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Guosen Yan

A Festschrift from Theoretical Chemistry Accounts



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Guosen Yan

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Contents

A tribute to Guosen Yan	1
Hua Guo, Daiqian Xie, Weitao Yang	
Transition between direct gap and indirect gap in two dimensional hydrogenated honeycomb Si _x Ge _{1-x} alloys	7
Nan Xia, Lan-Feng Yuan, Jinlong Yang	
A new ab initio potential energy surface and infrared spectra for the He–CS ₂ complex	13
ling Yuan, Hua Zhu	
Testing exchange–correlation functionals at fractional electron numbers Ali Malek, Degao Peng, Weitao Yang, Robert Balawender, Andrzej Holas	21
Stereoselective inclusion mechanism of ketoprofen into β-cyclodextrin: insights from molecular dynamics simulations and free energy calculations Mingsong Shi, Chunchun Zhang, Yani Xie, Dingguo Xu	29
State-to-state quantum dynamics of the N(⁴ S) + C ₂ ($\tilde{x}^{1}\Sigma^{+}$) \rightarrow CN($\tilde{x}^{2}\Sigma^{+}$) + C(³ P) reaction	41
Mechanism and regioselectivity of the cycloaddition between nitrone and dirhodium vinylcarbene catalyzed by Rh ₂ (O ₂ CH) ₄ : a computational study Xin Yang, Pan Xu, Ying Xue	49
Theoretical study on the working mechanism of a reversible light-driven rotary molecular motor	59
Lin wang, Ain Che, Humang Li, Jun Gao, Chengou Liu	
The combined CASPT2 and CASSCF studies on photolysis of 3-thienyldiazomethane and subsequent reactions Pei-Jie Guan, Wei-Hai Fang	67
Analysis of a failure of the CC2 coupled-cluster method for bond lengths of SnO and PbO Zhifan Wang, Fan Wang	75
Electrostatic potentials of camptothecin and its analogues Jianyong Lei, Yun Chen, Xuejun Feng, Jian Jin, Jiande Gu	83
A Comparison study of the H + CH ₄ and H + SiH ₄ reactions with eight-dimensional quantum dynamics: normal mode versus local mode in the reactant molecule vibration Yan Wang, Jun Li, Hua Guo, Minghui Yang	89
How the molecular face and the interaction vary as H atom approach H_2 molecule	99
Dong-Xia Zhao, Zhong-Zhi Yang	

State-to-state quantum versus classical dynamics study of the OH + CO \rightarrow H + CO ₂ reaction in full dimensions ($J = 0$): checking the validity	
of the quasi-classical trajectory method	111
Shu Liu, Jun Chen, Bina Fu, Dong H. Zhang	
NO adsorption and transformation on the BaO surfaces from density	
functional theory calculations	121
Nai-Xia Lu, Jing-Cong Tao, Xin Xu	
Microwave and infrared spectra of CO–(pH ₂) ₂ , CO–(oD ₂) ₂ , and mixed	105
CO–pH ₂ –He trimers	135
Xiao-Long Zhang, Hui Li, Robert J. Le Roy, Pierre-Nicholas Roy	
Numerical solution of solvent reorganization energy and its application	
in electron transfer reaction	149
Ting-Jun Bi, Mei-Jun Ming, Hai-Sheng Ren, Jian-Yi Ma, Xiang-Yuan Li	
Adsorption of water molecules on sodium chloride trimer	155
Cheng-Wen Liu, Gao-Lei Hou, Wei-Jun Zheng, Yi Qin Gao	
Thermal decomposition of FOX-7 studied by ab initio molecular dynamics simulations	165
Vong Liu Eong Li Hugi Sun	105
Tang Liu, reng Li, Huai Sun	
Hemibond complexes between H ₂ S and free radicals (F, Cl, Br, and OH) Benjamin Alday, Ryan Johnson, Jun Li, Hua Guo	177
DFT and TD-DFT studies on osmacycle dyes with tunable photoelectronic properties for solar cells Mingjun Sun, Zexing Cao	183
First-principles investigations on the anisotropic charge transport in 4,4'-bis((<i>E</i>)-2-(naphthalen-2-yl)vinyl)-1,1'-biphenyl single crystal Lili Lin, Xin Li, Guangjun Tian, Hua Geng, Zhigang Shuai, Yi Luo	191
Iron carbonyl thioboronyls: effect of substitution of sulfur for oxygen	
in the viability of binuclear complexes toward dissociation reactions	201
Xiaoli Gong, Liyao Zhu, Jing Yang, Xiumin Gao, Yaoming Xie, R. Bruce King	

INTRODUCTION

A tribute to Guosen Yan

Hua Guo · Daiqian Xie · Weitao Yang

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Professor Guosen Yan was born in 1930 in Nanchong, Sichuan Province, China. He graduated in 1951 from Chongqing University, majoring in Chemistry, and upon graduation joined the faculty in the same university. In 1953, he was transferred to Department of Chemistry in Sichuan University, rising through the ranks and became Associate Professor in 1962 and Full Professor in 1979. He served as the President of Sichuan University from 1984 to 1989. His leadership helped to secure Sichuan University as one of the top comprehensive universities in China. Professor Yan has also served as a director and executive director of the Chinese Chemical Society and in many other important posts in China, helping to lay a solid foundation for the ongoing expansion of the Chinese research enterprise.

Published as part of the special collection of articles celebrating the 85th birthday of Professor Guosen Yan.

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Key Laboratory of Theoretical Chemistry of Environment, Ministry of Education, School of Chemistry and Environment, South China Normal University, Guangzhou 510006, China Professor Yan is a pioneer in Theoretical Chemistry in China. He was one of eight original students of Professor Au-chin Tang, a Columbia-trained theoretical chemist, widely considered as the founding father of Theoretical Chemistry in China. Professor Yan's research interests spanned several areas, including crystal field theory, graph theory, quantum chemistry, and molecular spectroscopy. His major scientific contributions include (1) analysis of the symmetry and orthogonalization of the rotational group-point group coupling coefficients in ligand field theory, (2) graph theory for determinant expansions, (3) a vibrational self-consistent field/configuration interaction theory, and (4) molecular spectroscopy and potential energy surfaces. He has written three books and close to 200 journal articles.

Professor Yan has also been a great educator. His lectures were legendary. Whether it is quantum mechanics or group theory, he presented the materials in his characteristically meticulous fashion and filled the board with perfectly written derivations and equations, never miswriting or erasing anything. He has educated more than 20 graduate students (see list), many of whom have become accomplished scientists in China. His deep understanding of quantum mechanics and group theory greatly influenced his students.

In addition to being an accomplished scientist, Professor Yan also epitomizes a classical Confucius scholar. He read widely Chinese history, poetry, literature, and philosophy. Professor Yan is also an accomplished Chinese calligrapher. He has not only influenced many researchers through his scientific insights, but also helped many of us to shape our views of science, philosophy, and life.

On this occasion of Professor Yan's 85th birthday, we celebrate his accomplishments in science and education in this Special Collection. The breadth and depth of the work reported here are testaments of the vitality of theoretical

and computational chemistry in China, the fruits of the seeds sown by Professor Yan and his contemporaries.



List of students of Professor Guosen Yan

	Year	First name	Last name	中文姓名
M.S.	1978	Yaoming	Xie	谢尧明
	1981	Kaiqin	Lao	劳开勤
		Huai	Sun	孙 淮
	1982	Daping	Cao	曹大平
		Hua	Guo	郭 华
		Zhi	He	何 知
	1983	Gaoqing	Yuan	袁高清
		Chongde	Zheng	郑重德
	1984	Ruifeng	Liu	刘瑞峰
		Kehe	Su	苏克和
		Xingquan	Xiang	向兴全
		Xuefeng	Zhou	周学峰
Ph.D.	1987	Jie	Yang	杨捷
	1988	Changyuan	Tao	陶长元
	1989	Guoming	Liang	梁国明
	1990	Zexing	Cao	曹泽星
		Huabei	Zhang	张华北
	1991	Junkai	Xie	谢军楷
	1992	Xiangyuan	Li	李象远
		Jiande	Gu	顾建德
	1993	Hui	Xian	先 晖
	1994	Ying	Xue	薛 英
		Minghui	Yang	杨明晖
	1995	Yi	Ren	任 译
	1996	Xuejun	Xu	徐学军

1 Publications of Professor Guosen Yan

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REGULAR ARTICLE

Transition between direct gap and indirect gap in two dimensional hydrogenated honeycomb Si_xGe_{1-x} alloys

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Abstract Using first-principles calculations, we have explored the structural and electronic properties of fully hydrogenated honeycomb Si_rGe_{1-r}H alloys. Finite band gaps are opened by hydrogenation for x in the whole range from 0 to 1, while their nature and values can be tuned by x. When x is <0.7, the band gap is direct (from Γ to Γ). And when x is ≥ 0.7 , the gap turns into indirect (from Γ to M). For all the computed compositions, the two kinds of energy differences between valence band and conduction band, Γ - Γ and Γ -M, are described well by two polynomial functions of x. The smaller of the two functions gives a good prediction for the overall band gap at any x. The two curves cross at x = 0.7, leading to the change of band gap type. At PBE level, the values of band gap for different x spread from 1.09 to 2.29 eV. These findings give a new route to tune the electronic properties of these materials and may have potential applications in nanoscale optoelectronics.

Keywords Hydrogenated Si_xGe_{1-x} alloys · Virtual crystal approximation · Band gap engineering · First-principles calculation

Dedicated to Professor Guosen Yan and published as part of the special collection of articles celebrating his 85th birthday.

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1 Introduction

With the extensive applications of graphene in the electronic industry, in the past a few years more and more attentions have been paid to two dimensional (2D) atomically thin sheets, such as boron nitride (h-BN), transition metal dichalcogenides (e.g., MoS_2 and WS_2), and hybridized $B_xC_yN_z$ [1]. Similar to graphene, these 2D structures possess fantastic properties in various aspects (electronical, magnetical, optical, and mechanical, etc.) and exhibit great application prospects [2–4]. Silicene and germanene have been widely studied experimentally and theoretically [5–7], and it has been mentioned repeatedly that their band structures show linear band crossing at the Fermi level, and thus, the charge carriers behave like a massless Dirac fermion [8], similar to the case of graphene [9].

As two most important materials in the semiconductor industry, Si and Ge are both group 14 elements with similar structural properties; therefore, they can randomly mix to form alloys [10]. It has also been demonstrated that bulk Si_rGe_{1-r} alloys can be fabricated on Si substrates [11]. Not long ago, Padilha et al. [12] suggested that the electronic properties of $Si_x Ge_{1-x}$ random alloys in the honeycomb monolayer structure can be tuned by the value of x. To open a band gap and regulate the electronic properties of graphene-like materials, a variety of methods have been put forward, such as applying an external electric field [13], introducing hydrogen vacancies [14], uniaxial compression [15], biaxial compressive strain [16], cutting 2D materials into nanoribbons [17], and stacking into bilayer structures [18]. For silicene and germanene, hydrogenation is predicted to be a viable mean to open a finite gap [19, 20]. Therefore, it is highly probable that the 2D hydrogenated $Si_{r}Ge_{1-r}$ alloys also have finite gaps. If this is so, then we get a new parameter to manipulate this gap, i.e., the alloy





composition (*x*). In order to explore this route, we hereby investigate the evolution of the band structures of 2D disordered fully hydrogenated Si_xGe_{1-x} alloys (denoted as $Si_xGe_{1-x}H$), via first-principles calculations.

2 Methods and computational details

Our ab initio calculations are based on density functional theory (DFT) [21, 22] and are performed by the CASTEP software [23]. The generalized gradient approximation (GGA) with the Perdew-Burke-Ernzerholf (PBE) functional [24] is selected as the exchange correlation potential. For all the calculations, the ultrasoft pseudopotentials are used, with a 600 eV cutoff energy for the plane-wave basis. Both the lattice constants and the atomic positions are fully relaxed. The tolerances for geometry optimizations are 1.0×10^{-5} eV/atom for total energy, 0.02 eV/Å for maximum Hellman-Feynman forces, 0.1 GPa for maximum stress, and 0.002 Å for maximum ionic displacement. The periodic boundary condition is applied in the normal direction of the monolayer, with vacuum space of 20 Å to minimize the interlayer interaction. A primitive cell containing two Si (Ge) atoms and two H atoms is used, as shown in Fig. 1. Following the Monkhorst-Pack scheme [25], Brillouin zone integration is carried out at $9 \times 9 \times 1$ k-points for the geometry optimization and $15 \times 15 \times 1$ k-points for the static total energy calculations.

To simulate the electronic structure of these disordered alloys, the virtual crystal approximation (VCA) is used, which is a simple and reasonable approximation for this task. In this approximation, one studies a crystal with the primitive periodicity, but composed of fictitious atoms, which are hybrids of two or more element types [26]. The meaning of "hybrid" is that the potential of a virtual atom is treated as the concentration-weighted average of the constituent potentials, neglecting further compositional disorder effects [27]. VCA is widely used in the literature. For example, Bellaiche et al. [26] successfully employed VCA to study the dielectric and piezoelectric properties of the Pb($Zr_{0.5}Ti_{0.5}$)O₃ perovskite solid solution in its paraelectric and ferroelectric phases. Winkler et al. [28] showed that the VCA approach can be used to study Al/Si disorder in silicates, reproducing the experimental structural parameters. Employing VCA, a series of Si_xGe_{1-x}H alloys with

Employing VCA, a series of $Si_xGe_{1-x}H$ alloys with eleven different compositions have been simulated. The choices for *x* are from 0 (hydrogenated germanene) to 1 (hydrogenated silicene), with increment of 0.1, in the hope that the properties of an alloy with any *x* can be roughly estimated based on these eleven data points. There are two common configurations of fully hydrogenated silicene (germanene), i.e., boat like and chair like. It was reported that the chair configurations of silicene and germanene are more stable than the boat ones, by about 10 meV/atom [29]. In our calculations, for three typical alloys (x = 0.3, 0.5, and 0.8), the chair configurations are more stable than the boat ones as well, with the energy differences being 12.8, 12.8, and 13.9 meV/atom, respectively. Consequently, we will only consider the chair configurations of Si_xGe_{1-x}H alloys afterward.

3 Results and discussions

After geometry optimizations, the calculated lattice parameters *a* and atom bond lengths *d* of hydrogenated silicene and germanene are 3.88, 2.35, and 4.06, 2.45 Å respectively, which are consistent with the previous theoretical results 3.884, 2.356 Å [30] and 4.06, 2.46 Å [19]. Upon hydrogenation, the calculated buckling heights *h* are 0.72 and 0.75 Å, distinctly larger than those of silicene (0.45 Å) and germanene (0.64 Å). Namely, the hydrogenated configurations are more tortuous than the pristine structures. This can be understood in terms of variation in hybridization state along with hydrogenation. In silicene (germanene), the buckled structure gives rise to reduced π – π

Table 1 Structural and electronic parameters of Si_rGe_{1-r}H alloys in different composition of Si element (x)

a

Energy (eV)

с

Energy (eV)

x	a (Å)	$\Phi\left(\mathrm{eV}\right)$	$E_{\Gamma-\Gamma}$ (eV) (direct)	$\overline{E_{\Gamma-M}}$ (eV) (indirect)
0	4.058	4.809	1.089	1.912
0.1	4.046	4.855	1.435	2.035
0.2	4.035	4.870	1.710	2.135
0.3	4.023	4.972	1.905	2.207
0.4	4.009	4.932	2.058	2.250
0.5	3.998	4.950	2.157	2.271
0.6	3.983	4.992	2.234	2.285
0.7	3.962	5.015	2.297	2.285
0.8	3.941	5.028	2.327	2.262
0.9	3.915	5.012	2.340	2.224
1	3.884	5.029	2.332	2.174

The entries $a, \Phi, E_{\Gamma-\Gamma}$, and $E_{\Gamma-M}$ are the lattice constant, the effective work function, the gaps between conduction band and valence band from Γ to Γ and from Γ to M, respectively

overlap between p_z orbitals of adjacent Si (Ge) atoms, so the hybridization of each Si (Ge) atom is intermediate between sp² and sp³. After new σ bonds are formed with H atoms, the Si–Ge atoms in $Si_{,}Ge_{1,-}H$ become purely sp^{3} hybridized. Therefore, the bond angles in $Si_{y}Ge_{1-y}H$ are greater than those in silicene and germanene, resulting in higher degree of buckling.

For all the eleven compositions, the lattice parameters, the effective work functions (Φ), and two kinds of energy differences between valence band and conduction band $(\Gamma - \Gamma \text{ and } \Gamma - M)$ are listed in Table 1. The lattice parameter a decreases monotonically from 4.058 to 3.884 Å as x increases from 0 to 1, which is naturally owing to the larger atomic radius of Ge than that of Si. The effective work function, defined as the difference in potential energy of one electron between the highest occupied molecular orbital (HOMO) and the vacuum level, ranges from 4.81 to 5.03 eV.

To further illustrate how the composition affects the electronic properties of Si_xGe_{1-x}H, the band structures of four typical systems are presented in Fig. 2. In silicene and germanene, the π and π^* bands cross at the k-point around the Fermi level, rendering them gapless. On the other hand, all the 2D Si_xGe_{1-x}H alloys have finite band gaps, basically due to the interaction between the p_z orbitals of Si (Ge) and the 1 s orbitals of H atom. The valence band maximum (VBM) is always located at the Γ point. However, the conduction band minimum (CBM) can be either in Γ or in M, depending on the composition. Therefore, hydrogenated silicene is an indirect band gap (2.17 eV) semiconductor, while hydrogenated germanene is a direct band gap (1.09 eV) semiconductor. These results are qualitatively and quantitatively consistent with previous computational



Fig. 2 Band structures of $Si_xGe_{1-x}H$ alloys with four kinds of composition, i.e. x = 0 (a), 0.5 (b), 0.8 (c), 1 (d). The band gaps are noted in the picture. The Fermi levels are set to be energy zero

results, for instance, 2.0 eV [29], 2.19 eV [19] and 2.21 eV [14] for hydrogenated silicene, and 0.94 eV [19], 1.4 eV [29] and 1.5 eV [20] for hydrogenated germanene. Among the band gaps of our 11 systems, that of Si_{0.7}Ge_{0.3}H is found to be the biggest, which is an indirect gap of 2.29 eV, while the gap of hydrogenated germanene (1.09 eV) is the smallest. Thus, the band gap of hydrogenated Si_xGe_{1-x}H alloys can be tuned in the range from 1.09 to 2.29 eV by changing the composition. This is reminiscent of the result of Gao et al. [31] that the band gap of hydrogenated

x = 1



Fig. 3 Two sets of energy differences between valence and conduction bands of $\text{Si}_x\text{Ge}_{1-x}\text{H}$ as functions of Si composition (*x*): $\Gamma - \Gamma$ (*triangle dots*) and $\Gamma - M$ (*round dots*). The corresponding fourth order polynomial fitting curves are displayed as the *dash lines*

graphene can be continuously tuned by H coverage and configuration from 0 to 4.66 eV. That is a tuning in a wider range and with a constant nature of gap (direct), while our result is a tuning in a narrower range and with a variable nature of gap (from direct to indirect).

To gain further understanding on the transition of gap type, we list the Γ - Γ and Γ -M energy differences between valence band and conduction band for each composition in Table 1. We fit the two sets of data by two polynomial functions and find that fourth order polynomials are required to gain satisfactory match, as shown in Fig. 3. The curve for the Γ - Γ gap starts from the lowest value (1.089 eV) at x = 0 and increases quickly with x. When x approaches 1, this curve becomes flat. It reaches a maximum (2.340 eV) at x = 0.9 and drops a little bit (2.332 eV) eventually at x = 1. On the other hand, the curve for the Γ -M gap spreads in a much narrower range, from 1.912 to 2.285 eV. The maximum is reached at x = 0.6 and 0.7. In the beginning, i.e., for small values of x, the Γ - Γ gap is less than the Γ -M gap, so the system is a direct gap semiconductor. The two curves cross roughly at x = 0.7. After that, the order between the two gaps is reversed, and the system becomes an indirect gap semiconductor. Therefore, the band gap of any alloy can be represented by the smaller of the two polynomial functions, as listed below (in eV):

$$Eg = \begin{cases} 1.087 + 4.029x - 5.383x^2 + 3.875x^3 - 1.276x^4 (0 \le x < 7) \\ 1.911 + 1.465x - 1.948x^2 + 1.171x^3 - 0.425x^4 (0.7 \le x \le 1) \end{cases}$$

Using this piecewise function, one can describe well the band gap of $Si_xGe_{1-x}H$ for x along the whole range from 0 to 1.

At last, let us discuss the numerical uncertainty of our calculated gaps. There are two levels of errors, one from the treatment to disorder (i.e., VCA) and the other from the selection of functional (i.e., PBE). To check the viability of VCA in our systems, we compare the electronic properties by VCA and by conventional ordered PBE calculations for Si_{0.5}Ge_{0.5}H. We employ 6 kinds of supercells, i.e., $n \times n$ for n = 1-6, with different atomic arrangements of Si and Ge to simulate the randomized alloy. We test 8 arrangements for the 4×4 supercell and one arrangement for each of the other supercells. All the VCA and PBE calculations give rise to direct gaps. The band gaps from the supercell calculations are in the range of 1.68-1.72 eV, about 80 % of the VCA value 2.157 eV. Taking into consideration that a more realistic representation of the random alloy would require much larger supercells and many more atomic arrangements, this degree of agreement is acceptable. We acknowledge that VCA is a simple model, and the huge saving in computational cost is a major advantage and the main reason for us to adopt this method.

As for the second factor, GGA is known to tend to underestimate the band gaps of semiconductors and insulators and gives values typically 50-60 % smaller than those from better hybrid DFT methods. The gaps of fully hydrogenated silicene and germanene by HSE06 are reported to be 3.51 eV [32] and 1.56 eV [33], respectively. The optical properties of hydrogenated germanene were investigated by diffuse reflectance absorption (DRA) spectroscopy, and a linear approximation of the absorption edge suggests a band gap of approximately 1.59 eV [33]. There is no experimental data concerning the electronic properties of hydrogenated silicene. Therefore, the scaling factor for band gaps can be estimated as the ratio between the HSE06 to PBE values, which is 1.618 for hydrogenated silicene and 1.431 for hydrogenated germanene. So to compare quantitatively with experiments, it will be advisable to scale our PBE gaps by similar ratios, giving a range of 1.56-3.70 eV.

4 Conclusions

In summary, first-principles calculations based on the VCA have been applied to investigate the structural and electronic structures of 2D hydrogenated Si_xGe_{1-x} alloys. At PBE level, the band gap of these systems can be tuned in a range of 1.09–2.29 eV via varying their compositions. For x < 0.7, the alloys are direct band gap (Γ – Γ) semiconductors. And after x reaches 0.7, the band gaps transform to indirect (Γ –M). The origin of this transition is the change of the CBM location from Γ to M, or in another word, the crossing of the two curves of the Γ – Γ and Γ –M energy differences versus x. We fit the two curves by two polynomials, thus giving a tool to predict the band gap in the whole

range in terms of a piecewise function. Around the transition composition (x = 0.7), a small change in composition can lead to qualitative change in electronic and photonic properties for these alloys. Therefore, once combined with advanced Si/Ge nanotechnology, these materials have very valuable potential for band engineering and optical application.

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REGULAR ARTICLE

A new ab initio potential energy surface and infrared spectra for the He–CS₂ complex

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Abstract We report a new three-dimensional potential energy surface for the He–CS₂ complex including the Q_3 normal mode for the v_3 antisymmetric stretching vibration of the CS₂ molecule. The potential energies were calculated at the coupled-cluster singles and doubles with noniterative inclusion of connected triples level with augmented correlation-consistent quadruple-zeta basis set plus midpoint bond functions. Two vibrationally averaged potentials with CS_2 at both the ground (v = 0) and the first excited $(v = 1) v_3$ vibrational states were generated from the integration of the three-dimensional potential over the Q_3 coordinate. Both potentials have a T-shaped global minimum and two equivalent linear local minima. The radial discrete variable representation/angular finite basis representation method and the Lanczos algorithm were applied to calculate the rovibrational energy levels. Our calculated results show that the two potentials support eight vibrational bound states. The calculated band origin shift of the complex (0.1759 cm^{-1}) agrees very well with the observed one (0.1709 cm^{-1}) . The predicted infrared spectra and spectroscopic constants based on the two averaged potentials

Dedicated to Professor Guosen Yan and published as part of the special collection of articles celebrating his 85th birthday.

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T. Yuan · H. Zhu State Key Laboratory of Biotherapy, Sichuan University, Chengdu 610064, China are in excellent agreement with the available experimental values.

Keywords Ab initio potential energy surface \cdot Rovibrational energy levels \cdot Infrared spectra \cdot He–CS₂

1 Introduction

The weak intermolecular interactions and spectroscopy for the van der Waals (vdW) complexes of rare-gas (Rg) atoms with the CO₂ or CS₂ molecule have attracted considerable attention. Such studies have been aimed at understanding the nature of the intermolecular potential energy surface (PES) and dynamics of these weakly bound molecules. Carbon dioxide complexes have been thoroughly investigated experimentally [1–8] and theoretically [9–23], because CO₂ is one of the most important absorbers of infrared radiation in the earth's atmosphere and interstellar chemistry. As an analog of CO₂, CS₂ is also interesting because sulfur is a key element in chemistry and in the spectroscopy of giant planets [24]. It is interesting to explore the difference of structure and intermolecular dynamics between the CS₂ containing complexes and the CO₂ containing analogs.

The first spectra of the Ar–CO₂ complex were observed by Steed et al. [1] using the microwave spectroscopy. Subsequently, further experimental studies of microwave and infrared spectra for Rg–CO₂ (Rg = He, Ne, Ar, Kr) were reported [2, 3, 7, 8]. Along with these experimental studies, a number of theoretical researches have dealt with the construction of the ab initio potential energy surface and the calculation of the rovibrational energy levels. In the previous theoretical calculations [9–17], the CO₂ monomer was assumed as a rigid rotor, which is not sufficient for predicting the infrared spectra, since the transition in the intramolecular vibrational mode is usually involved [25]. In addition, the shifts of the band origin in the infrared spectra cannot be correctly reproduced using the rigid monomer model. One effective way is to take into account the influence of intramolecular vibration mode on the potential energy surface and dynamical calculations [26–32]. To date, we have successfully used this approach to construct the potentials including the dependence on the Q_3 normal coordinate of the CO₂ molecule and to reproduce the infrared spectra for Rg–CO₂ [20–23].

Until now, only a few studies of the complex involving CS₂ have appeared [33–38]. Mivehvar et al. [36] first reported the infrared spectra of He-, Ne-, and Ar-CS2 in the region of the CS_2 v_3 fundamental band around $1,535 \text{ cm}^{-1}$. The results showed that the Rg–CS₂ complexes have T-shaped structures. The vibrational band origin shifts and spectroscopic constants for Rg-CS₂ were also presented in their paper. Subsequently, Farrokhpour and Tozihi [37] constructed the ab initio potential energy surface of the Rg–CS₂ complexes (Rg = He, Ne, and Ar) using the coupled-cluster singles and doubles with noniterative inclusion of connected triple [CCSD(T)] theory with aug-cc-pVDZ basis set plus bond functions. Very recently, Zang et al. [38] reported the two-dimensional PESs for the three complexes, which were calculated using aug-ccpVTZ basis set at CCSD(T) level. Based on the ab initio potential energy surfaces, they further calculated the rovibrational bound states and pure rotational transition frequencies. In the above two theoretical studies, the CS₂ molecule was treated as a rigid rotor.

The purpose of this work was to present a reliable three-dimensional potential energy surface involving the antisymmetric Q_3 normal mode of CS₂ for the He–CS₂ complex and to provide more detailed theoretical information on the dynamics including the rovibrational energy levels, bound states, and transition line intensities. The potential-optimized discrete (PODVR) [39, 40] grid points were employed to construct the potential energy surface. The radial discrete variable representation (DVR)/angular finite basis representation (FBR) method and Lanczos algorithm were used to calculate the bound states and rovibrational energy levels. The transition frequencies and line intensities and the band origin shift were also calculated and compared with the available observed results.

2 Computational details

2.1 Ab initio calculations

The geometry of the He–CS₂ complex is defined in Jacobi coordinates: R, θ , Q_3 . R represents the distance from the CS₂ center-of-mass to the He atom. θ is the enclosed angle

between the *R* vector and the molecular axis of CS₂. The coordinate Q_3 denotes the normal mode for v_3 antisymmetric stretch of CS₂. Q_3 is defined as $Q_3 = (r_{cs1} - r_{cs2})/\sqrt{2}$, where rcs_1 and rcs_2 are the two C–S bond lengths of CS₂, respectively. We used one-dimensional potential for the Q_3 coordinate of CS₂ to determine the energy levels and wave functions for the Q_3 mode. A coordinate scaling method [41] was used to adjust the CCSD(T) potential curve to reproduce the experimental v_3 fundamental frequency value [42], and the scale factor is 0.964322 and 0.978711 for the ground and the v_3 excited states of CS₂, respectively. Thus, we generated five PODVR grid points of -0.24995, -0.11881, 0.0, 0.11881, and 0.24995 a_0 for the ground state, and -0.24989, -0.11878, 0.0, 0.11878, and 0.24989 a_0 for the v_3 excited state.

The intermolecular potential energies of He–CS₂ were calculated using the supermolecular approach at the CCSD(T) level [43] for a total of about 1,000 points at each of the ground and the first excited v_3 states of CS₂, with *R* ranging from 4.50 a_0 to 22.00 a_0 at 26 values, and 13 equidistant angles from $\theta = 0^{\circ}$ to 180° in a step of 15°. We employed the augmented correlation-consistent quadruple-zeta (aug-cc-pVQZ) basis set [44] for all atoms and the bond functions (3s3p2d1f1g) [45]. The full counterpoise procedure (FCP) [46] was used to correct the basis set superposition error (BSSE).

To test the convergence of the basis set, a few additional calculations using a extrapolation formula [47] were performed to estimate the complete basis set (CBS) limit for two points (R, θ) = (3.38 Å, 90.0°) and (4.50 Å, 45.0°) on the potential of He–CS₂ with CS₂ at the ground state. The ab initio calculated intermolecular energies are -54.06 and -16.44 cm⁻¹, respectively, whereas the extrapolation



Fig. 1 Contour plots (in cm⁻¹) of the averaged intermolecular potential energy surface for He–CS₂ with CS₂ at the ground (v = 0) state

Table 1 Calculated energy levels (in cm⁻¹) for the pure vdW vibrational bound states of the He–CS₂ complex for the v = 0 and 1 states of CS₂

N	Ground state	v_4 State
0	-21.352	-21.176
1	-10.536	-10.568
2	-10.444	-10.484
3	-9.125	-9.107
4	-6.909	-6.895
5	-4.198	-4.215
6	-2.849	-2.844
7	-0.185	-0.182

procedure gave the energies of -54.52 and -16.98 cm⁻¹. It is clear that the depth of the potentials presented in this work is only slightly underestimated because of the basis set incompleteness. All the ab initio calculations were carried out using the MOLPRO package [48].

2.2 Analytical fitting of the potential energy surface

The vibrationally averaged PESs $V_{\nu}(R, \theta)$ for the He–CS₂ complex can be defined as [49].



$$V_{\nu}(R,\theta) = \int_{-\infty}^{\infty} \psi_{\nu}(Q_3) V(R,Q_3,\theta) \psi_{\nu}(Q_3) dQ_3, \qquad (1)$$

where v denotes the quantum number for a specific vibrational state of isolated CS_2 molecule, and $\psi_v(Q)$ is the corresponding vibrational wave function.

The calculated ab initio potential energies can be fitted to an analytical form, [50–53]

$$V(R,\theta) = V_{\rm sh}(R,\theta) + V_{\rm as}(R,\theta).$$
⁽²⁾

The short-range part of the interaction is assumed to be

$$V_{\rm sh}(R,\theta) = G(R,\theta) e^{D(\theta) - B(\theta)R}$$
(3)

where $G(R, \theta)$, $D(\theta)$, and $B(\theta)$ are expanded as series in Legendre polynomials $P_1(\cos\theta)$,

$$G(R,\theta) = R^{-1} \sum_{i=1}^{2} R^{i} \sum_{l=0}^{8} g_{i,l} P_{l}(\cos \theta),$$
(4)

$$D(\theta) = \sum_{l=2}^{2} d_l P_l(\cos \theta), \tag{5}$$

$$B(\theta) = \sum_{l=0}^{8} b_l P_l(\cos \theta), \tag{6}$$



Table 2 Rovibrational energy levels (in cm⁻¹) of He–CS₂ for the v = 0 and 1 states of CS₂

Level	J	$E\left(J_{KaKc}\right)$					
		Even/even	Even/odd	Odd/even	Odd/odd		
Ground state	0	-21.352 (000)					
	1		-21.160 (1 ₀₁)	-20.924 (1 ₁₀)	-20.955 (111)		
	2	$-20.781(2_{02})$ $-19.894(2_{20})$	-20.510 (2 ₂₁)	-20.604 (2 ₁₂)	-20.510 (2 ₁₁)		
	3	-19.326 (3 ₂₂)	$-20.219(3_{03})$ $-19.310(3_{21})$	-19.892 (3 ₁₂) -18.230 (3 ₃₀)	$-20.079(3_{13})$ $-18.230(3_{31})$		
	4	$\begin{array}{r} -19.485~(4_{04})\\ -18.520~(4_{22})\\ -15.954~(4_{40})\end{array}$	$-18.566(4_{23}) \\ -15.954(4_{41})$	-19.382 (4 ₁₄) -17.468 (4 ₃₂)	$-19.073 (4_{13}) \\ -17.466 (4_{31})$		
	5	$-17.621\ (5_{24})\\-15.011\ (5_{42})$	$\begin{array}{c} -18.585\ (5_{05})\\ -17.518\ (5_{23})\\ -15.011\ (5_{41})\end{array}$	$\begin{array}{r} -18.055~(5_{14})\\ -16.507~(5_{32})\\ -13.085~(5_{50})\end{array}$	$-18.516(5_{15})$ $-16.513(5_{33})$ $-13.085(5_{51})$		
v_4 State	0	-21.176 (0 ₀₀)					
	1		-20.985 (1 ₀₁)	-20.750 (1 ₁₀)	-20.780 (111)		
	2	$-20.608(2_{02})$ $-19.723(2_{20})$	-19.726 (2 ₂₁)	-20.431 (2 ₁₂)	-20.339 (2 ₁₁)		
	3	-19.158 (3 ₂₂)	$-20.050(3_{03})$ -19.143(3 ₂₁)	-19.725 (3 ₁₂) -18.064 (3 ₃₀)	$-19.910(3_{13})$ $-18.064(3_{31})$		
	4	$\begin{array}{r} -19.320~(4_{04})\\ -18.358~(4_{22})\\ -15.795~(4_{40})\end{array}$	-18.404 (4 ₂₃) -15.795 (4 ₄₁)	$-19.217 (4_{14}) \\ -17.307 (4_{32})$	$-18.910 (4_{13}) \\ -17.306 (4_{31})$		
	5	$-17.464(5_{24})$ $-14.859(5_{42})$	$-18.425 (5_{05})$ $-17.362 (5_{23})$ $-14.859 (5_{41})$	$-17.899(5_{14})$ $-16.353(5_{32})$ $-12.936(5_{50})$	$-18.356(5_{15})$ $-16.358(5_{33})$ $-12.936(5_{51})$		

The long-range part $V_{\rm sh}(R, \theta)$ is expressed as,

$$V_{\rm as}(R,\theta) = -\sum_{n=6}^{12} \sum_{l=0}^{8} f_n(B(\theta)R) c_{n,l} R^{-n} P_l(\cos\theta)$$
(7)

where $f_n(B(\theta)R)$ is employed to damp the long-range force at short intermolecular distance, which is assumed to be in the Tang–Toennies form [54],

$$f_n(x) = 1 - e^{-x} \sum_{k=0}^n \frac{(x)^k}{k!}$$
(8)

2.3 Calculations of the rovibrational states

The vibrational averaged intermolecular Hamiltonian for the He–CS₂ complex in the Jacobi coordinates can be written as [55, 56]

$$\widehat{H} = -\frac{1}{2\mu} \frac{\partial^2}{\partial R^2} + \frac{\widehat{j}^2}{2I_v} + \frac{(\widehat{\mathbf{J}} - \widehat{j})^2}{2\mu R^2} + V_v(R,\theta),$$
(9)

where μ is the reduced mass of the He–CS₂ complex, $I_{\upsilon} = \langle \psi_{\upsilon}(Q_3) | I_{Q_3} | \psi_{\upsilon}(Q_3) \rangle$ is the vibrationally averaged rotational moment of inertia of CS₂ monomer, $\hat{\mathbf{J}}$, and \hat{j} are the angular momentum operators corresponding to the total and the CS₂ monomer rotations. $V_{\upsilon}(R, \theta)$ is the vibrationally averaged potential in a particular vibrational state v (v = 0, 1) of CS₂. The efficient radial DVR/angular FBR method [57, 58] was used to calculate the rovibrational energy levels. In our work, we used 120 sine-DVR [59] points for the *R* coordinate, 89 basis functions of associated Legendre polynomials, and 90 Gauss–Legendre grids for the angular coordinate. The Lanczos algorithm [60, 61] was employed to efficiently diagonalize the Hamiltonian matrix with 5,000 steps. The parity-adapted rotational basis is defined in the three Euler angles (α , β , γ) denoting the orientation of BF frame with respect to the SF frame,

$$C_{KM}^{Jp}(\alpha, \ \beta, \ \gamma) = [2(1+\delta_{K0})]^{-1/2} [D_{MK}^{J*}(\alpha, \ \beta, \ \gamma) + (-1)^{J+K+p} D_{M-K}^{J*}(\alpha, \ \beta, \ \gamma)], \qquad (10)$$
$$p = 0, \ 1$$

where $D_{MK}^{J}(\alpha, \beta, \gamma)$ is the normalized rotational function. The total parity is given by $(-1)^{J+p}$. We calculated the rovibrational energy levels with the total angular momentum $J \leq 8$.

3 Features of potential energy surface

The root mean square (rms) errors between the fitted and calculated points at v = 0 and 1 states of CS₂ are found to be about 0.04 cm⁻¹. The ab initio points and fitted potential

Table 3 Spectroscopic constants (in cm⁻¹, 1 cm⁻¹ = 29.979.25 MHz) for the He–CS₂ complex at both the ground and v_4 states

	Ground state		v_4 State	
	Experiment	This work	Experiment	This work
A	0.321565	0.316699	0.320765	0.315786
В	0.111322	0.111404	0.110604	0.110678
С	0.080321	0.079994	0.079845	0.079529
Δ_K	3.148×10^{-4}	6.231×10^{-5}	3.347×10^{-4}	6.205×10^{-5}
Δ_{JK}	1.077×10^{-4}	1.171×10^{-4}	1.078×10^{-4}	1.182×10^{-4}
Δ_{I}	6.701×10^{-7}	1.057×10^{-6}	6.701×10^{-7}	1.065×10^{-6}
δ_K	6.347×10^{-5}	7.302×10^{-5}	6.446×10^{-5}	7.407×10^{-5}

Experimental data are taken from Ref. [36]

parameters for He–CS₂ are presented in Tables S1 and S2 in the supplementary material, respectively. The contour plot of $V_{\nu}(R, \theta)$ ($\nu = 0$) is shown in Fig. 1. It is clear that the CCSD(T) potential is characterized by a global T-shaped minimum and two linear local minima. The global minimum is situated at R = 3.38 Å and $\theta = 90.0^{\circ}$ with a depth of 54.06 cm⁻¹. The two equivalent local minima of -32.04 cm⁻¹ occur at the linear geometry for R = 5.02 Å. A barrier separating the two minima is found at R = 4.50 Å and $\theta = 59.50^{\circ}$ with height of 10.62 cm⁻¹,

relative to the linear minimum. The barrier of 10.62 cm^{-1} for He–CS₂ is higher than that of 2.38 cm⁻¹ for He–CO₂ [18], which means that the linear minimum of the former is easier to trap the He atom, leading to a less wide delocalized angular distribution.

4 Rovibrational bound states

We follow Ref. [6] to label the v_3 antisymmetric stretch of CS_2 as the v_4 vibrational mode in the He-CS₂ complex. Table 1 presents the energy levels for the total pure vibrational bound states at the v = 0 and 1 states of CS₂. One can see that ab initio CCSD(T) potential supports eight vibrational bound states, while He-CO₂ only has five bound states [18]. The vibrational ground state of the He-CS₂ complex with CS₂ at the ground state is bound by 21.35 cm⁻¹, corresponding to a zero-point energy of 32.71 cm^{-1} . The zero-point energy is over half of the global well depth, similar to other He-containing vdW complexes. Our predicted vibrational band origin for the He–CS₂ complex is blue shifted by 0.1759 cm⁻¹, very close to the observed value [36] of 0.1709 cm^{-1} . Although the band origin shift for He-CS₂ is larger (0.1759 vs. 0.1034 cm^{-1}) than that for He–CO₂ [18], the CS₂ and CO₂ shifts are small in magnitude and positive in sign, which

Table 4 Calculated infrared transition frequencies (in cm⁻¹) and relative intensities for He–CS₂

$\overline{J'_{Ka'Kc'}-J''_{Ka''Kc''}}$	v_{cal}	$v_{cal} - v_{obs}$	Intensity	$J'_{Ka'Kc'} - J''_{Ka''Kc''}$	$v_{\rm cal}$	$v_{cal} - v_{obs}$	Intensity
5 ₀₅ 6 ₀₆	1,534.4569	0.0097	0.414	3 ₁₂ -2 ₁₁	1,536.1418	0.0048	0.787
4 ₂₃ -5 ₂₄	1,534.5733	0.0070	0.311	4 ₁₄ -3 ₁₃	1,536.2178	0.0031	0.899
3 ₁₂ -4 ₁₃	1,534.7040	0.0051	0.555	423-322	1,536.2783	0.0036	0.503
321-422	1,534.7333	0.0047	0.342	5 ₀₅ -4 ₀₄	1,536.4153	0.0016	0.884
3 ₀₃ -4 ₀₄	1,534.7904	0.0067	0.711	5 ₄₁ -4 ₄₀	1,536.4512	0.0030	0.606
221-322	1,534.9559	0.0056	0.280	5 ₃₂ -4 ₃₁	1,536.4700	0.0034	0.221
2 ₁₂ -3 ₁₃	1,535.0037	0.0062	0.640	5 ₂₃ -4 ₂₂	1,536.5137	0.0047	0.479
110-211	1,535.1170	0.0050	0.443	514-413	1,536.5294	0.0035	0.707
101-202	1,535.1513	0.0054	0.671	6 ₁₆ -5 ₁₅	1,536.5430	0.0012	0.660
4 ₁₄ -4 ₁₃	1,535.2115	0.0014	0.671	6 ₂₅ -5 ₂₄	1,536.6352	0.0022	0.394
6 ₂₅ -6 ₂₄	1,535.3146	-0.0010	0.004	6 ₃₄ –5 ₃₃	1,536.6517	0.0027	0.197
3 ₂₁ -3 ₂₂	1,535.5393	0.0054	0.382	7 ₀₇ –6 ₀₆	1,536.7182	0.0001	0.778
110-111	1,535.5615	0.0050	0.548	7 ₄₃ -6 ₄₂	1,536.8206	0.0023	0.020
5 ₂₃ -5 ₂₄	1,535.6147	0.0079	0.096	8 ₁₈ -7 ₁₇	1,536.8558	-0.0001	0.602
3 ₁₂ -3 ₁₃	1,535.7103	0.0068	0.139	7 ₃₄ –6 ₃₃	1,536.8603	0.0044	0.143
1 ₀₁ -0 ₀₀	1,535.7223	0.0045	0.443	7 ₁₆ -6 ₁₅	1,536.8872	0.0017	0.344
7 ₂₅ -7 ₂₆	1,535.8068	0.0130	0.016	7 ₂₅ 6 ₂₄	1,536.9253	0.0048	0.723
2 ₁₂ -1 ₁₁	1,535.8797	0.0043	0.548	8 ₂₇ -7 ₂₆	1,536.9761	0.0010	0.179
5 ₁₄ -5 ₁₅	1,535.9732	0.0100	0.043	845-744	1,537.0038	0.0017	0.045
3 ₀₃ -2 ₀₂	1,536.0864	0.0037	1.000	8 ₃₆ -7 ₃₅	1,537.0188	0.0020	0.259
321-220	1,536.1073	0.0046	0.367	Rms error	0.005		

Experimental data are taken from Ref. [36]

Fig. 3 Observed and simulated line intensities for the v_4 band of He–CS₂



shows a slight weakening of the vdW bond upon vibration excitation for the two complexes.

To shed some light on the dynamic of the complex, it is useful to visualize the wave functions. Figure 2 presents the contour plots of the wave functions for the ground and the first three lower excited states of He-CS₂ with CS₂ at the ground (v = 0) state. The vdW vibrational states are defined by two quantum numbers (n_s, n_b) , respectively, denoting the vdW stretching and bending modes. One can see that the ground state (0, 0) is localized at the T-shaped global minimum. Similar to He–CO₂, the wave functions for the lower excited states show predominantly bending character, which means that the potential is too shallow to support stretch vibrations along the R coordinate for the small reduced mass of the complex. However, some significant deviations do exist. For example, the wave function of the first excited state (0, 1) for He–CS₂ is distributed less widely than that for He–CO₂ along the angular coordinate. The average structures were, respectively, determined from the vibrational ground state wave function of the complex, $\langle R \rangle = \langle \psi_0 | R | \psi_0 \rangle$ and $\langle \cos^2 \theta \rangle = \langle \psi_0 | \cos^2 \theta | \psi_0 \rangle$. The calculated average distance *R* for the ground state is 3.766 Å, which is about 0.39 Å longer than its equilibrium value, and the angle θ is 83.53°, the latter is a measure of the average amplitude of the bending from the 90° average configuration, which can be interpreted as an effect of the zeropoint vibrational motion of CS₂ within the complex. The ground state of the system is quite far from rigid; the radial dispersion and angular dispersion are 0.287 Å and 6.88°, respectively.

The rovibrational energies were assigned by the antisymmetric rotor quantum numbers J_{KaKc} . *J* is the total angular momentum; K_a and K_c denote the projections of *J* onto the *a* and *c* axes in the principal axes of inertia. The rovibrational energy levels consist of four blocks, (even/ even), (even/odd), (odd/even), and (odd/odd) for different combination parity of (j/p). The calculated rovibrational energies for $J \leq 5$ of the vdW ground vibrational state

with CS₂ at both the ground (v = 0) and the first excited (v = 1) v_3 vibrational states are presented in Table 2.

5 Infrared spectra

The rovibrational energy levels of the vdW ground and v_4 states for $J \le 3$ were fitted to a Watson asymmetric top Hamiltonian [62] using the *a*-type reduction in the I^r representation. The fitted molecular spectroscopic constants are given in Table 3 together with the experimental results [36]. It is obvious that the rotational constants A, B, and C agree well with the experimental values. The inertial defect Δ_0 in the ground state of He–CS₂ was calculated to be 6.19 amu Å², whereas for the He–CO₂ complex, Δ_0 is 8.5 amu Å² [18], which indicates that the He–CS₂ complex is less floppy than the CO₂ analogs.

The calculated infrared transition is defined as the difference between the appropriate rovibrational energies on the upper (v = 1) and lower (v = 0) 2D surface, plus the experimental CS₂ band origin energy. The details regarding calculation of the transition intensities can be found in Ref. [63]. The calculated rovibrational transition frequencies and the relative intensities at the effective temperature of T = 3.0 K for He–CS₂ as well as the comparison with the observed infrared spectra are given in Table 4. The calculated transition frequencies are all in excellent agreement with the experimental values. Most of the calculated frequencies are within 0.005 cm⁻¹ of the observed values, and the root mean squares error of the 41 calculated transitions is only 0.005 cm^{-1} . Figure 3 plots a portion of the simulated infrared spectra, which is consistent with the observed spectra of Mivehvar et al. [36]. It is clear that both spectra have the 4_{14} - 3_{13} , 5_{05} - 4_{04} , 3_{12} - 2_{11} , and 7_{06} - 6_{06} , and the transition from 3_{03} to 2_{02} is found to be the strongest. Both the transition frequencies and intensities show that the new averaged potentials can reproduce the observed infrared spectra of the He-CS₂ complex well.

6 Conclusions

We have presented a three-dimensional PES for the He– CS₂ complex including the Q_3 normal mode of CS₂ at the CCSD(T) level with aug-cc-pVQZ basis set plus bond functions. The intermolecular potential was calculated over five PODVR grid points for the Q_3 normal mode. Based on the ab initio potential points, two vibrationally averaged PESs of the complex were generated. Each PES has a T-shaped global minimum and two equivalent local minima of linear geometry. The bound rovibrational energy levels of He–CS₂ were obtained by employing the radial DVR/angular FBR method and the Lanczos algorithm. The calculated spectroscopic constants and infrared spectra are in good agreement with the available observed values. The experimental band origin shift in the infrared spectra is reproduced very well. The v_3 antisymmetric and v_1 symmetric stretching vibrations of the CS₂ molecule will be both considered in the future work, and we will investigate the effect on the potential and spectroscopic properties of He–CS₂.

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REGULAR ARTICLE

Testing exchange–correlation functionals at fractional electron numbers

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Abstract Dependence of the ground-state energy and density on the fractional electron number should be linear, while that of the chemical potential and Fukui function should be constant within the region between integers. In practical calculations with commonly used approximate functionals, results deviate from these ideal conditions. Four indicators are defined here to measure such deviations: from the global and local linearity condition, and from the global and local constancy condition. These indicators are used to test the performance of DFT method with five exchange–correlation functionals, and also the Hartree–Fock method, on a set of high-symmetry atoms: He, Li, Be, Na, Mg. The rCAM-B3LYP functional, having all four indicators small, is found to be the best functional for fractional electron number applications.

Keywords Exchange–correlation potentials · Density functional · Fractional particle number

Dedicated to Professor Guosen Yan and published as part of the special collection of articles celebrating his 85th birthday.

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1 Introduction

Density functional theory (DFT) is, at present, the method of choice for the description of the ground-state (GS) properties of quantum systems with relatively low computational complexity. A practical and effective approach for electronic structure calculations is offered by Kohn-Sham (KS) method. The great success of KS-DFT is due to the fact that simple density functional approximations (DFAs) perform remarkably well for a wide range of problems in chemistry and physics, particularly for the prediction of the structure and thermodynamic properties of molecules and solids. DFT owes its power to an efficient formalism founded on the single-determinant state function that should in principle be able to yield the exact energy and density. All the complexity of this theory is hidden in one term-the exchange-correlation functional. This term holds 'the key to the success or failure of DFT' [1]. The main problem is that the explicit form of the exact functional is unknown, and only certain desired properties which should be satisfied by approximate functionals can be formulated [2, 3]. Failures in practical calculations are not the breakdown of the theory itself, but are only due to deficiencies in commonly used approximations [1]. Common approximations for the exchange-correlation functional have been found to give big errors for the linearity condition of fractional charges (electron numbers), leading to delocalization error, and for the constancy condition of fractional spins (the energy of the ensemble constructed from the components of the spin doublet is independent of the spin number), leading to static correlation error [1, 4, 5]. Violation of these conditions underlies the major failures of all known approximate functionals to describe the energy gaps and related properties in strongly correlated systems [6]. The linearity condition of the energy versus

the number of electrons N follows from the zero-temperature limit of the temperature extension of the DFT [7–9] or, alternatively, from the pure states for a collection of replicas at zero temperature [2, 10]. Standard local or semi-local DFAs such as local density approximation (LDA), generalized gradient approximation (GGA) and meta-GGA, result in a very poor description, giving a smooth, convex, almost parabolic, interpolation between the values at integers. Some corrections applied to these functionals straighten the curve for fractional charges but significantly worsen the description at integers. Hartree-Fock (HF) results have the opposite behavior, with a concave interpolation between the integers. However, the hybrid functionals which combine these two ingredients are still quite convex [11, 12]. There have been two strategies to tackle this problem. The first one is the scaling correction method [13–15] or the restoration scheme [16, 17] applied to existing functionals. The second is the construction of new functionals which satisfy the energy linearity condition [11, 18, 19] or the constancy condition of the frontier orbital energy (interpreted as the ionization energy) [20-22]. The "dilemma" of how to define the exchange-correlation potential for fractional N was discussed in early paper [23], while the papers [24, 25] represent the most recent investigations related to the fractional-N problem for density functionals. Recently, an orbital functional based on the particle-particle random phase approximation to many-body theory has been shown to exhibit minimal errors for fractional charges and fractional spins and have many promising features [26, 27].

In this article, we investigate four properties related to fractional number of electrons to test the quality of the exchange–correlation functional. As illustrative example, the results for DFT method with five exchange–correlation functionals (SVWN5, PBE, B3LYP, CAM-B3LYP, rCAM-B3LYP) and the Hartree–Fock method are presented.

2 Theoretical background

In the exact DFT [28], the total energy of the system is a functional of the electron density and the external potential $v(\mathbf{r})$,

$$E[\rho, v] = T[\rho] + E_{ee}[\rho] + \int v(\mathbf{r})\rho(\mathbf{r})d\mathbf{r}$$

= $T_s[\rho] + E_{es}[\rho] + E_{xc}[\rho] + \int v(\mathbf{r})\rho(\mathbf{r})d\mathbf{r}$ (1)

where $T[\rho]$ and $E_{ee}[\rho]$ are the kinetic energy and the electron-electron interaction energy functionals, respectively. Following the Kohn-Sham (KS) scheme, the sum of these two functionals can be represented as the sum of the non-interacting reference KS kinetic energy,

 $T_s[\rho]$, the electrostatic energy, $E_{es}[\rho]$, and the exchange– correlation energy, $E_{xc}[\rho]$. The GS energy, E[N, v], is the minimum $E[N, v] = \operatorname{Min}_{\rho \to N} E[\rho, v]$ with the constraint $N = \int \rho(\mathbf{r}) d\mathbf{r}$. The minimizer is the GS density. The main result of the fractional-number extension of the DFT at zero temperature is that the GS energy is continuous and piecewise linear function of the electron number (charge) [8]. Namely, for the system with fractional charge $N = J \pm \delta$, where J is an integer and $\delta \in [0, 1]$, the GS energy is

$$E_{\pm\delta}[J,v] \equiv E[J\pm\delta,v] = \delta E[J\pm1,v] + (1-\delta)E[J,v],$$
(2)

and the GS density is a linear mixture of the GS densities of the *J*- and the $(J \pm 1)$ -electron number systems,

$$\rho_{\pm\delta}(\mathbf{r}; J, v) \equiv \rho(\mathbf{r}; J \pm \delta, v) = \delta \rho(\mathbf{r}; J \pm 1, v) + (1 - \delta)\rho(\mathbf{r}; J, v).$$
(3)

The total energy is an implicit functional of the density via the dependence of the molecular orbitals and their occupancies on the density. The term representing the coupling between the electrons and the external potential is an exact, explicit functional of the electron density. The kinetic energy term, expressed as an explicit functional of spinorbitals and occupation numbers, is

$$\widetilde{T}_{s}[\{n_{p}\},\{\chi_{p}\}] = \sum_{p} n_{p} \int \chi_{p}^{*}(\mathbf{r}) \left(-\frac{1}{2}\nabla_{\mathbf{r}}^{2}\right) \chi_{p}(\mathbf{r}) d\mathbf{r}.$$
 (4)

 $E_{\rm es}$ represents the classical electrostatic energy

$$E_{\rm es}[\rho] = \frac{1}{2} \int \int \frac{\rho(\mathbf{r})\rho(\mathbf{r}')}{|\mathbf{r} - \mathbf{r}'|} d\mathbf{r} d\mathbf{r}'.$$
 (5)

Since such $E_{es}[\rho]$ is not linear in $\rho(\mathbf{r})$, it immediately follows that its nonlinearity has to be compensated by the appropriate exchange–correlation term $E_{\rm xc}[\rho]$, or the standard energy expression corrected by some additional terms [29, 30]. Alternatively, both $E_{\rm es}$ and $E_{\rm x}$ should be expressed in the ensemble terms [31]. In general, if the electrostatic energy for the $(J \pm \delta)$ system is defined as a linear combination of the electrostatic energies for the 'end' points, the form of the electrostatic energy as a functional of density is unknown [32]. Although no explicit form is available for the exact $E_{xc}[\rho]$, much is known about the "proper" way in which approximations for this energy term should be constructed. One of the main challenges for DFT is to keep, as its cornerstone, some element of simplicity [4]. Unfortunately, a simple density functional adopted from the solid-state physics, the LDA, does not perform well in many areas of chemistry. The introduction of the density gradient into the form of the exchange-correlation functional (the GGA or meta-GGA), allows to receive satisfactory results in the chemical applications. The next major advances came with the inclusion of a fraction of the HF exchange in the $E_{\rm xc}$ functional (hybrid functionals). One of the recent developments in functionals is due to the range separation. The idea is to separate the electron–electron interaction into two parts, the long-range one and the short-range one, and then to treat these parts with different functionals (see Ref. [4] for review).

In practice, the direct extension (to fractional *N*) of DFT is applied, for which the ground-state energy is evaluated as $E^{\text{DFA}}[N, y]$

$$\equiv \min_{\{n_p\},\{\chi_p\}} \left\{ \breve{E}[\{n_p\},\{\chi_p\};v] \middle| \sum_p n_p = N, n_p \in [0,1], \langle \chi_p \mid \chi_q \rangle = \delta_{pq} \right\}$$
(6)

where $E[\{n_p\}, \{\chi_p\}]$ is the energy functional defined in terms of commonly used approximations

$$\widetilde{E}[\{n_p\}, \{\chi_p\}, \nu] = \widetilde{T}_s[\{n_p\}, \{\chi_p\}] + \int \nu(\mathbf{r})\widetilde{\rho}(\mathbf{r}; \{n_p\}, \{\chi_p\})d\mathbf{r}
+ E_{\rm es}[\widetilde{\rho}[\{n_p\}, \{\chi_p\}]] + b_{\rm xc}E_{\rm xc}[\widetilde{\rho}[\{n_p\}, \{\chi_p\}]]
- \frac{1}{2}\sum_{pq} n_p n_q \Big\{ c_{\rm x} \langle \chi_p \chi_q \mid \chi_q \chi_p \rangle + b_{\rm x}^{\rm LR, \rm HF} \langle \chi_p \chi_q \mid \chi_q \chi_p \rangle^{\mu} \Big\}$$
(7)

with a set of orthonormal spinorbitals, $\{\chi_p\}$ and their occupation numbers, $\{n_p\}$. Here, the non-integer electron number *N*(fractional charge) is realized indirectly by the constraint on the spinorbital occupancies, $\{n_p\}$. The density of the system (the non-interacting KS density) is

$$\widetilde{\rho}(\mathbf{r}; \{n_p\}, \{\chi_p\}, \nu) = \sum_p n_p |\chi_p(\mathbf{r})|^2,$$
(8)

and can integrate to any nonnegative real number $N = \sum_{p} n_{p}$. The GS density $\rho(\mathbf{r}; N, v)$ is obtained from Eq. (8) with $\{\chi_p\}$ and $\{n_p\}$ being the minimizers in Eq. (6). $E_{\rm xc}[\tilde{\rho}]$ represents the part of the exchange–correlation energy which depends locally on the density (and on the density derivatives). If $b_{\rm xc} = 0$, there is no correlation part and all exchange is represented by the HF-like exchange. The parameters c_x and $b_x^{\text{LR,HF}}$ determine the portion of the HF exchange in the hybrid and the longrange correction functional, respectively. The parameter μ defines separation of regions. The spinorbitals $\{\chi_n\}$ and orbital energies $\{\varepsilon_n\}$ are the eigensolutions of the one-electron Hamiltonian with the effective potential $v_{\rm eff}(\mathbf{x})$. The $v_{\rm eff}(\mathbf{x})$ is a local (multiplicative) potential in the original KS method or a non-local potential in a generalized KS method (with hybrid functionals) and in the HF method.

As a consequence of the linearity condition, Eq. (2), the chemical potential (the first derivative of the GS energy with respect to N) is constant between the integers

$$\mu[J \pm \delta, \nu] \equiv \left(\frac{\partial E[N, \nu]}{\partial N} \right)_{\nu} \Big|_{N=J \pm \delta}$$
$$= \pm (E[J \pm 1, \nu] - E[J, \nu]), \ \delta \in (0, 1),$$
(9)

and it is related to the frontier (f) eigenvalue [8, 12, 33–35]

$$\mu[J \pm \delta, \nu] = \varepsilon_f^{\text{DFA}}[J \pm \delta, \nu]. \tag{10}$$

Now, we can formulate the first two indicators for testing the quality of functionals: the relative deviation from the global linearity condition (GLC), Eq. (2), as

$$\Delta_{\pm\delta}^{\text{GLC}} = \frac{E_{\pm\delta}^{\text{DFA}}}{\delta E_{\pm1}^{\text{DFA}} + (1-\delta)E_0^{\text{DFA}}} - 1, \tag{11}$$

and the relative deviation from the global constancy condition (GCC), Eqs. (9) and (10), as

$$\Delta_{\pm\delta}^{\text{GCC}} \equiv \frac{\varepsilon_f^{\text{DFA}}[J \pm \delta]}{\pm \left(E_{\pm 1}^{\text{DFA}} - E_0^{\text{DFA}}\right)} - 1.$$
(12)

Here, $E_{\pm\delta}^{\text{DFA}} = E^{\text{DFA}}[J \pm \delta, v].$

Besides the linearity condition for the total energy and the constancy condition for the energy of the highest (partially) occupied spinorbital, we can examine the local counterparts of these conditions related to the ground-state density and the Fukui function (FF) [28, 36]. Third indicator tests the fulfillment of the local linearity condition (LLC), Eq. (3). It is the integrated deviation from the LLC, defined as

$$\Delta_{\pm\delta}^{\text{LLC}} = (J \pm \delta)^{-1} \int \left| \rho_{\pm\delta}^{\text{DFA}}(\mathbf{r}) - \left(\delta \rho_{\pm1}^{\text{DFA}}(\mathbf{r}) + (1 - \delta) \rho_0^{\text{DFA}}(\mathbf{r}) \right) \right| d\mathbf{r},$$
$$\rho_{\pm\delta}^{\text{DFA}}(\mathbf{r}) = \rho(\mathbf{r}, J \pm \delta, \nu). \tag{13}$$

(The factor $(J \pm \delta)^{-1}$ is omitted in our illustrative example tests as it is irrelevant at comparisons).

The FF (the derivative of the ground-state density, Eq. (3), with respect to the electron number) is independent of the fraction δ (similarly as the chemical potential)

$$f(\mathbf{r}; J \pm \delta, v) \equiv \left(\frac{\partial \rho(\mathbf{r}; N, v)}{\partial N} \right)_{v} \Big|_{N=J\pm\delta}$$

$$= \pm (\rho(\mathbf{r}; J \pm 1, v) - \rho(\mathbf{r}; J, v))$$
(14)

(this form of FF was noted first in Ref. [37]), and the integrated deviation from this local constancy condition (LCC) is defined as

$$\Delta_{\pm\delta}^{\text{LCC}} = C^{\text{LCC}} \int \left| f_{\pm\delta}^{\text{DFA}}(\mathbf{r}) \mp \left(\rho_{\pm1}^{\text{DFA}}(\mathbf{r}) - \rho_0^{\text{DFA}}(\mathbf{r}) \right) \right| d\mathbf{r},$$
$$C^{\text{LCC}} = \left(\int \left| \left(\rho_{\pm1}^{\text{DFA}}(\mathbf{r}) - \rho_0^{\text{DFA}}(\mathbf{r}) \right) \right| d\mathbf{r} \right)^{-1}$$
(15)

	δ							
	0.01	0.1	0.3	0.5	0.7	0.9	0.99	
HF	-0.00 [-0.04]	-0.02 [-0.36]	-0.05 [-0.84]	-0.05 [-1.00]	-0.04 [-0.85]	-0.02 [-0.38]	-0.00 [-0.05]	
SVWN5	0.05 [0.27]	0.48 [2.51]	1.12 [6.15]	1.31 [7.66]	1.07 [6.65]	0.42 [2.80]	0.04 [0.25]	
PBE	0.05 [0.26]	0.48 [2.43]	1.11 [5.92]	1.31 [7.26]	1.08 [6.11]	0.44 [2.29]	0.04 [0.04]	
B3LYP	0.04 [0.21]	0.40 [1.95]	0.93 [4.8]	1.10 [6.01]	0.91 [5.29]	0.38 [2.28]	0.04 [0.21]	
CAM-B3LYP	0.02 [0.14]	0.20 [1.30]	0.46 [3.2]	0.55 [4.01]	0.47 [3.52]	0.20 [1.50]	0.02 [0.13]	
rCAM-B3LYP	0.00 [0.07]	0.02 [0.61]	0.04 [1.50]	0.05 [1.85]	0.05 [1.58]	0.02 [0.61]	0.00 [0.03]	

Table 1 The average (over the set: Li, Na, Be, Mg) of the test indicator for the global linear condition $\Delta_{-\delta}^{GLC}$

In brackets, the average over the set enlarged by He. All values are per thousand. Data for atoms are accessible from Supporting Information

where $f_{\pm\delta}^{\text{DFA}}(\mathbf{r}) \equiv \left(\partial \rho_{\pm\delta}^{\text{DFA}}(\mathbf{r}) / \partial \rho_{\pm\delta}^{\text{DFA}}(\mathbf{r}) \partial N . \partial N\right)\Big|_{N=J\pm\delta^{*}}$ (The factor C^{LCC} is omitted in our illustrative example tests as it is irrelevant at comparisons). Note that $(C^{\text{LCC}})^{-1}$ equals to one when the FF is not negative, but it is larger than one (even two) when FF is negative in some regions as in cases of redox-induced electron transfer [38, 39]. This Δ^{LCC} is the last indicator for testing the quality of functionals, proposed in this work. It should be noted that if the integrand in Eqs. (13) and (15) is taken without absolute value operation, the result of integration is exactly zero. According to the proposed definitions of indicators, high quality of a particular DFA is demonstrated when all inequalities

$$\left|\Delta_{\pm\delta}^{GLC}\right| \ll 1, \left|\Delta_{\pm\delta}^{GCC}\right| \ll 1, \Delta_{\pm\delta}^{LGC} \ll 1, \Delta_{\pm\delta}^{LCC} \ll 1, \quad (16)$$

are satisfied in the whole range $0 < \delta < 1$.

3 Illustrative examples and discussion

In this article, the performance of SVWN5 [40] (a LDAtype functional), PBE [41] (a GGA-type functional), B3LYP [42, 43] (a hybrid functional), CAM-B3LYP [44] (a long-range corrected functional), rCAM-B3LYP [11] (a long-range corrected functional with improved description of systems with fractional numbers of electrons), and HF method was tested. To eliminate complication due to the spatial degeneracy, our test set is restricted only to atomic systems with the total angular momentum equal zero, namely to He, Li, Be, Na, Mg atoms. All calculations were performed with cc-pVTZ basis set [45-47] in spinunrestricted formalism. The FF for the fractional electron system [48], $f_{\pm\delta}^{\text{DFA}}(\mathbf{r})$, was calculated using QM4D program [49] according to methods described in Ref. [48]. Only decreasing electron number process was considered for all systems (lower sign chosen in Eqs. (2), (3), (9)–(15), $\delta = 1$ means a cationic form of an atom). All results are presented per thousand. The results for atoms are accessible from Supporting Information.



Fig. 1 The average of $\Delta_{-\delta}^{\text{GLC}}$, Eq. (11), vs. fraction δ

To present our illustrative example results in a condensed form, the average value of an indicator over a set of four atoms, Li, Na, Be, Mg is displayed for each deviation. The results for He are not included, as significantly differing from those of other atoms in many cases: for Δ^{GLC} at all methods (see Table 1), for remaining indicators at rCAM-B3LYP. This peculiarity stems from the fact that the two-particle aspects of electron structure in comparison with the single-particle ones are more pronounced for the He atom than for remaining atoms. In all Figures, lines joining calculated points are drawn for eye guidance only.

The results of the global linearity condition testing, Eq. (11), are collected in Table 1 (zero values at $\delta = 0$ and $\delta = 1$ are omitted) and in Fig. 1. As expected, the standard local-potential functionals (SVWN5 and PBE) yield the largest, positive $\Delta_{-\delta}^{GLC}$. The HF method yields the weakest, negative $\Delta_{-\delta}^{GLC}$. The B3LYP functional combines these two effects, yielding $\Delta_{-\delta}^{GLC}$ slightly smaller than those generated by SVWN and PBE functionals,

but still presents a concave curve. A significant improvement is observed for functionals which contain some Coulombattenuated exchange (CAM). The rCAM-B3LYP functional yields the smallest GLC deviation. Reasonable explanation of observed tendencies is available, especially of the fact that extreme deviations are observed at $\delta = 0.5$ for all methods. Very weak GLC deviation for the HF method is the effect of the testing set (only atomic systems without spatial degeneracy) used in this article. All the occurring open shell systems at integral *N* are not spatially degenerate and the HF method for such systems is self-interaction free [31].

At first, it should be noted that the energy expression, Eq. (7), has no explicit N dependence. This dependence is implicit trough the dependence on $\{n_p\}$ and the condition $\sum_p n_p = N$. Namely, the trace of the first-order density matrix (DM), d_1 , is equal to electron number N

$$d_1(\mathbf{x}; \mathbf{x}'; \{n_p\}, \{\chi_p\}) \equiv \sum_p n_p \chi_p^*(\mathbf{x}) \chi_p(\mathbf{x}'), \quad \text{Tr}d_1 = N.$$
(17)

In the direct extension of the HF method, the hyper-HF one, the approximate second-order DM, d_2 , is constructed from the first-order DM as the following determinant

$$d_{2}(\mathbf{x}_{1}, \mathbf{x}_{2}; \mathbf{x}'_{1}, \mathbf{x}'_{2}; d_{1})$$

$$\equiv \frac{1}{2} (d_{1}(\mathbf{x}_{1}, \mathbf{x}'_{1}) d_{1}(\mathbf{x}_{2}, \mathbf{x}'_{2}) - d_{1}(\mathbf{x}_{1}, \mathbf{x}'_{2}) d_{1}(\mathbf{x}_{2}, \mathbf{x}'_{1}))$$

$$= d_{2}[N, \nu], \qquad (18)$$

(it is exact for the single Slater determinant). This d_2 leads to relations [50] $\operatorname{Tr} d_2[N, v] \ge \frac{1}{2}(N-1)N$, and [31]

$$\Delta E_{\text{ee},\pm\delta}^{\text{HF}} \equiv E_{\text{ee},\pm\delta}^{\text{HF}} - \left(\delta E_{\text{ee},\pm1}^{\text{HF}} + (1-\delta)E_{\text{ee},0}^{\text{HF}}\right) \ge 0.$$
(19)

where $E_{ee,\pm\delta}^{\text{HF}} = \text{Tr}d_2[J \pm \delta, v]\hat{V}_{ee}$ (the electron–electron energy due to 'generalized Fock' operator).

In the case of atoms considered in this work, the above deviation from the linearity for the electron–electron interaction energy is [31]

$$\Delta E_{\text{ee},\pm\delta}^{\text{HF}} = \delta (1-\delta) J_{\text{ff}} \ge 0, \qquad (20)$$

(here $J_{\rm ff}$ is the Coulomb integral for the frontier orbital) with the maximum at $\delta = 0.5$. The shape of this derivation, Eq. (21), was already observed in Fig. 1 of Ref. [51]. When combined with the negative denominator in Eq. (11), it leads to negative $\Delta_{-\delta}^{\rm GLC}$ for the HF method. Contrary to the hyper-HF method, compensation of the self-interaction errors of $E_{\rm es}$ and $E_{\rm x}$ for DFA methods is not complete or absent. Therefore, the deviation from linearity for $E_{\rm es}$, Eq. (5), is the dominating deviation. It can be easily evaluated as

$$\Delta E_{\text{es},\pm\delta}^{\text{DFA}} \equiv E_{\text{es}} \left[\rho_{\pm\delta}^{\text{DFA}} \right] - \left(\delta E_{\text{es}} \left[\rho_{\pm1}^{\text{DFA}} \right] + (1-\delta) E_{\text{es}} \left[\rho_{0}^{\text{DFA}} \right] \right) \\ = -\delta (1-\delta) E_{\text{es}} \left[\rho_{\pm1}^{\text{DFA}} - \rho_{0}^{\text{DFA}} \right] \le 0.$$
⁽²²⁾



Fig. 2 The average of $\Delta_{-\delta}^{GCC}$, Eq. (12), vs. fraction δ

When combined with the negative denominator in Eq. (11), it leads to positive $\Delta_{-\delta}^{GLC}$ for DFA methods, with the maximum at $\delta = 0.5$.

It should be pointed out that Eqs. (20) and (22) describe the GLC discrepancy of the E_{ee}^{HF} and E_{es}^{DFA} only approximately due to the following inconsistencies. All three terms $E_{ee,\pm\delta}^{HF}$, $E_{ee,\pm1}^{HF}$ and $E_{ee,0}^{HF}$ are constructed from a common set of orbitals, determined at $N = J \pm \delta$, appropriate for the first term only, while to be consistent, the orbitals determined at $(J \pm 1)$ and J should be applied for remaining two terms. In Eq. (22), a combination of $\rho_{\pm1}^{DFA}$ and ρ_{0}^{DFA} is taken for $\rho_{\pm\delta}^{DFA}$ [like in Eq. (3)], while this density should be constructed from orbitals determined at $N = J \pm \delta$.

The results of the global constancy condition testing, Eq. (12), are presented in Fig. 2. Since the constancy of $\mu[N] = (\partial E[N]/\partial N) = -\partial E_{-\delta}/\partial \delta$ is tested, the discrepancy for the HF method is accounted mainly by $\partial \Delta E_{\text{ee},-\delta}^{\text{HF}}/\partial \delta = -(1-2\delta)J_{\text{ff}}$, see Eq. (20). When combined with the negative denominator in Eq. (12), this produces $\Delta_{-\delta}^{GCC}$ linearly decreasing, zeroing at $\delta = 0.5$. Similarly, for DFA methods, Eq. (20) should be replaced by Eq. (22), leading to $\Delta_{-\delta}^{GCC}$ linearly increasing, zeroing at $\delta = 0.5$, and quite large at end points of δ range, with the smallest absolute values for the rCAM-B3LYP and HF methods. The zeroing at $\delta = 0.5$ is consistent with the transition state approach in which the ionization energy is determined by the highest occupied orbital energy of the intermediate state, $E[J] - E[J-1] \approx \varepsilon_{\text{HOMO}}[J-0.5]$ [52], see also Ref. [53].

As indicated in Eq. (9), the region of μ constancy is limited to $0 < \delta < 1$. Nevertheless, Fig. 2 includes results at



Fig. 3 The average of $\Delta_{-\delta}^{\text{GCC}}$, Eq. (13), vs. fraction δ

 $\delta = 0$. With $\varepsilon_f^{\text{DFA}}[J - \delta]$ defined to be the frontier (highest occupied) orbital energy, the value of Δ_0^{GCC} equals $\lim_{\delta \to 0} \Delta_{-\delta}^{\text{GCC}}$. It is consistent with $\Delta_{-0.01}^{\text{GCC}}$ for all atoms. However, the GCC indicator calculated formally at $\delta = 1$, Δ_{-1}^{GCC} , does not equal $\lim_{\delta \to 1} \Delta_{-\delta}^{\text{GCC}}$ and it differs significantly from $\Delta_{-0.99}^{\text{GCC}}$ (see Supporting Information). This difference is due to different calculational procedures for determination of $\varepsilon_f^{\text{DFA}}[J - \delta]$: at $\delta < 1$, the KS equations are solved self-consistently in the space of J occupied orbitals, while at $\delta = 1$ —in the space of (J - 1) occupied orbitals. The lowest virtual orbital energy is to be taken as $\varepsilon_f^{\text{DFA}}[J - 1]$.

The results of the local linearity condition testing, Eq. (13), are shown in Fig. 3. The shapes of plots are similar to those in Fig. 1, but the maximum discrepancies are about 50 times larger. This similarity can be understood with the help of the DFT mapping $\rho \rightarrow E[\rho]$ connecting the ground-state density and energy. The maximal discrepancy concerning the density in Fig. 3 correlates with the maximal discrepancy concerning the energy in Fig. 1. Of all DFA methods, the smallest discrepancy is observed for rCAM-B3LYP ($\Delta_{-0.5}^{LLC} = 0.015$). The HF method is even better ($\Delta_{-0.5}^{LLC} = 0.006$).

The results of the local constancy condition testing are shown in Fig. 4. The shapes of plots are similar to those in Fig. 2, provided $|\Delta_{-\delta}^{GCC}|$ would be plotted there instead of $\Delta_{-\delta}^{GCC}$. The magnitude of discrepancies is practically the same in both figures. This similarity can be viewed as an analog of similarity between the $\Delta_{-\delta}^{GLC}$ and $\Delta_{-\delta}^{LLC}$ plots, but now applied to derivatives with respect to *N*. Again, the



Fig. 4 The average of $\Delta_{-\delta}^{\text{GCC}}$, Eq. (15), vs. fraction δ

performance of the rCAM-B3LYP functional is the best among tested approximate functionals.

4 Conclusions

In concluding, our illustrative examples' calculations demonstrate that the performance of the rCAM-B3LYP functional is the best among the tested functionals. Moreover, satisfaction of inequalities (16) is quite reasonable $(\Delta_{-0.5}^{GLC} = 5 \times 10^{-5}, \Delta_{-0.3}^{GCC} = -4.5 \times 10^{-4}, \Delta_{-0.5}^{LLC} = 1.5 \times 10^{-2}, \Delta_{-0.5}^{LCC} = 3.8 \times 10^{-3})$. The especially small Δ^{GLC} is due to the fact that the rCAM-B3LYP functional was constructed to minimize just the discrepancy from the GLC. Another general conclusion can be reached: when the chemical potentials and the Fukui function are to be determined, the best results are found at $\delta = 0.5$ for all tested DFAs. It is surprising that the performance of the extended HF method is as good (or even better) as that of DFT with the rCAM-B3IYP functional. Probably, this is due to the particular set of systems (atoms) chosen for tests. In any case, the hyper-HF method should not be used for large systems because important correlation effects are missing.

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REGULAR ARTICLE

Stereoselective inclusion mechanism of ketoprofen into β -cyclodextrin: insights from molecular dynamics simulations and free energy calculations

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Abstract The host-guest inclusion mechanism formed between β-cyclodextrin and those poorly water-soluble drug molecules has important applications in supramolecular chemistry, biology and pharmacy. In this work, the chiral recognition ability of β-cyclodextrin to one of nonsteroidal anti-inflammatory drugs, ketoprofen, has been systematically investigated using molecular dynamics and free energy simulation methods. The *R*- and *S*-enantiomers of ketoprofen were explicitly bound within the cyclodextrin cavity in our simulations, respectively. In consistent with experimental observations, tiny structural difference between two isomers could be observed. Calculated absolute binding free energies using adapted biasing force (ABF) method and MM/GBSA approach for both isomers are comparable to experimental values. Significant binding fluctuations along the MD trajectory have been observed. The free energy profiles calculated using two different approaches reveal that the ketoprofen prefers binding in the cavity with the carboxylate group facing the wider edge of

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C. Zhang (⊠) · Y. Xie Analytical and Testing Center, Sichuan University, Chengdu 610064, Sichuan, People's Republic of China e-mail: cczh@scu.edu.cn β -cyclodextrin. Similar free energy profiles for two enantiomers obtained using ABF calculations indicate that it is very hard to separate and identify the chiral conjugates within the framework of the natural β -cyclodextrin.

 $\begin{array}{ll} \textbf{Keywords} & Molecular \ dynamics \cdot ABF \cdot \beta \ Cyclodextrin \cdot \\ Ketoprofen \cdot Chiral \ recognition \end{array}$

1 Introduction

The interaction or recognition mechanism of host–guest systems constitutes the core part of supramolecular chemistry, for which the main driving force of the formation of the systems is considered to be the hydrophobic interactions. Correct understanding or quantification of the intermolecular interaction would be particularly important for the "rational planning of new supramolecular systems, including intelligent materials, as well as for developing new biologically active agents" [1]. Among host–guest systems, the inclusion complexes formed between natural cyclodextrins (CDs) with small organic molecules, *esp.* those hydrophobic molecules, represents a typical example in this field and thus attracts a lot of attentions [2–9].

CDs are cyclic oligomers of glucose units. The most common natural CDs are composed of 6 (α), 7 (β) or 8 (γ) pyranose units connected by α -1,4 glycosidic linkage. Due to ${}^{4}C_{1}$ chair conformation of the pyranose units, the overall topology of these CDs can be described as a toroid/hollow truncated cone shape. The significant different hydrophilicity of their exterior environment and interior cavity makes the CDs very unique in the supramolecular chemistry. They can easily bind lipophilic molecules to form a host–guest inclusion complex and thus change the chemical, physical or even biological properties of these ligands. Such ability makes them widely applied in pharmaceutical development, drug delivery or food industry. People have pursued their usage in the synthesis of some certain mechanically interlocked molecular architecture, such as rotaxanes [10] and catenanes [11]. The photochemistry of molecules hosted by CDs is also of particular interests, e.g., arenes [12]. In this work, we will focus on the detailed chiral recognition mechanism of β -CD complexed with a widely used antiinflammatory drug, *ketoprofen*. The schematic representations of β -CD and ketoprofen are given in Scheme 1.

Ketoprofen (IUPACE name of (R/S)2-(3-benzoylphenyl) propionic acid) has long been considered to be the nonsteroidal anti-inflammatory drug (NSAID) with analgesic and antipyretic effects [13]. It has also served as one of pain relief drugs. Ketoprofen is usually marketed as 50:50 racemic mixtures of (R-) and (S-) enantiomers, but with poor solubility. It was demonstrated that only S-enantiomer has anti-inflammatory activity, while same functions can be found for other profen drugs such as ibuprofen and flurbiprofen. However, the chiral inversion of the R- to S-enantiomers of profen drugs has been observed in vivo, e.g., in pigs [14]. Based on this, the physiological functions of (R-)ketoprofen should not be totally neglected and might deserve some more researches. More importantly, some experimental studies have proved that (R-)flurbiprofen can be involved in the antitumor activity in mouse. Clearly, we cannot simply neglect either enantiomer. Due to the importance of the enantiomer drugs in the drug delivery, it might be interesting to pursue the synthesis, separation or chemical properties for different enantiomers. The recognition mechanism and binding structures of β -CD with ketoprofen have been reported in some detail from both experimental and theoretical sides [15–19]. However, little was known about the difference of the binding features, such as binding free energies or structures, for both enantiomers of ketoprofen. Recently, to tackle the photochemistry characteristics of (R-) and (S-)ketoprofen and their chiral recognition in β -CD, it was investigated by Marconi et al. [17] using various experimental techniques such as circular dichroism, isothermal titration calorimetry and NMR. For the association constants for both enantiomers, the experiments gave almost identical values considering experimental uncertainty. Based on circular dichroism and 2D ROESY spectra, tiny structural difference can be probed for both enantiomers. Therefore, it should be pointed out that it is very difficult to use experimental tools to get stereoselective recognition ability of β -CD versus (*R*-) and (*S*-)ketoprofen. Given the practically enantio-differentiated anti-inflammatory and analgesic activities of ketoprofen, it is then necessary to understand the detailed recognition mechanism for both chiral isomers from theoretical view.



Scheme 1 Schematic representations and definition of atoms of β -cyclodextrin and (*S*-) and (*R*-)ketoprofen molecules

Besides the stereoselective recognition mechanism of β -CD with ketoprofen, there is another interesting issue not quite well understood. For the profen drugs, there are two sides which bear different polarizability as shown in Scheme 1, with polar carboxylate group at one side and nonpolar aromatic group at the other side. So far there is no definite study to say which rim of the β -CD the carboxylate group is bound with. Indeed, some controversies were reported for other profen drugs, such as ibuprofen. The X-ray structure of the inclusion complex formed between β -CD and S-ibuprofen reported by Braga et al. [20] clearly indicates that the carboxylate group of the ligand faces the smaller rim of the β -CD. On the contrary, another crystallographic structure of S-ibuprofen complexed by a heptakis-substituted β -CD shows that the carboxylate group is facing the bigger rim [21]. In addition, theoretical simulations reported by Cai et al. [22] on steroid drugs recognized by β -CD indicate that the different binding orientations could lead to closed binding free energy profiles along the binding coordinates as well as the association constants. Such observations could further suggest that the formation of inclusion complex has some complicated processes. Therefore, to obtain a complete understanding of the β -CD bound with ketoprofen molecule or other small ligands, it is highly desired to perform extensive simulations on all of possible binding orientations for different enantiomers.

In this work, to obtain a complete understanding of the binding characteristics of ketoprofen/ β -CD inclusion complex, we then build four possible binding structures for two ketoprofen enantiomers as shown in Scheme 2. The absolute binding free energies will then be calculated based on molecular mechanics-generalized born surface area (MM-GBSA) method. To understand the detailed formation processes of each inclusion complex, the potential of mean force (PMF) profiles along the inclusion coordinates will be computed using adapted biasing forcing (ABF) method. Scheme 2 Schematic representations of two different inclusion directions of ketoprofen threading through the cavity of β -CD. The origin is set at the center of mass of seven glycosidic oxygen atoms of β -CD. The narrower edge is usually called as primary rim, while the wider edge as secondary rim



Orientation II

2 Computational details

2.1 Models of β-CD/ketoprofen inclusion complex

There is no published X-ray structure for β -CD/ketoprofen inclusion complex. To investigate the dynamics properties of this inclusion complex, it would be necessary to construct the model of inclusion complex firstly. It should be pointed out that since there are two isomers for ketoprofen molecule, we then docked R- and S-ketoprofen into β -CD cavity using AutoDock 4.2 program, respectively [23]. In addition, since there are also two possible binding orientations for each enantiomer as shown in Schemes 1 and 2, the structures for each orientation of different enantiomers with the lowest binding energy were chosen for the following extensive molecular dynamics simulation. For convenience, the glucose units of cyclodextrin are labeled from G1 to G7. We simply give the name for each binding orientation as **R-I**, **R-II** for (*R*-) ketoprofen, while S-I and S-II for (S-)ketoprofen, respectively. In binding orientation I, the carboxyl group is facing the secondary rim, while the carboxyl group stays at the primary rim in binding orientation II. Throughout the simulation, the stoichemistry of β-CD and ketoprofen is maintained to be 1:1 ratio, which has been suggested by many experimental works.

The obtained systems were firstly solvated in a pre-equilibrated TIP3P [24] water box. The typical size of the water box is calculated to be about 44 Å \times 44 Å \times 40 Å, consisting of 179 solute atoms and about 1800 solvent molecules. To prepare suitable force field for the ketoprofen molecule, we employed the standard Amber general amber force field (GAFF) generation procedure. First of all, the geometry optimizations of ketoprofen at HF/6-31G* level of theory were carried out using gaussian09 suite of program [25]. The partial atomic charges were then calculated using the restrained electrostatic potential (RESP) protocol after structure optimization and electrostatic potential calculations using B3LYP/6-31G* method. The force field parameters for the ligands generated using the Antechamber program were then described by GAFF module. The force field of β -cyclodextrin is expressed using Glycam06 carbohydrate parameters [26] in this work, which has been proved to accurately and efficiently represent carbohydrates. The periodic boundary conditions and an 8 Å cutoff for non-bond interactions were applied. The particle mesh ewald (PME) algorithm [27] was used to calculate the long-range electrostatic interactions. The positions of water molecules were relaxed by 500 steps of steepest descent (SD) and 1000 steps of conjugate gradient (CG) minimization approach with all of solute molecules fixed at their original positions. Further 10,000 steps of CG full minimization were carried out for total system. The obtained systems were gradually heated to 300 K in 20 ps in the NVT ensemble, followed by 20 ns equilibration simulation under 1 atm pressure. Subsequently, further 20 ns MD simulations in the isothermal-isobaric NPT ensemble were performed for data analysis. Newton's equations of atomic motion were integrated by the Verlet algorithm with a 2-fs time step. SHAKE algorithm [28] was applied to constraint bond stretching of the covalent bonds involving hydrogen atoms. All of MD simulations were performed using PMEMD module of CUDA version implemented in AMBER 12 software package.

2.2 Binding free energy calculations

To quantitatively assess the binding affinity of β -CD in complex with ketoprofen molecule, it is necessary to calculate the binding free energy for the inclusion complex.
Several methods have been proposed for calculating absolute binding free energy, e.g., linear response approximation (LRA) [29], linear interaction energy (LIE) [30–32], molecular mechanics Poisson Boltzmann (or generalized Born) surface area (MM-PB/GBSA) [33, 34], solvated interaction energy (SIE) [35], or free energy pathway method [36]. In this work, the binding free energy is calculated using MM-GBSA [33] method.

For the calculation of binding free energy in MM-GBSA framework, it has been discussed extensively [37–41]. Only a short description is summarized here:

 $\Delta G_{\text{binding}} = \Delta G_{\text{complex}} - \Delta G_{\beta - \text{CD}} - \Delta G_{\text{ketoprofen}}$ (1)

$$G = E_{\rm gas} + G_{\rm sol} - \mathrm{TS} \tag{2}$$

$$E_{\rm gas} = E_{\rm int} + E_{\rm vdW} + E_{\rm ele} \tag{3}$$

$$G_{\rm sol} = G_{\rm GB} + G_{\rm np} \tag{4}$$

 $\Delta G_{\text{complex}}, \Delta G_{\beta - \text{CD}}$ and $\Delta G_{\text{ketoprofen}}$ are free energies of the complex, the β -CD and ketoprofen, respectively. Each term can be obtained according to Eq. (2). Practically, they are calculated as the statistical averages over frames extracted from MD trajectories. The solvation free energy (G_{sol}) can be divided into polar (G_{GB}) and nonpolar (G_{np}) contributions. The polar solvation contribution is calculated by solving the Generalized Born (GB) equation [42]. Dielectric constants (ε) for solute and solvent were selected to be 1 and 80, respectively. The nonpolar contribution due to cavity formation and van der Waals interactions between the solute and the solvent can be estimated by the equation of $\gamma \cdot SA + b$, where $\gamma = 0.0072$ kcal/Å², b = 0.0 kcal/mol. The SA is defined as the solvent accessible surface area, which was estimated using the program MSMS [43]. For each complex system, binding energies were averaged over 1,000 frames of the 20-ns MD trajectory. It has long been recognized that the inclusion of entropic effect in the calculation of total binding free energy can largely reduce the difference between theoretical and experimental values [44]. Entropy contributions are from changes in the degrees of freedom including translation, rotation and vibration. The translational, rotational and vibrational entropy terms are functions of the mass and moments of inertia of the molecule and thus can be calculated using the standard equations of statistical mechanics [45]. In this work, vibrational entropy contributions were estimated using the normal mode analysis approach [46]. Due to the system size is just moderate, the $-T\Delta S$ was averaged over 200 snapshots of the MD trajectory.

2.3 Formation of inclusion complex

As suggested in Scheme 2, two possible binding models with different position of the propionic acid end have been

proposed for the molecules in profen family recognized by β -CD. Although the absolute binding free energy calculated using MM-GBSA method can provide some insights into the interaction affinity between β -CD and ligands, the detailed process to form the inclusion complex cannot be entirely revealed. Furthermore, the chiral recognition processes might be another interesting issue for the β -CD/ketoprofen inclusion complex. To address the formation of inclusion complexes, we further carried out extensive simulations of the inclusion processes using the adaptive biasing force (ABF) method [47]. MD simulations were performed using NAMD program [48], for which the initial system is constructed using the following strategy to avoid artificial impacts. As shown in Scheme 2, we randomly put the ketoprofen isomers outside the secondary (bigger) rim of the cavity of β -CD. We then simulate the translocation process through the cavity starting from the secondary rim, and finally moving out via the primary rim. The origin is defined as the center of mass of seven oxygen atoms (O4) of glycosidic linkage of the β -CD. In the present ABF calculations, we have two possible binding processes for each isomer: One is aromatic ring A enters the cavity first, and the second is the carboxylate group moves into the cavity first. Then, the order parameter or the inclusion coordinate should be defined correspondingly. For the first case, the orientation I, the order parameter (ξ) , which ranges from -16 to 14 Å, was chosen as the projection of the distance between the origin and the center of mass of the aromatic ring A of ketoprofen on the z-axis. For the second case, the orientation II, the order parameter (ξ) , which ranges from -16 to 14 Å, was chosen as the projection of the distance between the origin and the center of the mass of the carboxylate group of ketoprofen on the z-axis. The inclusion pathway was then divided into six consecutive windows to increase the computational efficiency. Each window was further divided into consecutive bins with bin width of 0.1 Å. For each window, up to 5 ns of MD trajectory was generated, resulted in a total of 30 ns simulation time. Instantaneous values of the forces, the first-order derivative of free energy with respect to the defined order parameter, were accrued in each bin to calculate the mean force value. Total of 1,500 force samples were accumulated in each bin prior to application of the adaptive bias. For the setup protocol of MD simulations, which is essentially the same as we did in previous section using Amber program, except the integration step is set as 1 fs time step in ABF calculations.

3 Results and discussion

3.1 Structures of the inclusion complexes

In this work, one of our major objectives is to understand the mechanism of chiral recognition of β -CD with Fig. 1 Distance between C4 atom and the center of mass of glycosidic oxygen atoms of β -cyclodextrin along the simulation time for four binding models



ketoprofen enantiomers. The phenomenon called chiral discrimination in the host–guest interaction has been observed in experimental way [49]. Here, we performed extensive MD simulations to examine the dynamic effects of β -CD interacting with the enantiomers of the ketoprofen.

Throughout the molecular dynamics simulations, all four inclusion complexes formed between ketoprofen and β -CD were maintained very well judged by the distance between the C4 atom of ketoprofen and the center of mass of glycosidic oxygen atoms as shown in Fig. 1. It looks like that for the *R*-ketoprofen, the binding orientation I (R-I) is much more stable than the other binding orientation (**R-II**) according to different binding fluctuation, while the difference between S-I and S-II models is smaller. Therefore, it could be deduced that a little smaller difference of binding affinity in S-conformation than that in R-conformation is expected. To get a direct view of the structural change along the simulation time, we further plotted corresponding snapshots from 20 ns to 40 ns for each binding conformer in Figs. 2 and 3. In most of simulation time, the ligand molecule is inserted into the cavity in tilted direction, which would ensure that the sugar ring has more interactions with aromatic rings of guest molecule. Such phenomenon

has been found in the simulation of β-CD/steroid drug systems investigated by Cai et al. [22]. At least one aromatic ring (A or B) of ketoprofen was found to be included in the cavity during the simulation, but some apparent fluctuations of the binding position could be observed. Such pattern could partially indicate the hydrophobic interaction plays important role in the guest molecule recognition by β -cyclodextrin. For clarity, the time course profiles of the distance between the center of mass of aromatic ring A (d_{cm}^A) or B (d_{cm}^B) and the center of mass of the glycosidic oxygen atoms of the host molecule are depicted in Fig. 4. It should be noted that this distance could only be used to elucidate the fluctuations for the binding status of the aromatic rings, but not for the estimation of binding stability, since this distance is different from the definition of order parameter (ξ) used in free energy pathway calculations as we gave above. Clearly, different binding performance for chiral isomers can be observed. For the orientation I, along the simulation trajectory, we can see that the carboxylate group always points outward and cannot be included into the cavity of β -CD. Such binding pattern requires that at least one of aromatic rings of ketoprofen is recognized by the hydrophobic interior environment of β -CD. According



Fig. 2 *Snapshots* of the inclusion complexes formed by the β -CD and (*R*-) and (*S*-)ketoprofen along the dynamics simulation time in orientation I, respectively. For clarity, the water molecules have been removed. The ketoprofen is plotted using stick style, while line style for β -CD



Fig. 3 Snapshots of the inclusion complexes formed by the β -CD and (*R*-) and (*S*-)ketoprofen along the dynamics simulation time in orientation II, respectively. For clarity, the water molecules have been removed. The ketoprofen is plotted using stick style, while line style for β -CD

to the snapshots given in Fig. 2 and the relative position analysis between the benzene rings and β -CD in Fig. 4, the substantial fluctuation of the binding of the benzene rings is quite clear. That means both aromatic rings have chance to be included into the cavity. It thus suggests that β -CD cannot identify any specific aromatic ring of included ligand. If we recall the NMR spectroscopy of the inclusion complex of β -CD/ketoprofen [17], experimental work indicated that both rings (*A* and *B*) can be deeply embedded in the cavity with very close chemical shifts. Our results are then in reasonably agreement with this observation. For the binding orientation **II**, a little different binding pattern could be observed. Especially, we can see that ring *A* of *R*-II and ring *B* of *S*-II seem to be dominantly recognized by β -CD cavity. Such binding feature might be inconsistent with the experimental observations. This feature should be further examined by binding free energy calculations.

It is also interesting to examine the possibility of hydrogen bond formed between host and guest molecules, since there are a lot of hydroxyl groups in β -cyclodextrin and one carboxylate group in ketoprofen. We summarized the hydrogen bond occupancy analysis results in Table 1 over the MD trajectories for all four inclusion complexes. Quite low hydrogen bond occupancies were found between host



Fig. 4 Distances between the aromatic rings of ketoprofen and the center of mass of seven glycosidic oxygen atoms of β -CD in binding orientation **I** (*panel I*) and in binding orientation **II** (*panel II*). For the binding of *R*-ketoprofen, d_{cm}^A is colored in *black*, and d_{cm}^B is colored in *green*. For the binding of *S*-ketoprofen, d_{cm}^A is colored in *red*, and d_{cm}^B is colored in *blue*

Table 1 Hydrogen bonds network analysis for interactions between β -CD and (*R*-) and (*S*-)ketoprofen (KET)

H-bond	Occupancy (%)					
	R-I	S-I	<i>R</i> -II	S-II		
KET:O1G7:O3	18.0	_	_	_		
KET:03…G1:02	13.8	16.1	_	_		
KET:03…G1:03	13.6	19.2	-	-		
KET:01G1:02	13.4	18.6	-	-		
KET:03G7:03	12.6	-	-	-		
KET:01G1:03	10.8	16.9	-	-		
KET:01…G5:06	-	-	21.6	-		
KET:01…G4:06	-	-	12.4	-		
KET:03…G5:06	-	-	11.1	_		
KET:01…G6:06	-	-	-	14.2		
KET:O3…G6:O6	-	-	-	13.3		
KET:O1…G7:O6	-	_	_	11.8		

Occupancy is in unit of percentage of the investigated time period (20 ns) during which specific hydrogen bonds are formed. The hydrogen bond is defined as the distance of acceptor and donor atoms shorter than 3.0 Å, and the internal angle of acceptor…H-donor is larger than 120°

and guest molecules. The biggest occupancy rate is just about 20 %. This supports from another angle that the hydrophobic interaction is possible the dominant factor in the stabilization of the inclusion complex. The electrostatic interactions should be considered to be a minor factor in the host–guest recognition. Indeed, the polar carboxylate group of ketoprofen is exposed to solvent molecules in almost most of simulation time no matter which rim it faces. However, the existence of the carboxyl group might be an issue to affect the formation of inclusion complex and the drug releasing, which we will discuss below using free energy calculations.

3.2 Binding free energy

Although binding models obtained from MD simulations can provide some insights into the guest molecule recognition, it is still difficult to identify which model is the favored one. Some quantitative measurements, such as absolute binding free energy calculation, can help us to get useful ideas. Due to chiral discrimination, different binding free energy for R- and S-enantiomers should be expected. Such results have been observed for the binding of ketoprofen, although researchers attributed the slight energetic difference into experimental uncertainty and thus cannot tell which the favored binding conformer is [17]. In this work, calculated absolute binding free energies are summarized in Table 2 for two binding orientations of both isomers, and the measured association free energies from different experimental sources [17, 18, 50-52] are included for comparison. The uncertainties for all of terms are included in the parentheses, which were calculated as the root mean square error for all of frames extracted in the MM/GBSA running. Overall, the calculated entropies for four binding models are negative, at the same time the enthalpies are also negative. This can suggest that the formation of inclusion complexes is an enthalpy driven process. For both R- and S-ketoprofen in binding orientation I, the calculated average binding free energy is around 3.8 kcal/mol, which is in excellent agreement with experimental values of ~4.0 kcal/ mol listed in Table 2. On the other hand, for the orientation II, a little larger difference between experimental and theoretical values could be observed. This could suggest that for the inclusion complexes formed by β -CD and ketoprofen, the binding orientation I, the propionic acid group of ketoprofen facing the wider rim of β -CD, is more likely the favored conformer, no matter R- or S-isomer.

On the other hand, since marked ketoprofen is often in a molar ratio of 50:50 racemic nature, little efforts have been applied to examine the binding difference between two chiral models. Recently, to understand the chiral photoreaction properties, Marconi et al. [17] examined the stereoselective binding affinity of β -CD bound with ketoprofen enantiomers. It is not surprised that only very tiny difference for the binding free energies can be observed, -3.95 kcal/mol (**S**) versus 3.76 kcal/mol (**R**). The experimental work by Marconi et al. [17] suggests that β -CD can bind *S*-ketoprofen slightly stronger than its *R*-chiral conjugate, although they put this to be within experimental uncertainty. Our

Table 2 Calculated absolute binding free energies for Image: Calculated absolute	Energy (kcal/mol)	<i>R</i> -ketoprofen		<i>S</i> -ketoprofen	
β -CD/(<i>R</i> -) and (<i>S</i> -)ketoprofen		Ι	II	I	II
MM/GBSA approach (comparison with experimental values)	$egin{aligned} & \langle \Delta E_{ m ele} angle \ & \langle \Delta E_{ m vdw} angle \ & \langle \Delta G_{ m sol,nonpolar} angle \end{aligned}$	-22.70 (3.21) -26.82 (16.71) -2.98 (0.31)	-25.34 (2.69) -19.56 (10.31) -3.06 (0.27)	-21.56 (3.47) -29.23 (16.70) -2.93 (0.32)	-25.09 (2.59) -20.18 (9.77) -3.08 (0.26)
We, in this work, calculated separately binding free energies of two binding orientations (I and II) for <i>R</i> - and <i>S</i> -ketoprofen isomers embedded into the β -CD cavity. Only Ref. [17] reports the binding free energies for two <i>R</i> - and <i>S</i> -ketoprofen enantiomers, other experimental work reports the binding free energies for the binding of ketoprofen with <i>R</i> and <i>S</i> isomers	$\begin{array}{l} \langle \Delta E_{\rm gas} \rangle \\ \langle \Delta G_{\rm sol,GB} \rangle \\ \langle \Delta E_{\rm gas} + {\rm G}_{\rm sol} \rangle \\ \langle -T\Delta S \rangle \\ \langle \Delta G_{\rm binding} \rangle \\ \Delta G_{\rm binding}^{\rm exp} \end{array}$	-49.53 (16.71) 28.75 (12.57) -20.78 (4.78) 17.11 (2.20) -3.67 -3.76 [17] -3.96 [50] -3.90 [51] -3.97 [18] -4.13 [52]	-44.90 (10.00) 26.28 (8.59) -18.62 (2.35) 16.26 (1.68) -2.36	-50.79 (15.81) 29.72 (12.28) -21.08 (4.37) 17.21 (1.94) -3.87 -3.95 [17]	-45.26 (10.36) 25.56 (7.81) -19.70 (3.36) 16.57 (1.89) -3.13

results are quite close this value, esp., the binding orientation I with the calculated absolute binding free energies are -3.87 kcal/mol for S-I, while -3.67 kcal/mol for R-I. For the binding orientation I, the difference of binding free energy between two enantiomers is calculated to be 0.2 kcal/mol, which agrees well with experimental difference of 0.19 kcal/mol [17] as shown in Table 2. Our results are also in consistent with a recent computational work, which suggests that the β -cyclodextrin binds (S-)ketoprofen a little tightly than (*R*-)ketoprofen [53].

Another interesting issue is that for both chiral isomers, the stronger binding affinity can be observed for orientation I than orientation II, esp. the R-isomer, which is consistent with our MD simulation result as described in above section. However, the negative binding free energies for the orientation II still suggest that such an inclusion conformation should not be totally neglected. Indeed, our simulation on the orientation II has confirmed that this conformer for both enantiomers is pretty stable as shown in Fig. 1. In addition, X-ray structure has proved the existence of the orientation **II** for the ibuprofen/β-CD inclusion complex [20]. It should be noted that the deviation of the calculated binding free energies of orientation II is a bit larger than those of orientation I. Based on the calculated absolute binding free energies, we then suggest that the preferred binding mode of ketoprofen by the natural β-cyclodextrin might take the orientation I for both isomers no matter it is in R- or S-isomer. Of course, such conclusion awaits more accurate experimental work to confirm. To this end, there is no contradiction between our results and a recent NMR study on the inclusion of R- and S-ketoprofen, in which Marconi et al. [17] concluded that the carboxyl group of ketoprofen exposed to the solvent from secondary edge of the host.

substrate binding, i.e., the electrostatic $(\Delta E_{ele} + \Delta G_{sol, GB})$ and nonpolar $(\Delta E_{\rm vdW} + \Delta G_{\rm sol, np})$ terms. It has long been accepted that the recognition of lipophilic molecules by β -CD is mainly due to hydrophobic interactions. The structural feature of aromatic included in the cavity revealed by MD simulation can support this proposal. Further, from Table 2, it can be seen that all four conformers agree this very well. For example, the electrostatic terms are calculated to be about 6.1, 1.0, 8.2 and 0.5 kcal/mol for R-I, R-II, S-I and S-II inclusion complexes, respectively. The electrostatic interaction is canceled out by the polar part of solvation. Positive values for electrostatic contribution indicate that polar interactions between host and guest molecules clearly disfavor the molecular recognition in the β -CD based inclusion complexes. In contrast, van der Waals interactions have dominantly favorable contributions to the binding affinity, as does the nonpolar part of solvation. The calculated total nonpolar terms are -29.8, -22.1, -31.2and -23.2 kcal/mol for R-I, R-II, S-I and S-II inclusion complexes, respectively. Quite interestingly, much larger nonpolar terms for the orientation I than the orientation II can be obtained in our simulations. This is also in consistent with our deduction of stronger binding affinity for orientation I than orientation II, since the hydrophobic interaction is the dominant factor in the formation of host-guest system we studied in this work.

Usually, there are two major factors that could affect the

3.3 ABF calculations

The binding free energies obtained using MM/GBSA method can explain the recognition driving force to some extent. However, it cannot tell us how the drug molecule is included into the cavity of host molecule. At the same time, it cannot identify the drug releasing direction and possible rate. To solve the problem, it is then necessary to examine the detailed energetic performance for the binding processes along the inclusion coordinates as depicted in Scheme 2. Calculated free energy profiles along the inclusion pathway for two binding orientations of *R*- and *S*-ketoprofen using ABF method are given in Figs. 5 and 6, respectively.

The free energy profile derived from ABF calculation could be integrated along the order parameters (ξ) to obtain a quantitative estimation of the association constant, K_a [54]. When ketoprofen penetrates through the β -CD cavity, the sampled volume was restricted to a cylinder defined by the area ($A_{\rm CD}$) available for in-plane movement of the guest molecule in the cavity ($A_{\rm CD} = \pi r^2$) and the linear trajectory perpendicular to this area [55].

$$K_a = N_A A_{\rm CD} \int \exp\left(-\frac{\Delta A(\xi)}{RT}\right) \mathrm{d}\xi,\tag{5}$$

where K_a is the system association constant with an unit of M⁻¹, $N_A = 6.02 \times 10^{23}$ is the Avogadro constant and the temperature T is chosen at 300 K. For the choice of radius of the cylinder, it could be calculated to be the average radius along the ligand penetrating the cavity along the z-axis direction. In this work, we simply take the value of r = 2Å as suggested by Auletta et al. [55]. The integration ranged over the whole sampled pathway. The association free energy can be obtained again according to the relationship of $\Delta G = -RT \ln K_a$. Calculated association constants and binding free energies of ΔG are listed in Table 3. The calculated association free energies are in excellent agreement with those values obtained using MM/GBSA method. Similarly, the association of S-isomer is systematically stronger than R-enantiomer, but with pretty small difference. For the binding orientation I, the results can be comparable to experimental values very well, with typical error less than 1.0 kcal/mol. These results could further indicate ABF method is one of suitable methods to address the energetic profiles for the formation of inclusion complexes. On the other hand, the results for the binding orientation II is not as good as those in orientation I. Considering the computational accuracy, our simulation results within both MM/GBSA and ABF frameworks seem to support that the inclusion complex formed in binding orientation I would be more likely the binding model observed in experimental way.

Since our simulations suggest that the model in orientation **I** would be the favored one, we will then focus on the free energy profiles depicted in Fig. 5. The overall topology of the free energy profiles for **R-I** and **S-I** is quite similar, which means that only a little difference can be observed in the chiral recognition. This is reasonable since there is only tiny difference for the chiral inclusion process revealed by



Fig. 5 Free energy profiles via ABF calculations for the inclusion of ketoprofen into β -CD along the *z*-axis in binding orientation **I**, *S*-ketoprofen (*solid line*), *R*-ketoprofen (*dash line*)



Fig. 6 Free energy profiles via ABF calculations for the inclusion of ketoprofen into β -CD cavity in binding orientation **II**, *S*-ketoprofen (*solid line*), *R*-ketoprofen (*dash line*)

 Table 3
 Calculated association constants and association free energies for all four binding models, and the experimental values from Ref. [17] are also included for comparison

Models	$K_a (\mathrm{M}^{-1})$	ΔG (kcal/mol)
S-I	632.5	-3.85
R-I	221.1	-3.22
S-II	176.5	-3.08
R-II	13.3	-1.53
Exp. [17]		-3.95 (S)
		-3.76(R)

S denotes the included compound is in *S*-enantiomer, while *R* represents the ligand is in its *R*-chiral conjugate

NMR experiments [17]. On the other hand, there are two distinguished energy minima along the order parameter, ca. $\xi = -3.6$ and 1.9 Å for S-I, while $\xi = -3.9$ and 1.1 Å for **R-I**, respectively. We then plotted corresponding snapshots in Fig. 7 for the inclusion of *S*-ketoprofen, and Fig. 8 for the inclusion of *R*-ketoprofen, which are all extracted from ABF trajectories. For the second minimum of S-I



Fig. 7 Snapshots of the inclusion complexes of S-ketoprofen with β -CD at the inflection points along the free energy curve in orientation I. **a** Near the first minimum, ca. $\xi = -3.6$ Å, **b** near the maximum, ca. $\xi = -2.6$ Å, **c** near the second minimum, ca. $\xi = 1.9$ Å



Fig. 8 Snapshots of the inclusion complexes of *R*-ketoprofen with β -CD at the inflection points along the free energy curve in orientation I. **a** Near the first minimum, ca. $\xi = -3.9$ Å, **b** near the maximum, ca. $\xi = -2.3$ Å, **c** near the second minimum, ca. $\xi = 1.1$ Å

 $(\xi = 1.9 \text{ Å})$ and **R-I** $(\xi = 1.1 \text{ Å})$ models, two aromatic rings are fully inserted into the β -CD cavity to maximum the hydrophobic interactions, while the carboxyl group is facing the wider rim just as we have observed in MD simulations. Surprisingly, instead of the ring A included in the cavity, we can see that for the first minima at $\xi = -3.6$ Å (S-I) and -3.9 Å (R-I), the ring B is included by β -CD cavity. One barrier at $\xi = -2.3$ Å (S-I) and -2.6 Å (R-I) with a moderate barrier height can be observed to connect these two free energy minima, and the structures for two isomers are presented in Figs. 7 and 8, respectively. It looks like that this barrier stays at the stage of the ring A beginning entering the cavity, but not fully bounded. Such geometry transformation characteristics could explain why we can see significant fluctuations in the ligand binding dynamics presented above. However, if we carefully examine the structures, we may partially attribute this barrier to the carboxyl group of ketoprofen, which could form hydrogen bond network with hydroxyl groups around the secondary rim. To understand this issue, we then calculated the hydrogen bond occupancy between the carboxylate group and sugar hydroxyl groups within the ABF simulation window of [-6.0 and -1.0 Å]. The hydrogen bond occupancy for this oxygen atom is calculated to be about 35 %. Clearly, albeit the ring A is not included by the cavity, the existence of hydrogen bond between the carboxyl group and β -CD could still anchor the ketoprofen a little while. That will make the ring *B* has the chance to be bound in the cavity. Therefore, although the polar interaction has been proved to be a minor factor to stabilize the host–guest system, it still should be considered as one of the key factors to affect the formation of the inclusion complex.

Although we pull the ketoprofen from the wide edge to narrow rim in the ABF calculations, we can still take a look at the free energy in a recursive way. Quite interestingly, no barrier can be located for the ligand migration from the narrow rim into the cavity, for which the carboxyl group enters the cavity firstly. However, the barrier occurs along the z-axis positive direction, which means to enter the cavity from the wider edge, and it has to overcome a barrier height of 3.0 and 2.7 kcal/mol for S- and R-ketoprofen according to our calculations, respectively. In other words, we might safely conclude that to form the ketoprofen/β-CD inclusion complex, the polar carboxyl group will enter the cavity first, but finally two aromatic rings will be bound in the cavity as shown in Fig. 7. It should be emphasized here that the hydrophobic interactions constitute the major factor in the final formation of inclusion complex. Accounting for the ketoprofen releasing direction, it depends on the binding position of the carboxyl group, i.e., where the carboxyl group occurs, and the releasing the drug could move out more easily through this direction. About 10 kcal/ mol energy has to be overcome for bounded ligand to migrate from the cavity through the narrow edge. We might dedicate this carboxyl group as one of manipulate factors for the drug releasing issue. This is quite smart of nature. According to our simulations, it is thus possible to design some new drugs with a desired releasing rate.

4 Conclusion

Correct understanding the inclusion mechanism of β-CD complexed with small drug molecules would be of particular importance in the drug delivery. We, in this work, systematically investigated the free energy profiles of the inclusion complexes formed between natural β-CD and ketoprofen enantiomers. Our results support that the R- and S-ketoprofen prefer binding β -CD with their carboxylate groups facing the wider edge of the cavity (binding orientation I). Additionally, both MM/GBSA and ABF approaches suggest that β -CD has stronger affinity with S- than R-ketoprofen, which is also consistent with NMR measurement. Calculated binding free energies for the model of S-I and R-I using MM/GBSA and ABF are very close to the experimental values, which further confirm that both methods are suitable to deal with the recognition problem related with host-guest supramolecular system. It has also been proved that natural β -CD might not be a good chiral separation matter due to similar binding performance of ketoprofen enantiomers into the cavity. Therefore, various substituted CDs might be good candidates to fulfill this object. Nonetheless, our simulations reveal that the ketoprofen might move out more easily from the cavity via the position of carboxyl group occurring. This carboxyl group can be denoted as one of manipulate factors in the drug releasing. It is our hope that this could be helpful for the further drug design with desired drug releasing rate.

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REGULAR ARTICLE

State-to-state quantum dynamics of the N(⁴S) + $C_2(\tilde{x}^1\Sigma^+) \rightarrow CN(\tilde{x}^2\Sigma^+) + C(^3P)$ reaction

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Abstract Quantum dynamics of the N(⁴S) + C₂(\tilde{X} ${}^{1}\Sigma^{+}) \rightarrow CN(\tilde{X}^{2}\Sigma^{+}) + C({}^{3}P)$ reaction was investigated on a global potential energy surface for the $\tilde{a}^4 A''$ state of C₂N using a Chebyshev real wave packet method. This reaction is barrierless and has two deep wells corresponding to the CCN and CNC species, respectively. The numerous oscillations in reaction probabilities and highly excited rovibrational states of the product CN reflect the large exothermicity of the reaction and the feature of the potential. As expected for a complex-forming reaction, the rotational state distributions of the CN product were highly excited. The calculated vibrational state distributions of the CN product are in reasonably good agreement with the experimental results. The differential cross sections were found to be dominated by scattering in both forward and backward directions with a forward bias, suggesting that the reaction is not completely statistic and the lifetime of the reaction intermediate is not particularly long.

Keywords Quantum dynamics · Wave packet propagation · Reaction probability · Cross section

Dedicated to Professor Guosen Yan and published as part of the special collection of articles celebrating his 85th birthday.

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1 Introduction

The CN radical is ubiquitous in the interstellar medium and has been observed in the dense clouds and around circumstellar envelopes [1-5], which is produced by the reactions such as [6, 7]

$$N + CH \to CN + H, \tag{1}$$

$$N + C_2 \rightarrow CN + C. \tag{2}$$

Both reactions also play important roles in the formation of CN (and N₂ indirectly) in interstellar clouds [8, 9]. For reaction (1), its rate constants were measured recently over the range of 56 K $\leq T \leq$ 296 K, which were excellently reproduced by our quantum mechanical calculations [10]. Based on the new rate constants, the CN abundance generated from the Nautilus gas-grain model is significantly less than the previous estimation, suggesting that this reaction is only a relatively minor source of the CN radical in dense clouds [10]. In addition, we further investigated quantum dynamics of reaction (1), which gave an intimate insight into the underlying reaction mechanism at a state-to-state level [11].

For reaction (2), both N(⁴S) + C₂($\tilde{X}^{1}\Sigma^{+}$) and N(⁴S) + C₂($\tilde{a}^{3}\Pi_{u}$) reactions lead to the products of C(³P) and CN($\tilde{X}^{2}\Sigma^{+}$). The rate constant of the N(⁴S) + C₂($\tilde{a}^{3}\Pi_{u}$) reaction at room temperature was found to be 2.8 × 10⁻¹¹ cm³ mol⁻¹ s⁻¹ from an experimental study by Becker et al. [12]. Very recently, Loison et al. [13] measured rate constants for the N(⁴S) + C₂($\tilde{X}^{1}\Sigma^{+}$) reaction, which turned out to be much larger than the estimated value for the N(⁴S) + C₂($\tilde{a}^{3}\Pi_{u}$) reaction. Astrochemical simulations showed that the title reaction should be dominant in the formation of CN in dense interstellar medium [13]. In addition, the vibrational distributions of the CN product produced by the exoergic indirect reaction (2) have

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been measured by Daugey et al. [14, 15] with laser-induced fluorescence, which were found to be non-statistical. High-resolution photoelectron spectra of C_2N^- obtained by using slow electron velocity-map imaging provided the first measurement of the low-lying $\tilde{a}^4\Sigma^-$ state of C_2N [16].

Theoretically, geometric and vibrational characterizations of the low-lying electronic states of the C₂N molecule have been calculated by several groups at various ab initio levels [17–23]. Near-equilibrium potential energy surface (PES) for the $\tilde{a}^4\Sigma^-$ state of the C₂N radical has been constructed using correlated CCSD(T) method to study the ro-vibrational spectra by Hill et al. [22]. Very recently, we constructed a global PES for the \tilde{a}^4A'' state of C₂N and calculated the rate constants of the title reaction, which showed good agreement with the recent experimental results [13]. To the best of our knowledge, there has been no theoretical study on the reaction dynamics of the title reaction, due to the lack of a global PES before.

In this work, we for the first time present an exact quantum mechanical study of the state-to-state integral and differential cross sections for the title reaction at the collision energies ranging from 0 to 0.16 eV, based on our new PES. This paper is organized as follows. In Sect. 2, we describe the ab initio calculations used to construct the PES and the details of the quantum mechanical calculations. The main characteristics of the PES, reaction probabilities, and cross sections are presented and discussed in Sect. 3. Finally, Sect. 4 gives a short summary.

2 Computational details

2.1 Potential energy surface

The title reaction can proceed adiabatically on the $\tilde{a}^4 A''$ state, which is $\tilde{a}^4 \Sigma^-$ in the linear geometries. The ab initio calculations were performed using internally contracted multi-reference configuration interaction method with Davidson correction (MRCI + Q) [24-26] employing the augmented correlation consistent polarized valence triple zeta (aug-cc-pVTZ) basis set, as implemented in the MOLPRO 2010 package [27]. All CI wave functions were based on the nature orbitals obtained from the stateaveraged completed active space self-consistent field (SA-CASSCF) calculations for equally weighted $\tilde{a}^4 A''$, $b^4 A''$, and $\tilde{c}^4 A'$ states. A full-valence active space involving 13 electrons in 12 orbitals was used in the CASSCF procedure, and three 1 s orbitals for both nitrogen and carbon atoms were kept doubly occupied but fully optimized. In the subsequent MRCI calculations, three 1s orbitals and one 2s orbital of the nitrogen atom were frozen. A total number of contracted configurations are typically 9795174.

					. 1.	
Table 1	Low-lying	vibrational	energy	levels	(cm^{-1})) of C ₂ N

(n_1, n_2, n_3)	E	(n_1, n_2, n_3)	Ε
(0, 0, 0)	0.0	(2, 1, 0)	2,851.4
(0, 1, 0)	522.5	(0, 6, 0)	2,897.4
(0, 2, 0)	1,014.7	(1, 0, 1)	2,984.5
(1, 0, 0)	1,165.3	(1, 4, 0)	3,150.3
	$(1,170 \pm 8)^{a}$	(0, 3, 1)	3,263.9
(0, 3, 0)	1,503.9	(2, 2, 0)	3,344.3
(1, 1, 0)	1,692.4	(0, 7, 0)	3,350.1
(0, 0, 1)	1,812.9	(3, 0, 0)	3,463.1
(0, 4, 0)	1,974.9	(1, 1, 1)	3,501.7
(1, 2, 0)	2,185.7	(0, 0, 2)	3,571.0
(2, 0, 0)	2,319.4	(1, 5, 0)	3,620.2
(0, 1, 1)	2,327.1	(0, 4, 1)	3,716.2
(0, 5, 0)	2,443.2	(0, 8, 0)	3,791.6
(1, 3, 0)	2,677.9	(2, 3, 0)	3,839.1
(0, 2, 1)	2,794.1	(1, 2, 1)	3,969.1

^a Experimental value taken from Garand et al. [16]

The internal coordinates ($R_{\rm NC}$, $R_{\rm CC}$, $\theta_{\rm N-C-C}$) were used to give a better description of the global PES. The PES was fitted with spline interpolation based on about 13,000 ab initio points ranging from $1.7 \le R_{\rm NC}/a_0 \le 15.0$, $1.75 \le R_{\rm CC}/a_0 \le 14.0$, and $0 \le \theta_{\rm N-C-C}/\deg \le 180$. The permutation symmetry of the two carbon atoms was kept by selecting the shorter NC bond as the NC coordinate in interpolation.

2.2 Vibrational energy levels

The vibrational energy levels of the C₂N intermediate were calculated by solving the time-independent Schrödinger vibrational equation in the Jacobi coordinates (R, r, γ) using the Lanczos algorithm [28]. All energy levels were generated by carrying out about 2000 Lanczos recursion steps. For the two Jacobi radial coordinates R and r, 120 potential-optimized discrete variable representation (PODVR) points [29] were used. And 110 Gauss–Legendre grid points [30] were used for the angular variable γ . The calculated energy levels, given in Table 1, are converged to be 0.01 cm⁻¹ or better.

2.3 Quantum dynamics

The reactive scattering dynamics were investigated using the Chebyshev real wave packet (CRWP) method. Since the details of the method can be found in earlier papers [31–34], only a brief description is given here. The Hamiltonian in the reactant Jacobi coordinates (R, r, γ) can be written as ($\hbar = 1$),

$$\hat{H} = -\frac{1}{2\mu_R} \frac{\partial^2}{\partial R^2} - \frac{1}{2\mu_r} \frac{\partial^2}{\partial r^2} + \frac{(\hat{J} - \hat{j})^2}{2\mu_R R^2} + \frac{\hat{j}^2}{2\mu_r r^2} + V(R, r, \gamma), \quad (3)$$

Table 2 Numerical parameters(in atomic units) used in thewave packet calculations

Grid/basis ranges and sizes Projection Initial wave packet Damping function	$\begin{split} R &\in [0.01, \ 19.0], \ N_R^1 = 511, \ N_R^2 = 269 \\ r &\in [1.7, \ 15.7], \ N_r^1 = 323, \ N_r^2 = 59 \\ \gamma &\in [0, 180^\circ], j = 0 \sim j_{\text{max}} = 300, N_j = 301 \\ R'_\infty &= 12.0 \\ R_0 &= 11.0, \ \delta = 0.1, \ E_0 = 0.05 \text{ eV} \\ D &= \begin{cases} \exp[-0.01(\frac{R-14.0}{5.0})^2], & 14.0 \le R \le 19.0 \\ \exp[-0.09(\frac{r-14.9}{1.0})^2 - 0.007], \ 14.9 < r \le 15.7 \\ \exp[-0.007(\frac{r-13.9}{1.0})^2], & 13.9 \le r \le 14.9 \\ 1, & \text{otherwise} \end{cases} \end{split}$		
Spectral range	Potential energy cutoff: 7.5 eV		
	Rotational kinetic energy cutoff: 5.5 eV		
Propagation step	60,000		

where μ_r and μ_R are the reduced masses of the diatomic C_2 and C_2N , respectively. *J* and *j* denote the total and C_2 rotational angular momentum. The reactant Jacobi coordinates (*R*, *r*, γ) represent the distance between the N atom and the C_2 center, the C_2 bond length, and the angle between *R* and *r*, respectively.

The Hamiltonian and wave packet were discretized in a mixed basis/grid representation [35]. To make the calculations efficient, an *L*-shaped grid was used for the radial coordinate [34]. For each total angular momentum J and parity p, the wave function in the body-fixed (BF) frame can be expanded as,

$$\left|\Psi^{Jp}\right\rangle = \sum_{\alpha_r \alpha_R j K} C^{Jp}_{\alpha_r \alpha_R j K} |\alpha_r\rangle |\alpha_R\rangle |jK, Jp\rangle, \tag{4}$$

where α_r and α_R denote the grid indices for the two radial coordinates, and *K* is the projection of *J* onto the BF *z*-axis. The parity-adapted angular basis can be written as

$$|jK; Jp\rangle = (2 + 2\delta_{K,0})^{-1/2} \times \left(|JK\rangle|jK\rangle + p(-1)^J|J-K\rangle|j-K\rangle\right),$$
(5)

where $|jK\rangle \equiv Y_j^K(\gamma, 0)$ are normalized associated Legendre functions and $|JK\rangle \equiv \sqrt{(2J+1)/8\pi^2}D_{KM}^J$ are the parityadapted normalized rotation matrix elements with the Wigner rotation functions D_{KM}^J [36].

The initial wave packet $|\Psi_i\rangle$ in the space-fixed frame is defined as [35],

$$|\Psi_i\rangle = N e^{-(R-R_0)^2/2\delta^2} \cos k_0 R \big|\varphi_{\nu_i j_i}\big\rangle |j_i l_i, Jp\rangle, \tag{6}$$

where *N* is a normalization factor, R_0 and δ represent, respectively, the initial position and width of the Gaussian function, $k_0 = \sqrt{2\mu_R E_0}$ is the mean momentum and $|\varphi_{v_i j_i}\rangle$ ($v_i = 0, j_i = 0$) is the eigenfunction of the initial rovibrational state of C₂. The initial wave function was transformed into the BF frame before propagation.

The propagation of the wave packet $|\Psi_k\rangle$ was performed using the modified Chebyshev recursion relation [37–39],

$$|\Psi_k\rangle = 2D\hat{H}_s|\Psi_{k-1}\rangle - D^2|\Psi_{k-2}\rangle, \ k \ge 2, \tag{7}$$

where *D* is a damping function. The propagation begins with $|\Psi_0\rangle = |\Psi_i\rangle$ and $|\Psi_1\rangle = D\hat{H}_s|\Psi_0\rangle$. The scaled Hamiltonian \hat{H}_s is defined as

$$\hat{H}_{\rm s} = (\hat{H} - H^+)/H^-,$$
(8)

in which H^+ and H^- are the spectral medium and half width of the Hamiltonian, respectively, which can be calculated from the upper and lower spectral limits of the Hamiltonian $(H_{\text{max}} \text{ and } H_{\text{min}})$ with the relation $H^{\pm} = (H_{\text{max}} \pm H_{\text{min}})/2$.

The coordinate transformation method [34] was used to carry out final state projection in the product channel, which yields the *S*-matrix elements $(S_{v_f j_f K_f}^J \leftarrow v_{ij_i K_i}(E))$. The state-to-state integral cross section (ICS) can be then evaluated as

$$\sigma_{\nu_f j_f \leftarrow \nu_i j_i}(E) = \frac{\pi}{(2j_i + 1)k_{\nu_i j_i}^2} \times \sum_{K_i} \sum_{K_f} \sum_J (2J + 1) \left| S_{\nu_f j_f K_f \leftarrow \nu_i j_i K_i}^J(E) \right|^2,$$
⁽⁹⁾

and the state-to-state differential cross section (DCS) is calculated as

$$\frac{\mathrm{d}\sigma_{v_{f}j_{f}\leftarrow v_{i}j_{i}}(\vartheta, E)}{\mathrm{d}\Omega} = \frac{\pi}{(2j_{i}+1)} \sum_{K_{i}} \sum_{K_{f}} \left| \frac{1}{2ik_{v_{i}j_{i}}} \sum_{J} (2J+1)d^{J}_{K_{f}K_{i}}(\vartheta)S^{J}_{v_{f}j_{f}K_{f}\leftarrow v_{i}j_{i}K_{i}}(E) \right|^{2},$$
(10)

in which $k_{v_i j_i}$ is the initial translational wave vector, ϑ is the scattering angle between the scattered CN + C product and the incoming N + C₂ reactant, and $d_{K_f K_i}^J(\vartheta)$ are the reduced Wigner rotational matrix elements [36].

The optimal numerical parameters listed in Table 2 were determined based on extensive convergence tests

Fig. 1 Energy diagram of the $N({}^{4}S) + C_2(X^{1}\Sigma^{+})$ reaction. The energies (in eV) are relative to the reactant. The geometries are presented with bond lengths (in bohr)



with respect to J = 0 reaction probability. A total of 60,000 Chebyshev propagation steps and large basis dimension were necessary for converging reaction probabilities due to deep wells in the reaction path. Partial waves for $J \le 94$ were included to converge the ICS to 0.16 eV. For $J \le 70$, a maximum K value of 30 was found to converge the probabilities well, whereas $K_{\text{max}} = 45$ was used for $70 < J \le 94$.

3 Results and discussion

3.1 Potential energy surface

The energies and geometries of the stationary points on the PES along the reaction pathways are shown in Fig. 1. The calculated exothermicity of the reaction is 1.322 eV including the zero point energies, which reasonably agrees with the experimental value of (1.48 ± 0.07) eV, estimated from the difference between the dissociation energies of $C_2(\tilde{X}^1\Sigma^+)$ and $CN(\tilde{X}^2\Sigma^+)$, which are (6.29 \pm 0.02) [40] and (7.77 \pm 0.05) eV [41], respectively. For the geometry of the global minimum, our result of $(2.253a_0, 2.520a_0, 180.0^\circ)$ is in good agreement with the CCSD(T)/VQZ-F12 result of $(2.25a_0, 2.51a_0, 2.55a_0, 2$ 180.0°) by Hill et al. [22]. The reaction path shows no barrier and has two deep wells, corresponding to the linear C₂N and CNC species, which are connected by two transition states and one intermediate. It is worth noting that there are conical intersections between the $\tilde{a}^4 A''$ and $b^4 A''$ states near the global minimum, which are not considered in the present work.

The vibrational energy levels, listed in Table 1, are labeled by three vibrational quantum numbers n_1 , n_2 ,



Fig. 2 Calculated reaction probabilities for J = 0, 30, 60, and 90 as a function of collision energy

and n_3 , representing the symmetric stretching, bending, and anti-symmetric stretching vibrations, respectively. The calculated fundamental frequencies of (1,165.3, 522.5, 1,812.9) cm⁻¹ are in good agreement with the anharmonic frequencies (1,178.5, 443.2, 1,821.8) cm⁻¹ of Hill et al. [22]. They are also consistent well with the harmonic fundamental frequencies reported by several groups, which are (1,111, 433, 1,915) cm⁻¹ at the CAS-SCF/cc-pVDZ level, (1,138, 434, 1,826) cm⁻¹ at the CCSD(T)/TZ2P level [19], and (1,101, 473, 1,910) cm⁻¹ at the CASSCF/6s4p2d2f level of theory [20]. Moreover, the fundamental frequency of 1,165.3 cm⁻¹ for the symmetric stretching vibrational mode is in excellent agreement with the experimental value of (1,170 ± 8) cm⁻¹ by Garand et al. [16].





Fig. 3 The effective potential energies including centrifugal potential as a function of Jacobi coordinate *R* with optimized *r* and γ in the N + C₂ entrance channel at *J* = 0, 30, 60, and 90



Fig. 4 Calculated total ICS for title reaction as a function of collision energy

3.2 Reaction probabilities

The reaction probabilities for J = 0, 30, 60, and 90 are shown in Fig. 2 as a function of the collision energy E_c . It is worth noting that the probabilities are basically around a value of 0.8, which shows the high reactivity of this reaction. There is no energy threshold for J = 0 reaction probability, consistent with the fact that the PES has no intrinsic barrier in the reactant channel. The threshold appears when J > 0 and gradually gets larger, since the effective potential turns to be a higher barrier as J increases due to the centrifugal potential $J(J + 1)/2\mu_R R^2$, as shown in Fig. 3. Moreover, the reaction probabilities consist of numerous sharp oscillations, which correspond to resonances supported by the two deep potential wells.



Fig. 5 Calculated product vibrational state distributions at $E_c = 0.04$, 0.08, and 0.15 eV. The experimental values over the room temperature are taken from Ref. [14]

3.3 Cross sections

The total ICS calculated by summing over partial waves $(J \le 94)$ is shown in Fig. 4. In consistent to the reaction probabilities, the total ICS shows no threshold. In general, the ICS decreases monotonically with the increase of the collision energy, due to the barrierless reaction path and complex-forming mechanism.

Figure 5 presents the vibrational state distributions of the CN product at collision energies of 0.04, 0.08, and 0.15 eV, obtained by summing all rotational state populations in each vibrational channel. It can be seen that the CN vibration is excited up to 6 quanta. At a collision energy of 0.04 or 0.08 eV, a slight population inversion appears, which implies the non-statistical nature of the reaction. At higher energies, the vibrational state populations decrease with the vibrational quantum number v_{f} consistent with the complex-forming mechanism. For comparison, the vibrational distribution measured at a distance of 4 mm from the reactant mixing zone by Daugey et al. [14] is also plotted in Fig. 5. It can be seen that their overall trends are similar. Since the reaction was investigated experimentally above room temperature, the reactant C₂, which is a mixture of C₂($\tilde{X}^1\Sigma^+$) and C₂ ($\tilde{a}^3\Pi_{\mu}$), contains both ground and excited ro-vibrational states, whereas we considered only the reaction between $N(^{4}S)$ and $C_2(\tilde{X}^1\Sigma^+)$ in the specific initial state $(v_i = 0, j_i = 0)$. This should be the major reason for the experiment-theory discrepancy.

So far, there has been no measured rotational state distribution of the CN product or theoretical prediction for the title reaction. The CN rotational state resolved ICSs at collision energy of 0.04, 0.08, and 0.15 eV are plotted in Fig. 6. The CN rotational state is highly excited up to



Fig. 6 Calculated product rotational state distributions at $E_c = 0.04$, 0.08, and 0.15 eV

more than 76 quanta in the ground vibrational state, since the reaction is largely exothermic. The number of rotational excited states decreases with the increase of the vibrational quantum number. The oscillatory structures in all populations are obviously caused by resonances.

The DCSs at collision energies of 0.04, 0.08, and 0.15 eV are presented in Fig. 7. It can be seen that the DCSs are dominated by scattering in both forward and backward directions, also consistent with the complex-forming mechanism. However, the DCSs are not entirely symmetric. The forward bias is similar to the N + CH and



Fig. 7 Calculated total DCSs for the title reaction as a function of scattering angle at $E_{\rm c}=0.04,\,0.08,\,{\rm and}\,0.15\,{\rm eV}$

O + NH reactions [11, 42], suggesting that the reaction is non-statistic and the lifetime of reaction intermediate is not particularly long.

4 Conclusions

In this work, we reported the state-to-state quantum dynamics of the $N(^4S) + C_2(\tilde{X}^1\Sigma^+)$ reaction, based on a new global PES of the \tilde{a}^4A'' state for C_2N . The PES was fitted from more than 13,000 ab initio points calculated at the MRCI + Q level using the aug-cc-pVTZ basis set. The reaction path is barrierless and has two deep wells corresponding to the linear CCN and CNC species. The resonances supported by these deep wells manifest in reaction probabilities as numerous oscillations. The vibrational and rotational states of the CN product are highly excited. The agreement with the available experimental vibrational state distributions of the product CN is reasonably good. The DCSs were found to be dominated by scattering in both forward and backward directions with a forward bias, providing evidence for the non-statistic behavior of this complex-forming reaction.

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REGULAR ARTICLE

Mechanism and regioselectivity of the cycloaddition between nitrone and dirhodium vinylcarbene catalyzed by $Rh_2(O_2CH)_4$: a computational study

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Abstract The B3LYP density functional studies on the cycloaddition between nitrone and dirhodium vinylcarbene with subsequent cascade carbenoid aromatic cycloaddition/N-O cleavage and rearrangement revealed the energetics and the geometries of important intermediates and transition states in the catalytic cycle. The reaction initially occurs through a stepwise [3+2]-cycloaddition in which the dirhodium carbene activates the adjacent vinyl group for [3+2]-cycloaddition by the nitrone. Driven by the reactivity of the carbone carbon, intramolecular cyclopropanation takes place to release the catalyst, followed by the electrocyclic opening of the cyclopropane ring to form a seven-membered ring intermediate. N-O cleavage and [1,7]-oxygen migration to final product proceed in a single step with a high activation energy. With meta-substituent (Cl) in the N-aryl group, the charge density of chlorine's ortho-site decreases due to the negative inductive effect of chlorine. Therefore, as compared to the electrophilic addition on the C^8 atom, the electrophilic addition on the C^9 site owns a significantly incensement on the energy barrier of cyclopropane (TSIII and TSIX). In addition, according to the calculations, the basicity of the product 4 is stronger than that of the product 5. So an acidic compound

Dedicated to Professor Guosen Yan and published as part of the special collection of articles celebrating his 85th birthday.

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1,1,1,3,3,3-hexafluoro-2-propanol added in the experimental study (Wang et al. in Angew Chem Int Ed 51:5907– 5910, 2012) more easily reacts with **4** than **5** and thus promotes the formation of product **4**.

KeywordsDirhodium vinylcarbene \cdot DFT calculation \cdot Transition state \cdot Reaction mechanism \cdot Regioselectivity

1 Introduction

Diazo compounds have been extensively studied during the last few decades, and their value in organic synthesis is well known [1-5]. Transition-metal carbenoids derived from reactions of diazo compounds with a variety of metal complex are versatile intermediates in organic synthesis [6]. For several years, the rhodium-catalyzed reactions of donor/acceptor-substituted carbenoids have been explored, and the results showed that those carbenoids display much greater chemoselectivity when compared to the more conventional acceptor-substituted carbenoids [7-10]. A clear example of this effect is seen in a Hammett study of intermolecular cyclopropanation of para-substituted styrenes (Scheme 1) [11]. Electronic character of styrenes [ρ value of $-0.9 \ (\sigma^+ \text{ scale})$] can strongly affect the relative rates of reaction of methyl phenyldiazoacetate with various styrenes, while ethyl diazoacetate shows virtually no selectivity [11]. The greater chemoselectivity of the donor/ acceptor carbenoids has opened up a number of reactions that were previously not viable with conventional carbenoids. In particular, vinylcarbene intermediates derived from vinyldiazoacetates exhibit a high level of selectivity in cyclopropanation [12-14] and carbon-hydrogen insertion reaction [15–17] and can participate in a diverse array of more elaborate transformation [15, 18, 19], such as the

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Scheme 1 Competition reaction with styrene. Using metal carbenoids. Reproduced with permission from Ref. [11] Copyright 2000, ScienceDirect



Scheme 2 Dirhodium vinylcarbene-induced nitrone cycloaddition with subsequent cascade carbenoid aromatic cycloaddition/N–O cleavage and rearrangement (CACR)

combined C–H insertion/cope rearrangement [18], and the ylide formation/[2,3]-sigmatropic rearrangement/oxy-cope rearrangement/ene reaction cascade sequence [19]. Furthermore, their propensity for formal [3+2] [20–22] and [4+3] [23–26] cycloaddition reactions has been demonstrated.

Recently, Doyle et al. reported an efficient and highly regioselective cycloaddition reaction between the vinylcarbene from methyl 2-diazo-3-butenoate **1** and diverse nitrones catalyzed by rhodium octanoate [Rh₂(Oct)₄] (Scheme 2) [27]. The dirhodium vinylcarbene induced nitrone cycloaddition with subsequent cascade carbenoid aromatic cycloaddition/N–O cleavage, and rearrangement (CACR) gave products which have both oxygen and nitrogen-fused rings and a quaternary carbon in the middle. The compound structure obtained by X-ray diffraction of a single crystal reveals that extensive rearrangement has occurred, and the carboxylate group from the vinyldiazoacetate is bound to the quaternary carbon that connects the tricycle. Even though the reactions were carried out at room temperature, the products of the process were formed with remarkable specificity, and this type of reaction can be used as a general and selective method for preparation of multi-functionalized tricyclic heterocycles through an abnormal cascade process. A further surprising feature of the reactions is that, with *meta-* and *ortho*-substituents on the *N*-aryl group, cyclopropanation could have occurred on either side of the *N*-aryl bond, which would have led to the formation of two regioisomers; however, only a single regioisomer was formed in high yields. Therefore, this multi-step cascade process occurred with very high regiocontrol. This class of reaction has already become a standard method for laboratory and is being considered as a promising protocol for commercial process [28–30].

In light of the experimental results and for further development of the CACR chemistry, it was felt that the deeper understanding of the reaction mechanism, in particular the origins of the observed selectivity on the molecular scale, was indispensable. We report herein the pathways of the rhodium-catalyzed reactions of methyl 2-diazo-3-butenoate and nitrone by using the density functional calculations, which provide the first qualitative account of the experimental data, as well as the structures of important intermediates and transition states involved in the catalytic cycle. The calculations explain how the CACR happens and disclose the reason why the reaction can have such high regioselectivity. It is worthwhile to mention that this is the first report on reaction mechanism for the CACR reaction in detail. The theoretical analysis of the properties of intermediates and transition states revealed unique roles of the rhodium carbene. The studies and mechanistic insights obtained herein can ultimately provide value in accelerating the design and optimization of catalytic processes of transition-metal-catalyzed carbene procedures.

2 Computational methods

2.1 Models

In the present work, we have studied a catalytic model reaction between vinylcarbenoid complex 2 and nitrone 3 to form two different products 4 and 5 using dirhodium tetrakis (formate), $Rh_2(O_2CH)_4$, as a model of a common catalyst [31–34] (Scheme 3). Product 4 is in accord with the actual synthetic compound. The reaction to generating product 5 has not been reported in the literature and was investigated herein for comparison in regioselectivity. These model reactions serve to investigate the fundamental properties of the dirhodium carbene complex, as well as the mechanism and regioselectivity of CACR reactions. The vinylcarbenoid model 2 has been described in previous

Scheme 3 Model reactions for the computational studies



computational studies on cyclopropanation chemistry [31, 34]. This study is focused on an analysis of how the rhodium carbenoid reacts with the nitrone. For detailed discussions on the formation of the reactive vinylcarbenoid complex, see Refs. [31, 34].

2.2 Computational details

All of the calculations were performed with the Gaussian 09 software package [35]. Density functional theory (DFT) was employed using the three-parameter hybrid functional B3LYP [36, 37] to locate stationary points on the potential energy surface (PES). Structures were optimized with a basis set consisting of the 1997 stuttgart relativistic smallcore effective core-potential and basis set [Stuttgart RSC 1997 ECP] [38-40] for Rh, augmented with a 4f-function ($\zeta_{4}(Rh) = 1.350$) [31] and the split valence basis set 6-31G(d,p) for all other atoms (C, H, N, Cl and O). The method and the basis sets used here are known to give reliable results for rhodium (carbene) reactions [5]. This composite basis set is abbreviated 6-31G(d,p)[Rh-RSC+4f]. The main discussion is based on single-point energies calculated at the B3LYP/6-311+G(2d,2p)[Rh-RSC+4f]// B3LYP/6-31G(d,p)[Rh-RSC+4f] level of theory, with zero-point energy corrections calculated at the B3LYP/6-31G(d,p)[Rh-RSC+4f] level and Gibbs free energies calculated at the B3LYP/6-31G(d,p)[Rh-RSC+4f] level of theory. Heavy-atom basis set definitions and corresponding pseudopotential parameters were obtained from the EMSL basis set exchange library [41, 42]. Each stationary point was adequately characterized by normal coordinate analysis (only one imaginary for a transition state and no imaginary frequencies for an equilibrium structure). The intrinsic reaction coordinate (IRC) analysis [43] was carried out throughout the reaction pathways to confirm that all stationary points are smoothly connected to each other.

To evaluate the effect of the solvent polarity on the energetics of CACR reactions, a single-point energy calculation was performed with self-consistent reaction field (SCRF) method [based on the SMD model [44], $\varepsilon = 10.125$ for dichloroethane (DCE)]. Natural population analysis and natural bond orbital (NBO) analyses were carried out at the same level as the one used for geometry optimization [45, 46]. All charge distribution analyses discussed throughout this article are made on the basis of the nature population analysis.

3 Results and discussion

3.1 Reaction to produce the real product 4

First, we investigated the reactions of 3 with vinylcarbenoid complex 2 to generate product 4 as a basic model reaction. Scheme 4 shows the reaction course that the catalytic cycle involves five steps. The free energy profile of this CACR reaction in the gas phase is given in Fig. 1. First, 3 and 2 interact through a stepwise [3+2] cycloaddition to form the complex III. The formation of C-O bond proceeds via TSI with an activation free energy of 2.8 kcal/mol, which is followed by the formation of C-C bond with the activation free energy barrier of 4.5 kcal/mol. Subsequent intramolecular cyclopropanation by the rhodium carbene on the nitrogen-bound aryl group is proposed to form intermediate V via two transition states TSIII and TSIV with the free energy barriers of 10.2 and 7.6 kcal/mol, respectively, and the catalyst is regenerated. The next step toward the final product first involves an electrocyclic opening of the cyclopropane ring (aromatic cycloaddition) to form intermediate VI via TSV with a free energy barrier 4.5 kcal/mol. Unexpected and unique N-O bond cleavage and [1,7]-oxygen migration then take place through TSVI with the largest free energy barrier of 22.2 kcal/mol followed by a downhill path to the final product with large exothermicity (64.7 kcal/mol). In TSVI, the cleavage of N-O bond and migration of oxygen to conjugated olefinic carbon atom take place simultaneously in a single step. No intermediates were located near TSVI along the intrinsic reaction coordinate.

To probe the solvent effect on the reaction energetics, SCRF calculations (SMD model) were performed based on the gas-phase geometry. Dichloroethane (DCE, $\varepsilon = 10.125$) was investigated as the solvent because it is used for the CACR reaction in practice. The relative free energies of the stationary points on the potential energy surface in DCE are also given in Fig. 1 (in blue). The free energy profile did not change very much from the one in the gas-phase calculations. The N–O cleavage and [1,7]-oxygen migration processes have the largest free energy barrier (19.6 kcal/mol in DCE). Therefore, this step can be regarded as the ratedetermining step (RDS). Moreover, the free energy barrier for **TSVI** has a small decrease, and the energy barrier for cyclopropanation has a slight increase. As compared to the free energy profile in gas phase, the [1,7]-oxygen migration





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Fig. 1 Free energy profile of the $Rh_2(O_2CH)_4$ -catalyzed CACR reaction of 2 and 3 to produce 4 at the B3LYP/6-311+G(2d,2p)[Rh-RSC+4f]/B3LYP/6-31G(d,p)[Rh-RSC+4f] level. The result of the free energies in gas phase is given in*red*, and the result of the free energies in DCE is shown in*blue*

step has the highest energy barrier, which suggests that how to overcome the energetic bottleneck for the [1,7]-oxygen migration becomes very crucial for accelerating the reaction rate.

The 3D structures of representative stationary points are shown in Fig. 2. The Rh¹–Rh² bond of I has no significant change toward the formation of the complex III (2.479 \rightarrow 2.474 Å). The Rh¹–C¹ bond of I (2.036 Å) has a significant increase in first to form the intermediate II (2.103 Å) and then shortens through the C–C bond formation stage TSII (2.063 Å) to form the complex III (1.981 Å, 5.8 % decrease), which indicates that II barely has the character of carbene complex. To probe the function of the nuclear structure of the dirhodium catalyst, we analyzed the nature of the Rh–Rh–C π and σ -bond systems in **II** (carbene complex). Important orbital of **II** is shown in Fig. 3. LUMO of the complex **II** was found to be exactly what is expected and composed mainly of the carbene 2p orbital, which accepts rather small back-donation form the Rh¹ 4d_{xz} orbital to form an extended π^* -system that possesses a strongly electrophilic carbene carbon.

Analysis of the charge distribution during the reaction course was found to be informative and showed strong parallelism between this realistic model to form 4 (Fig. 4a), and the model to form 5 discussed later (Fig. 4b). The nitrone moiety releases negative charge (C⁴: +0.03 \rightarrow +0.16, O^6 : $-0.51 \rightarrow -0.42$) upon formation of II, and the negative charge moves largely into the vinylcarbene carbon C^1 $(+0.04 \rightarrow -0.12)$ and the N⁵ atom $(+0.09 \rightarrow +0.01)$. In the C^2-C^4 bond formation stage (TSII), the nitrogen moiety becomes less positive (N⁵: +0.01 \rightarrow -0.15, C⁴: $+0.16 \rightarrow -0.06$), and the carbon becomes cationic $(-0.12 \rightarrow +0.16)$. In the complex IV, the benzene ring attached to N⁵ atom significantly becomes more positive $(C^7: +0.15 \rightarrow +0.31)$, because the negative charge moves largely from the p-type orbital of the C^7 into the C^1 atom. The Charge of the Rh¹ and Rh² atoms changes just a little during the C¹–C⁸ bond formation (Rh¹: +0.44 \rightarrow +0.47; Rh²: +0.51 \rightarrow +0.55). However, in the C¹-C⁷ bond formation. Rh^1 and Rh^2 lose a lot of negative charge (Rh^1 : $+0.47 \rightarrow +0.68$; Rh²: $+0.55 \rightarrow +0.68$) and C⁷ becomes less positive ($+0.31 \rightarrow +0.13$), which is in accord with the removing of the donation from carbene to Rh. In the electrocyclic reaction TS (**TSV**), the C^1-C^7 and C^1-C^8 bond significantly shorten (1.580-1.503 and 1.537-1.490 Å, respectively.), and C^7 and C^8 atom lose negative charge $(C^{7}: +0.13 \rightarrow +0.20, C^{8}: -0.26 \rightarrow -0.19)$. The negative



Fig. 2 B3LYP/6-31G(d,p)[Rh-RSC+4f] structures of stationary points in CACR reaction of 2 and 3-4. The numbers refer to bond length (Å)



Fig. 3 Electronic structure and shape of the frontier electronic level (LUMO) of complex II

charge moves largely into the C¹ atom ($-0.16 \rightarrow -0.22$). In [1,7]-oxygen migration step, the C⁷–N⁵ bond shortens (1.449 Å \rightarrow 1.282 Å), and the benzene ring attached to N⁵ loses negative charge (C⁷: +0.20 \rightarrow +0.25, C⁸: $-0.19 \rightarrow$ +0.06). The nitrone moiety becomes negatively charged (N⁵ decrease by +0.21) because of forming of the C⁷–N⁵ π -bonding electrons. Some negative charge is also transferred to the O⁶ atom (-0.13).

3.2 Reaction to produce isomer 5

Furthermore, to prove the reasonableness of the mechanism proposed for the above catalytic system and gain insight into the influencing factors for highly regioselectivity for the present reaction, the other reaction of nitrone **3** with dirhodium vinylcarbene **2** to produce **5** was investigated. The free energy profile along the reaction progress is shown in Fig. 5. To make a concise expression, the detailed reaction mechanism to form **5** is shown in Fig. 6.



Fig. 4 Charge distribution along the reaction coordinate for a the reaction to 4 and b the reaction to 5



Fig. 5 Free energy profile of the $Rh_2(O_2CH)_4$ -catalyzed CACR reaction of 2 and 3 to produce 5 at the B3LYP/6-311+G(2d,2p)[Rh-RSC+4f]/B3LYP/6-31G(d,p)[Rh-RSC+4f] level. The result of the free energies in gas phase is given in*red*, and the result of the free energies in DCE is shown in*blue*

The reaction pathway and the energetics obtained for this reaction are essentially the same as those of the reaction to form 4 (Fig. 1). The CACR reaction also starts with the [3+2]-cycloaddition. This step goes through stepwise mechanism via **TSVII** and **TSVIII** with low free energy

barriers of 3.2 and 3.8 kcal/mol, respectively, in the gas phase. The charge distribution shows the same tendency as found for the reaction to form 4. In the next cyclopropanation step, carbene atom C^1 reacts with C^9 instead of C^8 to form the intermediate IX. Unlike the result in the reaction to obtain 4, the $C^1 - C^9$ bond formation step process via transition state TSIX with a significantly higher activation free energy (18.1 kcal/mol in gas phase). In the next electrocyclic opening of the cyclopropane ring, the free energy barrier for overcoming **TSXI** is much lower ($\Delta G^{\neq} = 5.7$ kca/ mol). Downhill from TSXI, seven-membered ring intermediate XI is formed. Since the activation of the N-O bond is poor, the electrophile attack of O to C^9 via a concerted transition state **TSXII** requires a higher Gibbs free energy of activation of 22.0 kcal/mol in gas phase, which is nearly identical with that of the reaction to form 4 (22.2 kcal/ mol). When taking the solvent effect into consideration, the energy profile has a little different from that in gas phase. The step of [1,7]-oxygen migration also bears a large free energy barrier (19.2 kcal/mol), which indicates that the oxygen migration is also difficulty in all cases. However, in the reaction to form 5, the C^1-C^9 bond formation bears the largest energy barrier (19.8 kcal/mol) and the cyclopropanation step has become the rate-determining step (RDS).

Two reactions mentioned above affording **4** and **5** are two parallel. Based on our calculational results, although the free energy barrier of RDS to **5** is larger than that to **4**, the difference between those two reactions is small. So the reaction dynamics is not the main factor responsive to the fact that the amount of product **4** dominates over **5** in DCE. We noted that an acidic additive 1,1,1,3,3,3-hexafluoro-2-propanol (HFIP) was used to capture the basic product in the experiment research [27]. So the proportion of alkaline dissociation equilibrium constant K_b values of products **4** and **5** was also calculated in our study according to the Eqs. (1) and (2).

$$B + H^+ \stackrel{\Delta G_B^{\mu}}{\rightleftharpoons} BH^+ \tag{1}$$

$$\frac{K_{b,4}}{K_{b,5}} = \exp\left(-\frac{\Delta G_4^{\theta} - \Delta G_5^{\theta}}{RT}\right) \\
= \exp\left(\frac{G_4^{\theta} - G_{H4^+}^{\theta} - G_5^{\theta} + G_{H5^+}^{\theta}}{RT}\right)$$
(2)

where ΔG_4^{θ} and ΔG_5^{θ} are the free energy changes of reactions corresponding to producing **4** and **5** in the standard state, respectively. *R* is mole gas constant, and *T* equals to 298.15 K. Table 1 lists the Gibbs free energies of product **4** and **5**, as well as their corresponding protonated substances **H4⁺** and **H5⁺**. It is easy to find that the basicity of the product **4** is stronger than that of the product **5**. As a



Fig. 6 B3LYP/6-31G(d,p)[Rh-RSC+4f] structures of stationary points in CACR reaction of 2 and 3-5. The numbers refer to bond length (Å)

Table 1 Gibbs free energy (in kcal/mol) of product 4 and 5, and their corresponding protonated substances $H4^+$ and $H5^+$ in DCE

Substance B	4	H4 ⁺	5	H5 ⁺
G^{θ}_{B} /kcal mol ⁻¹	-1,435.3	-1,437.0	-1,437.2	-1,438.1
		at	R ₁	
L _x	⁸ М=С(б⁺	L _x M	=C _δ -	
	R_2		R_2	

Fig. 7 Fischer and Schrock-type carbenoids

result, HFIP more easily reacts with 4 than 5, and thus, the reaction equilibrium will move to the formation of product 4, which is consistent with the experimental results that the reaction between 1 and 3 has high regioselectivity.

It is well known that rhodium carbenoids are Fischertype carbenoids (see Fig. 7). From the charge distribution along the reactions, it is clear that the carbene C^1 atom occurs electrophilic addition, along with a significant reduction of its charge. When there is an electronwithdrawing group such as chlorine at C^{10} of nitrone,the charge density of chlorine's *ortho*-site decreases due to the negative inductive effect of chlorine. On the other hand, the electron density on C^8 is slightly affected by the negative inductive effect of chlorine. Therefore, the electrophilic addition is more likely to occur at C^8 site, rather than C^9 site.

4 Conclusion

On the basis of our theoretical computations, we have developed a mechanistic proposal for the CACR reaction. The present studies have revealed, for the first time, the energetics, the electronic nature, and the 3-D structures of the intermediates and the transition states in the catalytic cycle of the dirhodium vinylcarbene-induced nitrone cvcloaddition with subsequent cascade carbenoid aromatic cycloaddition/N-O cleavage and rearrangement. The reaction of vinyldiazoacetate-induced nitrone starts from the stepwise [3+2] cycloaddition. Subsequent intramolecular cyclopropanation and electrocyclic opening of the cyclopropane ring by the rhodium carbene on the nitrogen-bound aryl group take place to form a heptatomic ring intermediate. A [1,7]-oxygen migration to final product occurs via a concerted transition state. The reactions to form 4 and 5 have similar mechanisms, and the step of cyclopropanation is the regioselectivity determining step. With meta-substituent (Cl) on the N-aryl group, the charge density of chlorine's ortho-site decreases due to the negative inductive effect of chlorine. Therefore, as compared to the electrophilic addition on the C^8 atom, the electrophilic addition on the C^9 site owns a significantly incensement on the energy barrier of cyclopropane (TSIII and TSIX). Since the basicity of the product 4 is stronger than that of the product 5, the acidic additive HFIP promotes the formation of product 4.

5 Supporting information

The optimized Cartesian coordinates and geometrical structures of all stationary points along the potential energy profiles. This material is available free of charge via the Internet at http://springerlink.lib.tsinghua.edu.cn/content/.

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REGULAR ARTICLE

Theoretical study on the working mechanism of a reversible light-driven rotary molecular motor

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Abstract Recent progress in the experimental design and synthesis of artificial light-driven molecular motors has made it possible to reverse the light-powered rotation of a molecular motor during the rotary process with multilevel control of rotary motion. With the interconversion between the photochemically generated less-stable isomers in one unidirectional rotary cycle with the stable isomers in the other cycle, the clockwise and anticlockwise rotations can be regulated with configuration inversion at the stereogenic centre. In this work, we report theoretical calculations by using density functional theory and time-dependent density functional theory, revealing the working mechanism of the clockwise and anticlockwise rotations. Specifically, by locating the relevant stationary points along the excitedstate and ground-state reaction paths, the clockwise and anticlockwise rotary cycles are explored in great detail,

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State Key Laboratories of Transducer Technology, Shanghai Institute of Microsystem and Information Technology, Chinese Academy of Sciences, Shanghai 200050, People's Republic of China which is of complementary importance in understanding the overall rotary process of the reversible rotation.

Keywords Light-driven molecular motor · Rotation reversal · Working mechanism · TD-DFT

1 Introduction

Molecular motor, in general, is a kind a nano-machine which can perform continuously mechanical functions by converting external energy to repetitive 360° motion. There are various molecular motors in nature, existing in the biological systems such as ATPase, dimeric kinesin, myosin, and dynein, which accomplish the extremely delicate and intricate microscopic works that compose the brilliant biological world. According to the different driven forces, it can be sorted as chemical [1], electrical, thermal, and light-driven molecular motors. On the other hand, due to the rapid development of the 'bottom-up' approach for experimental chemists, studies focusing on the design and synthesis of molecular-level machines that mimic the basic functions in nature have increased dramatically in the past decade. Although it is still far to construct artificial molecular motor that can perform integrated functions for realistic works, people have been able to design and to synthesis some relatively simple molecular motors which possess linear or rotary movement working under ambient conditions.

An eligible biological or artificial molecular motor that can be operated effectively at molecular level should work against the random thermal Brownian motion, which means the unidirectionality and controllability of the motor are essential criteria. The artificially designed molecular motors in laboratory that are powered by different energy sources and possess various mechanical motions have been inspired extensively by the pioneering synthetic unidirectional rotary molecular motors presented by Feringa [2, 3] and Kelly [4, 5] and their co-workers over the last 15 years.

One of the most interesting characteristics of the rotary motors existing in biological systems is the reversal of the direction of rotation [6]. Recently, a new molecular motor (henceforth denoted motor 1) driven by light, whose rotation direction can be controlled to be capable of performing reversible mechanical works by certain chemical catalysis epimerization, has been reported by Feringa et al. [7]. The design of motor 1 is based on one of the typical second-generation molecular motor (shown in Scheme 1) by replacing the methyl substituent at the stereogenic centre 2'position with electron-withdrawing-CONMe₂ group at the 3' position [7, 8]. In this way, the bidirectional 360° rotary cycle of motor 1 can be achieved, and the concerted working mechanism was experimentally observed, as shown in Fig. 1. According to the relative rotating between the upper rotor and lower stator, the two rotating cycles are named clockwise rotation and anticlockwise rotation, respectively.

Computational studies complement these experimental efforts and provide more insights into the rationale of the working mechanism, which could be helpful for the design and operation of these high rotational frequency motors in functional devices [9-12]. Moreover, quantum chemical studies could be regarded as the first step before exploring the rotary cycle of such motors by making them work as practical molecular machines. Some theoretical studies focusing on the working mechanisms of different light-driven molecular motors have been reported widely. Grimm et al. [13] have investigated the photochemical steps of a Feringa alkene-based molecular motor by using Car-Parrinello molecular dynamics simulations and thermal steps by semiempirical AM1 method, which has revealed the important role of methyl groups in determining the unidirectionality of the rotation. Excited-state dynamics studies of fluorene-based molecular motor also include TD-DFT simulation [14] and semiclassical trajectory surface hopping (TSH) simulations combined with orthogonalization-corrected OM2 Hamiltonian and multireference configuration interaction treatment (OM2/MRCI) by Filatov et al. [15]. Recently, the detailed photoisomerization processes of a redesigned stilbene-based light-driven molecular motor were specified by Liu and Morokuma [16] with CASPT2//CASSCF calculations. On the contrary, Perez-Hernandez and González [17] mainly focused on the possible reaction paths and corresponding transition states in the ground state.

As for the working mechanism of motor $\mathbf{1}$, despite the experimental observation has shown the bidirectional rotations, a detailed understanding of the excited-state photoisomerization and ground-state thermal helix inversion following the rotary motion is still of essential importance



Scheme 1 Second-generation (*left panel*) and redesigned reversible rotary motor 1 (*right panel*). With dihedral angles defined as follows, $\theta = C11'-C1'-C1-C10a$, $\alpha = C1-C1'-C11'-C10'a$, $\alpha' = C1'-C1-C10a-C10$, $\beta = C1-C1'-C2'-C3'$, and $\beta' = C1'-C1-C1a-C2$

considering improved design of reversible light-driven rotary molecular motors in the future.

We report quantum chemical calculations in this work by using density functional theory (DFT) [18–24] and timedependent density functional theory (TD-DFT) [25–29], which illustrate the overall rotary process including both the excited-state photoisomerization and the ground-state thermal helix inversion steps at the same level of theory. It is found that the long-range corrected (LC) density functional CAM-B3LYP [30] is able to demonstrate the unidirectionality of the photoisomerizations in the *clockwise* and *anticlockwise* rotary processes. By computing the potential energy curves (PECs) of the ground state and three lowest excited states, the different profiles of these states in the power-stroke photochemical reactions have been revealed.

2 Computational methods

The complete understanding of the rotary cycle of motor 1 involves the description of both photoisomerization reaction and thermal helix inversion steps. Therefore, the selection of a rational computational method that is equally good at describing excited-state and ground-state potential energy surfaces at the same level of theory is of fundamental importance. Although the widely used globally hybrid density functional B3LYP [18, 21, 31] is recognized as one of the most accurate for ground-state structures, energies, and frequencies [18, 21], there are still some limitations for handling partial π -bond breaking or interactions dominated by medium-range correlation energy such as overestimate of rotation barriers [32], failure to give correct excitation energies [33-35], and shortcomings in some other cases [36]. Hence, CAM-B3LYP was employed in the work motivated by its merit in describing the excited-state potential energy surfaces as well as relatively accurate ground-state energy barriers as demonstrated by Rostov et al. [34, 35].



Fig. 1 Schematic *clockwise* and *anticlockwise* rotary cycles of motor 1

Table 1 CAM-B3LYP and B3LYP ground-state geometric parameters and relative energetic differences of two most stable structures (3'S)-(M)-1 and (3'R)-(P)-1 with different basis sets

Isomer	Method	Basis set	θ	α	α'	β	β'	C1′–C1	ΔG
(3'S)-(M)-1	CAM-B3LYP	6-31G+(d,p)	0.2 (177.3)	-63.4	-50.5	-107.1	52.4	1.348	0.0
		6-311G++(d,p)	0.5 (177.3)	-63.6	-51.0	-107.2	52.9	1.343	0.0
	B3LYP	6-31G+(d,p)	-0.9 (176.5)	-63.1	-49.9	-107.0	51.7	1.359	0.2
		6-311G++(d,p)	-0.6 (176.7)	-63.4	-50.4	-107.1	52.2	1.354	0.2
(3'R)-(P)-1)	CAM-B3LYP	6-31G+(d,p)	-177.2 (-0.2)	63.5	50.4	107.0	-52.5	1.348	0.0
		6-311G++(d,p)	-177.3 (-0.5)	63.6	51.0	107.2	-52.9	1.343	0.0
	B3LYP	6-31G+(d,p)	-176.3 (1.1)	63.0	50.0	106.6	-51.9	1.359	0.0
		6-311G++(d,p)	-176.4 (0.7)	63.4	50.6	106.7	-52.5	1.354	0.0

The definition of dihedral angles is listed in Scheme 1. Dihedral angles in degrees, bond lengths in angstrom, and relative energies in kJ mol⁻¹. ΔG is the relative energy differences between (3'S)-(M)-1 and (3'R)-(P)-1 computed at the same level of theory

In order to explore the rotary cycle of motor 1 with clarity and simplicity, each 360° rotation process of *clockwise* and *anticlockwise* is divided into two identical rotary halves due to the symmetric chemical structure of the lower stator, i.e., every 180° half includes a photochemical and a thermal step (step 1 and step 2 for *clockwise* rotation and step 1' and step 2' for *anticlockwise* rotation as shown in Fig. 1). The two most stable stationary points ((3'S)-(M)-1) and (3'R)-(P)-1) on the ground state were located by geometry optimizations and frequency calculations using long-range corrected hybrid density functional CAM-B3LYP in combination with the 6-31+G(d,p) basis set. As a complementary comparison, geometry optimizations and frequency calculations with B3LYP functional and 6-31+G(d,p)basis set were also performed to ensure the reliability of CAM-B3LYP for ground-state properties (see Table 1 in following section).

Starting from (3'S)-(M)-1 and (3'R)-(P)-1) as the most stable structures on the ground state in each rotary cycle, constrained geometry optimizations along the dihedral angle C11'-C1'-C1-C10a (θ in Table 1) with an interval of 10° were carried out with the same DFT methodology to identify the ground-state PECs. Then, based on the ground-state minima and PECs, the excited-state PECs in each rotary cycle ((3'S)-(P)-1' and (3'R)-(M)-1') including the three lowest electronic excited states $(S_1, S_2, and$ S₃) were evaluated by TD-DFT single-point calculations at CAM-B3LYP/6-311++G(d,p) level of theory, where triple- ζ quality and more diffuse functions are included (basis set effect on the ground-state geometries also discussed in Table 1). It should be noticed that during the Franck-Condon (FC) relaxation, i.e. analytic TD-DFT gradients [28, 37] calculations from (3'S)-(M)-1and (3'R)-(P)-1 structures, 6-31+G(d,p) basis set and

Tamm–Dancoff approximation (TDA) [38] were employed in order to give balanced results between accuracy and efficiency. Afterwards, TD-DFT single-point excitation energies of the TDA optimized geometries were calculated at 311++G(d,p) level.

The less-stable structures (3'S)-(P)-1 and (3'R)-(M)-1can be recognized as products of photoisomerizations, whereafter the transition state search was carried out by constrained geometry optimization along the relevant dihedral angle α shown in Scheme 1. The saddle-point structure for each curve was taken out and was optimized to a transition structure at B3LYP/6-31G+(d,p) level of theory using the standard transition state search algorithms implemented in Gaussian 09 [39]. After a transition structure is optimized, the geometries and energies were refined using B3LYP/6-311G++(d,p) and CAM-B3LYP/6-311G++(d,p) together with the frequency calculations in order to confirm all the transition structures have only one imaginary frequency and to obtain thermal corrected free energies. Also, intrinsic reaction coordinate calculations were done to make sure the reactants and products do connected by a given transition structure.

All calculations in this work were performed with Gaussian 09 package of programs [39].

3 Results and discussions

3.1 The (3'S)-(P)-1 and (3'R)-(M)-1 isomers

Taken as the preliminary photochemical reactants in each rotary cycle, the ground-state geometric parameters and relative energy difference of (3'S)-(M)-1 and (3'R)-(P)-1 were computed by CAM-B3LYP and B3LYP with different basis sets, respectively, as shown in Table 1.

It can be known from Fig. 1 the stable mirror opposite enantiomer (3'R)-(P)-1 which can be obtained by base-catalysed epimerization of the less-stable (3'S)-(P)-1' should be of the same stereogenic structure and energetic stability with (3'S)-(M)-1. It is confirmed by the comparison in Table 1, in which the maximum disparities are 0.1°, 0.0 Å, and 0.0 kJ mol⁻¹ for CAM-B3LYP/6-31G+(d,p), 0.0°, 0.0 Å, and 0.0 kJ mol⁻¹ for CAM-B3LYP/6-311G++(d,p), 0.4°, 0.0 Å, and 0.2 kJ mol⁻¹ for B3LYP/6-31G+(d,p), and 0.4°, 0.0 Å, and 0.2 kJ mol⁻¹ for B3LYP/6-311G++(d,p) in dihedral angles, C1'-C1 bonds, and relative energies. Moreover, the increase in the size of the basis set from 6-31G+(d,p) to 6-311G++(d,p) has similar and minor effect on the chemical structures, with maximum disparities of 0.5° and 0.005 Å for CAM-B3LYP, and 0.6° and 0.005 Å for B3LYP.

From this point of view, both functionals combined with each of two basis sets give reasonably good description of the ground-state geometry and energy. Importantly, CAM-B3LYP predicts (3'S)-(M)-1 and (3'R)-(P)-1 to be exactly identical, which proves that this functional is capable of yielding accurate ground-state results needed for the calculations in this work.

3.2 FC relaxation and potential energy curves

As the first power stroke in the rotary cycle, the irradiation of light promotes the two initial structures in each cycle, (3'S)-(M)-1 and (3'R)-(P)-1, to the higher energy excited electronic states, followed by an ultrafast relaxation to the S_1 minimum region, which can be evidenced by the electronic structure analyses for the three lowest electronic excited states. (For detailed discussion, please refer to the Supporting Information.) The energy difference between vertical and adiabatic excitation energy is the driving force of the photoisomerization process, and the isomer is considered to go down to the ground state near the twisted perpendicular conical intersection. Therefore, the directionality of the photoinitiated rotation around C1'-C1 is controlled by the FC relaxation and can be reflected by the shape of PECs on the excited state. Some key parameters of excitation energy and geometry variation during this process are collected and shown in Table 2.

Table 2 S_1 vertical (VEE) and adiabatic (AEE) excitation energies (eV) of (3'S)-(M)-1 and (3'R)-(P)-1 calculated at CAM-B3LYP with 6-31G+(d,p) and 6-311G++(d,p) level of theory

Isomer	Basis set	λ_{max}^{b}	VEE	AEE	Geometry change	
					$\Delta \theta$	ΔC1′–C1
(3'S)-(M)-1	6-31G+(d,p)	3.97	4.14	2.99	$0.2 \rightarrow -57.5$	$1.348 \rightarrow 1.447$
	$6-311G++(d,p)^{a}$		4.14	3.01	_	-
(3'R)-(P)-1	6-31G+(d,p)	3.97	4.14	2.99	$-177.2 \rightarrow -126.4$	$1.348 \rightarrow 1.447$
	$6-311G++(d,p)^{a}$		4.14	3.01	-	_

^a The excitation energies are computed from the geometries optimized with 6–31G+(d,p) basis set

^b Experimental absorption maximum taken from Ref. [7]

It can be seen from Table 2 how well CAM-B3LYP can describe the excitation energies of the excited electronic states. The vertical excitation energies (~0.15 eV higher than the experimental values) are in reasonable agreement with the experimental absorption maxima which should correspond to the vertical approximation, as the calculated adiabatic excitation energies are ~ 0.90 eV lower than the experimental values. The basis set effect was evaluated by single-point calculations increasing basis set from 6-31G+(d,p) to 6-311G++(d,p) and only shows marginal difference (~0.02 eV in adiabatic excitation energies). With regard to the geometry changes during the FC relaxation of the two isomers, there are both similarity and difference. While a pronounced elongation of ~ 0.1 Å in C1'-C1 bond is observed for both isomers, the driving forces on the relative rotation between rotor and stator of (3'S)-(M)-1 and (3'R)-(P)-1 are of opposite directions which can be observed clearly in Fig. 2.

In Fig. 2, the S_0 minimum energy path (MEP) and the single-point excitation energies of the first three excited states for (3'S)-(M)-1 and (3'R)-(P)-1 are shown. It should be pointed out that the aim of the research is to elucidate the reversible rotation processes and to understand the working mechanism of the motor; therefore, the expensive excited-state potential energy surface calculation was not performed. Furthermore, although CAM-B3LYP functional performs well for the excitation energy calculation, it is a single-reference method and is well known not as good as multireference methods in describing conical intersections. Instead, the nature of the excited-states MEPs was estimated by computing the single-point excitation energies of the first three excited states from the optimized geometries on S_0 MEP, and still some interesting features are revealed.

As shown in Fig. 2, the key feature is the inversion symmetry of the PECs of the two isomers. As for the (3'S)-(M)-1 isomer, the FC relaxation on the excited state follows the S₁ PEC after the molecule is excited, which results from the barrierless feature of the first excited state and high energy barriers on the second and third excited states.

The descent of energy from the FC point on the S₁ PEC drives the excited state isomerization towards the twisted intermediate conformation $(3'S)-(M)-1^*$. The dihedral angle θ changes from $\sim 0^\circ$ to -60° roughly, suggesting an efficient *clockwise* initial rotation. While for the (3'R)-(P)-1 isomer, the PECs feature similar characteristics with that of the (3'S)-(M)-1 isomer, except the dihedral angle θ changes from $\sim -180^\circ$ to $\sim -120^\circ$ towards $(3'R)-(P)-1^*$ and is in an *anticlockwise* fashion.

As indicated in the discussion above, both S_1 and S_0 PECs in Fig. 2 cannot be capable of locating the conical intersections between the excited state and ground state. Nevertheless, an excitation energy of approximately zero is shown at -130° for (3'S)-(M)-1 and at -50° for (3'R)-(P)-1, which indicates that the conical intersection is in the range of $(-60^{\circ}, -130^{\circ})$ for (3'S)-(M)-1 and $(-120^{\circ}, -130^{\circ})$ -50°) for (3'R)-(P)-1, respectively. Such assumption can be further confirmed by carrying out ground-state geometry optimizations from twisted (3'S)-(M)-1* and (3'R)-(P)-1* conformations to locate the less-stable photochemical products. Interestingly, it was observed that in the intermediate (3'S)-(M)-1* and (3'R)-(P)-1* conformations, the orientation of phenyl rings on the stator has changed from pointing-in the plane to pointing-out direction. Furthermore, when optimized to the ground-state minima from the excited-state intermediates $(3'S)-(M)-1^*$ and $(3'R)-(P)-1^*$, the orientation of phenyl rings on the stator was preserved and less-stable structures (named as (3'S)-(P)-1 and (3'R)-(P)-1(M)-1) different with that in the original working mechanism (shown in Fig. 1) were obtained. The photochemical products (3'S)-(P)-1 and (3'R)-(M)-1 suggest that motor 1 should work in a slightly different manner that does lead to the experimentally observed thermal barriers.

3.3 Working cycles and transition states

Taken as the rate-determining steps in the overall rotary cycle, the thermal helix inversions of step 2 and step 2' start from the photoisomerized (3'S)-(P)-1 in the clockwise

Fig. 2 FC relaxation directions and excitation energies with S_0 MEPs for (3'S)-(M)-1 (left panel) and (3'R)-(P)-1 (right panel). The photoisomerization directions indicated by green arrows



rotation and from (3'R)-(M)-1 in the anticlockwise rotation, during the first half of each rotary cycle. The location of transition states is of utmost importance in determining the rotary frequency and in understanding the working mechanism of the motor. To this aim, two-dimensional relaxed potential energy scans are carried out by changing torsional angle α and α' in order to get suitable starting geometries for the geometry optimizations of transition states. After the corresponding transition states are optimized, all of the requisite frequency and intrinsic reaction coordinate calculations are performed to get the free energies and to verify the reactants and products mediated by the transition states. (The atomic coordinates of optimized ground and excitedstate stationary points are provided in Supporting Information.) By means of locating all the possible stationary points, the proposed feasible working cycles for the clockwise and anticlockwise rotations are shown in Fig. 3. The exclusive *syn-fold* structures (3'S)-(P)-1 and (3'R)-(M)-1 of the photochemical products have been theoretically predicted, which means that there is an *anti-fold* \rightarrow *syn-fold* geometrical transformation during the rotary process. All the five isomers of motor 1 in step 1 and step 2 of clockwise rotation and another five isomers in step 1' and step 2' are depicted in Fig. 4, where also given are the relative stabilities of different isomers with respect to the *global* stable isomer in each rotary cycle.

As shown clearly in Fig. 4, the clockwise and anticlockwise rotary cycles show highly symmetric characteristic, which has an opposite rotary direction but nearly equivalent relative free energies for all the corresponding isomers



Fig. 3 Proposed feasible working cycles for the clockwise and anticlockwise rotations



Fig. 4 Relative free energies (in kJ mol⁻¹) at CAM-B3LYP/6–31G+(d,p) level of theory with respect to the most stable isomers of motor 1 in the *clockwise* and *anticlockwise* rotations

in each cycle. The step 1 of the clockwise rotation and step 1' of the anticlockwise rotation are the power stroke photochemical steps, which go through the excited-state intermediates (3'S)-(M)-1* and (3'R)-(P)-1*, respectively. The subsequent concerted thermal helix inversion steps are accomplished via the transition states TS1 and TS1'. The Gibbs free energies for the thermal isomerizations were found experimentally to be 107.9 kJ mol⁻¹ for the energetically downhill isomerization and 120.1 kJ mol⁻¹ for the uphill isomerization [7]. Accordingly, the obtained activation barriers for the downhill isomerization (from (3'S)-(P)-1 to TS1 for step 2 and from (3'R)-(M)-1 to TS1' for step 2') and the uphill isomerization (from (3'S)-(M)-1 to TS1 for step 2 and from (3'R)-(P)-1 to TS1' for step 2') are about 111.9 and 135.2 kJ mol⁻¹, which is in good agreement with the experimental ones. Also compared with the experimental results is the temperature effect on the rate constant of the thermal downhill isomerization process which gives the same trend as provided by the experiment (See Supporting Information for more detail).

As for the different photoisomerization products compared with that observed from the experiment, our calculations do predict the *syn-fold* conformers in both clockwise and anticlockwise rotary cycles. Although there is lack of evidence for the existence of *syn-fold* isomers of motor 1, both experimental and theoretical studies have identified the *syn-fold* conformers as possible chemical constitutions for certain second-generation light-driven molecular motors with analogous structure [17, 40].

4 Conclusions

In summary, using long-range corrected density functional CAM-B3LYP, we have explored the working mechanism of a recently synthesized reversible light-driven molecular motor in detail by locating all the stationary points along the reaction paths. The different profiles of the groundstate and three lowest excited-states PECs were evaluated, by which the unidirectional clockwise and anticlockwise rotary motions of the photoisomerizations can be well understood. Although the appreciable structures of the conical intersections of S₀/S₁ are not accurately located, we are able to obtain the less-stable photochemical products by optimizing the highly twisted molecular structures within a large range to the ground state. Interestingly, the photoisomerization products show syn-fold conformations which are slightly different with that in the experiments, for which there are possible evidences both from experimental and theoretical studies of light-driven molecular motors with similar structures. Finally, by comparing the excitation energies and calculated free-energy barriers with the experimental values, it has been found that the CAM-B3LYP function cannot only give good description of excited states, but also show good performance for the thermal helix reactions.

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REGULAR ARTICLE

The combined CASPT2 and CASSCF studies on photolysis of 3-thienyldiazomethane and subsequent reactions

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Abstract The triplet 3-thienvlcarbene and α -thialmethylenecyclopropene were observed as the main products upon photolysis ($\lambda > 540$ nm) of 3-thienyldiazomethane (3-TD) under the matrix-isolated conditions. The underlying mechanism was explored by the combined CASPT2 and CASSCF calculation in the present work. The S₁, T₂, and T₁ surfaces were found to be quasi-degenerate over a wide structural region. Meanwhile, there is a relatively strong spin-orbit interaction between the S₁ and T₂ states, due to the sulfur-atom effect. As a result, the triplet 3-thienylcarbene can be formed with a noticeable efficiency. The S₁ minimum and S₁/S₀ intersection structures for 3-TD were predicted to be undistinguishable by the present CASPT2//CASSCF calculations. Therefore, the $S_1 \rightarrow S_0$ internal conversion that occurs in the vicinity of the S_1/S_0 intersection (S_1 minimum) is in competition with the $S_1 \rightarrow T_1$ intersystem crossing through the three-surface $(S_1, T_2, and T_1)$ intersection region. Once the excited 3-TD molecule returns to the S₀ state, the singlet 3-thienylcarbene can be generated with a considerable efficiency, which is followed by the carbene rearrangement, leading to the final product of α -thial-methylenecyclopropene (α -TMC). The concerted N₂ loss and rearrangement in the excited state was proposed for the α -TMC formation in previous experimental study, which is not supported by the present CASPT2//CASSCF calculation. The reason for this

Dedicated to Professor Guosen Yan and published as part of the special collection of articles celebrating his 85th birthday.

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Key Laboratory of Theoretical and Computational Photochemistry, Ministry of Education, College of Chemistry, Beijing Normal University, Beijing 100875, China e-mail: Fangwh@bnu.edu.cn might come from the matrix effect, which is open for further study.

Keywords 3-Thienyldiazomethane · Photodissociation · Ab initio calculation

1 Introduction

Diazomethane and its derivatives have been widely used as reagents in synthetic organic chemistry. For example, they were employed as methylating agents in cyclopropanation reactions with olefins [1-4] and for one-carbon chain elongation by insertion into the primary C-H bond [5]. Diazomethane derivatives can function as photochemical precursors of singlet carbenes, which are the reactive intermediates and play important role in many branches of chemistry [6-10]. Mechanistic study on how to form the singlet carbene from these precursors has a long and venerable history [6, 11] and is still active area in the fields of chemical kinetics and physical organic chemistry [12-16]. The photolysis of diazomethane has been extensively studied from experimental and theoretical viewpoints [13, 17–21]. Upon irradiation of diazomethane (CH_2N_2) in the gas phase, methylene (CH_2) was observed as the initial product in different singlet states that are dependent on excitation wavelengths (a weak $n \rightarrow \pi^*$ transition at 300–500 nm and a strong $\pi \rightarrow \pi^*$ transition at 200-260 nm) [17-20]. It has been argued whether the dissociation of CH_2N_2 to $CH_2 + N_2$ occurs in the excited electronic state [13, 21].

Ultrafast time-resolved spectroscopic techniques have been used to probe the excited-state dynamics of aryl diazocompounds and the formation of reactive carbene intermediates [22–29]. The excited-state lifetimes of *p*-biphenylyldiazomethane (BpCN₂H) and *p*-biphenylyldiazoethane (BpCN₂CH₃) were inferred to be within 300 fs by femtosecond time-resolved UV-vis transient absorption studies [23–27]. The experimentally observed absorption bands at ~360 nm were assigned to the singlet carbenes (¹BpCH and ¹BpCCH₃) with their lifetimes changed from a few hundreds of picoseconds in acetonitrile to ~ 10 ps in a carbene scavenging solvent of methanol. Meanwhile, the quantum yield for formation of ¹BpCCH₃ was determined to be much lower than that for formation of ¹BpCH. It was proposed that the excited state of BpCN₂CH₃ underwent 1,2-H shift in concert with N₂ extrusion [24, 25]. Subsequently, the direct experimental evidence for the rearrangement in the excited states was obtained by ultrafast time-resolved IR spectroscopic study of alkyl diazoester (CH₃CN₂CO₂CH₃) in chloroform [29].

It is well known that singlet carbenes are primarily released by photodissociation of aryl diazocompounds [23–29]. The nascent singlet carbene can undergo either intramolecular or intermolecular reactions to form final products. As organic reactive intermediates, heteroaryl carbenes have been extensively investigated in the past several decades, due to their importance in astronomical environments [30]. A major channel for formation of heteroaryl carbene involves pyrolysis and photolysis of heteroaryl diazocompounds, such as thienyldiazomethanes. Because of the interaction between the heteroaryl ring and electronrich diazo groups, thienyldiazomethanes were expected to exhibit some unique characters. Early in 1978, Shechter and co-workers studied pyrolysis of 2-thienyldiazomethane and its methyl-substituted analogs [31]. Heteroaryl carbenes were inferred to be reactive intermediates, and the main products were observed to be unsaturated thiocarbonyls and 1,2-di(2-thienyl)ethylenes. The primary product has been proposed to be 3-thienylcarbene upon photolysis of 3-thienyldiazomethane in argon at 10 K with $\lambda > 450$ nm [32]. In order to determine whether the triplet 3-thienylcarbene is formed initially, a spectroscopic study has been performed on photochemistry of furyl- and thienyldiazomethanes by IR, UV/vis, and EPR spectroscopy [16]. The conformational isomers of 3-thienylcarbene exhibit a weak electronic absorption of triplet characteristic in the visible spectrum ($\lambda_{max} = 467$ nm) and an unusually large difference in zero-field splitting parameters in the EPR spectrum, which afford strong evidence for formation of the triplet 3-thienylcarbene. However, the underlying mechanism is still unclear up to now.

Photolysis of diazomethane and its aryl-substituted derivatives has been a subject of numerous ab initio calculations and dynamics simulations in order to explore the mechanism for formation of singlet carbene [13, 15, 20, 21, 33]. However, how the triplet 3-thienylcarbene is produced from the photolysis of 3-thienyldiazomethane was not explored from theoretical viewpoint up to now. In the present work, we performed the combined CASPT2//CASSCF study on structures and reactivity of 3-thienyldiazomethane in the ground and low-lying excited states, which enable us to understand mechanism of 3-thienylcarbenes generated by the photolysis of 3-thienyldiazomethane at $\lambda > 540$ nm. Meanwhile, 3-thienylcarbene rearrangements in singlet and triplet states were determined on the basis of the CASSCF optimized stationary and intersection structures as well as the CASPT2 calculated relative energies. We believe that the present study provides new insights into mechanism for formation of 3-thienylcarbene, which is very helpful for understanding mechanistic photochemistry of heteroaryl diazocompounds.

2 Computational details

Stationary structures in the ground and low-lying excited states were fully optimized at the level of the complete active space self-consistent field (CASSCF), which was confirmed to be appropriate for the potential energy surface (PES) topology analysis of diazocompounds [13, 20]. All the optimized stationary structures were characterized as minima or saddle points on PESs by frequency analysis. The minimum-energy structures on intersection seams of different electronic states were determined with the state-averaged CASSCF wave functions to ensure a balanced description of multi-states at the intersection area. On the basis of the CASSCF optimized structures for the stationary and intersection points, the energies were recomputed with the CASPT2 method (the complete active space perturbation theory to second order). The combined CASPT2//CASSCF calculation was extensively used to study excited-state behavior of different systems [34-37] and proven to be well suited for the accurate description of the lowest-lying electronic states. The active space of the present CASSCF calculation for 3-thienyldiazomethane is comprised of twelve electrons distributed in eleven orbitals, assigned as CAS(12,11) hereafter, which is composed of all π and π^* orbitals. Since the C–N fission occurs in the process of N₂ extrusion, the C–N σ and σ^* orbitals are included in the active space. As for 3-thienylcarbene and its rearrangement reactions, we chose eight electrons in eight orbitals as the active space, which originate from four π and π^* orbitals in the thiophene ring, the C–S σ and σ^* orbitals, and the two singly occupied orbitals of the carbene C atom. The 6-31G* basis set was used for all CASSCF and CASPT2 calculations, which were carried out with Gaussian 09 and Molcas 7.4 software packages [38, 39].

3 Results and discussion

3.1 The lowest-lying electronic states

Photochemistry of 3-thienyldiazomethane (3-TD) was investigated with excitation wavelength of 543 nm or longer [16]. In order to identify the initial state populated by irradiation at ~540 nm, geometric structures of 3-TD in the lowest-lying electronic states were optimized at the CAS(12,11)/6-31G* level and the obtained results were shown in Fig. 1, along with the selected bond parameters.

As expected, the C=N=N moiety has a linear configuration in the ground-state (S_0) equilibrium structure of 3-TD (3-TD_S₀_min). However, the C–N=N angle becomes ~126° in the minimum-energy structures of the lowest excited singlet (S_1) and triplet (T_1) states, which are referred to as 3-TD_S₁_min and 3-TD_T₁_min, respectively. Meanwhile, the C–N bond is mainly of a

Fig. 1 CASSCF optimized stationary and intersection structures, along with the selected bond lengths in angstrom

single-bond character in 3-TD_S₁-min and 3-TD_T₁-min. On the basis of the CAS(12,11)/6-31G* calculation for 3-thienyldiazomethane in the low-lying electronic states, eleven active orbitals are schematically shown in Fig. 2 and are labeled by AOi (i = 1-11). Two singly occupied orbitals of the S₁ or T₁ state are found to be AO6 (the out-ofplane C–N bonding π orbital) and AO8 (the in-plane N=N anti-bonding π^* orbital). It is evident that the S₁ or T₁ state originates from $\pi \rightarrow \pi^*$ transition localized in the C–N–N structural region. The second excited singlet (S₂) and triplet (T₂) states are of the $\pi\pi^*$ nature and are mainly localized in the thiophene ring with a considerable delocalization into the C–N–N region, since two unpaired electrons for the S₂ and T₂ states are mainly distributed in AO5, AO9, and AO10 in Fig. 2.

On the basis of the CAS(12,11)/6-31G* optimized 3-TD_ S₀_min structure, the vertical excitation energies to the S₁ and S₂ states were calculated to be 58.4 and 108.8 kcal/mol




Fig. 2 Active orbitals of the CASSCF calculations for 3-thienyldiazomethane (a) and 3-thienylcarbene (b)

at the CASPT2//CAS(12,11)/6-31G* level of theory. In comparison with the absorption peak of 494 nm (57.9 kcal/mol) for 3-thienyldiazomethane in acetonitrile solution [16], we come into conclusion that the S_1 state is initially populated upon photolysis of 3-TD at 490 nm and longer wavelengths.

3.2 Formation of the singlet 3-thienylcarbene

It has been established that 3-thienylcarbene (3-TC) is energetically lower in the lowest triplet state than in the lowest singlet state [16], which are, respectively, referred to as 3-TC_T_1 and 3-TC_S_1 hereafter. Once the 3-TD molecule is excited to the S₁ state, the C–N bond cleavage might occur along the S₁ pathway. Direct extrusion of molecular nitrogen from 3-TD on the S₁ state was traced by the CAS(12,11)/6-31G* stepwise optimizations, followed by the CASPT2 single-point energy correction. The barrier height for the C–N bond cleavage on the S₁ state was estimated to be 69.5 and 39.7 kcal/mol with respect to the S₀ and S₁ minima, respectively. It is evident that the direct

 S_1 C–N fission is inaccessible in energy upon photolysis of 3-TD in the wavelength range of 490–540 nm. Therefore, there is little possibility that the singlet 3-thienylcarbene (3-TC_S₁) is directly formed along the S₁ pathway.

From the S_1 state, there are mainly two non-radiative pathways for the 3-TD molecule to deactivate: internal conversion (IC) to the S_0 state and intersystem crossing (ISC) to a triplet state. Here, we pay attention to the $S_1 \rightarrow S_0$ IC process and the subsequent formation of the singlet 3-thienylcarbene along the S_0 pathway. The minimum-energy conical intersection between S1 and S0 was determined by the two state-averaged CAS(8,7)/6-31G* calculations, referred to as $3-TD_S_1/S_0$. It was found that the $3-TD_$ S_1 _min and 3-TD_ S_1/S_0 structures are not distinguishable from each other. As a result, the IC process from S_1 to S_0 takes place very effectively, leading to the 3-TD molecule in a hot S₀ state. A transition state, 3-TD_S₀_TS, was determined on the S₀ pathway from 3-TD_S₀_min to the singlet 3-thienylcarbene and N2. The barrier height was predicted to be 32.1 kcal/mol at the CASPT2//CAS(12,11)/6-31G* level of theory. The present CASPT2//CASSCF calculation reveals that the singlet 3-thienylcarbene $(3-TC_S_1)$ can be formed with a considerable efficiency upon irradiation of 3-thienyldiazomethane at ~540 nm.

3.3 Formation of the triplet 3-thienylcarbene

In addition to the $S_1 \rightarrow S_0$ IC process, the $S_1 \rightarrow T_1$ intersystem crossing is another important pathway for the S_1 deactivation, which is probably responsible for the triplet 3-thienylcarbene observed experimentally [16]. The minimum-energy crossing point between singlet and triplet states was optimized at the state-averaged CAS(12,11)/6-31G* level. The optimized structure was confirmed to be the crossing point between the S_1 and T_2 states by the CASSCF wave functions and electron populations, referred to as $3-TD_S_1/T_2$ hereafter. As shown in Fig. 1, the 3-TD_S₁/T₂ structure is still in the S₁ Franck–Condon (FC) region and has its relative energy of 52.5 kcal/mol at the CASPT2//CAS(12,11)/6-31G* level. As pointed out before, the two unpaired π electrons in the T₂ state are mainly distributed in the thiophene ring and the C=N bond is mainly of a double-bond character. In addition, the C=N bond fission from the T₂ state is adiabatically correlated with an excited triplet state of 3-thienylcarbene, which is not favorable in energy. Thus, there is little possibility for the C=N bond to break along the T_2 pathway to the triplet 3-thienylcarbene observed experimentally.

To explore the possibility of the $T_2 \rightarrow T_1$ internal conversion, a conical intersection between the T_1 and T_2 states was determined by the state-averaged CAS(12,11)/6-31G* optimization, referred to as 3-TD_T₂/T₁ hereafter. As shown in Fig. 1, 3-TD_T₂/T₁ is similar to 3-TD_S₁/T₂ in

structure. It is found that a small difference in 3-TD $T_2/$ T_1 and 3-TD_S₁/T₂ structures originates mainly from the redistribution of the conjugated π electrons, which will not result in a large change in energy. The energy difference between $3-TD_T_2/T_1$ and $3-TD_S_1/T_2$ was predicted to be 2.6 kcal/mol by the CASPT2/CAS(12,11)/6-31G* calculations. Test single-point energy calculations at the CASPT2/CAS(12,11)/6-31G* level for a series of the structures show that the S₁, T₂, and T₁ states are nearly degenerate in the structural region from $3-TD_S_1/T_2$ to $3-TD_T_2/T_2$ T₁. As discussed before, the T₂ state is of the ${}^{3}\pi\pi^{*}$ character localized in the thiophene ring. Because of a heavyatom effect of sulfur, there is a relatively strong spin-orbit interaction between the S1 and T2 states, which enhances considerably the rate of the $S_1 \rightarrow T_2$ process. The spinorbit coupling (SOC) between S₁ and T₂ was calculated to be 35.1 cm⁻¹ in the vicinity of the 3-TD_S₁/T₂ structure by using a one-electron operator, and effective nuclear charges for spin-orbit interaction implemented in Gaussian 03 software package. In addition, the time scale for the $T_2 \rightarrow T_1$ internal conversion process in the vicinity of the $3-TD_T_2/T_1$ conical intersection is expected to be on the order of a vibrational period [40]. Thus, the $S_1 \rightarrow T_1$ intersystem crossing occurs efficiently in the structural region from $3-TD_S_1/T_2$ to $3-TD_T_2/T_1$, due to a function of the T_2 state as a bridge [41, 42].

The C–N bond cleavage of 3-thienyldiazomethane was determined to be the most probable pathway on the T_1 state, leading to formation of the triplet 3-TC and N_2 . A transition state of 3-TD_T₁_TS was obtained on the T_1 pathway, which was confirmed to be the saddle point on the T_1 pathway from 3-TD_T₁_min to N_2 and 3-TC_T₁_min (the equilibrium geometry of 3-thienylcarbene in the T_1 state). The barrier height was calculated to be 13.1 and 34.8 kcal/mol with respect to the T_1 and S_0 vibrational zero levels, respectively. Formation of the triplet 3-thienylcarbene along the T_1 pathway is favorable in energy upon irradiation of 3-thienyldiazomethane at ~540 nm.

3.4 Formation of α-thial-methylenecyclopropene

Upon irradiation of 3-thienyldiazomethane (3-TD) at $\lambda > 534$ nm [16], α -thial-methylenecyclopropene (α -TMC) was observed as one of the main products in Ar or N₂ at 10 K. There are two possible mechanisms responsible for the α -TMC formation, the concerted N₂ loss and rearrangement in the excited state of the diazo precursor [16, 24, 29] and the stepwise process of molecular nitrogen extrusion followed by carbene isomerization in the ground state [27, 32, 43]. The concerted process from 3-TD to α -TMC involves formation of the C–C bond and fissions of the C–N and C–S bonds simultaneously. Because the initial relaxation from the S₁ Franck–Condon (FC) structure



Fig. 3 Schematical potential energy profiles for formation of the singlet 3-thienylcarbene (**a**), for formation of the triplet 3-thienylcarbene (**b**), and for rearrangement of the singlet 3-thienylcarbene (**c**). The

stationary structures on the reaction pathways and their relative energies (kcal/mol) are given, together with some key bond lengths (Å) in the related structures

to 3-TD_S₁_Min is downhill process, the concerted process takes place in the S₁ state with little possibility. In order to explore the possibility that the concerted process occurs in the S₀ state, the S₀ potential energy surface was scanned as a function of the C–N and C–S distances with the other bond parameters optimized fully. The scanned S₀ potential energy surface shows clearly that the reaction from 3-TD to α -TMC proceeds in a stepwise way. The first step involves the C–N bond fission of 3-TD_S₀ yielding singlet 3-thienylcarbene (3-TC_S₁), which is followed by the C–S bond fission and isomerization to α -TMC.

As pointed out before, the direct S_1 C–N fission is inaccessible in energy upon photolysis of 3-TD at $\lambda > 540$ nm. Since the ultrafast $S_1 \rightarrow S_0$ internal conversion occurs at the vicinity of the 3-TD_S₁/S₀ conical intersection, the singlet 3-thienylcarbene (3-TC_S₁) was formed from 3-TD_S₀ with a barrier of 32.1 kcal/mol, which is considered as the first step (3-TD_S₀ \rightarrow 3-TC_S₁ + N₂) for formation of α -TMC from the 3-TD photolysis. An intermediate was determined on the pathway from 3-TC_S₁ to α -TMC, which is assigned as 1H-2-thiabicyclo[3.1.0]hexa-3,5-diene (THD) observed experimentally [16]. As shown in Fig. 3, the barrier height was calculated to be 7.7 kcal/mol at the CASPT2//CAS(8,8)/6-31G* level for the second step of 3-TC_S₁ \rightarrow THD, while the third step of THD $\rightarrow \alpha$ -TMC has a barrier of 17.7 kcal/mol at the same level of theory.

On the basis of the CASSCF calculated frequencies for the stationary structures on the pathway from 3-TD_ S_0 to α -TMC + N_2 and the CASPT2 calculated barrier heights, the RRKM rate theory [43] was used to estimate rate constants for each step of the sequential reaction with the CASPT2//CAS(12,11)/6-31G* calculated vertical excitation energies of 58.4 kcal/mol as the available energies. The calculated rate constants indicate that the 3-TC_ S₁ formation is the rate-determining step and the rate constant of the THD formation is about 100 times larger than that on the third step from THD to α -TMC. As a result, the THD intermediate might be observed experimentally, when the sequential reaction is in its stationary state. However, there is little possibility to have the singlet 3-thienylcarbene observed under the experimental condition. A minor amount of a species was assigned as THD, which is based on a comparison of experimental and the computed IR spectra [16].

4 Summary

In the present work, we performed the combined CASPT2//CASSCF study on photodissociation of 3-thienyldiazomethane (3-TD) and the subsequent reactions. Upon irradiation of 3-TD at ~540 nm, the $S_1(^1\pi\pi^*)$ state is initially populated, which originates from one-electron excitation from the out-of-plane C–N π orbital to the inplane N=N π^* orbital. From the S₁ state, there are three possible non-radiative pathways for the 3-TD molecule to deactivate: the direct S_1 reactions, internal conversion (IC) to the S₀ state, and intersystem crossing (ISC) to a triplet state. The CASPT2//CASSCF calculated results provide a strong evidence that the C-N bond fission and the concerted N_2 loss take place in the S_1 state with little possibility. On the basis of CASSCF optimized structures for the S1/T2 and T2/T1 intersections as well as the CASPT2//CASSCF calculated relative energies, it is found that the S_1 , T_2 , and T_1 surfaces are near degenerate over a wide structural region. Because of the heavy-atom effect of sulfur, there is a relatively strong spin-orbit interaction between the S_1 and T_2 states, which enhances considerably the rate of the $S_1 \rightarrow T_2$ process. In addition, the time scale for the $T_2 \rightarrow T_1$ internal conversion process is expected to be on the order of a vibrational period in the vicinity of T_2/T_1 conical intersection. Thus, the $S_1 \rightarrow T_1$ intersystem crossing occurs efficiently with a function of the T₂ state as a bridge, once the 3-TD molecule is populated in the S₁ state. A barrier of 13.1 kcal/ mol was determined by the CASPT2//CASSCF calculation on the T_1 pathway from the triplet 3-thienyldiazomethane $(3-TD_T_1)$ to the triplet 3-thienylcarbene $(3-TC_T_1)$. All of these enable us to understand mechanism of 3-thienylcarbenes generated by the photolysis of 3-thienyldiazomethane at $\lambda > 540$ nm [16].

The 3-TD_S₁-min and 3-TD_S₁/S₀ structures were found to be undistinguishable by the present CASPT2//CASSCF calculations. Therefore, the S₁ \rightarrow S₀ IC process takes place very effectively after irradiation of the 3-TD molecule to the S₁ state, which is at least in competition with the $S_1 \rightarrow T_1$ ISC process. Once the 3-TD molecule returns to the hot S₀ state, the singlet 3-thienylcarbene $(3-TC_S_1)$ can be formed with a barrier of 32.1 kcal/ mol on the pathway. Since the singlet 3-thienylcarbene is very reactive, the α -thial-methylenecyclopropene (α -TMC) can be formed easily, which has been observed as one of the main products upon photolysis of 3-TD in Ar or N₂ at 10 K. An intermediate was determined on the pathway from 3-TC_{S_1} to α -TMC, which is assigned as 1H-2thiabicyclo[3.1.0]hexa-3,5-diene (THD). The calculated rate constant for formation of THD from 3-TC_S₁ is about 100 times larger than that for the reaction of THD $\rightarrow \alpha$ -TMC, which might be responsible for a minor amount of a species observed experimentally [16]. Finally, it should be pointed out that the concerted N2 loss and rearrangement in the excited state was proposed for the α -TMC formation upon photolysis of 3-TD in Ar or N₂ at 10 K [16], which is not supported by the present CASPT2//CASSCF calculations. The reason for this might come from the matrix effect of Ar or N2 at 10 K, and the related studies are still in progress and will be reported in due time.

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REGULAR ARTICLE

Analysis of a failure of the CC2 coupled-cluster method for bond lengths of SnO and PbO

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Abstract CC2 model is found to overestimate bond lengths of SnO and PbO by about 0.25 Å, while both second-order Møller-Plesset perturbation theory and coupledcluster singles and doubles give reasonable results. Previously, analysis shows that the $[[U, T_1], T_1]$ term in the doubles equation of CC2 is the origin of failure for CC2 and some truncated CC models have been suggested to achieve reasonable result for ozone, where CC2 is unable to obtain a stable structure. However, these remedies are unable to afford reasonable bond lengths of SnO and PbO. Based on a term-wise analysis, our results indicate that the $[U, T_1]$ term results in failure of CC2. CC2 model by removing this term will provide results that agree well with those of MP2. Furthermore, the $[[U, T_2], T_1]$ term absent in the CC2 while present in doubles equation of CCSD can balance this $[U, T_1]$ term and CC2 model augmented with this term is able to afford reasonable results for PbO, SnO and ozone.

Keywords Coupled-cluster theory \cdot CC2 \cdot Unexpected failure

Dedicated to Professor Guosen Yan and published as part of the special collection of articles celebrating his 85th birthday.

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1 Introduction

Coupled-cluster (CC) theory [1–3] is one of the most commonly used non-variational methods in quantum chemistry which shows good performance on correlation energies and properties for single-reference systems. Accuracy of results improves rapidly within the hierarchy of truncated coupled-cluster methods: CCS, CCSD [4] and CCSDT [5]. However, computational costs also increase rapidly and scalings of CCS, CCSD and CCSDT are N⁴, N⁶ and N⁸, respectively, where N represents system size. On the other hand, some intermediate models like CC2 [6], CCSD(T) [7] and CC3 [8], which scale as N⁵, non-iteratively N⁷ and iteratively N⁷, respectively, have been developed to achieve results with similar accuracy to those of the more expensive CC methods.

The CC2 model was developed by Christiansen et al. [6] as an approximation to the CCSD method and is mainly used for excited states [9-14]. Some benchmark calculations indicate that excitation energies with CC2 are even in better agreement with those high-level approaches such as CC3 and CCSDR(3) [15] than those of CCSD especially for valence-type excitations of single-reference molecules [16–19]. Due to its success and relatively low computational cost, various variants of CC2 by employing resolution of identity (RI) [20, 21], density fitting [10, 11], spincomponent scaling [22, 23], localized orbitals [20, 21, 24], reduction of virtual spaces [25], etc. have been proposed to reduce computational effort even further. In addition, CC2 with a scaling of N⁴ based on tensor hypercontraction [26, 27], scaled opposite-spin [28], pair natural orbital [29, 30] or Laplace transformation [24, 31] has also been developed. These developments enable applications of CC2 to groundand excited-state energies of large molecules. Besides energies, analytical gradients [32-34] and Hessian [35] as well as properties for ground and excited states based on CC2 have also been implemented.

CC2 can be understood as a generalization of secondorder Møller-Plesset perturbation theory with an iterative treatment on single excitations. Equation for single excitation amplitudes in CC2 is the same as that in CCSD, and it is expected to provide results with similar accuracy to MP2. Recently, we calculated bond lengths and harmonic frequencies of some closed-shell diatomic heavy-element compounds with CC approaches and MP2 [36]. CC2 is found to perform poorly for PbO and SnO, and their bond lengths are calculated to be about 0.25 Å longer than those with CCSD and MP2 as well as experimental data. Bond lengths for monoxides of this group by various methods as well as experimental data [37] are listed in Table 1, and it can be seen from this table that bond lengths for CO and SiO with CC2 are somewhat larger than those of MP2, while performance of CC2 deteriorates for heavy-element compounds. It has been noticed that CC2 is sensitive to strong electron correlation effects [22]. In fact, strong correlation does not necessarily mean large deviation between CC2 and MP2 according to our results [36]. Bond lengths of AuH and Au2 with CC2 agree reasonably with those of MP2, although electron correlation reduces bond lengths of these two molecules by about 0.1 Å at MP2 level [36]. One may argue that CC2 is mainly developed to calculate excitation energies. Pilot calculations indeed show that transition energies to the lowest several excited states of PbO at experimental bond length with CC2 agree reasonably with those based on equation-of-motion CCSD (EOM-CCSD) [38] and differences are <0.1 eV. However, equilibrium bond length for the lowest spin-singlet excited state with CC2 is still about 0.15 Å longer than that with EOM-CCSD and EOM-CCSDT. This shows poor performance of CC2 on the ground state structure will inevitably affect its performance on structures of excited states. It should be noted that bond lengths with CCSD(T) for SnO and PbO in Table 1 are overestimated by about 0.08 Å, which is due to the employed basis set as well as the frozen inner-shell electrons.

Recently, Pabst et al. [39] reported that CC2 leads to a severe failure in calculation of the equilibrium structure and vibrational frequencies of ozone where MP2 and CCSD provide reasonable results. They carried out a term-wise analysis on CC2 equations and found that the $[[H, T_1], T_1]$ term in the doubles equation was the origin of this failure. Unfortunately, our calculations indicate that modified CC2 approaches suggested by them that could achieve reasonable structure for ozone fail for SnO and PbO. In this work, a term-wise analysis of CC2 on potential energy surface of PbO is performed following closely the work by Pabst et al. An alternative remedy is suggested to achieve reasonable results for bond lengths of PbO and SnO.

Theor Chem Acc (2014) 133:1579

Table 1 Bond lengths (in Å) for monoxides of group IV

	0			8 1			
	MP2	CC2	CCSD	CCSD(T)	Exp. ^a		
СО	1.138	1.149	1.129	1.136	1.128		
SiO	1.534	1.563	1.515	1.527	1.510		
GeO	1.658	1.726	1.634	1.649	1.625		
SnO	1.908	2.148	1.875	1.900	1.833		
PbO	2.027	2.247	1.952	2.015	1.922		

^a Ref. [37]

2 Theory

CC2 is an approximation to CCSD and we will start from CCSD equations. Total energy in CCSD can be written as the following:

$$E_{\text{CCSD}} = \langle \Phi_0 | H \exp(T_1 + T_2) | \Phi_0 \rangle, \tag{1}$$

where *H* is the Hamiltonian, Φ_0 is the HF wavefunction of the corresponding *H*, T_1 and T_2 are single and double excitation operator, respectively, and are defined as the following

$$T_1 = \sum_{i,a} t_i^a a_a^+ a_i, \quad T_2 = \frac{1}{4} \sum_{a,b,i,j} t_{ij}^{ab} a_a^+ a_i a_b^+ a_j, \tag{2}$$

where i, j, ..., (a, b, ...) are indices for occupied (virtual) spin orbitals in Φ_0 . The cluster amplitudes t_i^a and t_{ij}^{ab} in T_1 and T_2 are determined based on the following equations in CCSD:

$$\left\langle \Phi_{i}^{a} \right| \exp\left(-T_{1} - T_{2}\right) H \exp\left(T_{1} + T_{2}\right) |\Phi_{0} \right\rangle = 0,$$
 (3)

$$\left\langle \Phi_{ij}^{ab} | \exp\left(-T_1 - T_2\right) H \exp\left(T_1 + T_2\right) | \Phi_0 \right\rangle = 0,$$
 (4)

where Φ_i^a and Φ_{ij}^{ab} are singly and doubly excited determinant with respect to Φ_0 , respectively. The Hamiltonian *H* can be divided into the one-electron Fock operator *F* and the two-electron perturbation *U*: H = F + U, where *F* is treated as zeroth order and *U* as first order. The *F* operator will be diagonal when canonical molecular orbitals (MO) are used. With this division and canonical MO, the above equations can be written as

$$E_{\text{CCSD}} = \langle \Phi_0 | H + [H, T_2] + 1/2[[H, T_1], T_1] | \Phi_0 \rangle, \qquad (5)$$

$$\langle \Phi_i^a | H + [F, T_1] + [U, T_1] + [H, T_2] + 1/2[[H, T_1], T_1] + [[H, T_2], T_1] + 1/6[[[H, T_1]T_1]] \Phi_0 \rangle = 0,$$
(6)

$$\left\langle \Phi_{ij}^{ab} \middle| U + [F, T_2] + [U, T_1] + 1/2[[H, T_1], T_1] \right. \\ \left. + 1/6[[[H, T_1], T_1], T_1] + 1/24[[[[H, T_1], T_1], T_1], T_1] \right. \\ \left. + [U, T_2] + [[H, T_2], T_1] + 1/2[[[H, T_2], T_1], T_1] \right.$$
(7)

$$\left. + 1/2[[H, T_2], T_2] \middle| \Phi_0 \right\rangle = 0.$$

Based on the Brillouin theorem, it can readily be shown that T_1 is of second order and T_2 is of first order in U. However, orbital relaxation is described mainly by T_1 and it will be of zeroth order if non-HF wavefunction is employed. In CC2 approach, T_1 is set as zeroth order in U and T_2 is approximated to be correct through first order. Total energy in CC2 is thus correct to second order in U. Total energy and singles equation in CC2 are the same as those in CCSD in Eqs. (5) and (6), while the following doubles equation is employed in CC2:

$$\left\langle \Phi_{ij}^{ab} \middle| U + [F, T_2] + [U, T_1] + 1/2[[H, T_1], T_1] \right. \\ \left. + 1/6[[[H, T_1]T_1]T_1] \right.$$

$$+ 1/24[[[[H, T_1], T_1]T_1]T_1]] \Phi_0 \right\rangle = 0.$$

$$(8)$$

By comparing Eqs. (7) and (8), it can be seen that the last four terms in the doubles equation of CCSD are neglected. In the present work, CC2 was implemented by calculating each term in Eqs. (5, 6) and (8) separately through an interface to the CFOUR program package [40] so that a term-wise analysis on CC2 equation can be carried out. Furthermore, the last four terms in Eq. (7) were also implemented in a term-wise way.

3 Results and discussion

All calculations in this paper were performed by the CFOUR program package. The energy-consistent pseudopotential developed by the Stuttgart/Cologne groups, i.e., ECP28MDF and ECP60MDF [41] are employed for Sn and Pb and the corresponding cc-pVTZ-PP basis set [42] is adopted. For C, Si, Ge and O, the cc-pVTZ basis set is used [43]. $4s^24p^64d^{10}$ electrons of Sn, $5s^25p^65d^{10}$ electrons of Pb and $1s^2$ electrons of O are kept frozen in correlation calculations. Equilibrium bond length of SnO and PbO was obtained by fitting total energies at five points with a space of 0.02 Å around the equilibrium bond length with a polynomial function up to fourth order.

Total energies of PbO, norms of the T_1 - and T_2 -amplitude vectors ($||T_1||$ and $||T_2||$) at different bond lengths with MP2, CC2, CCSD and CCSDT are demonstrated in Fig. 1. It can be seen from this figure that total energies with CC2 are much too low compared with those of MP2, CCSD and CCSDT. In addition, underestimation of CC2 energies with respect to the other approaches is more pronounced at larger bond length which results in a too long bond length with CC2. Significant underestimation of total energies with CC2 is accompanied by large $||T_1||$ and $||T_2||$ of CC2. $||T_1||$ and $||T_2||$ with CCSD and CCSDT increase slightly with bond length while those with CC2 grow much more rapidly.

Performance of CC2 on PbO is similar to that on ozone as in Ref. [39], where $||T_1||$ and $||T_2||$ are overestimated and



Fig. 1 Potential energy curve of PbO with MP2, CC2, CCSD, CCSDT as well as $||T_1||$ and $||T_2||$ for CC2, CCSD and CCSDT at different bond lengths

the total energy is underestimated significantly by CC2. After a term-wise analysis, the $[[U, T_1], T_1]$ term in doubles equation is identified as the main origin of failure of CC2 on ozone. The following truncated CC2 models have been suggested to achieve reasonable results for structure of ozone: (1) CC2-noT₁q: deleting the $[[U, T_1], T_1]$ term in Eqs. (5–7), i.e., the third term on r.h.s. of Eq. (5) and the fourth term on l.h.s. of Eqs. (6) and (7): (2) CC2-T₁l(μ_2): Total energy and singles equation are the same as those in CC2, while only the linear terms in T_1 and T_2 , i.e., the first three terms on l.h.s. of Eq. (7) are retained in doubles equation.

These truncated CC2 models are applied to potential energy curve of PbO and the results together with $||T_1||$ and $||T_2||$ of these models are illustrated in Fig. 2. It can be seen from this figure that none of these two models is able to provide a reasonable equilibrium bond length of PbO. Energies



Fig. 2 Potential energy curve of PbO as well as $||T_1||$ and $||T_2||$ for CC2, CC2-T₁lu2, CC2-noT₁q, CC2T2S and CCSD with respect to bond lengths

of CC2-T₁l(μ_2) are slightly higher than those with CC2, while CC2-noT₁q gives rise to energies that are even lower. Performance of these models on $||T_1||$ and $||T_2||$ is consistent with their performance on energies, where large $||T_1||$ and $||T_2||$ correspond to lower energies. Furthermore, it is difficult to achieve converged results at large bond length with CC2noT₁q. One may suspect that this is because electron correlation effect is more pronounced in PbO than that in ozone. However, as we have shown in our previous work [36] that large correlation effect does not necessarily mean failure of CC2. Electron correlation reduces bond length of Au₂ by about 0.17 Å at MP2 level, while bond length with CC2 is only about 0.005 Å shorter than that with MP2. According to our calculations, the largest absolute T_1 amplitude for ozone at CCSD level is 0.073 and it is indeed smaller than that for PbO, i.e., 0.103. On the other hand, the largest T_2 amplitude for ozone at CCSD level is 0.213, while it is only 0.077 for PbO. This indicates that the origin of failure of CC2 for ozone may be different from that for PbO.



Fig. 3 Potential energy curve of PbO as well as $||T_1||$ and $||T_2||$ for MP2, CC2, CCSD, CC2-noT₁, CC2-p1, CC2-p2, CC2-p3, CC2-p4

One can see from these results that the failure of CC2 on PbO is closely related to the abnormally large $||T_1||$ and $||T_2||$. T_1 is treated as zeroth order in CC2, and the maximum absolute T_1 amplitude is indeed larger than the maximum absolute T_2 amplitude. However, contribution to correlation energy from T_1 is smaller than that of T_2 by two orders of magnitude. This indicates that failure of CC2 should mainly come from the too large $||T_2||$, instead of $||T_1||$. However, it is still unclear whether this overestimation of $||T_2||$ stems from large $||T_1||$ or from some unbalanced terms in the doubles equation of CC2. To clarify this point, the converged T_1 amplitudes from CCSD calculation are plugged into the doubles equation of CC2, and this method is termed as CC2T₂S. Results of CC2T₂S are also plotted in Fig. 2. It can be seen from this figure that $||T_2||$ with CC2T₂S is indeed much smaller. Unfortunately, bond length with CC2T₂S is still too long, which means $||T_2||$ from CC2 equation even with reasonable T_1 amplitudes is still overestimated, especially at long bond length range. This implies some unbalanced term in the doubles equation should be the origin of failure of CC2.

Table 2Bond lengths (in Å) ofSnO and PbO

Method	MP2	CC2	CC2-noT ₁	CC2-p2	CCSD	CCSDT
SnO	1.908	2.148	1.903	1.922	1.875	1.898
PbO	2.027	2.247	2.014	2.091	1.952	2.004

The fourth term in the doubles equation in CC2, i.e., Eq. (8), was identified as the origin of failure of CC2 previously. However, removing this term could not afford improved bond length for PbO. The last two terms should only have minor contribution to T_2 amplitudes since they are cubic and quartic terms in T_1 , respectively. The first two terms on l.h.s. of Eq. (8) give rise to MP2 results. Besides these two terms, the most important contribution to T_2 amplitudes should thus be the $[U, T_1]$ term. To investigate importance of this term, we proposed the truncated CC2 noT_1 model where this $[U, T_1]$ term is removed from the doubles equation of CC2. Results of this model are plotted in Fig. 3 together with those of MP2 and CCSD, and one can see from this figure that $CC2-noT_1$ is able to achieve reasonable bond length for PbO. Furthermore, energies with CC2-noT₁ are in close agreement with those of MP2 and $||T_1||$ and $||T_2||$ of CC2-noT₁ are even smaller than those with CCSD. The $[U, T_1]$ term is the most important coupling term between T_1 and T_2 operators in doubles equation, and omitting this term in CC2 will thus give rise to results that closely resemble those of MP2. Bond lengths of PbO and SnO with CC2-noT₁ are listed in Table 2 together with results of MP2, CCSD and CCSDT. It can be seen from this table that bond length with $CC2-noT_1$ is about 0.01 Å shorter than that with MP2. CC2 has been developed mainly for excitation energies, and the $[U, T_1]$ term is critical to couple single and double excitations. Excitation energies based on CC2-noT1 formulation will give rise to excitation energies of similar accuracy to that of CIS and CC2-noT1 should thus not be used for excitation energies.

Our result indicates that failure of CC2 is due to the unbalanced $[U, T_1]$ term in doubles equation. Considering that CCSD is able to provide reasonable bond length of PbO, the $[U, T_1]$ term must be balanced by one or any combination of the addition four terms in doubles equation of CCSD: $[U, T_2]$, $[[U, T_2], T_1]$, $[[U, T_2], T_2]$ and $[[[U, T_2], T_2]]$ T_2], T_1], T_1]. To further investigate this point, we proposed four models: CC2-pn with n = 1,2,3,4, and CC2-pn means the CC2 model with the additional n-th term from these four additional members included in the doubles equation. Results of these four models are demonstrated in Fig. 3. One can see from this figure that energies with CC2-p1 are rather close to those of $CC2-noT_1$ for bond lengths below 2.10 Å, but performance of this model deteriorate at longer bond lengths. On the other hand, energies of CC2p3 are only slightly higher than those of CC2 and converged results cannot be obtained with the CC2-p4 model when bond length exceeds 2.0 Å. It is interesting to note that although energies with CC2-p4 are even lower than those of CC2, $||T_1||$ and $||T_2||$ with this model are smaller than those with CC2. According to this figure, the CC2-p2 model improves upon CC2 model to a large extent and reasonable bond length of PbO can be achieved with this model. These results show that the $[U, T_1]$ term in doubles equation which result in large error on bond length of PbO in CC2 can somehow be balanced by the $[[U, T_2], T_1]$ term in CCSD. Bond lengths of PbO and SnO with CC2-p2 are



Fig. 4 Potential energy curve as well as $||T_1||$ and $||T_2||$ of ozone with MP2, CC2, CCSD, CC2-noT₁q, CC2-T₁l(μ_2), CC2-noT₁ and CC2-p2

Table 3 Bond lengths (in Å) of Bi₂, TlH, Sb₂ and InH

	MP2	CC2	CC2-noT1	CC2-p2	CCSD	CCSD(T)
InH	1.860	1.861	1.860	1.864	1.871	1.873
TlH	1.926	1.927	1.927	1.926	1.944	1.946
Sb_2	2.559	2.585	2.558	2.532	2.506	2.532
Bi ₂	2.733	2.765	2.732	2.696	2.661	2.691

also listed in Table 2. Compared with results of MP2, CC2p2 overestimates this bond length by about 0.06 Å for PbO and 0.01 Å for SnO.

To further demonstrate the reliability of the CC2-noT₁ and CC2-p2 models, potential energy curve of ozone is calculated with these two models and results are plotted in Fig. 4 together with those given by CC2-noT₁q, CC2-T₁l(μ_2), CC2, MP2 and CCSD. It can be seen from this figure that CC2-noT₁q and CC2-T₁l(μ_2) models yield reasonable result for ozone, which is consistent with results in Ref. [39]. On the other hand, results of the two models suggested in this work also agree qualitatively with those of MP2 and CCSD. Unsurprisingly, energies with CC2-noT₁ are in good agreement with those of MP2.

Bond lengths of InH, TlH, Sb₂ and Bi₂ have also been calculated with CC2-noT1, CC2-p2, MP2, CC2, CCSD and CCSD(T), and results are listed in Table 3. Basis set and ECPs employed in these calculations are the same as those for SnO and PbO. Electron correlation effect for InH and TIH is insignificant where bond lengths with CC2 for these two molecules are in good agreement with those with MP2. On the other hand, electron correlation effect somewhat more pronounced for Sb₂ and Bi₂ and CC2 bond lengths are about 0.03 Å larger than that of MP2. Bond lengths with CC2-noT₁ closely resemble those with MP2 as expected and differences are about 0.001 Å. On the other hand, results with CC2-p2 for InH and TlH are in good agreement with those of CC2, while they are shorter than those with CC2 and are in better agreement with CCSD and CCSD(T) results.

4 Conclusion

CC2 model fails to achieve equilibrium structure for ozone, and bond lengths with CC2 are overestimated by about 0.25 Å for PbO and SnO. In a previous analysis, the $[[U, T_1], T_1]$ term in the doubles equation is identified as the origin of the failure of CC2 and some truncated CC2 models have been proposed to provide reasonable results for ozone. Unfortunately, these models are unable to give rise to reasonable bond length for PbO. Based on a termwise analysis of CC2, we show that the $[U, T_1]$ term in the doubles equation result in this failure of CC2. This term is the most important term that couples T_1 amplitudes with

 T_2 amplitudes and truncated CC2 model by removing this term from the doubles equation provide results that are in close agreement with those of MP2. Furthermore, we also found that the $[[U, T_2], T_1]$ term that is absent in CC2 equation can balance this $[U, T_1]$ term. CC2 model augmented with this term in the doubles equation is able to afford reasonable bond lengths of PbO and SnO as well as equilibrium structure for ozone.

It should be noted that the computational scaling of the $[[U, T_2], T_1]$ term is N⁶, and performance of CC2-noT₁ closely resembles that of MP2. The purpose of this work is to provide an understanding on failure of CC2, instead of proposing some improved models. On the other hand, CC2 is developed mainly to calculate excitation energies, while MP2 should be preferred to calculate ground state energy. Indeed, excitation energies for PbO at equilibrium bond length of PbO with CC2 are in good agreement with those of EOM-CCSD. However, its poor performance on ground-state structure leads to its significant overestimation of excited state bond length. Reasonable energy for excited state may be obtained by summation of excitation energy with CC2 and ground state energy from MP2. The suggested approach is similar to CIS(D) [44] and ADC(2) [45], where the MP2 ground state energies are employed. However, it is well known that CC2 usually provides better excitation energies than CIS(D) and ADC(2) and the suggested approach may still outperform CIS(D), ADC(2) and even CC2 itself in determining excited state structures.

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REGULAR ARTICLE

Electrostatic potentials of camptothecin and its analogues

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Abstract In order to understand the influences of the modification of E-ring on the main frame of camptothecin (CPT), studies of the E-ring modification resulted changes in the stability and the electrostatic potential around the main frame of CPT were performed by the density functional theory. The results of present study indicate that the stability of the close-ring lactone form of CPT and homocamptothecin (hCPT) is similar to their open-ring hydroxylate forms, especially when in aqueous solutions. As an E-ring-modified CPT analogue, hCPT has essentially the same electrostatic potential (ESP) as CPT around the main frame (from A- to D-ring). However, the electrostatic potentials of the open-ring compounds are more negative around the main frame than that of CPT. The changes in the ESP of the CPT derivatives are found to be correlated with the corresponding dipole moments. Since electrostatic potential could influence the $\pi - \pi$ stacking pattern between CPT (and its analogues) and the DNA bases, present study

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suggests that this $\pi - \pi$ interaction of the open-ring forms of CPT and hCPT might be different from that of the closering lactone form. The information revealed in this study sheds light on the developments of new CPT-type antitumor drugs.

Keywords Electrostatic potentials \cdot Camptothecin $\cdot \pi - \pi$ Interaction

1 Introduction

Camptothecin (CPT, see Fig. 1) and its analogues are antitumor drugs, which strongly inhibit nucleic acid synthesis and initiate DNA strand breaks [1–4]. Structure-activity studies indicate that their activity is due to an action on topoisomerase I [5, 6].

CPT and its analogues are found to intercalate between DNA bases of both strands at the enzyme-induced nick in the eukaryotic DNA topoisomerase I. The binding pattern consists of stacking interactions with the bases of DNA, H-bonding with residual Asp, and water-bridged H-bonding interactions with the active site phosphotyrosine and Asn [7]. Based on the crystal structure of the topoisomerase I and DNA complex, Kerrigan and Pilch predicted that the closed lactone form (E-ring) of CPT might also bind to residuals Arg and Asp of the topoisomerase and guanine and ribose moieties of DNA through H-bonding [8]. Molecular modeling studies suggested that both stacking and H-bonding interactions played important roles in stabilizing the CPT-topoisomerase I–DNA complex [9].

The lactone moiety (a six-membered α -hydroxylactone) within CPT can be hydrolyzed at physiological condition, forming the open-ring carboxylate structure [2]. Compared with the closed lactone form, the open carboxylate form of

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Fig. 1 The optimized structures of the lactone form of CPT and its hydrolyzed open-ring hydroxylate forms. Calculated at the B3LYP/6-311G(d,p) level of theory. Numbering scheme is in black. The values in parenthesis were evaluated at the M06-2X/6-311G(d,p) level of theory. Color representations: red for oxygen, gray for carbon, blue for nitrogen, and white for hydrogen. Atomic distances are in Å. Overlay-1 is the overlay structures of CPT, CPT-1, and CPT-2. Hydrogen atoms have been omitted in Overlay-1 for clarity



CPT (and its analogues) provides more H-bonding sites. Therefore, the open carboxylate form is expected to bind more tightly to the topoisomerase I. Furthermore, opening of the E-ring may influence the electrostatic potential around main frame of CPT.

As an E-ring-modified CPT, homocamptothecin (hCPT, see Fig. 2) bears a seven-membered β -hydroxylactone ring. This E-ring-expanded compound exhibits enhanced lactone stability due to the reduced electrophilicity of a β -hydroxylactone [10–12]. However, there are still 20 % of hCPT adopt the open-ring form [12]. Moreover, the expanded E-ring in hCPT provides more H-bonding sites than that of CPT.

Theoretical study at different levels of theory suggested that the alteration of the electrostatic potential might affect the π - π stacking with the DNA bases [13]. Since the orientation of the π - π stacking between the main frame of CPT and the DNA bases could be a main factor contributing to the stability of the CPTtopoisomerase I–DNA complex [14], the knowledge of the influences of the modification of E-ring on the main frame of CPT is crucial for improving the selectivity of CPT-related compounds. Here, we report the density functional theory study of the influences of the E-ring modifications on the electrostatic potential of the main frame of CPT. The information revealed in this study sheds light on the developments of new CPT-type antitumor drugs.

2 Methods

The B3LYP [15, 16] and M06-2X [17] functionals with basis sets of triple-ζ quality plus polarization functions (denoted 6-311G(d,p)) [18, 19] were used to determine the local minimal geometries, energetics, and vibrational frequencies of CPT and hCPT (and their corresponding open-ring forms as well). For better description of longrange interactions, diffuse functions were also included (6-311 + G(d,p)) in the determination of the electrostatic potentials. These compounds were also been optimized using the polarizable continuum model (PCM) [20] with a dielectric constant of water ($\varepsilon = 78.39$) in order to evaluate the energies of the systems in a polarizable medium. It should be noted that this PCM model used in the present study only accounts for the influences of the polarizable surroundings; the important effects of the micro-hydration could not be included in this approach. Therefore, the PCM model only to some extent approximates the real situation of aqueous solution. The GAUSSIAN 09 program [21] was used for all computations.



Fig. 2 The optimized structures of the lactone form of hCPT and its hydrolyzed open-ring hydroxylate forms. Calculated at the B3LYP/6-311G(d,p) level of theory. The values in parenthesis were evaluated at the M06-2X/6-311G(d,p) level of theory. Color representations: red

3 Results and discussions

3.1 Geometry and energy properties

Both B3LYP and M06-2X functionals predict the similar geometries for all the species. The main difference in the geometric parameters is that the intra-molecular H-bond length predicted by the M06-2X functional is 0.05 Å longer than that predicted by the B3LYP approach at most. Moreover, the geometries of the CPT and hCPT derivatives optimized with the PCM model are also close to the corresponding compounds optimized in the gas phase. Therefore, the discussion below is mainly based on the results obtained at the B3LYP/6-311G(d,p) level of theory in the

for oxygen, gray for carbon, blue for nitrogen, and white for hydrogen. Atomic distances are in Å. Overlay-2 is the overlay structures of CPT, hCPT, and hCPT-1. Hydrogen atoms have been omitted in Overlay-2 for clarity

gas phase. The optimized geometries of CPT and the corresponding open-ring compounds (CPT-1 and CPT-2, see Fig. 1) reveal that the geometric variations due to the E-ring opening mainly limited on the E-ring. In general, the C–C bond distances of the E-ring moiety in the open-ring forms increase slightly as compared to those in the lactone form. Specifically, C15–C20 distance increases 0.03 Å in CPT-1 (1.55 Å) and 0.02 Å in CPT-2 (1.54 Å), and C16–C22 distance elongates 0.02 Å in CPT-1 (1.52 Å) and 0.01 Å in CPT-2 (1.51 Å), respectively, as compared to those in CPT. Moreover, C20–O bond length also increases in the open carboxylate forms (1.42 Å in CPT-1 and 1.44 Å in CPT-2 vs. 1.41 Å in CPT). The main difference between CPT-1 and CPT-2 is that there is an intra-molecular H-bond in the opened E-ring moiety of CPT-1. To form this intra-molecular H-bond, the OH group at the C20 position in CPT-1 has to rotate around C15–C20 bond to the opposite position after the hydration of CPT. Compared with CPT-1, the relative positions of C20, C21, and C22 of the opened E-ring moiety in CPT-2 are close to those of CPT. The geometric feature of CPT-2 indicates that CPT-2 is the direct product of the hydration of CPT. On the other hand, the overlay of the structures of CPT, CPT-1, and CPT-2 indicates that the geometries of the A-, B-, C-, and D-rings of CPT do not change significantly in response to the hydrolysis of the lactone form.

Similarly, the optimized geometries of hCPT and its hydrolysis products (hCPT-1, hCPT-2, hCPT-3, and hCPT-4, Fig. 2) demonstrate that the introduction of CH₂ group at the α -position of the lactone moiety does not lead to geometric variations in the main frame (A-, B-, C-, and D-rings) of CPT (see Overlay-2 in Fig. 2). However, the C-C bond distances of the E-ring moiety increase in the open-ring carboxylate forms as compared to those in the lactone form of hCPT. It is important to note that the intramolecular H-bond in the α-hydroxylactone moiety of CPT does not exist in hCPT. Nevertheless, in the hydrolysis products of hCPT, intra-molecular H-bond can be identified at various sites of the E-ring moiety. In hCPT-1, the intramolecular H-bond through the proton of the β -hydroxyl group and the carbonyl oxygen of carboxylate group forms a six-membered ring structure as can be seen in Fig. 2. For other conformers of the open-ring carboxylate forms of hCPT, the intra-molecular H-bond is found to be involved in the ω -hydroxyl group and in the carboxylate group. The formation of intra-molecular H-bond is expected to be one of the important factors that stabilize the systems.

The relative energies (relative to the close-ring lactone form) summarized in Table 1 reveal that the stability of the open-ring form of CPT is similar to that of the lactone form. The energy of CPT-1 is about 2.5 kcal/mol lower than that of CPT, while the energy of CPT-2 is about 2.5 kcal/mol higher than that of CPT at the B3LYP/6-311G(d,p) level of theory. Therefore, the balance between the open-ring carboxylate form and the close-ring lactone form could be easily shifted in different environments. The higher stability of CPT-1 as compared to CPT-2 should be attributed to the existence of the intra-molecular H-bond in the former.

On the other hand, two of the open-ring forms of hCPT (hCPT-1 and hCPT-2) are considerably more stable than the lactone form of hCPT. The energy of hCPT-1 and hCPT-2 is about 10.1 kcal/mol and 5.7 kcal/mol lower than that of hCPT, respectively. These two structures should dominate the open carboxylate form of hCPT. Although the structures of hCPT-3 and hCPT-4 are similar, the total energy of hCPT-3 is about 3 kcal/mol lower than that of hCPT-4. The stronger intra-molecular H-bondingin hCPT-3 (with O…H distance of 1.92 Å in hCPT-3 vs. 2.02 Å in hCPT-4) should be the main contribution of this energy difference.

Including the polarizable medium slightly raises the stability of the close-ring form of camptothecin. Using the PCM model, the energy variations between the open-ring (CPT-1) and close-ring form of CPT are <0.1 kcal/mol. On the other hand, CPT-2 in the aqueous solutions is approximately 4 kcal/mol less stable than CPT (predicted by the B3LYP functional). Due to the existence of the polarizable surroundings, the energy variations between the open-ring and close-ring form of hCPT are <5.0 kcal/mol. All of the open-ring forms of hCPT derivatives are more stable than the close-ring form with the exception of hCPT-4, which is about 3 kcal/mol less stable than hCPT.

Compared with the B3LYP functional, the M06-2X functional predicts that the open-ring structures are more stable. The relative energies of the open-ring forms of CPT and hCPT evaluated by the M06-2X approach are about

	ΔΕ	ΔE^0	ΔG	$\Delta E(PCM)^{a}$	μ
$CPT + H_2O$	0.0	0.0	0.0	0.0	6.5 (6.5)
CPT-1	-2.49 (-6.73)	1.17 (-3.09)	10.27 (6.04)	-0.06 (-4.13)	6.9 (6.8)
CPT-2	2.47 (-2.86)	5.40 (0.18)	14.70 (9.53)	4.09 (-1.47)	2.1 (2.0)
$hCPT + H_2O$	0.0	0.0	0.0	0.0	7.1 (7.0)
hCPT-1	-10.14 (-12.29)	-6.94 (-9.15)	1.83 (-0.26)	-4.84 (-7.09)	1.4 (1.5)
hCPT-2	-5.69 (-9.14)	-2.50 (-5.92)	6.48 (3.40)	-0.87 (-4.03)	4.4 (4.1)
hCPT-3	-3.091 (-5.82)	0.12 (-2.53)	9.14 (6.64)	-0.02 (-2.72)	4.8 (5.0)
hCPT-4	0.162 (-2.46)	3.45 (0.93)	12.59 (9.94)	3.03 (0.38)	5.0 (5.0)

Table 1 Energy properties (in kcal/mol) and dipole moments (μ , in Debye) of the optimized structures of CPT and hCPT

Calculated at the B3LYP/6-311G(d,p) level of theory. The values in parenthesis were evaluated at the M06-2X/6-311G(d,p) level of theory. ΔE is the energy relative to the total energy of the close-ring lactone form and one water molecule; ΔE^0 is the zero-point-energy corrected ΔE ; ΔG is the free energy difference at 298 K

^a Using the polarizable continuum model with a dielectric constant of water ($\epsilon = 78.39$)



Fig. 3 The electrostatic potential (ESP) map of CPT. The ESP is mapped on the iso-surface of electron density with iso-value of 0.0005. The ESP color ranges from -0.01 (*blue*) to 0.01 (*red*). Unit is in au

2–5 kcal/mol lower than those by the B3LYP method. In the PCM-modeled aqueous solutions, the open-ring derivatives are more stable than the corresponding close-ring compounds in the M06-2X studies (except for hCPT-4, which is 0.4 kcal/mol higher than hCPT in energy). Since all the energy differences between the close-ring lactone form of CPT and hCPT are close to the corresponding open-ring hydroxylate forms, one may expect that both forms exist in aqueous solutions.

3.2 Electrostatic potential and dipole moment

Electrostatic potential (ESP) around the molecular surface is one of the crucial properties of activity of a drug. Changes in electrostatic potential might result in changes in the interaction patterns between a drug and the corresponding reaction site. Moreover, changes in electrostatic potential could influence the transportation process of a drug. The electrostatic potential of CPT mapped on its electron density iso-surface (with iso-value of 0.0005 au) is depicted in Fig. 3. On this iso-surface, negative ESP (colored in blue) regions spread around the oxygen and the nitrogen atoms. Meanwhile, a slightly positive ESP (in orange) area locates on the vicinity of the edge of the carbon side of the B- and C-ring. Electrostatic potential is essentially zero (in green) on the iso-surface of the A-ring moiety of CPT.

As expected, the alteration of the E-ring of CPT could result in dramatic changes in the electrostatic potential around its neighborhood. The electrostatic potential



Fig. 4 The electrostatic potential difference (ESPD) maps of the E-ring-modified compounds of CPT. The ESPD is mapped on the iso-surface of electron density with iso-value of 0.0005. The ESPD color ranges from -0.005 (*blue*) to 0.005 (*red*). Unit is in au

difference (ESPD) between CPT-1 and CPT mapped on the electron density iso-surface in Fig. 4 demonstrates that the change in the electrostatic potential only limited on the isosurface around the E- and D-ring. The electrostatic potential of the main frame from A- to C-ring is effectively unaffected in CPT-1. However, the other form of the open-ring hydroxylate compound, CPT-2, has substantial changes in its electrostatic potential due to the structural variations of the E-ring of CPT. The ESPD map of CPT-2 reveals that the electrostatic potential is notably more negative around the main frame of CPT-2 as compared to that of CPT.

The influence of the E-ring modification on the electrostatic potential is primarily limited on the E-ring moiety of hCPT, as revealed in its ESPD map. Therefore, the binding pattern of stacking interactions with the bases of DNA should be the same for CPT, CPT-1, and hCPT. On the other hand, the ESPs of the hydrolyzed products of hCPT decrease over the main frame, ranging from A- to D-ring, as compared to that of CPT. ESPD maps depicted in Fig. 4 show that the decrease of ESP around the B-, C-, and D-ring moiety of the main frame of hCPT-1 is the most obvious among the four open-ring structures of hCPT. Meanwhile, the ESPD map of hCPT-2 demonstrates moderate decrease in the ESP around the main frame. The reduction of ESP over the A- and B-ring of hCPT-3 and hCPT-4 is less significant as illustrated by the corresponding ESPD maps.

It is interesting to note that the changes in the ESP of the CPT derivatives can be correlated with the corresponding dipole moments. Similar dipole moments are found for CPT, CPT-1, and hCPT (6.5, 6.9, and 7.1 Debye, respectively), which matches the near-zero ESPD around the main frame for the corresponding compounds. Meanwhile, small dipole moments of hCPT-1 (1.4 Debye) and CPT-2 (2.1 Debye) correlate the significant ESP reduce on their iso-surfaces around B-, C-, and D-rings. Moderate dipole moments of hCPT-2 (4.4 Debye), hCPT-3 (4.8 Debye), and hCPT-4 (5.0 Debye) associated with the observable ESP lessen on the C- and D-rings as shown in the ESPD maps.

4 Concluding remarks

The results of present study indicate that the stability of the close-ring lactone form of CPT and hCPT is similar to their open-ring hydroxylate forms, especially when in aqueous solutions. E-ring modification of hCPT essentially does not alter the ESP around the main frame (from A- to D-ring) as compared to that of CPT. However, the electrostatic potential of the open-ring compounds around is more negative

around the main frame than that of CPT. The changes in the ESP of the different structures of the CPT derivatives are found to be correlated with the corresponding dipole moments. Significant decreases of the ESP around the A-, B-, C-, and D-ring of CPT-2 and hCPT-1 are found to be associated with the dramatic reduce of their dipole moments, while minor changes in the ESP over the A-, B-, C-, and D-ring region of CPT, CPT-1, and hCPT can be correlated with their similar dipole moments.

Since electrostatic potential could influence the π - π stacking pattern between CPT (and its analogues) and the DNA bases, present study suggests that this π - π interaction of the open-ring forms of CPT and hCPT might be different from that of the close-ring lactone form.

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REGULAR ARTICLE

A comparison study of the $H + CH_4$ and $H + SiH_4$ reactions with eight-dimensional quantum dynamics: normal mode versus local mode in the reactant molecule vibration

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Abstract While molecular vibration of CH_4 is well described by the normal-mode paradigm, the local mode picture is more suitable for understanding the SiH₄ stretching vibrational motion. To compare the roles of the two types of molecular vibration in reaction dynamics, the $H + CH_4 \rightarrow H_2 + CH_3$ and $H + SiH_4 \rightarrow H_2 + SiH_3$ reactions have been investigated using an eight-dimensional (8D) quantum dynamics method in which the nonreacting XH₃ (X = C, Si) group keeps its C_{3v} symmetry in the reaction. The reaction probabilities, integral cross sections and thermal rate constants in the temperature range of 200– 2,000 K were calculated for both reactions. Strong mode specificity was found in both reactions, and the differences were rationalized by the vibrational characteristics of the CH₄ and SiH₄ reactants.

Keywords Quantum dynamics $\cdot H + CH_4 \cdot H + SiH_4 \cdot$ Normal mode \cdot Local mode

Dedicated to Professor Guosen Yan and published as part of the special collection of articles celebrating his 85th birthday.

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1 Introduction

Based on observations in atom-diatom reactions, Polanyi summarized the role played by various forms of energy in promoting reactivity. For a reaction with an "early" (reactant-like) barrier, translational energy is more effective than vibrational energy. However, for a reaction with a "late" (product-like) barrier, vibrational energy has higher efficacy in promoting reaction [1]. Much effort has been made to test Polanyi's rules in reactions involving polyatomic molecules, both experimentally [2–6] and theoretically [7–10]. The results indicated that Polanyi's rules are mostly upheld, but the capacity of various reactant vibrational modes for promoting reactivity may be different. These studies have stimulated new models to understand mode specificity in chemical reactions [11–15].

The vibration of a polyatomic molecule is traditionally described with the normal-mode paradigm, in which each normal-mode vibration involves a particular form of synchronized atomic motion [16, 17]. For example, the vibration of CH₄ is classified into four normal modes: the symmetric stretching mode (v_1), the asymmetric bending mode (v_2), the asymmetric stretching mode (v_4 , also known as umbrella mode). These synchronized motions involve all atoms in the molecule, and the energy deposited in each vibrational mode can be used to promote reactivity in a reaction. Previous studies have indicated that all excitations of the CH₄ vibration could enhance the H + CH₄ reaction with different levels of efficacy [18–20].

In some molecules, the normal-mode picture is inadequate and a local mode description of molecular vibration may be more suitable. A good example is the SiH₄ molecule, in which the four Si–H bonds vibrate nearly independently due to weak kinetic coupling among these stretching

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modes [21–24]. The ro-vibrational spectrum of SiH₄ has been extensively investigated as a model to understand the local mode vibration by Zhu et al. [25, 26]. The SiH₄ molecule is a spherical top in its vibrational ground state. Upon the Si–H vibrational excitation, however, its rotational spectrum displayed symmetric top characters. The differences observed in the SiH₄ ro-vibrational spectra could be used to assign the stretching excited states, which are better described in the local mode regime. Theoretically, Halonen and Child [23] have developed a theory to calculate the different kinetic and potential couplings and analyzed the conditions under which the local mode vibration could be observed.

As a prototype of polyatomic reactions, the dynamics of the $H + CH_4$ reaction has attracted significant interests over the past decades, both experimentally [2, 27, 28] and theoretically [19, 20, 29-38]. As a full-dimensional quantum dynamics calculation is still difficult to perform, a number of reduced-dimensional models have been proposed. For example, Takayanagi [38] reported a three-dimensional study by approximating $H + CH_4$ with a collinear fouratomic system. Yu and Nyman [37] included the umbrella mode in their four-dimensional treatment within the rotating bond approximation (RBA). Wang and Bowman [36] performed a six-dimensional calculation by approximating CH₄ with a triatomic molecule. A semi-rigid vibrating rotor target (SVRT) model proposed by Zhang has been applied to this reaction in which the reacting polyatomic molecule CH_4 is treated as a semi-rigid vibrating rotor [35]. Last but not least, Palma and Clary [39] have suggested a quantum mechanics model for the $X + YCZ_3$ reactions, in which the nonreacting CZ_3 group is restricted to C_{3v} symmetry. Based on this model, Yang and Zhang [20] reported a seven-dimensional initial state-selected time-dependent wave packet study, and Zhang et al. [32] have carried out an eight-dimensional transition-state wave packet study of $H + CH_4$ reaction. Very recently, Manthe et al. [31] studied initial state-selected reaction probabilities for this reaction with a full-dimension MCTDH (multi-configuration timedependent Hartree) method, and Zhang et al. reported a sixdimensional state-to-state quantum dynamics study of this reaction [29].

Experimentally, the dynamics of the F + SiH₄ [40, 41] and Cl + SiH₄ [42, 43] reactions have been studied recently by Yang et al. However, no experimental study of the H + SiH₄ reaction has been reported. Theoretically, Espinosa-Garcia et al. [44] have developed an analytic potential energy surface (PES) for the H + SiH₄ reaction. Based on this PES, rate constants and kinetic isotope effects were studied using variational transition-state theory by the same authors [44] and later using a semi-classical quantum instanton method by Zhao et al. [45]. Recently, Bian et al. constructed a global twelve-dimensional ab initio PES for the $H + SiH_4$ reaction [46] and studied the dynamics with quasi-classical trajectory calculations [47, 48] and the kinetics with variational transition-state theory [49].

In this work, a reduced-dimensional quantum dynamics study of the $H + SiH_4$ reaction is performed to explore the role of local mode in the reaction dynamics, using the PES of Espinosa-Garcia et al. [44] (denoted as SiH5-EGSC-1998). For comparison, the quantum dynamics calculations for the $H + CH_4$ reaction have also been carried out on a PES also developed by Espinosa-Garcia [50] (denoted as CH5-EG-2002). This PES is completely symmetric with respect to the permutation of the four CH₄ hydrogen atoms and has been carefully calibrated [50]. This permutation symmetry is absent in the PES developed by Jordan and Gilbert [51] (denoted as JG-PES), and the calculated potential energy is dependent on the order of four hydrogen atoms, although the JG-PES has been used in most previous quantum dynamics studies. The SiH5-EGSC-1998 and CH5-EG-2002 PESs have recently been employed to calculate the thermal rate constants using variational transition-state theory [44, 50] and semi-classical methods [45, 52], thus providing theoretical data to compare with. For these reasons, these PESs are selected in this work, although several more accurate PESs exist [53-55]. For the quantum dynamical calculations, a recently developed eight-dimensional (8D) quantum mechanical (QM) Hamiltonian is employed [19]. This 8D QM model is similar to the pioneering work of Palma and Clary [39] for the $X + YCZ_3 \rightarrow XY + CZ_3$ reaction and has recently been applied to the $H/O(^{3}P) + CH_{4}$ reactions [9, 19] and the CH_{4} dissociative chemisorption on Ni(111) [8], and the results were in good agreement with experimental observations.

The paper is organized as follows. In Sect. 2, the 8D QM method is briefly described. The reaction probabilities, integral cross sections and thermal rate constants for both reactions are presented in Sect. 3. This is followed by a discussion of these results in Sect. 4. Finally, conclusions are given in Sect. 5.

2 Methodology

90

2.1 The coordinate system

The eight-dimensional model for the X + YCZ₃ \rightarrow XY + CZ₃ reaction is described in a Jacobi coordinate system as shown in Fig. 1: *R* is the vector from the center of mass of YCZ₃ to X; *r* is the vector from the center of mass of CZ₃ to Y. The CZ₃ group could be defined with polar coordinates (ρ , χ) [32, 55] or Cartesian coordinates (x, s) [39, 56]. Here, ρ is the bond length of CZ, and χ is the angle between a CZ bond and vector *S*, which is the symmetry axis of CZ₃; x denotes the distance between atom Z



Fig. 1 a Jacobi coordinate system of the $X + YCZ_3$ model; b Cartesian coordinates and polar coordinates of CZ_3 group

and the symmetry axis *S* and *s* the distance between atom C and the center of three Z atoms. θ_1 is the bending angle between the vectors *R* and *r* and φ_1 is the azimuthal angle of the rotation of YCZ₃ around the vector *r*; θ_2 is the bending angle between vectors *r* and *S* and φ_2 is the azimuthal angle of the rotation of CZ₃ around the vector *S*.

Four frames were introduced to describe the angular coordinates and rotation of the system: the space-fixed frame, the body-fixed frame (XYCZ₃-fixed frame), the YCZ₃-fixed frame and the CZ₃-fixed frame. The *z*-axis of the body-fixed frame lies along the vector \mathbf{R} , and the vector \mathbf{r} is always in the *xz*-plane of the frame. The *z*-axis of the YCZ₃-fixed frame lies along the vector \mathbf{r} , and the vector \mathbf{S} is always in the *xz*-plane of the frame. The *z*-axis of the CZ₃-fixed frame lies along its symmetry axis, the vector \mathbf{S} , and the first Z atom is always in the *xz*-plane of the frame. The four frames form three pairs of related space and body-fixed frames.

2.2 The model Hamiltonian

By restricting the nonreacting CZ_3 group in C_{3v} symmetry, the eight-dimensional model Hamiltonian for $XYCZ_3$ system is given by

$$\hat{H} = -\frac{1}{2\mu_{\rm R}}\frac{\partial^2}{\partial R^2} - \frac{1}{2\mu_{\rm r}}\frac{\partial^2}{\partial r^2} + \frac{(\hat{J}_{\rm tot} - \hat{J})^2}{2\mu_{\rm R}R^2} + \frac{\hat{l}^2}{2\mu_{\rm r}r^2} + \hat{K}_{\rm CZ_3}^{\rm vib} + \hat{K}_{\rm CZ_3}^{\rm vib} + V(R, r, x, s, \theta_1, \varphi_1, \theta_2, \varphi_2),$$
(1)

where $\mu_{\rm R}$ is the reduced mass of the X + YCZ₃ system and $\mu_{\rm r}$ is the reduced mass of Y + CZ₃. The first two terms are the kinetic energy operators for *R* and *r*, respectively; $\hat{J}_{\rm tot}$ is the total angular momentum operator of the system. \hat{J} is the rotational angular momentum operator of atom Y with respect to CZ₃. $\hat{K}_{\rm CZ_3}^{\rm vib}$ and $\hat{K}_{\rm CZ_3}^{\rm rot}$ are the vibrational and rotational kinetic energy operators of CZ₃, respectively. Due to the symmetry requirement and definition of the CZ₃-fixed frame, no vibration–rotation coupling exists in this eight-dimensional Hamiltonian. The last term $V(R, r, x, s, \theta_1, \varphi_1, \theta_2, \varphi_2)$ in Hamiltonian is the potential energy.

The vibrational and rotational kinetic operators of the CZ₃ group could be expressed with a new scaledpolar coordinate system (q, γ) , with $q = \sqrt{x^2 + y^2}$ and $\gamma = \arctan(-x/y)$, here $y = \sqrt{\frac{m_C}{m_C + 3m_Z}}s$. The explicit form of the operators is

$$\hat{K}_{\text{CZ}_3}^{\text{vib}} = \frac{1}{2\mu_x} \left(\frac{\partial^2}{\partial q^2} + \frac{1}{q^2} \frac{\partial^2}{\partial \gamma^2} + \frac{1}{4q^2} \right), \tag{2}$$

with $\mu_x = 3m_Z$ and the integral element as $\partial q \partial \gamma$, and

$$\hat{K}_{CZ_3}^{\text{rot}} = \frac{1}{2I_A}\hat{j}^2 + \left(\frac{1}{2I_C} + \frac{1}{2I_A}\right)\hat{j}_z^2,\tag{3}$$

with \hat{j} as the rotational angular momentum of CZ₃ and \hat{j}_z its z-component. The rotational inertia I_A and I_C are defined as [39],

$$I_{\rm A} = \frac{3}{2} m_{\rm Z} \left[x^2 + \frac{2m_{\rm C}}{m_{\rm C} + 3m_{\rm Z}} s^2 \right],\tag{4}$$

$$I_{\rm C} = 3m_Z x^2. \tag{5}$$

2.3 Wavefunction and reaction flux

For the case of $J_{\text{tot}} = 0$, the wave packet is expanded as a linear combination of the product of the basis sets for *R*, *r*, *q*, γ and the parity-adapted rotational basis functions.

$$\Psi^{\varepsilon} = \sum_{n_{Z}, n_{r}, n_{q}, n_{\gamma}} \sum_{Jljk} c^{\varepsilon}_{n_{Z}n_{r}n_{q}n_{\gamma}Jljk}(t) G_{n_{R}}(R) F_{n_{r}}(r)$$

$$Q_{n_{q}}(q) H_{n_{\gamma}}(\gamma) \Phi^{\varepsilon}_{Jljk}(\hat{r}, \hat{S}), \qquad (6)$$

k is the projection of *j* onto the C_{3v} symmetry axis *S*, and ε is the parity under space inversion. The parity-adapted rotational basis function is expressed as

$$\Phi_{Jljk}^{\varepsilon}(\hat{r},\hat{S}) = \sqrt{\frac{1}{2(1+\delta_{k0})}} \times \left[\Phi_{Jljk}(\hat{r},\hat{S}) + \varepsilon(-1)^{J+l+j+k} \Phi_{Jlj-k}(\hat{r},\hat{S}) \right],$$
(7)

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with

$$\Phi_{Jjlk}(\hat{r},\hat{S}) = \sum_{m} \bar{D}_{0m}^{J}(\hat{r}) \sqrt{\frac{2l+1}{2J+1}} \langle jml0|Jm \rangle \bar{D}_{mk}^{j}(\hat{S}), \quad (8)$$

where D'_{MK} is the normalized Wigner rotational matrix and $\langle jmj'm'|JM \rangle$ the Clebsch-Gordan coefficient [57].

The time-dependent wave packet method was used to solve the Schrödinger equation with an initial wave packet consisting of a Gaussian wave packet and a particular rovibrational state of XH_4 (X = C, Si)

$$\Psi^{\varepsilon}(t=0) = N e^{-\left[(R-R_0)/\alpha\right]^2} e^{i\kappa_0 R} \psi_{\rm XH_4},\tag{9}$$

where N, R_0 , α , κ_0 are, respectively, the normalization factor, position, half-width and central energy of the Gaussian wave packet. ψ_{XH_4} is the ro-vibrational wavefunction of XH₄, which has been predetermined in the reactant asymptote. In this work, the rotationless reactants were used.

2.4 Analysis of the reaction path Hamiltonian

The reaction path Hamiltonian [58] for both reactions was determined using POLYRATE 9.7 [59], starting from the saddle point geometry and going downhill to both directions in mass-weighted Cartesian coordinates using the Page and McIver's method [60] with a step size of $1.9 \times 10^{-4} \,\mathrm{u}^{1/2} \cdot a_0$. Along the minimum energy path (MEP), the reaction coordinate (s) is defined as the signed distance from the saddle point (s = 0), with s > 0 referring to the product side and s < 0 to the reactant side. The reaction path was followed between s = -1.5 and s = 1.5 u^{1/2}· a_0 , and the Hessian matrix was calculated every 10 points. These parameters have been tested for convergence. All calculations were carried out in mass-scaled coordinates with a reduced mass equal to 1 u. Along the MEP, a generalized normal-mode analysis was performed by projecting these modes along the reaction path as well as overall rotations and translations. With such information, the vibrationally adiabatic potentials were determined as:

$$V_{\rm a}(n,s) = V_{\rm MEP}(s) + \varepsilon_{\rm int}(n,s), \tag{10}$$

where $V_{\text{MEP}}(s)$ is the potential energy of the MEP with its zero at the reactants, and $\varepsilon_{\text{int}}(n,s)$ is the vibrational energy at *s* from the generalized normal-mode vibrations orthogonal to the reaction coordinate. $V_a(n = 0, s)$ corresponds to the ground vibrational state adiabatic potential, $V_a^G(s)$.

3 Results

3.1 Basis set

In this work, an *L*-shaped wavefunction expansion for R and r was used to reduce the size of the basis set [61].

	$H + CH_4$	$\mathrm{H}+\mathrm{SiH}_4$
$\overline{N_R}$	85/26	160/80
(R_{\min}, R_{\max})	$(2.5, 15.0)^{a}$	$(3.0, 11.0)^{a}$
N _r	5/25	6/28
(r_{\min}, r_{\max})	$(1.0, 5.0)^{a}$	$(1.5, 5.5)^{a}$
N_q	4	4
N_{γ}	5	5
J _{max}	45	72
l _{max}	27	48
$\dot{J}_{\rm max}$	18	24
k _{max}	6	6
N _{rot}	18,654	62,048
N _{tot}	3.5×10^8	3.4×10^9

^a Unit in a_0

For the $H + SiH_4$ reaction, a total of 160 sine basis functions ranging from 3.0 to 11.0 a_0 were used for the R basis set expansion with 80 nodes in the interaction region, and 6 and 28 basis functions of r were used in the asymptotic and interaction regions, respectively. For the vibration of the CZ₃ group, 4 and 5 basis functions were used for coordinates q and γ , respectively. The size of the rotational basis functions is controlled by the parameters, $J_{\text{max}} = 72$, $l_{\text{max}} = 48$, $j_{\text{max}} = 24$ and $k_{\text{max}} = 6$. After considering parity and C_{3v} symmetry, the size of rotational basis functions is 62,048 and the size of the total basis functions is \sim 3.4 billion. These parameters are listed on Table 1. For comparison, the corresponding values for the $H + CH_4$ reaction are also listed on this table. Due to the small mass of C, the size of basis set for the $H + CH_4$ reaction is smaller than that for the $H + SiH_4$ reaction.

3.2 Total reaction probabilities

For the H + CH_4 reaction, total reaction probabilities were calculated with nonrotating CH_4 initially in 5 vibrational states which labeled as (v_1, v_2, v_3, v_4) : (1) the ground vibrational state (0, 0, 0, 0) with a zero point energy of $4,872.29 \text{ cm}^{-1}$, (2) the excited states of the umbrella mode (0, 0, 0, 1) and (0, 0, 0, 2) with excitation energies of $1,288.18 \text{ cm}^{-1}$ (0.160 eV) and $2,543.35 \text{ cm}^{-1}$ (0.315 eV), (3) the symmetric stretching excited state (1, 0, 0, 0) with an excitation energy of 2,822.18 cm⁻¹ (0.350 eV) and (4) the asymmetric stretching vibrational state (0, 0, 1, 0) with an excitation energy of 2,949.29 cm⁻¹ (0.366 eV). These excitation energies are listed in Table 2 and compared with the experimental values [62]. Note that the relatively large errors are due to inaccuracies in the PES. It is also clear that the frequency for the (0, 1, 0, 0) state has a large discrepancy with the corresponding experimental value, due

Table 2 Vibration frequencies of CH_4 and SiH_4 (cm⁻¹)

	CH ₄		SiH ₄		
	Cal	Exp ^a	Cal	Exp ^b	
(0, 0, 0, 0)	4,872.29		3,364.02		
(0, 0, 0, 1)	1,288.18	1,367	851.22	914	
(0, 1, 0, 0)	2,517.06	1,583	1,677.36	975	
(0, 0, 0, 2)	2,543.35		1,686.89		
(1, 0, 0, 0)	2,822.16	3,026	1,992.27	2,187	
(0, 0, 1, 0)	2,949.29	3,157	2,075.69	2,191	

^a Reference [62]

^b Reference [63]

presumably to the fact that the bending motion is not compatible with the C_{3v} symmetry of the CZ_3 moiety [9]. As a result, the dynamics calculations for this state were not attempted.

Figure 2 shows the total reaction probabilities (J = 0) as a function of the translational energy and total energy

relative to $H + CH_4(0, 0, 0, 0)$, respectively. From the figure, all vibrational excitations of CH_4 appear to enhance the reactivity and the excitation energies were partially utilized to reduce the reaction threshold. At low collision energies, the reactivity is apparently enhanced by the symmetric stretching (1, 0, 0, 0) and asymmetric stretching (0, 0, 1, 0) excited states. However, the reaction probabilities as a function of total energy show that the translational energy is the most effective in promoting the reaction. These observations are consistent with our previous results on the earlier JG-PES [19].

The calculated energies of the six low-lying vibrational states of SiH₄ are also listed in Table 2 and compared with the experimental values [63]. Again, the (0, 1, 0, 0) state is poorly described by the 8D model and is not included in the dynamics calculations. The total reaction probabilities for the H + SiH₄ reaction were calculated with nonrotating SiH₄ initially in these five vibrational states and are shown in Fig. 3. The efficacies of the two umbrella excitations are very similar for both the H + SiH₄ and H + CH₄ reactions:





Fig. 2 Total reaction probabilities for the $H + CH_4$ reaction from different initial vibrational states of CH_4 as a function of translational energy (*upper panel*) and as a function of total energy measured with respect to $H + CH_4$ (0, 0, 0, 0) (*lower panel*)

Fig. 3 Total reaction probabilities for the H + SiH₄ reaction from different initial vibrational states of SiH₄ as a function of translational energy (*upper panel*) and as a function of total energy measured with respect to H + SiH₄ (0, 0, 0, 0) (*lower panel*)

(0, 0, 0, 2) > (0, 0, 0, 1). For the two stretching excitations of SiH₄, although the frequencies of v_1 (2,187 cm⁻¹) and v_3 (2,191 cm⁻¹) modes are close in energies, they show very different efficacies in reducing the reaction threshold. From Fig. 3, the (1, 0, 0, 0) excitation pronouncedly enhances the reactivity while the (0, 0, 1, 0) excitation has a minor effect on the reactivity. In the H + CH₄ reaction, two stretching excitations were found to have roughly the same efficacy in promoting the reaction. Similar behaviors have also been found in the previous studies of the O + CH₄ reaction with a four-dimensional model by Palma and Clary [64] and an eight-dimensional model by Liu and coworkers [9].

3.3 Integral cross sections and thermal rate constants

The integral cross section (ICS) for a specific initial state is obtained by summing the total reaction probabilities over all the partial waves,

$$\sigma_i(E) = \frac{1}{2J+1} \frac{\pi}{2\mu E} \sum_{J_{\text{tot}}} (2J_{\text{tot}} + 1) P_i^{J_{\text{tot}}}(E).$$
(11)

In this work, the $J_{tot} > 0$ partial waves were calculated with the centrifugal-sudden (CS) approximation [65, 66]. For H + CH₄, the maximum value of J_{tot} needed to converge the ICS is $J_{tot} = 40$. And for H + SiH₄, the corresponding value is $J_{tot} = 50$. The ICSs have been multiplied by a factor of 4 to account for all possible reaction channels. The results for ICS are plotted on Figs. 4 and 5.

The ICS curves are very similar to the reaction probability curves for both reactions. As shown in Fig. 4, all vibrational excitations of CH_4 promote the H + CH_4 reaction, consistent with the previous calculations [19]. It should be noted that the calculated values of the ICS are close to those obtained on the JG-PES [19] but almost 10 times larger than that obtained using recently developed ab initio PESs [55]. The lower panel of Fig. 4 shows that the groundstate reaction yields the largest ICS throughout the energy range. For the H + SiH₄ reaction, the H + SiH₄(1, 0, 0, 0) reaction has the largest contribution to the reaction when the total energy is above 0.33 eV and the ground-state reaction dominates the reactivity when the total energy is below 0.33 eV. The efficacy of the SiH₄(1, 0, 0, 0) excitation for the promotion of reactivity is rather pronounced. From the upper panels of Fig. 5, the ICS for the $H + SiH_4(1, 1)$ (0, 0, 0) reaction is overall 2 times larger than that for the $H + SiH_4(0, 0, 0, 2)$ reaction. Overall, the ICS values for the (0, 0, 0, 1), (0, 0, 1, 0) and (0, 0, 0, 2) excitations of SiH₄ are very close to each other over the entire collision energy range studied here.

The rate coefficients are calculated from the Boltzmann average of the cross sections for the five vibrational states considered in this work,



Fig. 4 Integral cross sections for the $H + CH_4$ reaction from different initial vibrational states of CH_4 as a function of translational energy (*upper panel*) and as a function of total energy measured with respect to $H + CH_4$ (0, 0, 0, 0) (*lower panel*)

$$k_{i}(T) = \left(\frac{8kT}{\pi\mu}\right)^{1/2} (k_{\rm B}T)^{-2} \int_{0}^{\infty} E_{\rm t} \exp(-E_{\rm t}/kT)\sigma_{\rm i}(E_{\rm t}) dE_{\rm t},$$
(12)

where $k_{\rm B}$ is the Boltzmann constant and μ is the reduced mass of the scattering coordinate. These rate coefficients are presented in Figs. 6 and 7. For comparison purposes, the calculated values with two other theoretical, the canonical variational theory (CVT) [44, 50] and quantum instanton (QI) methods [45, 52], as well as experimental values [67-71] are also showed in the figures. These theoretical calculations have been performed on the same PESs. For the $H + CH_4$ reaction, the 8D QM calculated values satisfactorily described the rate constants in the temperature range of 200-2,000 K and are in good agreement with the results calculated using other theoretical methods. For the $H + SiH_4$ reaction, on the other hand, the 8D QM rate coefficients are overall smaller than the experimental and QI values, which are quite close, by a factor of 1.5, and are slightly larger at low temperature but about 2 times smaller



Fig. 5 Integral cross sections for the $H + SiH_4$ reaction from different initial vibrational states of SiH₄ as a function of translational energy (*upper panel*) and as a function of total energy measured with respect to $H + SiH_4$ (0, 0, 0, 0) (*lower panel*)

at high temperatures than the CVT values. The discrepancy between 8D QM and CVT rate constant at high temperatures indicates that higher vibrational excited state may have considerable contributions to the reactivity.

4 Discussion

The differences in mode specificity between the two reactions can be explained by the vibrational character of the XH₄ reactant molecules. The (1, 0, 0, 0) and (0, 0, 1, 0) stretching modes of CH₄ are best described by the normalmode picture, while those of SiH₄ have different characters from the CH₄ molecule, as the vibrational excitation is localized in each Si–H bond. These differences are clearly shown in Fig. 8, where the corresponding wavefunctions are displayed in the (r, q) coordinates. For CH₄, the wavefunction is normal as the node is along either the (r + q) or (r - q) normal coordinate. In the classical picture, this corresponds to synchronized vibrational motions involving



Fig. 6 Arrhenius plot of thermal rate coefficients for the $H + CH_4$ reaction, in comparison with other theoretical [50, 52] and experimental rate constants [67–69]



Fig. 7 Arrhenius plot of thermal rate coefficients for the $H + SiH_4$ reaction, in comparison with other theoretical [44, 45] and experimental rate constants [70, 71]

all atoms. On the other hand, the wavefunctions for SiH_4 are largely local, as the node is along either the *r* or *q* coordinate. Classically, this represents vibrational motion in either the C–Y or C–Z₃ bond. It is important at this point to stress that the 8D model used here does not have full permutation symmetry. As a result, the C–Y and C–Z bonds are not treated as equals. In particular, the wavefunction of

Fig. 8 The contour plots of $|\psi_{XH_4}(r, q)|^2$ with integration over other coordinates. **a**, **b** (1, 0, 0, 0) and (0, 0, 1, 0) states of CH₄ and **c**, **d** (1, 0, 0, 0) and (0, 0, 1, 0) states of SiH₄, respectively



 SiH_4 in full dimension should be a linear combination of all four local modes with equal weights [23]. Consequently, the description of the reactant vibrational and reaction dynamics within the 8D model is approximate in nature. Nevertheless, this reduced-dimensional model captures the essence of the normal versus local vibrations, as evidenced by the reasonably good agreement with experimental vibrational frequencies in Table 2.

The results in Sect. 3 clearly demonstrated that the energy deposited into vibration can be used to overcome the reaction barrier. From Fig. 2, for example, 0.212 eV out of 0.350 eV for the (1, 0, 0, 0) excitation energy and 0.212 eV out of 0.366 eV for the (0, 0, 1, 0) excitation energy are used to reduce the reaction threshold for the H + CH₄ reaction, respectively. The corresponding percentages are 60 % and 58 %, clearly larger than the average value of 25 %. The dominance of the local mode features in the SiH₄ helps to promote its reaction with H even more effectively. The effect of the symmetric stretching mode is much more prominent in the $SiH_4 + H$ reaction than that in the $CH_4 + H$ reaction. This is clearly seen in Fig. 5, where the $SiH_4(1, 0, 0, 0)$ excited reactivity is much larger than both the ground and other excited vibrational states.

A word of caution concerning the limitations of the reduced-dimensionality model used in this work is in order. As mentioned above, the lack of permutation symmetry in this model prevents the proper treatment of the four equivalent X–H bonds in XH₄. As a result, the local mode wavefunctions for both the (1, 0, 0, 0) and (0, 0, 1, 0) states of SiH₄ are not properly symmetrized. In particular, the

full-dimensional (1, 0, 0, 0) state should also include excitation of the three C–Z bonds by symmetry. On the other hand, the reduced-dimensional wavefunction has only a node in the *r* coordinate along the C–Y bond. Furthermore, the attribution of the vibrational excitation to all three C–Z bonds for the (0, 0, 1, 0) state in our reduced-dimensional model is probably not a realistic representation of the local mode vibration in this molecule. Hence, the strong mode specificity observed in our reduced-dimensional model calculations is expected to be much less pronounced in fulldimensional dynamics.

To further illustrate the influence of the local mode vibration of SiH_4 during the H + SiH_4 reaction, we have computed the vibrationally adiabatic reaction paths for the corresponding vibrational excitation in SiH₄ in full dimension, which are presented in Fig. 9. For comparison, the corresponding results for the CH_4 + H reaction are also included in the same figure. Despite the fact that the reaction paths are qualitatively similar for these two reactions, the extent of localization is different. As shown in the figure, the symmetric stretching mode of both the isolated CH₄ and SiH₄ molecules involves all four bonds. However, as the H collisional partner approaches, the (1, 0, 0, 0) vibration localized to the proximal C-H or Si-H bond involved in the reaction, but with different extents. At s = -0.95, the SiH₄ vibration is already exclusively in the proximal Si-H bond, while the CH₄ vibration has still some residual normal-mode character. This is readily understood as the former has very weak coupling among the four Si-H stretching vibrations, while the coupling is moderate in CH₄ [22].



Fig. 9 Vibrationally adiabatic reaction paths for both the H + CH₄ (*upper panel*) and H + SiH₄ (*lower panel*) reactions on the PESs of Espinosa-Garcia et al. [44, 50]. The vibrational modes of the CH₄ or SiH₄ reactant in the reactant asymptote are labeled. The normal-mode vibrational vectors are also given for the isolated reactant ($s = -\infty$) and the reactive system at s = -0.95

At this point, it is interesting to consider isotopically substituted methane as a prototype for local excitation. Due to the larger differences in the vibrational frequencies in CHD₃, the C–H and C–D stretches do not couple anymore. The v_1 mode of CHD₃ is the C–H local mode, and the v_3 mode is localized in the C–D stretches. A recent study [72] of the H + HCD₃ \rightarrow H₂ + CD₃ reaction has showed that the C–H excitation led to a much large enhancement than the bending or umbrella mode. Similar results have also been obtained in the studies of the Cl + CHD₃ \rightarrow HCl + CD₃ reaction [7, 73]. These results together with the current study of the H + SiH₄ reaction demonstrate that the high efficiency promotes the reaction by local mode excitation.

Very recently, we have proposed the sudden vector projection (SVP) model [14, 15], which complements the adiabatic picture described in Fig. 9. Assuming the collision process is much faster than the intramolecular vibrational energy redistribution (IVR), the mode specificity in a bimolecular reaction is attributed to the projection of the reactant normal-mode vectors on to the reaction coordinate vector at the transition state. Although the $H + CH_4$ reaction has been analyzed in another publication [74] based on ab initio data, we repeat here the calculation on the CH5-EG-2002 PES. The SVP values are 0.51, 0.00, 0.28 and 0.038 for the symmetric stretching, bending, asymmetric stretching and umbrella modes of CH₄. These values are similar to those reported elsewhere [74], where the validity of the SVP predictions has been discussed. The same SVP calculations have been done on the (SiH5-EGSC-1998) PES as well for the $H + SiH_4$ reaction, and the corresponding SVP values are 0.47, 0.001, 0.27 and 0.001. In both systems, the SVP model predicts a larger enhancement effect for the symmetric stretching mode than that for the asymmetric mode, and small efficacies of the bending modes for enhancing the reactions. Both predictions are consistent with the 8D results presented above, although the SVP model seems to underestimate the bending modes.

5 Conclusions

In this work, an eight-dimensional quantum dynamics study has been carried out for both the H + CH₄ and H + SiH₄ reactions on two analytic PESs developed by Espinosa-Garcia and coworkers [44, 50]. The quantum dynamics are based on a reduced-dimensional model originally proposed by Palma and Clary [39] and reformulated by introducing a coordinate transform [19]. The 8D model includes representatives of all XH₄ vibrational modes, thus presenting a realistic reduced-dimensional model for the $X + YCZ_3 \rightarrow XY + CZ_3$ type reactions.

The total reaction probabilities and integral cross sections were obtained for reactants initially in the ground and four excited vibrational states. Strong mode specificity was found in both reactions. The mode specificity was analyzed and found to be consistent with the SVP predictions. In addition, the ability to enhance the reactivity is also correlated with the normal or local mode characters of the reactant molecules. For both reactions, the thermal rate coefficients were also calculated by averaging the contributions from all five initial vibrational states. The quantum dynamical rate coefficients are in reasonably good agreement with the experiment measurements and other theoretical calculated values.

This work demonstrates that the effects of reactant local modes on reaction dynamics could be studied using the 8D quantum dynamics method. On the other hand, it should be noted that the results are not expected to be quantitative due to the limitations of the reduced-dimensional model. An ultimate understanding of the mode specificity in these reactions can only be achieved with full dimensionality. Acknowledgments This work was supported by National Science Foundation of China (Projects No. 21221064 and 21373266 to MY) and US Department of Energy (DE-FG02-05ER15694 to HG).

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REGULAR ARTICLE

How the molecular face and the interaction vary as H atom approach H₂ molecule

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Abstract First, we briefly introduce the potential acting on one electron in a molecule (PAEM). Second, based on PAEM, the molecular intrinsic characteristic contours (MICC) is defined uniquely and intrinsically, and then, the electron density distribution is mapped on the MICC which is called molecular face (MF), an identification card, which is an intrinsic characteristic "face" or "fingerprint" for a molecule. Third, the polarization phenomena have been quantitatively demonstrated, i.e., the changing features of the spatial characteristic and the frontier electron density on the MF surface have been shown for reaction $H_2 + H \rightarrow H + H_2$, denoted as H1 - H2 + H3 \rightarrow H1 + H2 - H3, along the linear reaction path, which is compared with those quantities of the reaction $H + H \rightarrow H_2$ process. Furthermore, the conversion between the van der Waals interaction and chemical bonding during the process of $H1 - H2 + H3 \rightarrow H1 + H2 - H3$ reaction is shown by the PAEM-MO diagram, which provides insight into the characteristic of the interaction between two atoms: At the starting step of the reaction, the interaction between H1 and H2 atoms is chemically bonded, while the interaction between H2 and H3 is van der Waals interaction, and the contour of H2 atom in H₂ molecule does not overlap with that of the H3 atom, which means that the electron interflow between H1 and H2 atoms is free and the electron interflow between H2

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D.-X. Zhao e-mail: zhaodx@lnnu.edu.cn and H3 atoms is blocked by a PAEM barrier or must tunnel the barrier. At the transition state, both the pair of H1 and H2 and the pair of H2 and H3 are chemically bonded.

Keywords Molecular face \cdot The potential acting on one electron within a Molecule (PAEM) \cdot H₂ + H \rightarrow H + H₂ reaction \cdot PAEM-MO diagram \cdot Conversion between van der Waals interaction and chemical bonding

1 Introduction

One is used to say and to visualize an object with shape, size and color. Scientists often do the same things for molecules. Molecular shape and size are basic and fundamental concepts [1-12]. It is just through their boundary surfaces that molecules recognize, communicate and interact with each other. Therefore, molecular contour is essential in describing molecular properties, particularly in solutions [6-9]. Most molecular properties, even the chemical processes of life and physiology, are strongly dependent on molecular shape and size.

In this paper, we will demonstrate how the molecular face (MF) and the interaction character (van der Waals or chemical bonding) vary during the process of an H atom approaching a H_2 molecule as an example.

Usually, a molecular shape is visualized by a set of overlapping hard spheres with the corresponding van der Waals (vdW) radii. There are mainly three kinds of surfaces based on this idea: the vdW surface (vdWS), the solventaccessible surface (SAS) [13] and the molecular (or solvent excluded) surface (MS or SES) [14, 15]. The vdWS is formed by centering the spheres with properly chosen vdW radii at the positions of atomic nuclei. The SAS and MS are often used for dealing with solvation of solutes and

macromolecules. The SAS was first proposed by Lee and Richards as the locus of the center of an imaginary hard probe sphere when it rolls over the vdWS of the macromolecule. Later. Richards defined MS as the part of the surface of the solvent probe sphere that faces the macromolecule. As the simplest model, the vdWS and the volume it enclosed (van der Waals volume) are trivial to be programmed and able to be calculated analytically [6, 16, 17]. A large number of articles on algorithms for computing SAS and MS and the corresponding volumes can be found. The researches until 1996, including Connolly's dot surface [18], the parametric representations introduced by Duncan and Olson [19] and GEPOL proposed by Silla et al. [20], have been reviewed by Connolly in detail [21]. Many new approaches have also been reported recently [22-35]. An analytic method for reproducing Lee-Richards molecular volume [10] was proposed to yield an efficient solvent representation in simulations of biopolymers by Lee et al., Pomelli and Tomasi et al. [22, 23] presented the DefPOL program, which is more efficient to describe solvent effects on molecular system of large size. Wiser and coworkers presented the NLR optimization to accelerate the computation of hard-sphere molecular surfaces [24]. However, a problem for all of these approaches is the choice of an appropriate set of vdW radii or an assumed probe radius. The vdW radius is not a unique and fixed parameter characteristic of an atom. In general, vdW radii of same atom accepted by different authors vary by about 10 % [26]. This ambiguity can lead to a difference of about 12 % in calculating the vdW volume [29].

As a quantum chemical approach, electron density has been employed to give a sort of molecular boundary surface in many studies [1, 30-33]. Boyd [30] discussed the relative sizes of atoms from H to Xe based on the electron density distribution by using a cutoff value 10^{-4} of electron density. Bader and coworkers proposed the surface of constant electron density (usually typical values of 0.001 or 0.002 a.u.) to describe the shape and size of diatomic molecules [31-33]. They also demonstrated that the 0.002 a.u. isosurface of electron density is closely related in size to Corley-Pauling-Koltun (CPK) surfaces for hydrocarbons. Mezey proposed a method for topological analysis of contour surfaces represented by electron isodensity-fused spheres, which is called the molecular isodensity contour (MIDCO), and the MIDCO surfaces of biopolymers are constructed by a set of molecular density fragments [1]. Recently, isosurface function of promolecule electron density has also been presented by Mitchell and Spackman [34]. The molecular surface areas and volumes based on electron density are, of course, sensitive to selection of the cutoff value of electron density, and the choice of the cutoff value could not be determined intrinsically. Moreover, a single cutoff value is found not to be suitable for all molecular species [35]. For example, Stefanovich used an isodensity cutoff value of 0.001 a.u. to calculate the solvation energies of neutral and ionic species. However, the solvation energies of anions were found to be underestimated, while solvation energies of cations overestimated, and the assignment of a cutoff value to be employed for a zwitterionic system becomes ambiguous, i.e., the cutoff value applicable for neutral species would not appear to be suitable for both positive and negative ones. Thus, the previous methods and models, including the hard-sphere models and the electron density approach, have not yet provided an intrinsic and unique representation to molecular shape and size.

Recently, by using the classical turning point of electron movement in a molecule, the molecular intrinsic characteristic contour (MICC) [36-39] has been proposed by Yang and his workers. Furthermore, we define the MF that is the MICC mapped with the electron density [40]. MF is such a unique and intrinsic 3D picture that provides both molecular spatial appearance and molecular frontier electron density, an intuitive and instructive identification card for a molecule. Studies on some small molecules show that the MF is a reasonable representation [41–44]. In the previous researches, we have defined and calculated a set of atomic and ionic characteristic boundary radii in the spirit of this model [45–47]. The atomic characteristic radii [45, 48, 49] show a close correlation with the commonly used vdW atomic radii, and the surface derived by this sort of radius is taken as the accurate classical turning point surface to resemble by Ayers [50] and is called the atomic turning radius by Dekock et al. [51] in their investigation and classification of the atomic radii. The ionic characteristic radii [47, 52] also correlate quite well with Pauling ionic radii, as well as Shannon and Prewitt ionic radii. Moreover, the MF model was used to study the shape changing and polarization of some reactions, H + H, H + F, and $HCl + CH_2CH=CH$, and so on [37, 39, 53–58]. Very recently, we proposed and constructed the PAEM-MO diagram which is used to give a rule of distinguishing chemical bonding from van der Waals interaction [59]. Herein, both MF model and PAEM-MO diagram will be used to investigate the process of $H_2 + H \rightarrow H + H_2$ reaction to show the conversion between the chemical bonding and vdW interaction.

The following sections describe the MF model, computational algorithm, results and a brief summary.

2 Formalism of the molecular face (MF)

2.1 The potential acting on one electron in a molecule and its 3D representation

The potential acting on one electron in a molecule (PAEM) [36, 37, 59, 60] was formulated in previous

studies. It should be emphasized that the electron considered in this kind of the potential belongs to this molecule rather than an external test electron. As we know, however, the positive unit charge in defining the molecular electrostatic potential (MEP) [61–63] is an external test charge that does not belong to the molecule and only electrostatically interacts with all electrons and nuclei of the molecule. In addition, there are various potentials in atomic and molecular descriptions and studies [64–83], which are of some significance that will not be introduced here.

If one electron in a molecule locates at position \mathbf{r}_1 , it feels a potential coming from the remaining particles of the molecule, the *N*-1 remaining electrons and all nuclei. This potential acting on this electron in a molecule (PAEM), the $V(\mathbf{r}_1)$ at a position \mathbf{r}_1 can be expressed

$$V(\mathbf{r}_1) = -\sum_A \frac{Z_A}{r_{1A}} + \frac{1}{\rho(\mathbf{r}_1)} \int \frac{\rho_2(\mathbf{r}_1, \mathbf{r}_2)}{|\mathbf{r}_1 - \mathbf{r}_2|} d\mathbf{r}_2$$
(1)

in which the first term is the attractive potential provided by all nuclei, and the second term is the potential which is the interaction energy of this electron of interest with all the remaining electrons of the molecular system. In Eq. (1), Z_A is the charge on nucleus A, r_{1A} stands for the distance of nucleus A to the electron considered, summation involving index A is over atomic nuclei, \mathbf{r}_1 and \mathbf{r}_2 denote the electronic coordinates, $\rho(\mathbf{r}_1)$ is the one-electron density function of finding an electron at position \mathbf{r}_1 , and $\rho_2(\mathbf{r}_1, \mathbf{r}_2)$ is the two-electron density function of finding an electron at position \mathbf{r}_1 and at the same time another electron at position \mathbf{r}_2 .

In the ab initio configuration interaction (CI) method, the one-electron density function can be deduced as,

$$\rho(\mathbf{r}_{1}) = \sum_{I} \sum_{J} \sum_{i_{1} \in I} \sum_{j_{1} \in J} c_{I} c_{J}^{*} (-1)^{s-t} \delta_{\{I-i_{1}\}\{J-j_{1}\}} \times \left[f_{i_{1}}^{\alpha}(\mathbf{r}_{1}) f_{j_{1}}^{*\alpha}(\mathbf{r}_{1}) + f_{i_{1}}^{\beta}(\mathbf{r}_{1}) f_{j_{1}}^{*\beta}(\mathbf{r}_{1}) \right]$$
(2)

Here c_1 is the configuration interaction coefficient summation, $f_{i_1}^{\alpha}(\mathbf{r}_1)$ is the i_1 -th molecular orbital (MO) with α spin corresponding to the *s*-th row in the determinant *I*, and $f_{j_1}^{*\alpha}(\mathbf{r}_1)$ is the complex conjugate of the j_1 -th molecular orbital (MO) with α spin corresponding to the *t*-th row in the determinant *J*. Summation involving index *I* and *J* is over the configurations. Summation involving index i_1 and j_1 is over the molecular orbitals. The Kronecker delta, $\delta_{\{I-i_1\}\{J-j_1\}}$, is defined to be unity when $\{I - i_1\}$ equals $\{J - j_1\}$ and zero otherwise.

The spatial two-electron density function can practically be expressed by the following formula,

$$\rho_2(\mathbf{r}_1,\mathbf{r}_2)$$

$$= \sum_{I,J} c_I c_J^* \sum_{\substack{i_1 \in I \\ j_1 \in J \\ i_1 < i_2 \\ j_1 < j_2}} (-1)^{s+t-u-v} \delta_{\{I-i_1-i_2\}\{J-j_1-j_2\}} \cdot$$

$$\begin{cases} f_{i_{1}}^{\alpha}(\mathbf{r}_{1})f_{j_{1}}^{\alpha}(\mathbf{r}_{1})f_{i_{2}}^{\alpha}(\mathbf{r}_{2})f_{j_{2}}^{\alpha}(\mathbf{r}_{2}) + f_{i_{1}}^{\alpha}(\mathbf{r}_{1})f_{j_{1}}^{\alpha}(\mathbf{r}_{1})f_{i_{2}}^{\beta}(\mathbf{r}_{2})f_{j_{2}}^{\beta}(\mathbf{r}_{2}) \\ - f_{i_{1}}^{\alpha}(\mathbf{r}_{1})f_{j_{2}}^{\alpha}(\mathbf{r}_{1})f_{i_{2}}^{\alpha}(\mathbf{r}_{2})f_{j_{1}}^{\alpha}(\mathbf{r}_{2}) - f_{i_{1}}^{\alpha}(\mathbf{r}_{1})f_{j_{2}}^{\beta}(\mathbf{r}_{1})f_{j_{2}}^{\beta}(\mathbf{r}_{2})f_{j_{1}}^{\beta}(\mathbf{r}_{2}) \\ - f_{i_{2}}^{\alpha}(\mathbf{r}_{1})f_{j_{1}}^{\alpha}(\mathbf{r}_{1})f_{i_{1}}^{\alpha}(\mathbf{r}_{2})f_{j_{2}}^{\alpha}(\mathbf{r}_{2}) - f_{i_{2}}^{\alpha}(\mathbf{r}_{1})f_{j_{1}}^{\alpha}(\mathbf{r}_{1})f_{i_{1}}^{\beta}(\mathbf{r}_{2})f_{j_{2}}^{\beta}(\mathbf{r}_{2}) \\ + f_{i_{2}}^{\alpha}(\mathbf{r}_{1})f_{j_{2}}^{\alpha}(\mathbf{r}_{1})f_{i_{1}}^{\alpha}(\mathbf{r}_{2})f_{j_{1}}^{\alpha}(\mathbf{r}_{2}) + f_{i_{2}}^{\alpha}(\mathbf{r}_{1})f_{j_{2}}^{\beta}(\mathbf{r}_{1})f_{i_{1}}^{\beta}(\mathbf{r}_{2})f_{j_{1}}^{\beta}(\mathbf{r}_{2}) \\ + f_{i_{1}}^{\beta}(\mathbf{r}_{1})f_{j_{1}}^{\beta}(\mathbf{r}_{1})f_{i_{2}}^{\alpha}(\mathbf{r}_{2})f_{j_{2}}^{\alpha}(\mathbf{r}_{2}) + f_{i_{1}}^{\beta}(\mathbf{r}_{1})f_{j_{2}}^{\beta}(\mathbf{r}_{1})f_{i_{2}}^{\beta}(\mathbf{r}_{2})f_{j_{2}}^{\beta}(\mathbf{r}_{2}) \\ - f_{i_{1}}^{\beta}(\mathbf{r}_{1})f_{j_{2}}^{\beta}(\mathbf{r}_{1})f_{i_{2}}^{\alpha}(\mathbf{r}_{2})f_{j_{1}}^{\alpha}(\mathbf{r}_{2}) - f_{i_{1}}^{\beta}(\mathbf{r}_{1})f_{j_{2}}^{\beta}(\mathbf{r}_{1})f_{i_{2}}^{\beta}(\mathbf{r}_{2})f_{j_{1}}^{\beta}(\mathbf{r}_{2}) \\ - f_{i_{2}}^{\beta}(\mathbf{r}_{1})f_{j_{1}}^{\beta}(\mathbf{r}_{1})f_{i_{1}}^{\alpha}(\mathbf{r}_{2})f_{j_{1}}^{\alpha}(\mathbf{r}_{2}) - f_{i_{1}}^{\beta}(\mathbf{r}_{1})f_{j_{2}}^{\beta}(\mathbf{r}_{1})f_{i_{2}}^{\beta}(\mathbf{r}_{2})f_{j_{1}}^{\beta}(\mathbf{r}_{2}) \\ - f_{i_{2}}^{\beta}(\mathbf{r}_{1})f_{j_{1}}^{\beta}(\mathbf{r}_{1})f_{i_{1}}^{\alpha}(\mathbf{r}_{2})f_{j_{1}}^{\alpha}(\mathbf{r}_{2}) - f_{i_{2}}^{\beta}(\mathbf{r}_{1})f_{j_{1}}^{\beta}(\mathbf{r}_{1})f_{i_{1}}^{\beta}(\mathbf{r}_{2})f_{j_{2}}^{\beta}(\mathbf{r}_{2}) \\ + f_{i_{2}}^{\beta}(\mathbf{r}_{1})f_{j_{1}}^{\beta}(\mathbf{r}_{1})f_{i_{1}}^{\alpha}(\mathbf{r}_{2})f_{j_{1}}^{\alpha}(\mathbf{r}_{2}) - f_{i_{2}}^{\beta}(\mathbf{r}_{1})f_{j_{1}}^{\beta}(\mathbf{r}_{1})f_{i_{1}}^{\beta}(\mathbf{r}_{2})f_{j_{2}}^{\beta}(\mathbf{r}_{2}) \\ + f_{i_{2}}^{\beta}(\mathbf{r}_{1})f_{j_{2}}^{\beta}(\mathbf{r}_{1})f_{i_{1}}^{\alpha}(\mathbf{r}_{2})f_{j_{1}}^{\alpha}(\mathbf{r}_{2}) - f_{i_{2}}^{\beta}(\mathbf{r}_{1})f_{j_{2}}^{\beta}(\mathbf{r}_{1})f_{i_{1}}^{\beta}(\mathbf{r}_{2})f_{j_{1}}^{\beta}(\mathbf{r}_{2}) \\ + f_{i_{2}}^{\beta}(\mathbf{r}_{1})f_{j_{2}}^{\beta}(\mathbf{r}_{1})f_{i_{1}}^{\alpha}(\mathbf{r}_{2})f_{j_{1}}^{\alpha}(\mathbf{r}_{2}) - f_{i_{2}}^{\beta}(\mathbf{r}_{1})f_{$$

where $\delta_{\{I-i_1-i_2\}\{J-j_1-j_2\}}$ is equal to 1, if the cofactor $\{I-i_1-i_2\}$ of the determinant *I* is equal to the cofactor $\{J-j_1-j_2\}$ of the determinant *J*, and otherwise is zero.

Based on the expressions of the electron density $\rho(\mathbf{r}_1)$ Eq. (2), and the two-electron density function $\rho_2(\mathbf{r}_1, \mathbf{r}_2)$, Eq. (3), we can obtain the concrete expression of PAEM defined in Eq. (1) and can then calculate it by using an ab initio CI method and an our in-house program.

2.2 Definition of molecular intrinsic characteristic contour (MICC)

The idea for this definition can be considered as follows [36, 37, 40]. When an electron moves in a molecule, its kinetic energy varies with its relative position to the other particles of the molecule. If its energy is equal to its felt potential, i.e., the average kinetic energy of this electron at this point equals zero, then this position \mathbf{r} is a classical turning point of this electron movement. Supposing that the one-electron energy at \mathbf{r} is equal to minus of the first ionization potential of the molecule, then we have the formula, $V(\mathbf{r}) = -I$, i.e., the classical turning point equation of this electron movement, where $V(\mathbf{r})$ is the PAEM defined by Eq. (1) and I is the first ionization potential of the molecule. We assume that such a point \mathbf{r} is termed as an intrinsic characteristic contour point of this molecule. The set of all the intrinsic characteristic points defines the molecular intrinsic characteristic contour (MICC), which can be expressed as:

$$\boldsymbol{G}(-I) = \{ \mathbf{r} : V(\mathbf{r}) = -I \}$$
⁽⁴⁾

in which G denotes the MICC, i.e., the MICC is composed of all the classical turning points of electron movement in

the molecule and is an iso-PAEM contour where PAEM (or one-electron energy) equals the negative value of the ionization potential of the molecule. In other words, if one electron located on the MICC obtains an equivalent energy of the molecular first ionization potential, it becomes a free electron.

2.3 Program and calculation

By using the ab initio MELD program [84] developed by Davidson et al. and a separated program developed by our own, the PAEM was calculated. The SDCI calculation in MELD program with the near Hartree-Fock quality Gaussian type orbital basis sets [85] was used to calculate the molecular integrals in the expression of the PAEM. At the same level of ab initio method, the vertical ionization potentials of a molecular system in its ground state were also calculated if there is no experimental value can be utilized. According to the classical turning point of electron movement, the molecular intrinsic characteristic contour (MICC) points were calculated one by one for a molecule, then collected, and thereby a MICC was drawn. All of the numerical computations were carried out on a SGI O-300 server. Based on this model, we have, respectively, studied the MICCs of a number of molecular systems in their ground states.

The calculation of the PAEM and MICC in our model for a molecule can be summarized in the following three steps:

- (a) Calculate the molecular integrals in Eqs. (2) and (3). Here, we utilize the MELD program [84] developed by Davidson et al. to perform the ab initio calculations. All calculations were carried out on an O300 server (16 \times 500 MHz MIPS R14000 CPU) and a SGI Octane2 workstation (2 \times 400 MHz MIPS R12000 CPU). Then, by using a program of our own, calculate the PAEM and physical quantities expressed by Eqs. (2) and (3), including the electron density.
- (b) Calculate the MICC points. According to the classical turning point equation, Eq. (4), the MICC points are obtained by a large number of calculations. The vertical ionization potential *I* that we have used is the experimental vertical ionization potential or the calculated value by the accurate ab initio CI method.
- (c) Sketch the MICC. Collecting all the MICC points, a 3D picture of MICC can be drawn. Besides this, significantly, the electron density distribution on the MICC is also mapped out. Matlab [86] was used to depict and to obtain visualization plots of the MF.

3 Results and discussion

3.1 3D Representations of the PAEM and MF: HI molecule as an example

Using the formulism and computational method mentioned above, we have studied the potentials acting on an electron for some molecules. Here, take HI molecule as an example to demonstrate the procedure of obtaining the PAEM and MF as follows.

We put I atom at the origin of the coordinate system and H atom at the *X*-axis. The equilibrium bond length of the HI molecule is 3.041 a.u. (experimental value) [87]. Since HI has an axial symmetry around its molecular axis that contains the two nuclei, it is sufficient to only represent the PAEM $V(\mathbf{r}_1)$ where \mathbf{r}_1 is on a plane that passes through the molecular axis.

First of all, the calculations of the $V(x_1)$ were carried out along the molecular axis, the *x*-axis. When \mathbf{r}_1 runs on the molecular plane, the *xy* plane, i.e., taking a lot of net points on the *xy* plane, the calculated $V(\mathbf{r}_1)$ is displayed in the *z*-axis, then a 3D representation of the PAEM $V(\mathbf{r}_1)$ is brought out and drawn in Fig. 1. The near-nucleus region is the area that is located around a nucleus of a molecule. For the PAEM of HI molecule, there are two deep potential wells which originate from the nuclear attraction, a



Fig. 1 a The 3D graph of the PAEM for HI molecule on xy plane (Molecular Plane). All quantities are in atomic units in this paper. **b** The molecular intrinsic characteristic contour with electronic density (MICCED or MF) of HI molecule

wider potential well corresponding to the I atom of the larger nuclear charge and a narrower potential well relating to the H atom of the smaller nuclear charge. The PAEM surface between the two wells has a saddle shape with a saddle point at $x_1 = 2.10$, $y_1 = 0.0$, V(2.10,0.0) = -1.5726hartree which is called PAEM saddle point (psp). The corresponding electronic position of the PAEM saddle point between the H and I atoms is named the bond center (bc), marked by star. The corresponding PAEM of this point is called the height of PAEM barrier. The vast remaining part is the out-bond region of the PAEM, the rest area subtracting the near-nuclei and bond regions from the total PAEM. In this region, the potential gradually ascends with enlarging the distance between the electron and nucleus and then approaches to zero at the infinite, which is a correct asymptotic behavior.

Then, Eq. (4), called equation of the classical turning point, is performed to obtain the MICC points of HI molecule. At the same time, the electron density at every MICC point is calculated. Thus, we collect the MICC point and their corresponding electron density. As result, a colorful picture of molecular intrinsic characteristic contour with electron density (MICCED or MF) is obtained, as shown in Fig. 1b, where the color scale indicates the electron density distributed on the MICC, i.e., the red color area denotes lower electron density and the blue area has larger one. It can be found that the MF contains not only the spatial shape and size but also the electron distribution on the MICC. Therefore, it provides insight into a significant feature of both molecular spatial appearance and frontier electron density mapped on the MICC, an intuitive picture of molecular fingerprint or face, which is denoted as MF.

3.2 The changing pictures of the molecular face (MF) for the reaction $H_2 + H \rightarrow H + H_2$ at different internuclear distances along the energy-favorite linear reaction path

Now, the typical three-atomic exchange reaction $H_2 + H \rightarrow H + H_2$ is taken as an example to show both its spatial and electronic MF varying pictures, along the energy-favorite collinear reaction path [88–90], the route of the intrinsic reaction coordinate [91]. The calculations were performed by a program at the ab initio CI level based on the MELD program package developed by Davidson et al. [84] and a separate program with 18s15p3d basis set for each H atom.

The coordinate system is chosen as follows: along the collinear approaching line (the chemical bond direction in H_2 molecule), the atoms from left to right are the two H atoms of the hydrogen molecule and the approaching H atom, denoted as H1, H2 and H3 atoms. The structural formula, $H1(R_1)H2(R_2)H3$, where R_1 is the distance between H1 and H2 atoms, R_2 is the distance between H2 and H3

atoms, is used to denote the corresponding state at each point of the IRC route.

The vivid changing pictures of MFs during the process of the H atom approaching hydrogen molecule drawn along their IRC route are shown in Fig. 2b-h, in which the dark blue region has large electron density while the dark red region has small electron density, and the value of MFED or MF is denoted by the color scale (marked by Den) on the right side of the picture. Figure 2 shows a series of MF pictures at different distances of the process of H atom approaching H₂ molecule, a reaction of H₃ system. As the H atom, going along the linear reaction path, is 4.33 a.u. close to the adjacent H atom of the H₂ molecule that has a nuclear separation 1.4011 a.u., i.e., the geometry H1(1.4011)H2(4.33)H3, both the H_2 part and the H part have stretched their hands and increased their electron density around the active areas, even with a weak green signal from the H₂ right hand, as depicted in Fig. 2c. About the transition state structure, as shown in Fig. 2h, the H₃ linear reaction is symmetric. After this step, the right H3 atom and the middle H2 atom are combining while the left H1 atom leaving, whose MF pictures are just the images of Fig. 2a–g about the middle vertical plane.

In order to quantitatively describe and compare the changing features of different MFs along the IRC route, several characteristic parameters of the molecular intrinsic characteristic contour (MICC) and its corresponding electron density (MFED) are defined and exhibited in Fig. 2b. For example, in the molecular cross section including atoms H1, H2 and H3, the horizontal straight line passing through all nuclei has four intersection points with the molecular contour of H₃ system, so R_1 (H1), R_r (H2) and $R_1(H3)$ and $R_r(H3)$ are in turn defined as the distances from the left side of H1 atom, the right side of H2 atom, the left side of H3 atom, the right side of H3 atom to the intersection points, and the corresponding electron density on these points are denoted by *Den*₁(H1), *Den*_r(H2) and *Den*₁(H3) and $Den_r(H3)$, respectively. The vertical straight line passing through nucleus H1 has two intersection points with the MF, and then, the distance from nucleus H1 to one of the intersection points is denoted $R_{\nu}(H1)$, the corresponding electron density is represented by $Den_{v}(H1)$. The calculated numerical values of some characteristic quantities of MICC and MFED at different geometries for the H₃ system are listed in Table 1.

The polarization phenomena, including both the deformation of spatial shape and the charge of the frontier electron density, are clearly demonstrated by the characteristic quantities, as given in Table 1, for the H₃ reaction, particularly the $R_r(H2)$ and $R_l(H3)$ that denote the reaction sites of H₂ molecule part and H atom part, respectively. The streching of both parts for the reaction process is obviously seen from the following change of $R_r(H2)$ values (from

Fig. 2 3D MF of the energy-favorite linear $H_2 + H \rightarrow H + H_2$ reaction system at the nuclear arrangement along IRC reaction route during the process of H atom gradually approaching to H_2 molecule. **a** The isolated H_2 molecule and H atom; **b**-**h** corresponding to the nuclear separation of H(1.4009)H(5.0)H, H(1.4010)H(4.33) H, H(1.4011)H(4.27)H, H(1.4070)H(3.57)H, H(1.421)H(3.57)H, H(1.421)H(2.971)H, H(1.518)H(2.179)H and H(1.757)H(1.757)H of the linear H_3 reaction system, respectively

1.8012 to 1.9793 a.u.) and $R_{\rm l}({\rm H3})$ values (from 2.0532 to 2.2899 a.u.). Correspondingly, the values of the frontier electron densities have the following changes: $Den_{\rm r}({\rm H2})$ from 5.921 \times 10⁻³ to 6.541 \times 10⁻³ a.u. and $Den_{\rm l}({\rm H3})$ from 5.664 \times 10⁻³ to 6.533 \times 10⁻³ a.u. listed in Table 1. The following four paragraphs are given for more detailed descriptions.

If the H atom and H₂ molecule are separate, the MF of the H atom is round with a boundary radius of 2.00 a.u. and the electron density of 5.832×10^{-3} a.u., and the MF of the H₂ part is like an egg with nearly uniform electron distribution on the MF frontier surface, which is shown in Fig. 2a.

As the H atom along the linear reaction path is 5.00 far from the adjacent H atom of the H₂ moiety that is at nuclear separation 1.4009, or simply denoted by H(1.4009) H(5.0)H, the 3D MF is shown in Fig. 2b, where the color indicates the electron density on the MF surface (the red or blue area has lower or higher electron density). Obviously, there are H₂ molecular and H atomic regions, and between them there is a medium that is a classically forbidden region for the electronic motion. The H atom looks still round with nearly uniform electron density on MF, while the MF region of the H₂ part is clearly polarized in the electron distribution although its equilibrium nuclear separation is still 1.4009, around the middle of the H₂ region, the electron density (blue) is larger than that around the two end sides, and the left one being red and the right one being vellow (yellow has higher electron density than red but blue has the highest). It seems that the H_2 part defends its covalent bond concentrating its electrons around the middle of the H₂ bond but giving a bit warning with its yellow right hand to the attacking H atom.

The Fig. 2d of the geometry H(1.4011)H(4.302)H shows a critical step along the H₃ reaction path, where only one touching point appears between the two MF regions. Here, the two regions are more significantly deformed and polarized, particularly around the touching tip. It seems that both the H₂ molecule and H atom regions stretch their body to welcome each other with the green hands, implying that they both intensify their electron density on the adjacent active areas, which just belong to the bonding domain described by the Hellmann–Feynman theorem [92]. This means both regions have strengthened their interaction.



Table 1The molecularintrinsic characteristic quantities(MICC and MFED) of the H_3 systems, distances in a.u.,electron densities in $\times 10^{-3}$ a.u

R_1 R_2 $R_v(H1)$ $R_l(H1)$ $R_v(H2)$ $R_r(H2)$ $R_v(H3)$ $R_l(H3)$ $R_r(H3)$ 1.40095.0001.97281.82221.96061.80122.01782.05322.01341.40104.5001.98901.82631.97621.83042.03352.12212.02831.40114.3301.99021.82931.97901.90102.04052.21252.03561.40114.3021.99621.83281.98611.99722.04312.29942.03751.40114.2001.99241.83391.98352.04382.030401.40114.2701.98921.83101.97882.04072.03401.40103.5712.04121.88492.03632.11442.10061.42102.9712.17941.98282.14782.22092.19011.51802.1792.44592.24952.41202.44612.36211.75701.7572.54842.40682.51172.54842.4068 R_1 R_2 $Den_v(H1)$ $Den_v(H2)$ $Den_v(H2)$ $Den_v(H3)$ $Den_t(H3)$ 1.40104.5007.3375.0647.5186.5685.4446.0035.4901.40114.3027.2205.0247.3396.5415.3315.4091.40114.3007.3335.0537.4525.3655.4351.40114.2007.2755.0147.3765.3315.4091.40									
1.4009 5.000 1.9728 1.8222 1.9606 1.8012 2.0178 2.0532 2.0134 1.4010 4.500 1.9890 1.8263 1.9762 1.8304 2.0335 2.1221 2.0283 1.4011 4.302 1.9902 1.8293 1.9790 1.9010 2.0405 2.2125 2.0356 1.4011 4.302 1.9962 1.8328 1.9861 1.9972 2.0431 2.2994 2.0375 1.4011 4.270 1.9892 1.8310 1.9788 2.0407 2.0438 2.0380 1.4011 4.270 1.9892 1.8310 1.9788 2.0407 2.0438 2.0340 1.4010 4.270 1.9892 1.8310 1.9788 2.0407 2.0412 2.0340 1.4070 3.571 2.0412 1.8849 2.0363 2.1144 2.1006 1.4210 2.971 2.1794 1.9828 2.1478 2.2209 2.1901 1.5180 2.179 2.4459 2.2495 2.4120 2.4461 2.3621 1.757 1.757 2.5484 2.4068 2.5117 2.5484 2.4068 R_1 R_2 $Den_v(H1)$ $Den_t(H1)$ $Den_v(H2)$ $Den_v(H2)$ $Den_v(H3)$ $Den_t(H3)$ 1.4010 4.500 7.337 5.064 7.518 6.568 5.444 6.003 5.490 1.4011 4.302 7.220 5.024 7.339 6.541 5.3365 5.435 1.4011 4.290	R ₁	R_2	$R_{\rm v}({ m H1})$	<i>R</i> _l (H1)	$R_{\rm v}({\rm H2})$	<i>R_r</i> (H2)	$R_{\rm v}({\rm H3})$	<i>R</i> _l (H3)	<i>R_r</i> (H3)
1.40104.5001.98901.82631.97621.83042.03352.12212.02831.40114.3301.99021.82931.97901.90102.04052.21252.03561.40114.3021.99621.83281.98611.99722.04312.29942.03751.40114.2901.99241.83391.98352.04382.20942.03801.40114.2701.98921.83101.97882.04072.03401.40703.5712.04121.88492.03632.11442.10061.42102.9712.17941.98282.14782.22092.19011.51802.1792.44592.24952.41202.44612.36211.75701.7572.54842.40682.51172.54842.4068 R_1 R_2 $Den_v(H1)$ $Den_t(H2)$ $Den_r(H2)$ $Den_v(H3)$ $Den_t(H3)$ 1.40105.0007.5965.1097.7605.9215.6165.6645.6461.40104.3007.3375.0647.5186.5685.4446.0035.4901.40114.3027.2205.0247.3396.5415.3315.4351.40114.2007.2755.0147.3765.3655.4351.40114.2007.2755.0147.3765.3655.4351.40114.2007.2755.0147.3765.3655.4351.40114.2007.2755.014 <th>1.4009</th> <th>5.000</th> <th>1.9728</th> <th>1.8222</th> <th>1.9606</th> <th>1.8012</th> <th>2.0178</th> <th>2.0532</th> <th>2.0134</th>	1.4009	5.000	1.9728	1.8222	1.9606	1.8012	2.0178	2.0532	2.0134
1.40114.3301.99021.82931.97901.90102.04052.21252.03561.40114.3021.99621.83281.98611.99722.04312.29942.03751.40114.2901.99241.83391.98352.04382.03802.03801.40114.2701.98921.83101.97882.04072.03401.40703.5712.04121.88492.03632.11442.10061.42102.9712.17941.98282.14782.22092.19011.51802.1792.44592.24952.41202.44612.36211.75701.7572.54842.40682.51172.54842.4068R1R2Denv(H1)Dent(H1)Denv(H2)Denv(H2)Denv(H3)Dent(H3)1.40105.0007.5965.1097.7605.9215.6165.6645.6461.40104.5007.3375.0647.5186.5685.4446.0035.4901.40114.3027.2205.0247.3396.5415.3396.5335.4171.40114.2007.2755.0147.3765.3655.4355.4351.40114.2707.3335.0537.4525.3655.4355.4351.40114.2907.2755.0147.3765.3655.4355.4351.40114.2907.2755.0147.365.3655.4355.4351.40114.290 </th <th>1.4010</th> <th>4.500</th> <th>1.9890</th> <th>1.8263</th> <th>1.9762</th> <th>1.8304</th> <th>2.0335</th> <th>2.1221</th> <th>2.0283</th>	1.4010	4.500	1.9890	1.8263	1.9762	1.8304	2.0335	2.1221	2.0283
1.40114.3021.99621.83281.98611.99722.04312.29942.03751.40114.2901.99241.83391.98352.04382.03801.40114.2701.98921.83101.97882.04072.03401.40703.5712.04121.88492.03632.11442.10061.42102.9712.17941.98282.14782.22092.19011.51802.1792.44592.24952.41202.44612.36211.75701.7572.54842.40682.51172.54842.4068R_1R_2Den_v(H1)Den_v(H2)Den_r(H2)Den_v(H3)Den_t(H3)1.40105.0007.5965.1097.7605.9215.6165.6645.6461.40104.5007.3375.0647.5186.5685.4446.0035.4901.40114.3027.2205.0247.3396.5415.3396.5335.4171.40114.2007.2755.0147.3765.3315.4091.40114.2707.3335.0537.4525.3655.4351.40703.5716.5614.5006.5604.6324.8311.42102.9714.8433.6965.0603.7624.1581.51802.1792.7012.2392.7832.4863.1241.75701.7572.1642.3102.2262.1642.310	1.4011	4.330	1.9902	1.8293	1.9790	1.9010	2.0405	2.2125	2.0356
1.40114.2901.99241.83391.98352.04382.03801.40114.2701.98921.83101.97882.04072.03401.40703.5712.04121.88492.03632.11442.10061.42102.9712.17941.98282.14782.22092.19011.51802.1792.44592.24952.41202.44612.36211.75701.7572.54842.40682.51172.54842.4068 R_1 R_2 $Den_v(H1)$ $Den_t(H1)$ $Den_v(H2)$ $Den_r(H2)$ $Den_v(H3)$ $Den_t(H3)$ 1.40105.0007.5965.1097.7605.9215.6165.6645.6461.40114.3007.3375.0647.5186.5685.4446.0035.4901.40114.3027.2205.0247.3396.5415.3396.5335.4171.40114.2707.3335.0537.4525.3655.4355.4351.40114.2707.3335.0537.4525.3655.4351.40703.5716.5614.5003.7624.1581.51802.1792.7012.2392.7832.4863.1241.57701.7572.1642.3102.2262.1642.310	1.4011	4.302	1.9962	1.8328	1.9861	1.9972	2.0431	2.2994	2.0375
1.40114.2701.98921.83101.97882.04072.03401.4070 3.571 2.04121.88492.03632.11442.10061.42102.9712.17941.98282.14782.22092.19011.51802.1792.44592.24952.41202.44612.36211.75701.7572.54842.40682.51172.54842.4068 R_1 R_2 $Den_v(H1)$ $Den_t(H1)$ $Den_v(H2)$ $Den_r(H2)$ $Den_v(H3)$ $Den_t(H3)$ 1.4010 5.000 7.596 5.109 7.760 5.921 5.616 5.664 5.646 1.4011 4.300 7.337 5.064 7.518 6.568 5.444 6.003 5.490 1.4011 4.302 7.220 5.024 7.339 6.541 5.339 6.533 5.417 1.4011 4.200 7.275 5.014 7.376 5.365 5.435 1.4070 3.571 6.561 4.500 6.560 4.632 4.831 1.4210 2.971 4.843 3.696 5.060 3.762 4.158 1.5180 2.179 2.701 2.239 2.783 2.486 3.124 1.7570 1.757 2.164 2.310 2.226 2.164 2.310	.4011	4.290	1.9924	1.8339	1.9835		2.0438		2.0380
1.4070 3.571 2.0412 1.8849 2.0363 2.1144 2.1006 1.4210 2.971 2.1794 1.9828 2.1478 2.2209 2.1901 1.5180 2.179 2.4459 2.2495 2.4120 2.4461 2.3621 1.7570 1.757 2.5484 2.4068 2.5117 2.5484 2.4068 R_1 R_2 $Den_v(H1)$ $Den_1(H1)$ $Den_v(H2)$ $Den_r(H2)$ $Den_v(H3)$ $Den_1(H3)$ $Den_r(H3)$ 1.4010 5.000 7.596 5.109 7.760 5.921 5.616 5.664 5.646 1.4010 4.500 7.337 5.064 7.518 6.568 5.444 6.003 5.490 1.4011 4.302 7.220 5.024 7.339 6.541 5.339 6.533 5.417 1.4011 4.290 7.275 5.014 7.376 5.365 5.435 1.4011 4.270 7.333 5.053 7.452 5.365 5.435 1.4070 3.571 6.561 4.500 6.560 4.632 4.831 1.4210 2.971 4.843 3.696 5.060 3.762 4.158 1.5180 2.179 2.701 2.239 2.783 2.486 3.124 1.7570 1.757 2.164 2.310 2.226 2.164 2.310	.4011	4.270	1.9892	1.8310	1.9788		2.0407		2.0340
1.42102.9712.17941.98282.14782.22092.19011.51802.1792.44592.24952.41202.44612.36211.75701.7572.54842.40682.51172.54842.4068 R_1 R_2 $Den_v(H1)$ $Den_1(H1)$ $Den_v(H2)$ $Den_r(H2)$ $Den_v(H3)$ $Den_1(H3)$ $Den_r(H3)$ 1.40105.0007.5965.1097.7605.9215.6165.6645.6461.40104.5007.3375.0647.5186.5685.4446.0035.4901.40114.3027.2205.0247.3396.5415.3396.5335.4171.40114.2907.2755.0147.3765.3655.4355.4351.40114.2707.3335.0537.4525.3655.4351.40703.5716.5614.5006.5603.7624.1581.51802.1792.7012.2392.7832.4863.1241.75701.7572.1642.3102.2262.1642.310	.4070	3.571	2.0412	1.8849	2.0363		2.1144		2.1006
1.51802.1792.44592.24952.41202.44612.36211.75701.7572.54842.40682.51172.54842.4068 R_1 R_2 $Den_v(H1)$ $Den_l(H1)$ $Den_v(H2)$ $Den_r(H2)$ $Den_v(H3)$ $Den_l(H3)$ $Den_r(H3)$ 1.40105.0007.5965.1097.7605.9215.6165.6645.6461.40104.5007.3375.0647.5186.5685.4446.0035.4901.40114.3027.2205.0247.3396.5415.3396.5335.4171.40114.2907.2755.0147.3765.3655.3655.4351.40114.2707.3335.0537.4525.3655.4351.40703.5716.5614.5006.5604.6324.8311.42102.9714.8433.6965.0603.7624.1581.51802.1792.7012.2392.7832.4863.1241.75701.7572.1642.3102.2262.1642.310	1.4210	2.971	2.1794	1.9828	2.1478		2.2209		2.1901
1.7570 1.757 2.5484 2.4068 2.5117 2.5484 2.4068 R_1 R_2 $Den_v(H1)$ $Den_l(H1)$ $Den_v(H2)$ $Den_r(H2)$ $Den_v(H3)$ $Den_l(H3)$ $Den_r(H3)$ 1.4010 5.000 7.596 5.109 7.760 5.921 5.616 5.664 5.646 1.4010 4.500 7.337 5.064 7.518 6.568 5.444 6.003 5.490 1.4011 4.330 7.313 5.060 7.448 6.697 5.366 6.382 5.436 1.4011 4.302 7.220 5.024 7.339 6.541 5.339 6.533 5.417 1.4011 4.290 7.275 5.014 7.376 5.365 5.435 1.4011 4.270 7.333 5.053 7.452 5.365 5.435 1.4070 3.571 6.561 4.500 6.560 4.632 4.831 1.4210 2.971 4.843 3.696 5.060 3.762 4.158 1.5180 2.179 2.701 2.239 2.783 2.486 3.124 1.7570 1.757 2.164 2.310 2.226 2.164 2.310	1.5180	2.179	2.4459	2.2495	2.4120		2.4461		2.3621
R_1 R_2 $Den_v(H1)$ $Den_1(H1)$ $Den_v(H2)$ $Den_r(H2)$ $Den_v(H3)$ $Den_1(H3)$ $Den_r(H3)$ 1.40105.0007.5965.1097.7605.9215.6165.6645.6461.40104.5007.3375.0647.5186.5685.4446.0035.4901.40114.3307.3135.0607.4486.6975.3666.3825.4361.40114.3027.2205.0247.3396.5415.3396.5335.4171.40114.2907.2755.0147.3765.3655.3655.4351.40114.2707.3335.0537.4525.3655.4351.40103.5716.5614.5006.5604.6324.8311.42102.9714.8433.6965.0603.7624.1581.51802.1792.7012.2392.7832.4863.1241.75701.7572.1642.3102.2262.1642.310	1.7570	1.757	2.5484	2.4068	2.5117		2.5484		2.4068
1.40105.0007.5965.1097.7605.9215.6165.6645.6461.40104.5007.3375.0647.5186.5685.4446.0035.4901.40114.3307.3135.0607.4486.6975.3666.3825.4361.40114.3027.2205.0247.3396.5415.3396.5335.4171.40114.2907.2755.0147.3765.3655.4351.40114.2707.3335.0537.4525.3655.4351.40103.5716.5614.5006.5604.6324.8311.42102.9714.8433.6965.0603.7624.1581.51802.1792.7012.2392.7832.4863.1241.75701.7572.1642.3102.2262.1642.310	R ₁	<i>R</i> ₂	Den _v (H1)	Den _l (H1)	$Den_v(H2)$	Den _r (H2)	$Den_v(H3)$	$Den_{l}(H3)$	Den _r (H3)
1.40104.5007.3375.0647.5186.5685.4446.0035.4901.40114.3307.3135.0607.4486.6975.3666.3825.4361.40114.3027.2205.0247.3396.5415.3396.5335.4171.40114.2907.2755.0147.3765.3315.4091.40114.2707.3335.0537.4525.3655.4351.40703.5716.5614.5006.5604.6324.8311.42102.9714.8433.6965.0603.7624.1581.51802.1792.7012.2392.7832.4863.1241.75701.7572.1642.3102.2262.1642.310	1.4010	5.000	7.596	5.109	7.760	5.921	5.616	5.664	5.646
1.40114.3307.3135.0607.4486.6975.3666.3825.4361.40114.3027.2205.0247.3396.5415.3396.5335.4171.40114.2907.2755.0147.3765.3315.4091.40114.2707.3335.0537.4525.3655.4351.40703.5716.5614.5006.5604.6324.8311.42102.9714.8433.6965.0603.7624.1581.51802.1792.7012.2392.7832.4863.1241.75701.7572.1642.3102.2262.1642.310	1.4010	4.500	7.337	5.064	7.518	6.568	5.444	6.003	5.490
1.40114.3027.2205.0247.3396.5415.3396.5335.4171.40114.2907.2755.0147.3765.3315.4091.40114.2707.3335.0537.4525.3655.4351.40703.5716.5614.5006.5604.6324.8311.42102.9714.8433.6965.0603.7624.1581.51802.1792.7012.2392.7832.4863.1241.75701.7572.1642.3102.2262.1642.310	1.4011	4.330	7.313	5.060	7.448	6.697	5.366	6.382	5.436
1.40114.2907.2755.0147.3765.3315.4091.40114.2707.3335.0537.4525.3655.4351.40703.5716.5614.5006.5604.6324.8311.42102.9714.8433.6965.0603.7624.1581.51802.1792.7012.2392.7832.4863.1241.75701.7572.1642.3102.2262.1642.310	.4011	4.302	7.220	5.024	7.339	6.541	5.339	6.533	5.417
1.40114.2707.3335.0537.4525.3655.4351.40703.5716.5614.5006.5604.6324.8311.42102.9714.8433.6965.0603.7624.1581.51802.1792.7012.2392.7832.4863.1241.75701.7572.1642.3102.2262.1642.310	.4011	4.290	7.275	5.014	7.376		5.331		5.409
1.40703.5716.5614.5006.5604.6324.8311.42102.9714.8433.6965.0603.7624.1581.51802.1792.7012.2392.7832.4863.1241.75701.7572.1642.3102.2262.1642.310	.4011	4.270	7.333	5.053	7.452		5.365		5.435
1.42102.9714.8433.6965.0603.7624.1581.51802.1792.7012.2392.7832.4863.1241.75701.7572.1642.3102.2262.1642.310	1.4070	3.571	6.561	4.500	6.560		4.632		4.831
1.51802.1792.7012.2392.7832.4863.1241.75701.7572.1642.3102.2262.1642.310	.4210	2.971	4.843	3.696	5.060		3.762		4.158
1.75701.7572.1642.3102.2262.1642.310	1.5180	2.179	2.701	2.239	2.783		2.486		3.124
	1.7570	1.757	2.164	2.310	2.226		2.164		2.310

Obviously, the other areas on the MF of the H_3 system vary accordingly.

The picture for the transition state structure, H(1.757)H(1.757)H, that has the highest energy for the linear H_3 reaction has been shown in Fig. 2h. Remarkably, the middle of the original H_2 part becomes dark red, which means that the electron density strongly decreases making its bond much weaker. This is consistent with that this transition structure has the highest energy along the reaction path. It is also seen that the electrons increase around the two end sides of the system. About this transition state structure, the H_3 linear reaction is symmetric. After this stage, the right and the middle H atoms combine while the left H atom leaves, whose MF pictures are just the images of Fig. 2a–c about the middle vertical plane.

In addition, we have also investigated the MF changing pictures for $H + H \rightarrow H_2$ reaction, which provides a comparison with the H_3 system. For an easy comparison with H_3 system, the two H atoms in H_2 system are marked by H2 and H3 atoms. When two isolated H2 and H3 atoms approach each other, the MFs have investigated. The calculated numerical values of some characteristic quantities of MICC and MFED for the chosen H_2 geometries are listed in Table 2. Figure 3 only depicts the MFs where the distances between two H atoms are in turn 7.0, 5.0, 4.314, 1.757 and 1.4009 a.u. (equilibrium geometry) with the same color bar, the scale of electron density. When the distance between two H atoms is 4.314 a.u., the two atoms touch each other and begin to form a body (contacting point).

It can be seen from Tables 1 and 2 that the contacting distance between the H2 and H3 atoms in H₂ system is 4.314 a.u., but that is 4.302 a.u. for H₂ + H system. Before they arrive at their respective contacting point, for two H atomic system, the MICCs in three directions gradually shrink as a result of a decrease in the distance between two H atoms in H₂ system, while the MICCs in three directions gradually swell as a result of a decrease in the distance between the H₂ and H parts in H₃ system.

3.3 Conversion between van der Waals interaction and chemical bonding based on PAEM-MO diagram

We have constructed the PAEM-MO diagram, i.e., the local PAEM curve between two atoms inserted with the molecule orbitals with their main atomic components, from which a rule is provided for distinguishing chemical bonding from van der Waals interaction between two atoms [59]. Here, for $H_2 + H \rightarrow H + H_2$ reactions, we just take three points on the reaction route, namely H(1.4009)H(5.0) H, H(1.4011)H(4.302)H and H(1.757)H(1.757)H, to demonstrate the conversion between van der Waals interaction and chemical bonding.

As the H atom along the linear reaction path is 5.00 a.u. far from the adjacent H atom of the H_2 molecule whose

Table 2The molecularintrinsic characteristic quantities(MICC and MFED) of the H_2 systems, distances in a.u.,electron densities in $\times 10^{-3}$ a.u

$\overline{R_2}$	<i>R_r</i> (H2)	Den _r (H2)	<i>R</i> _{<i>l</i>} (H3)	Den _l (H3)	<i>R</i> _v (H3)	Den _v (H3)
5.0	2.0760	4.957	1.9904	7.387	2.0739	5.073
4.5	2.0901	4.565	2.0176	9.975	2.0840	4.962
4.4	2.0919	4.499	2.0498	10.733	2.0834	4.958
4.35	2.0937	4.479	2.0744	11.151	2.0829	4.961
4.33	2.0942	4.470	2.1214	11.326	2.0826	4.963
4.32	2.0945	4.464	2.1455	11.415	2.0824	4.964
4.315	2.0945	4.462	2.1573	11.461	2.0823	4.965
4.314	2.0946	4.461			2.0823	4.965
4.313	2.0946	4.461			2.0823	4.965
4.31	2.0946	4.459			2.0822	4.966
4.3	2.0946	4.449			2.0817	4.965
4.2	2.0947	4.385			2.0791	4.983
4.1	2.0973	4.301			2.0773	5.113
4.0	2.0953	4.268			2.0736	5.137
3.5	2.0605	4.363			2.0352	5.532
3.0	1.9881	4.746			1.9661	6.468
2.5	1.8945	5.182			1.8675	8.251
2.0	1.7452	6.728			1.7631	10.886
1.4009	1.5089	10.575			1.6409	15.561



Fig. 3 3D MF of the H_2 molecule at the nuclear arrangement. The nuclear separation of H(7.0)H, H(5.0)H, H(4.3314)H, H(1.757)H, H(1.4009)H (equilibrium) of the H + H \rightarrow H₂ reaction, respectively

nuclear separation is 1.4009 a.u., named H(1.4009)H(5.0) H, the PAEM-MO diagram is described in Fig. 5a, along the line of the three H atoms. The PAEM barrier top between two H atoms in the H_2 molecule is -1.9127 hartree, while the PAEM barrier top between isolated H atom and the adjacent H atom of the H_2 molecule is -0.4097hartree. The 3 electrons of the H₃ system distribute 2 molecular orbitals, 1σ and 2σ , whose energies are -0.5971and -0.4995 hartree, respectively. The electrons of 1σ MO are dominantly located on the H-H chemical bond of the H_2 molecule with bonding character, and one electron in 2σ MO is dominantly located on the H₃ atom. The 1σ and 2σ in energy are higher than the barrier top of -1.9127 hartree between H1 and H2 atoms, while they are lower than the barrier top -0.4097 hartree, which implies that is a chemical bonding between the H1 and H2 atoms, while the interaction between H2 and H3 atoms belongs to a van der Waals interaction.

As shown in Fig. 2d, when the distance between H_2 molecule and H atom is 4.302 a.u., denoted by H(1.4011) H(4.27)H, there is one touching point between the H_2 molecule and H atom along the H_3 reaction path, based on their MF. For this critical point, the PAEM-MO diagram is correspondingly depicted in Fig. 4b where the PAEM barrier top between two H atoms in the H_2 molecule is -1.9174 hartree, while the PAEM barrier top between the coming H atom and the adjacent H atom of the H_2 molecule




Fig. 4 The PAEM-MO diagrams along the nuclear line of the energy-favorite linear $H_2 + H \rightarrow H + H_2$ reaction system at the nuclear arrangement. **a** From the left to the right the H–H distances are, respectively, 1.4009 and 5.0 a.u., denoted as H(1.4009)H(5.0)H. **b** At the nuclear separation of H(1.4011)H(4.302)H. **c** The structure of transition state H(1.757)H(1.757)H

is -0.4958 hartree. The three electrons of the H₃ system distribute two molecular orbitals, 1 σ and 2 σ , whose energies are -0.6015 and -0.4968 hartree, respectively. The electrons of 1 σ MO are dominantly located on the H₁-H₂ region, and one electron in 2 σ MO is dominantly located on the H3 atom. The 1 σ and 2 σ in energy are higher than the barrier top of -1.9174 hartree, while 1 σ MO in energy is



Fig. 5 The PAEM-MO diagrams of H_2 system at the nuclear arrangement. **a** H(7.0)H, **b** H(4.314)H, **c** H(1.757)H, H(1.4009)H

lower than the barrier top of -0.4958 hartree and 2σ MO in energy (-0.4968 hartree) is almost at the same level of the barrier top (-0.4958 hartree). It is shown that the interaction between H1 and H2 atoms still belongs to the chemical bonding, but the interaction between H₂ molecule and isolated H3 atom start chemical bonding from van der Waals, in terms of PAEM-MO diagram.

As H₂ molecule gradually approaches to H atom, the distance between H1 and H2 atoms gradually enlarges and separation between H2 and H3 atoms shortens. Along the IRC of the linear H_3 reaction, the geometry H(1.757) H(1.757)H in energy is highest, being the transition state. The three H atoms have formed one body, as described by the MF in Fig. 2h. Figure 4c shows its corresponding PAEM-MO diagram where the PAEM barrier tops between two adjacent H atoms are -1.5149 hartree. The three electrons of the H₃ system occupied 1σ and 2σ MOs whose energies are -0.7065 and -0.3869 hartree, respectively. 1σ and 2σ MOs in energy in this case are higher than the PAEM barrier top (-1.5149 hartree). It is shown that the interaction between H2 and H3 has become chemical bonding. It is demonstrated from the Fig. 4a, b and c that there is conversion between van der Waals interaction and chemical bonding in the linear H₃ reaction $(H_2 + H \rightarrow H + H_2)$.

In addition, Fig. 5a–d displays the PAEM-MO diagrams of the H(7.0)H, H(4.314)H, H(1.757)H and H(1.4009) H for H + H \rightarrow H₂ reaction, respectively. Corresponding, four out of their PAEM-MO diagrams are shown in Fig. 5a–d, where the PAEM barrier tops are, respectively, -0.2925, -0.4720, -1.4628 and 1.9435 hartree, and the 1 σ MO energies are in turn -0.3099, -0.3695, -0.5442 and -0.5945 hartree. As a consequence of comparing their respective PAEM barrier and the MO energy, H(7.0)H is van der Waals interaction, while H(4.314)H, H(1.757)H and H(1.4009)H are chemical bonding.

4 Summary

The method for defining molecular intrinsic characteristic contour (MICC) and then the MF that is the MICC mapped by the electron density (MFED) is formulated. This is based on the intrinsic motion of an electron within a molecule and rigorously calculated by an accurate ab initio method. MF is a molecular turning surface for electron motion and provides a unique and intrinsic description of both the molecular boundary surface and the frontier electron density.

To sum up, some features can be abstracted for the MF change and PAEM-MO diagram for a molecular reaction system: (a) In the respect of spatial polarization, the adjacent active areas of MF regions of two reactants may stretch to welcome each other; the nearer they approach,

the larger stretching they make, and then they touch and bond together while the rest part of the reaction system may have the concerted changing. (b) In the respect of polarization of the electron cloud, during the approaching process of reactants, the electron density on the MF contour usually concentrates onto the reaction active areas to strengthen the interaction of two reactants while accordingly lessening the electron density of other areas, especially from the bondbreaking region. This may provide a pattern of recognition strategies between molecules. (c) At the critical point, two reactant MF regions have only one touching point with the largest stretching in shape and the most polarization in the electron density, particularly on the active reaction areas. It might practically be assumed that only one touching point reached by two reacting MF regions represents the starting point of forming or breaking the chemical bonding between reactants. (d) In addition, it was shown that the chemical bonding or van der Waals interaction between two adjacent atoms can be distinguished by the barrier height of the potential acting on an electron (PAEM) compared with the valence bounding MO energy levels, i.e., by the PAEM-MO diagram. (e) In most cases, the MF and PAEM-MO diagram give consistent distinction of chemical bonding from van der Waals interaction, but MF gazes at the shape, while the frontier electron density and PAEM-MO diagram focuses on the electronic structure, namely the potential, bonding and the electron interflow.

Really, we need to do more studies on the PAEM, MF and PAEM-MO diagram for more molecular systems later on.

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REGULAR ARTICLE

State-to-state quantum versus classical dynamics study of the $OH + CO \rightarrow H + CO_2$ reaction in full dimensions (J = 0): checking the validity of the quasi-classical trajectory method

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Abstract We report full-dimensional state-to-state quantum mechanical (QM) and quasi-classical trajectory (QCT) calculations on the title reaction for the ground rovibrational initial state with total angular momentum fixed at zero on the accurate potential energy (PES) constructed recently by using permutation-invariant polynomial-neural network method (Li et al. in J Chem Phys 140:044327, 2014), to check the validity of the QCT method for the reaction. It is found that the QM state-tostate results strongly depend on the resonance structures in reaction, but the collision energy-averaged results show a smooth change with the increase of collision energy. Overall, the agreement between collision energy-averaged QM and QCT state-to-state results is satisfactory, in particular at high collision energy region, indicating that the QCT method is rather accurate on describing dynamics of the reaction on the PES. On the other hand, because earlier studies revealed the QCT results on the PES do not agree very well with the experimental measurements available, more theoretical and experimental studies should be carried out to achieve a full understanding on the dynamics of this benchmark complex-forming reaction.

Keywords Complex-forming reaction · State-to-state · Quantum dynamics · Quasi-classical trajectory

Dedicated to Professor Guosen Yan and published as part of the special collection of articles celebrating his 85th birthday.

S. Liu \cdot J. Chen \cdot B. Fu \cdot D. H. Zhang (\boxtimes)

1 Introduction

Because of its crucial role in the conversion of CO to CO₂, the OH + CO \rightarrow H + CO₂ reaction is important to both atmospheric [1] and combustion chemistry [2]. This exothermic reaction proceeds via two pathways with the formation of HOCO intermediate complex in trans and cis deep wells, representing a prototype for complex-forming four-atom reactions, just as $H_2 + OH \rightarrow H + H_2O$ is for direct four-atom reactions. Over the years, a large number of experimental studies have been devoted to this reaction, both for kinetic and dynamical properties. The measured thermal rate constants are nearly independent of temperature between 80 and 500 K, but increase sharply with temperature above 500 K [3-5]. This strong non-Arrhenius behavior has been attributed to the intricate minimum energy paths (MEP), which have near-isoenergetic barriers in the entrance (OH + CO) and exit $(H + CO_2)$ channels. The molecular beam experiment carried out by Alagia et al. showed peaks both in the forward and backward directions, indicating the existence of reasonable long-lived intermediate species [6, 7].

Theoretically, the OH + CO reaction presents a great challenge to both potential energy surface (PES) construction and quantum dynamics. Much of the earlier work on the development of the global PES has been pioneered by Schatz and coworkers [8–10], by using a relatively small number of ab initio points. Following that, a few better PESs have been constructed and widely used, such as the YMS, LTSH, and VvHK PES [11–13], but they were still not sufficiently accurate for dynamical studies. The situation has been greatly improved, thanks to new PESs based on large numbers of high-level ab initio points. In 2012, Li et al. developed a global PES [14] and its modified version [15] using the permutation-invariant polynomial (PIP)

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method [16]. Subsequently, Chen et al. reported another PES using the neural network (NN) method [17], with data points spreading in a more complete configuration space [18]. It was shown that the NN PES fits the ab initio points much better than the PIP PES. Very recently, Li et al. reported a new full-dimensional global PES based on the same ab initio data set generated by Chen et al. [18], but fit using the newly proposed permutation-invariant polynomial–neural network (PIP–NN) method [19, 20], which can be considered as the most accurate PES so far for the title reaction [21].

Because of the presence of three heavy atoms and a relatively long-lived intermediate complex, the rigorous quantum scattering calculations of this reaction is difficult. Extensive quantum mechanical (QM) studies were only carried out for the total reaction probabilities and rate constants on different PESs [21-26]. In 2011, we performed the first full-dimensional state-to-state quantum dynamics calculation for total angular momentum J = 0 on LTSH PES for the title reaction, for the ground and two vibrationally excited initial states [27, 28]. It was found that the initial CO vibrational excitation essentially has the same effect on the product energy partition as the reagent translational excitation, while the initial OH excitation leads to slightly more internal energy of CO₂. However, because the total reaction probabilities for J = 0 obtained on the LTSH PES were found to be substantially larger than those on the more accurate NN [18] and PIP-NN [21] PESs, state-to-state dynamics calculations on the NN or PIP-NN PES are highly desired to theoretically study product state distribution of this important reaction.

Although the quantum wave packet methods have been developed to compute differential cross sections (DCS) for direct four-atom reactions, such as the $HD + OH \rightarrow H_2O + D$ [29, 30], which achieved excellent agreement with high-resolution crossed-molecular beam experiments, it is still not feasible at present to compute final state resolved cross sections for complexforming four-atom reactions due to the long wave packet propagation time and huge basis sets required. Quasi-classical trajectory (QCT) calculations of the title complexforming reaction were carried out extensively on different PESs [12, 15, 21, 31-34]. Detailed dynamics features, i.e., DCS, product translational energy, vibrational and rotational state distributions, as well as the excitation functions and rate constants were calculated and compared with available experimental results. The recent QM and QCT study on the new PIP-NN PES indicated better agreement with both experimental kinetic and dynamical data than on previous PESs, but the differences still remain between QCT and experimental DCS [21]. Therefore, it is important to test the validity of the QCT method to describe chemical reaction dynamics for the title reaction.

In this article, we perform the full-dimensional (6D) state-to-state QM and QCT studies from the ground rovibrational initial state for the OH + CO \rightarrow H + CO₂ reaction on the new PIP–NN PES. Both QM and QCT calculations were carried out at the total angular momentum (impact parameter) fixed at zero. The rest of this paper is organized as follows. In Sec. II, we present the theory of the full-dimensional state-to-state quantum dynamics treatment to diatom–diatom reactions and some computational details. Sec. III contains our results, including QM and QCT total reaction probabilities, product distributions and energy partitioning information for this reaction. In Section IV we summarize our conclusions.

2 Theory

The QM calculations were carried out using the multistep reactant-product decoupling (MRPD) scheme based time-dependent wave packet (TDWP) method [30, 35– 37]. The basic strategy of the RPD scheme is to partition the full time-dependent (TD) wavefunction into a sum of reactant component (Ψ_r) and all product components (Ψ_p , p = 1, 2, 3, ...) that satisfy the following decoupled equations,

$$i\hbar\frac{\partial}{\partial t}|\Psi_{r}(t)\rangle = H|\Psi_{r}(t)\rangle - i\sum_{p}V_{p}|\Psi_{r}(t)\rangle$$

$$i\hbar\frac{\partial}{\partial t}|\Psi_{p}(t)\rangle = H|\Psi_{p}(t)\rangle + iV_{p}|\Psi_{r}(t)\rangle,$$
(1)

where $-iV_p$ is the negative imaginary potential (absorption potential) employed to prevent the wavefunction $\Psi_r(t)$ from entering the *p*th product arrangement. The solution for $\Psi_r(t)$ can be propagated in the reactant Jacobi coordinates just as for total reaction probability calculations,

$$\Psi_r(t+\Delta) = e^{-V_p \Delta/\hbar} e^{-(i/\hbar)H\Delta} \Psi_r(t).$$
(2)

Every product wave function, Ψ_p , can be propagated in its own coordinates as in a normal wave packet propagation except for a source term, ξ_p , prepared by Eq. 2,

$$\Psi_p(t+\Delta) = e^{-(i/\hbar)H\Delta}\Psi_p(t) + \xi_p(t) = e^{-(i/\hbar)H\Delta}\Psi_p(t) + \left(1 - e^{-V_p\Delta/\hbar}\right)\Psi_r(t+\Delta).$$
(3)

In the original RPD approach, the source term $\xi_p(t) = (1 - e^{-V_p \Delta/\hbar}) \Psi_r(t)$ is saved and transformed from reactant to product Jacobi coordinates at every propagation time step. To reduce the computational cost, we use MRPD by saving and transforming the source term at every *M* time steps in the actual calculation,



Fig. 1 The reagent Jacobi coordinates $(R, r_1, r_2, \theta_1, \theta_2, \varphi)$ for the AB + CD diatom-diatom arrangement, and product Jacobi coordinates $(R', r'_1, r'_2, \theta'_1, \theta'_2, \varphi')$ for the A + BCD atom-triatom arrangement. Δ is the angle between \hat{R} and \hat{R}'

$$\Psi_{r}(t+\Delta) = e^{-V_{p}M\Delta/\hbar}e^{-(i/\hbar)H\Delta}\Psi_{r}(t),$$

when $mod((t+\Delta) - t_{0}, M\Delta) = 0,$
$$\Psi_{p}(t+\Delta) = e^{-(i/\hbar)H\Delta}\Psi_{p}(t) + \left(1 - e^{-V_{p}M\Delta/\hbar}\right)\Psi_{r}(t+\Delta),$$

(4)

where t_0 is the starting point for performing wave function transformation. At other time steps, we carried out the standard split-operator propagation for $\Psi_r(t)$ and $\Psi_p(t)$ without the absorption potential, V_p , related terms. In this way, we can cut the computational time for wave function transformation from reactant coordinates to product coordinates by a factor of M. Finally, one can extract the final state information, such as state-to-state S matrix elements or reaction probabilities, from the Fourier transformation of $\Psi_p(t)$.

The coordinate systems and wavefunctions representations for both diatom-diatom and atom-triatom scattering studies are outlined here. For details, please refer Refs. [38, 39]. Figure 1 shows the reagent Jacobi coordinates $(R, r_1, r_2, \theta_1, \theta_2, \varphi)$ for the OH + CO diatomdiatom arrangement, and product Jacobi coordinates $(R', r'_1, r'_2, \theta'_1, \theta'_2, \varphi')$ for the H + CO₂ atom-triatom arrangement. As can be seen r_2 and r'_2 share the same vector.

The system Hamiltonian in the OH + CO arrangement for a given total angular momentum *J* can be written as

$$H = -\frac{\hbar^2}{2\mu} \frac{\partial^2}{\partial R^2} + \frac{(\mathbf{J} - \mathbf{j}_{12})^2}{2\mu R^2} + \frac{\mathbf{j}_1^2}{2\mu_1 r_1^2} + \frac{\mathbf{j}_2^2}{2\mu_2 r_2^2} + V(R, r_1, r_2, \theta_1, \theta_2, \varphi) + h_1(r_1) + h_2(r_2),$$
(5)

where μ is the reduced mass between the center of mass of OH and CO, **J** is the total angular momentum operator, \mathbf{j}_1 and \mathbf{j}_2 are the rotational angular momentum operators of OH and CO, which are coupled to form \mathbf{j}_{12} . The diatomic reference Hamiltonian $h_i(r_i)$ (i = 1, 2) is defined as

$$h_i(r_i) = -\frac{\hbar^2}{2\mu_i} \frac{\partial^2}{\partial r_i^2} + V_i(r_i).$$
(6)

The time-dependent wave function can be expanded in terms of the translational basis of R, the vibrational basis $\phi_{v_i}(r_i)$, and the BF rovibrational eigenfunction as

$$\Psi_{\nu_{0}j_{0}K_{0}}^{JM\epsilon}(\mathbf{R},\mathbf{r_{1}},\mathbf{r_{2}},t) = \sum_{n,\nu,j,K} F_{n\nu_{j}K,\nu_{0}j_{0}K_{0}}^{JM\epsilon}(t)u_{n}^{\nu_{1}}(R)\phi_{\nu_{1}}(r_{1})\phi_{\nu_{2}}$$
$$\times (r_{2})\mathscr{Y}_{jK}^{JM\epsilon}(\hat{R},\hat{r}_{1},\hat{r}_{2}).$$
(7)

The BF total angular momentum eigenfunctions can be written as,

$$\mathscr{Y}_{jK}^{JM\epsilon} = (1+\delta_{K0})^{-1/2} \sqrt{\frac{2J+1}{8\pi}} \Big[D_{K,M}^{J*} Y_{j_1j_2}^{j_{12}K} + \epsilon (-1)^{j_1+j_2+j_{12}+J} D_{-K,M}^{J*} Y_{j_1j_2}^{j_{12}-K} \Big]$$
(8)

$$\equiv \mathscr{D}_{K,M}^{J*} Y_{j_{1}j_{2}}^{j_{12}K} + \epsilon (-1)^{\tilde{j}} \mathscr{D}_{-K,M}^{J*} Y_{j_{1}j_{2}}^{j_{12}-K}$$
(9)

where $D_{K,M}^{J}(\Theta \Phi \Psi)$ is the Wigner rotation matrix [40] with three Euler angles $(\Theta \Phi \Psi)$, ϵ is the parity of the system defined as $\epsilon = (-1)^{j_1+j_2+L}$ with *L* being the orbital angular momentum, and $Y_{j_1j_2}^{j_12K}$ is the angular momentum eigenfunction of **j**₁₂,

$$Y_{j_1j_2}^{j_{12}K} = \sum_{m_1} \langle j_1 m_1 j_2 K - m_1 | j_{12} K \rangle y_{j_1m_1}(\theta_1, 0) y_{j_2K-m_1}(\theta_2, \phi)$$
(10)

where y_{jm} are spherical harmonics.

For the $H + CO_2$ arrangement, similarly to the discussion about OH + CO arrangement, we have the same forms of equations, except the quantities with unprimed indices replaced by primed ones. The system Hamiltonian for the $H + CO_2$ arrangement, corresponding to Eq. (5),

$$H = -\frac{\hbar^2}{2\mu'}\frac{\partial^2}{\partial R'^2} + \frac{(\mathbf{J} - \mathbf{j'}_{12})^2}{2\mu' R'^2} + \frac{\mathbf{j'}_1^2}{2\mu'_1 r'_1^2} + \frac{\mathbf{j'}_2^2}{2\mu'_2 r'_2^2} + \frac{$$

where μ' is the reduced mass between the center-of-mass of H and CO₂, $\mathbf{j'}_{12}$ is the total angular momentum operator of CO₂, $\mathbf{j'}_2$ is the rotational angular momentum operator of CO, and $\mathbf{j'}_1 = \mathbf{j'}_{12} - \mathbf{j'}_2$ is the orbital angular momentum of CO₂. The reference vibrational Hamiltonian $h_i(r'_i)$ (i = 1, 2) is defined as

$$h_i(r'_i) = -\frac{\hbar^2}{2\mu'_i} \frac{\partial^2}{\partial r'_i^2} + V_i(r'_i).$$
 (12)

Corresponding to the Eq. 7, we have

$$\Psi_{i}^{JM\epsilon}(\mathbf{R}',\mathbf{r}_{1}',\mathbf{r}_{2}',t) = \sum_{n',\nu',j',K'} F_{n'\nu'j'K',i}^{JM\epsilon}(t) u_{n'}^{\nu'_{1}}(R')\phi_{\nu'_{1}}(r_{1}')\phi_{\nu'_{2}} \times (r_{2}')\mathscr{Y}_{j'K'}^{JM\epsilon}(\hat{R}',\hat{r}_{1}',\hat{r}_{2}'),$$
(13)

It should be pointed out here that the functional form for the product arrangement basis (primed) are different with the one for the reagent arrangement basis (unprimed).

The BF total angular momentum basis $\mathscr{Y}_{j'K'}^{JM\epsilon}(\hat{R'}, \hat{r'}_1, \hat{r'}_2)$ in Eq. 13 are defined as,

$$\mathscr{Y}_{j'K\hat{a}L^{TM}}^{JM\epsilon} = (1 + \delta_{K'0})^{-1/2} \sqrt{\frac{2J+1}{8\pi}} \Big[D_{K'M}^{J*} Y_{j_1j_2}^{j_{12}K'} + \epsilon (-1)^{j_1'+j_2'+j_{12}'} D_{-K'M}^{J*} Y_{j_1j_2}^{j_{12}-K'} \Big]$$
(14)

$$\equiv \mathscr{D}_{K'M}^{J*} Y_{j_1' j_2'}^{j_{12}'K'} + \epsilon (-1)^{\tilde{j}'} \mathscr{D}_{-K'M}^{J*} Y_{j_1' j_2'}^{j_{12}'-K'}$$
(15)

 ϵ is the total parity of the system defined as $\epsilon_0(-1)^J$ with ϵ_0 being the parity of the system. $Y_{j_1j_2}^{j_1'Z'}$ is the angular momentum eigenfunction of \mathbf{j}'_{12} defined as,

$$Y_{j'_{1}j'_{2}}^{j'_{12}K'} = \sum_{m} D_{K'm}^{j'_{12}*}(\chi,\theta_{1}',\varphi') \sqrt{\frac{2j'_{1}+1}{4\pi}} \langle j'_{2} m j'_{1} 0 | j'_{12}m \rangle y_{j'_{2}m}(\theta_{2}',0)$$
(16)

and y_{jm} are spherical harmonics. Note that in Eq. 15, the restriction $\epsilon (-1)^{j'_1+j'_2+j'_{12}} = 1$ for K' = 0 partitions the whole rotational basis set into even and odd parities. Thus a K' = 0 initial state can only appear in one of these two parity blocks. For K' > 0, however, there is no such restriction, the basis set is the same for even and odd parities. Hence a K' > 0 initial state can appear in both parities.

We used the same parameters in Ref. [28], except for a larger asymptotic region with 60 sine basis functions for the R' from 1.0 to 13.0 a_0 in continuous propagation atom-triatom coordinates. A total number of 45,000 time steps with a time increment of 10 a.u. were used to accomplish the wave packet propagation from reagent channel to product channel. To minimize the computational cost, the coordinate transformation, which transfers the no-return part of reacted wave packets from the diatom-diatom coordinates to atom-triatom coordinates, was carried out at every 12 time steps. Even so, it is very time consuming to carry out the wave packet propagation in the reactant and product arrangements and to make the coordinate transformation.

The QCT calculations used essentially the same method as in the previous work [21]. Briefly, reactive scattering at the collision energy below 0.4 eV was simulated using the VENUS code [41]. The initial OH and CO reactants in their respective ground rovibrational states are separated by 6.0 Å and the trajectories are terminated when products (or reactants for non-reactive trajectories) reached a separation of 6.0 Å. The propagation time step was selected to be 3 a.u., and exceptionally long trajectories were halted if the propagation time reached a pre-specified value (36 ps). The scattering parameters (vibrational phases, and spatial orientation of the initial reactants) were selected with a Monte Carlo approach. The gradient of the PES was obtained numerically by the central-difference algorithm.

3 Results

Figure 2 shows the OM total reaction probabilities for J = 0 ground initial state on the PIP–NN PES from the summation of the state-to-state reaction probabilities obtained by using the RPD approach, in comparison with those from the initial state selected wave packet (ISSWP) approach only using reactant Jacobi coordinates. As can be seen, the agreement between the total reaction probabilities from RPD and ISSWP calculations is very good, indicating the RPD partition of the wave function as well as the continuous propagation in atom-triatom coordinates is of high accuracy. The QCT reaction probabilities for some collision energies on the same PES are represented by red dots, which show the same trend as the OM curve, although many narrow but overlapping resonances exist on QM probabilities apparently due to the long-lived HOCO complex. Obviously, the reaction probabilities are quite small, indicating this bimolecular reaction is very inefficient, despite its low energy barriers along the MEP. The present QM probabilities are substantially different from those obtained on the LTSH PES, not only for the magnitude of the reaction probabilities but also for the position of narrow resonance peaks. Based on the accuracy of the ab initio method used in generating energy points for the PIP-NN PES and the extremely small fitting errors [21], the current QM probabilities on the PIP-NN PES should be much more reliable as compared to the results on the LTSH PES.

Figure 3 shows the CO_2 product vibrational state distributions at $E_c = 0.1$ and 0.4 eV obtained by the QM calculation on the PIP-NN PES. For CO₂, because the symmetric stretching vibrational frequency v_1 is very close to the double of the bending frequency v_2 , there are strong Fermi resonances between levels (100) and (020), and their corresponding overtones. In addition, with the increase of CO₂ rotational angular momentum, the number of vibrational states proliferates rapidly. Therefore, it is extremely hard to make a complete assignment of all the vibrational states, and we only managed to assign some of the vibrational states with low excitation energies. As seen from Fig. 3a at $E_c = 0.1$ eV, the CO₂ products are dominantly excited in bending and symmetric modes, but essentially have no excitation in the antisymmetric stretching mode. It is interesting that quite large population is seen from the



Fig. 2 Total reaction probabilities for J = 0 ground initial state on the PIP–NN PES, including QM results obtained from state-to-state RPD and ISSWP calculations, and QCT results for some collision energy, in comparison with QM probabilities on the LTSH PES



Fig. 3 Product vibrational state distributions of CO₂ for total angular momentum J = 0 at $E_c = 0.1$ (a) and 0.4 eV (b) obtained by the QM calculation

simultaneous excitations of combined bending and symmetric modes, such as (110), (12²0) and (210). With the increase of collision energy, higher vibrationally excited states of CO₂ become populated as shown in Fig. 3b. The largest population for a single vibrational state decreases substantially, from 11 % at $E_c = 0.1$ eV to merely 3.5 % at $E_c = 0.4$ eV. Contrary to highly bending and symmetric



Fig. 4 Product vibrational state distributions below 0.6 eV for total angular momentum J = 0 at $E_c = 8.6$ kcal/mol obtained by adding the sublevels of each bending mode together

stretching excitation, the population of the antisymmetric stretching states, even with one quantum number, is still small at $E_c = 0.4 \text{ eV}$. Because the CO bond is substantially excited vibrationally in the complex wells as found in Ref. [25], it clearly reflects the fact that the reaction system has to put much of the available energy on the reaction coordinate to overcome the exit channel barrier. In the mean time, the substantial excitation of bending vibration are consistent with the geometry of the exit channel cis-H-OCO transition state, where O-C-O has an angle of 157°. The excitation in the symmetric stretching modes can likely be attributed to the Fermi-link with bending mode. The vibrational distributions on the LTSH PES in Ref. [27] are less excited than the present results. At $E_c = 0.1 \text{ eV}$, the LTSH distribution peaks at (000) state with 36 % population, and is limited to a small number of states with energy below 0.25 eV. The peak moves slightly to $(02^{2}0)$ state at $E_c = 0.4$ eV. We also found more populations with antisymmetric stretching excited on the LTSH PES. As we have discussed in Ref. [18], in contrast to the tight exit cis-H-OCO transition state on the PIP-NN PES, the exit channel of LTSH PES is much wider, resulting in larger probabilities for the complex decaying into product channel and larger total reaction probabilities. And the bending angle of O-C-O changes slower to the linear product on the LTSH PES, leading to the cooler vibrational excitation patterns.

The QCT product vibrational state distribution at $E_c = 8.6$ kcal/mol for this reaction has been determined using a normal-mode analysis method on PIP (CCSD-2/d) PES in an earlier work, as shown in Fig. 3 of Ref. [33]. To compare with that, we added the sublevels of each bending mode together in QM distribution. Because of the difficulty



Fig. 5 Normalized product CO_2 rotational state distributions for total angular momentum J = 0 at some collision energy, obtained by QCT (a) and QM (b) calculations

on assigning vibrational states in high energy region, Fig. 4 only includes the treated QM distribution below 0.6 eV. Although the QCT and QM calculations were carried out on different PESs and different impact parameters, the trends of the two distributions are quite similar: the populations in each group (distinguished by symmetric stretching mode) firstly increase and then decrease with the increase of bending quantum number in general, despite the slight differences in the population of a single state and the position of peaks. However, the ratio of symmetric stretching excitations in the QM distribution is larger than that in the QCT one.

Figure 5 shows the normalized CO₂ rotational state distributions at some collision energies, with all vibrational states included. As seen, the reaction leads to substantial rotational excitations. Overall, the two sets of distributions calculated by QCT and QM methods are similar except some details. With the increase of collision energy, the QCT distribution becomes slightly broader, with the peak nearly fixed at the rotational angular moment of roughly 20. In contrast, the variation of QM distributions with the collision energies is more complicated. The behavior of QM rotational state distribution at $E_c = 0.1$ eV is similar to that of QCT results, except slightly higher peak for the QM distribution. However, as the collision energy increases, the QM distribution turns to show oscillatory structures which can be attributed to resonances during the reaction.

To further investigate this, we depict in Fig. 6a the QM rotational state distributions, respectively, at two pairs

Theor Chem Acc (2014) 133:1558



Fig. 6 Normalized QM product CO₂ rotational state distributions for total angular momentum J = 0, at peak and dip positions of total probability (a), and after Gaussian function-weighted averaging (b)

of collision energies (0.122/0.132 and 0.312/0.318 eV), where the total reaction probability exhibits a pronounced peak and a dip nearby as shown in Fig. 2. The behavior of these distributions is quite different, indicating the QM rotational state distributions depend strongly on the reaction resonances. To eliminate the influence of resonances, we averaged the QM populations using a Gaussian function weighting for each collision energy. As a result, the resonance-smoothing QM distributions shown in Fig. 6b are in very good agreement with QCT results. In addition, the rotational state distributions shown in Fig. 5b are quite different from those on the LTSH PES in Ref. [27], where the CO₂ products were shown to be more rotationally excited as the collision energy increases. The present PES is substantially more accurate and thus the current dynamics results are more reliable. The small variation of rotational state distributions with collision energy in this study reflects the fact that the HOCO intermediate complex lives sufficiently long in the wells in the collision energy region we studied.

Figure 7a, b compare the normalized product translational energy distributions on the PIP–NN PES calculated by QM and QCT methods at the collision energies of 0.1 and 0.4 eV, respectively. The zero point energy (ZPE) problem is an important issue in QCT calculations. As for this reaction, some CO_2 products can emerge with the vibrational energies below its ZPE level. Hence, we have shown the QCT results without (blue curve) and with ZPE constraint (magenta curve). The ZPE constraint results are Fig. 7 Normalized product translational energy distributions on the PIP–NN PES at $E_c = 0.1$ (a) and 0.4 eV (b) obtained by both QM and QCT calculations, in comparison with QM results on the LTSH PES (c, d). The maximum translational energy determined by the ZPE level of CO₂ is indicated by the *arrow*



analyzed for those trajectories in which CO₂ has at least the corresponding ZPE. It can be seen that the ZPE constraint QCT distributions are similar to the original QCT distributions, except at high translational energy region. At $E_c = 0.1$ eV, the QM distribution is lower in translational energy by about 0.1 eV than the QCT distributions, indicating the QM calculation yields a higher internal energy distribution. This discrepancy possibly results from the tunneling and ZPE effects that QCT calculation does not consider in the reaction. While $E_c = 0.4$ eV, the agreement between the QM and QCT distributions are quite good, except that the QCT distributions slightly overestimate populations at high translational energy region.

To investigate the influence of the reactant translational energy on the product translational energy, we show the QM product translational energy distributions on the PIP-NN PES at $E_c = 0.1$ and 0.4 eV in Fig. 7(c), together with the previous QM results on the LTSH PES in Fig. 7(d). On the whole, the two batches of results on the two different PESs are rather different. At $E_c = 0.1 \text{ eV}$, the product translational energy obtained on the PIP-NN PES peaks roughly at 0.72 eV, whereas the peak of translational energy distribution on the LTSH PES nearly reaches the maximum of available translational energy, in accord with the previous calculation that only the CO₂ vibrational states with low excitation energies are mainly populated [27]. With the increase of collision energy, the product translational energy distribution on the PIP-NN PES shifts substantially to higher energy, although more rovibrational states are populated as discussed above, indicating the reactant collision energy transforms substantially into the translational motion of products. Similar trend is seen from the



Fig. 8 The fraction of the total available energy in product channel going into internal rovibrational motion of CO_2 as a function of collision energy, together with the QCT results on the same PES and QM result on the LTSH PES

product translational energy distributions on the LTSH PES in Fig. 7(d).

Figure 8 shows the fraction of the total available energy in product channel (with respect to the ground state CO_2) going into internal rovibrational motion of the product as a function of collision energy from the QM and QCT calculations. It can be seen that although the QM curve fluctuates rapidly just like the total reaction probability shown in Fig. 2, the average value increases steadily with the collision energy from 28 % at $E_c = 0.1$ eV to 37 % at $E_c = 0.4$ eV. The QCT fractions for some collision energies on the same PES and their ZPE constrained results are represented by blue and magenta dots, with the latter agreeing well with the QM results except in low energy region. As we know the partial wave average will wash out most oscillatory structures exhibited in the total reaction probability as well as in the fraction of the total available energy going into CO₂ rovibrational excitation, and thus the QM energy partition will be much smoother if seen from the integral cross section, which results in better agreement with the QM and OCT result. The OM results calculated on the LTSH PES, indicated by red curve, underestimated this fraction in the entire energy region considered here, implying that the total available energy goes into the translational motion of the product more efficiently. At $E_c = 8.6$ kcal/mol, the QM fraction of the available energy released into product translation, $\langle f_T \rangle$, is 0.618 on the PIP–NN PES. It is lower than both the molecular beam experimental value of 0.70 [7], and the QM value of 0.738 on the LTSH PES, suggesting more excitation in the internal degrees of freedom of CO₂ on the PIP–NN PES. Furthermore, if we assume the CO2 rotational constant does not change substantially for various vibrational states, we can get the approximate fraction of energy going into product rotation, only 1.72 % on the PIP–NN PES at $E_c = 8.6$ kcal/mol due to the tiny rotational constant of CO₂, much smaller than the QCT value of 9.36 % from VENUS code. However, the VENUS code also provided an uncertainty in rotational energy close to the rotational energy itself, indicating that QCT result given by VENUS code excessively overestimated product rotational energy by including part of vibrational energy in bending modes in particular when CO₂ vibrates close to linear geometry. The QCT fraction can be reduced to 3.92 % by using the minimum rotation energy of each trajectory instead of the average value, but is still larger than the quantum result. It is worthwhile to point out that our QM and QCT calculations are for J = 0, while rotational energy in product may increase with the increase of J.

4 Conclusions

We performed full-dimensional (6D) state-to-state QM and QCT calculations for the ground rovibrational initial state for the OH + CO \rightarrow H + CO₂ reaction on the new PIP–NN PES, with the total angular momentum (impact parameter) fixed at zero. It is found that the QM rovibrational distributions of CO₂ strongly depend on the resonance structures of the reaction, but the collision energy-averaged distributions show a smooth change with the increase of collision energy. It turns out that the vibrational excitations in CO₂ bending and symmetric stretching modes are

quite substantial and increase with the increase of collision energy. Very little excitation in CO_2 antisymmetric stretching states is observed. The rotational distributions of CO_2 do not change substantially with the collision energy, peaking at angular momentum of about 20. Overall, the agreement between collision energy-averaged QM and QCT state-to-state results is satisfactory, in particular at high collision energy region, indicating that the QCT method is rather accurate on describing dynamics of this complexforming reaction on the PIP–NN PES.

On the other hand, since the QCT results on this PIP– NN PES do not agree very well with the experimental measurements available [21], the present PIP–NN PES may not be sufficiently accurate for the dynamics studies of the reaction at the state-to-state level. More efforts therefore should be devoted to the construction of more accurate PESs for this benchmark complex-forming reaction. In the meanwhile, because the PIP–NN PES was fitted very well to such a large number of ab initio energy points calculated at a high-level theory [18], it is highly desirable to carry out more experimental investigations on the reaction to eliminate possible uncertainties from experiment.

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REGULAR ARTICLE

NO adsorption and transformation on the BaO surfaces from density functional theory calculations

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Abstract Density functional theory combined with embedded cluster model calculations have been used to investigate the NO adsorption and transformation reactions on the BaO(100) surfaces. NO is found to adsorb on the anion sites to form a NO_2^{2-} species, which can then couple with another NO to form a $N_2O_3^{2-}$ species. These surface species provide an alternative explanation for the infrared bands that were used to be assigned to the nitrite/ nitrate and hyponitrite species. The calculations suggest a large intrinsic barrier for the transformation from $N_2 O_3^{2-1}$ to $N_2O_2^{2-}$. The latter species acts as a chemisorbed N_2O_2 , which is envisioned as a key intermediate for further NO reduction. The present study provides a detailed description at the molecular level for the NO/BaO(100) system, which shed some light on the NO_x storage-reduction systems, as well as NO direct decomposition.

Keywords NO_x storage and reduction $\cdot BaO \cdot NO_x$ trap \cdot Lean-burn $\cdot DFT \cdot Vibrational properties$

Dedicated to Professor Guosen Yan and published as part of the special collection of articles celebrating his 85th birthday.

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1 Introduction

The need for fuel economy and consequently a reduction in CO₂ emissions for environmental protection has brought about the lean-burn engine technology with high air/fuel ratio for motor vehicles. In such a circumstance, however, the conventional three-way catalysts (TWC) in removing engine exhausts, notably CO, hydrocarbons (HC) and nitrogen oxides (NO_x), are no longer effective in removal of NO_x. One promising solution to this problem is to use the so-called NO_x storage-reduction (NSR) catalyst [1-5]. The concept of NSR is that NO_x is first stored under the lean condition over the catalyst, which is regenerated during the short periods of rich conditions, where NO_x is released and subsequently reduced with HC, H₂ and CO to form N₂, H₂O and CO₂ [6-8]. A typical NSR catalyst consists of noble metals (generally Pt) for the purposes of NO oxidation/reduction, and materials with Lewis basicity (notably a metal oxide like BaO) for the purposes of NO_x storage, which are dispersed on a high surface area support like γ -Al₂O₃ [1, 6–8]. To further improve the performance of the catalysts in regard of NO_x storage ability, temperature stability, regeneration times, and sulfur resistance, a good knowledge about the storage and transformation mechanisms is vital.

Despite the fact that huge efforts have been dedicated to kinetic measurements and spectroscopic characterizations [e.g., 1–31], certain key aspects concerning the mechanistic features of the NSR processes remain elusive, such that no clear understanding of this catalytic system has emerged [1, 5]. The difficulty arises not only from the variations of the operation conditions, but also from the complexity of both the reactant gas mixture (i.e., NO, NO₂, O₂, H₂O, CO₂, etc.) and the catalyst material (e.g., Pt/Ba/ γ –Al₂O₃) with the abundance of the different adsorption and reaction sites. To

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build a comprehensive understanding of the mechanism and to set up a predictive kinetic model concerning how different reactant gases adsorb, react and even compete against each other for active sites, and how different components of catalyst materials interplay to enhance or depress the reaction activity, model catalysts and model reactant gases have to be employed to individually probe the adsorption and chemistry on different sites [10–14, 17–28]. Along this line, the present work focuses only on NO, which is the main form of NO_x in exhaust gas, adsorbed on and reacted with BaO, which is the primary storage material, with embedded cluster model calculations using density functional theory. The results shall be best compared with the experiments concerning NO adsorption and reactions on BaO surfaces by surface chemistry approaches [12, 13, 15, 23, 26].

In a NSR catalytic system, it is generally believed that NO is first oxidized to NO_2 on noble metal sites,

$$NO + \frac{1}{2}O_2 \stackrel{Pt}{\rightleftharpoons} NO_2 \tag{1}$$

which is then stored on BaO as nitrites (NO_2^{-}) and nitrates (NO_3^{-}) [6–11]. However, such a conversion was shown to be kinetically limited at lower temperature and thermodynamically limited at higher temperature [1, 11]. For example, Muncrief et al. [11] reported a <30 % steady-state NO to NO₂ conversion at temperatures below 250 °C and above 450 °C, with a maximum conversion of 60 % around 350 °C on a Pt/Ba/Al₂O₃ catalyst for a gas feed of 500 ppm NO and 5 % O₂ flowing at 200 cm³/min. Hence, understanding the NO adsorption and transformations is an important issue in unraveling the NO_x storage mechanism.

In experiments, the ad-species of NO exposure on barium-containing catalysts is complicated and controversial. The NO storage on monolith experiments showed that negligible conversion of NO to NO2 occurred with the Ptfree catalyst, and catalysts with >1 % Pt loading had significantly improved the trapping efficiencies [21]. These experiments suggested the importance of a Pt-Ba couple and associated spillover processes that enhanced the storage [10, 18, 21]. Temperature-programmed desorption (TPD) experiments were carried by Mahzoul et al. [19], which also confirmed the role of platinum. Nevertheless, they showed that NO storage was already significant even if no oxygen was present in the gas phase and even in the absence of platinum [19]. Hence, their experiments suggested that BaO alone is reactive toward pure NO. The X-ray photoelectron spectroscopy (XPS) has been used to characterize the ad-species of NO on a 15-20 Å BaO film deposited on the aluminum substrate [12], where the influence of the substrate was actually minimized. The nitrogen and oxygen core level spectra acquired following NO exposure strongly suggested that new ad-species had formed, which were primarily ascribed as a nitrite-like species (NO_2^{-}) , accompanied by the formation of a small amount of nitrate-like ad-species (NO₃⁻) [12]. These results were, however, not supported by a later XPS and TPD investigation on BaO layers deposited on Cu(111) surfaces [23], which concluded that BaO itself did not readily trap NO. A systematic study of the Pt/Ba/Al₂O₃ system was performed by Prinetto et al. [13]. The evolutions of the surface species were monitored with FT-IR where Pt/Ba/Al₂O₃, Pt/Al₂O₃, BaO/Al₂O₃, and Al₂O₃ were exposed independently to NO or NO2 in O2. A couple of sharp bands at 1,375 and 1,310 cm⁻¹, a complex absorption at 1,050–950 cm^{-1} with maximum at 1,025 cm^{-1} and bands of minor intensity at 1,600, 1,470, and 1,220-1,180 cm⁻¹ were observed upon NO adsorption on either Ba/Al₂O₃ or Pt/Ba/Al₂O₃ [13]. Regardless of the presence of platinum or not, there was little difference on the IR spectra for these two systems [13]. Based on these results, the formation of nitrites (NO_2^{-}) , nitrates (NO_3^{-}) , and hyponitrite $(N_2O_2^{2-})$ species was assumed, and a mechanism was postulated [13]

$$2 \operatorname{NO} + \operatorname{O}_{(s)}^{2-} \longrightarrow \operatorname{NO}^{-} + \operatorname{NO}_{2(s)}^{-}$$
(2)

$$4 \text{ NO} + 2 \text{ O}_{(s)}^{2-} \longrightarrow 3 \text{ NO}_{(s)}^{-} + \text{ NO}_{3(s)}^{-}$$
 (3)

$$2 \operatorname{NO}_{(s)}^{-} \longrightarrow \operatorname{N}_2 \operatorname{O}_{2(s)}^{2-} \tag{4}$$

However, it should be noted that these seemingly simple steps are actually sum of several elementary reaction steps, which already includes some uncertainties. For example, the suggested formation of the hyponitrite species involves coupling of two negative species (NO⁻), which is suspicious due to the large electrostatic repulsion.

In NSR, conversion to N_2 is also thought to occur over noble metal sites, which generally involves NO dissociation to form atomic N and O on platinum, where N₂O was found to be an important product at lower temperature when the NO coverage is higher [1, 11, 18, 21, 29]. Although the primary focus of this study is the NSR system, direct decomposition of NO is also of interest. In fact, decomposition of NO over metal oxides in no presence of noble metals has been actively studied [31–35]. For example, Lunsford and co-workers have studied the BaO supported MgO systems using in situ Raman spectroscopy. This system was claimed to be particularly active for NO decomposition at Ba loadings of 11 mol% or greater, where a nitro species was assumed to be an intermediate in the catalytic cycle [31]. Recently, NO direct decomposition on the BaO/Y2O3 catalyst was reported by Ishihara et al. [35]. N₂O, nitrite and nitrate species were supposed to form based on IR spectra. Thus, the disproportional reactions of NO were suggested to occur [35]:

 $3 \text{ NO} \longrightarrow \text{N}_2\text{O} + \text{NO}_2$ (6)

(5)

Clearly, Eqs. 5 and 6 do not refer to elementary steps and the mechanistic details remain to be clarified.

Identification of reaction pathways and intermediates by means of experiments represents a great challenge, while theoretical calculations can help to gain a more clear-cut answer for a given model system. So far, only a few theoretical calculations, based on either periodic-slab models [36, 37] or cluster models [38–40] have been carried out to investigate the interaction of NO with the BaO surfaces. It was generally agreed that NO adsorbs on the oxygen anion sites to form a NO_2^{2-} species [38, 39]. However, the subsequent reactions of the surface species and the possible reaction paths that lead to the formation of the possible hyponitrite $(N_2O_2^{2-})$ species and the release of N_2O have not yet been explored.

Thermodynamics for NO decomposition on MgO has been studied using cluster model calculations, where no transition states have been characterized [41]. A $N_2O_3^{2-}$ species was proposed to be an important intermediate in the NO decomposition process [41]. Unlike the neutral N_2O_2 dimer, it was noticed that anionic dimeric species such as $N_2O_2^{-}$ and $N_2O_2^{2-}$ present a much shorter N–N bond, typical of the N-N double bond [42, 43]. Therefore, charge transfer from the adsorption site to the ad-species was essential to activate the adsorbed precursors for the production of molecular N₂O [42]. On MgO, the lattice sites were found to interact with NO very weakly [41], while chemisorbed species was formed only at low-coordination anions (steps, edges, or corners) or oxygen vacancies [42, 43]. As it is well known that the surface basicity of the cubic alkaline earth oxides (MgO, CaO, SrO, and BaO) increases going down the series as a result of the reduction of the Madelung potential with the crystal lattice expansion [44], the more basic anions on the BaO surfaces are expected to facilitate the NO adsorption and subsequent reactions.

The present theoretical study deals with NO adsorption and transformation on BaO surfaces. It aims, on the one side, at providing a detailed understanding of NO and BaO interactions at the molecular level, and on the other side, at complementing the available experimental work to gain, in particular, a better assignment of the species observed in the IR spectra.

2 Computational details

Density functional theory (DFT) calculations were carried out using the gradient-corrected Becke three-parameter hybrid exchange functional [45–49] in combination with the correlation functional of Lee, Yang, and Parr [50] (B3LYP). Recently, the inter-conversion between NO₂ and N_2O_4 has been studied using the B3LYP functional and several ab initio methods (MP2, CCSD(T), etc.). The results indicated that the B3LYP functional led to good energetics for these conversions [51].

An embedded cluster model approach was adopted to describe the BaO(100) surface as well as the low-coordination surface sties. A Ba₁₆O₁₆ cluster was chosen to represent the BaO(100) terrace sites (O_{5c}), while Ba₁₂O₁₂ and Ba₁₀O₁₀ were used as models for the four-coordinated edge sites (O_{4c}) , and the three-coordinated corner sites (O_{3c}) , respectively (see Fig. 1). These cluster models were constructed in fulfillment of some general requirements, namely the neutrality principle, the stoichiometry principle, and the coordination principle [52-56]. All these clusters were embedded in a large array of ± 2 point charges (PCs), which were aimed to reproduce the Madelung potential at the adsorption sites. A Ba–O lattice distance of 2.76 Å was assumed [57]. To avoid the artificial polarization of the O^{2-} anions at the cluster border induced by the PCs, the positive PCs at the interface were replaced by the total ion effective core potentials (TIPs) [58-60] to ensure a proper description of the Pauli repulsion. No basis functions were associated with the TIPs. The Ba atoms in the clusters were also treated with the effective core potentials (ECPs) originally developed by Hay and Wadt [61]. These were the small core ECPs with the contraction scheme generally known as LANL2DZ. The O and N atoms were described with a $6-31+G^*$ basis set [62, 63].

Geometry optimizations were performed by means of analytical gradients with no symmetry constraints. The adsorbed NO molecules and atoms in clusters which were not in direct contact with the TIPs were included in the geometry optimization. Vibration frequencies were computed by determining the second derivatives of the total energy with respect to the internal coordinates. No scaling factors were applied when comparing the calculated frequencies with the experimental ones.

The adsorption enthalpy changes at room temperature were calculated using the formula:

$$\Delta H_{ad} = H_{tot}^{clus + ad} - H_{tot}^{clus} - H_{tot}^{ad}$$
(7)

where $H_{tot}^{clus + ad}$, H_{tot}^{clus} , and H_{tot}^{ad} refer to the total enthalpy of the full system (BaO cluster + adsorbate), the enthalpy of the bare BaO cluster, and the enthalpy of the gas-phase molecules, respectively. According to this definition, $\Delta H_{ad} < 0$ stands for an exothermic adsorption.

Cluster models have generally been recognized as a useful tool to facilitate analysis and to be able to treat the neutral (e.g., NO and NO dimer) and charged adsorbates (e.g., nitrite and nitrate) on metal oxide surfaces even-handedly [64]. Previously, the same methodology has been used to study the O_2 dissociation mechanisms over the BaO(100)



Fig. 1 Cluster models used in the present work: **a** $(BaO)_{16}$ for terrace sites, **b** $(BaO)_{12}$ for edge sites, and **c** $(BaO)_{10}$ for corner sites. *Red spheres* stand for O^{2–}, while *large white spheres* for Ba²⁺. *Small gray*

surface [65]. Other related studies, in particular, regarding the cluster size and embedding method [64–67], have well documented the validity of the present methodology [64–77].

All calculations were performed with the Gaussian 03 suite of program [62].

3 Results and discussion

3.1 One NO adsorption

On the terrace sites of the BaO(100) surfaces, our calculations predict that NO binds, with the N-down orientation, quite strongly to the O_{5c} anion sites [see Fig. 2(1)]. The adsorption energy is calculated to be $\Delta H_{ad} = -28.4$ kcal/ mol (see Table 1). Mulliken analysis shows that the unpaired spin density is almost entirely localized on the adsorbed NO (0.95, as shown in Table S1 in the supporting information, SI). Upon adsorption, the NO moiety is negatively charged (-0.89), such that partial charge is transferred from the surface anion (O_s) to the antibonding π^* orbital of the adsorbed NO, resulting in an elongation of the N–O bond length from 1.158 Å in the gas phase to 1.291 Å on the surfaces. As a relatively short O_s–N bond (1.516 Å) is formed, the adsorption of NO on the O_s site can be formally regarded as the formation of a NO₂²⁻ species:

$$NO + O_{(s)}^{2-} \longrightarrow NO_{2(s)}^{2-}$$
(8)

spheres refer to the point charges (PCs) that are in close contact with O^{2-} in the clusters as described by the total ion effective core potentials (TIPs). The rest of PCs are not shown

This was first postulated by Lunsford based on the results of the electron paramagnetic resonance (EPR) technique on MgO [78] and was later supported by the observation on CaO [79]. Certainly, $NO_2^{2^-}$ is unstable as an isolated species. It is stabilized on the surfaces by the Madelung potential.

As expected, the reactivity on the low-coordination sites for such an adsorption is much higher than that on the terrace sites due to the increased basicity of O_s [44]. For instance, The NO adsorption energies on the O_{4c} edge sites, or the O_{3c} corner sites, are calculated to be -38.8 and -46.1 kcal/mol, respectively (Table 1). The stronger interactions are also reflected in the shorter O_{s} -N distances of 1.461 Å, and 1.423 Å on the O_{4c} and O_{3c} sites, respectively (Table S1). Spin densities (0.92) are nearly localized on the NO moieties which carrier more negative charges (-0.98), as compared to that on the terrace site, in accord with the increased adsorption energies.

Previously, a detailed comparison between the slab model and the embedded cluster model has been carried out in the study of the O_2 dissociation mechanisms over the BaO(100) surface [65]. A general agreement on the reaction trend was observed with these two methods. The calculations for NO adsorption are also generally in agreement with previous slab model [36, 37] and cluster model [38, 39] calculations. For example, with cluster models [38], Pacchioni et al. calculated the adsorption energies for NO on the O_{5c} terrace and O_{4c} step sites to be -18.4, and -38.0 kcal/mol, respectively. It is interesting to note that

Fig. 2 Optimized structures. **1** for one NO molecule adsorbed on an O_{5c} anion site at the terrace of BaO (*left, side view; right, top view*); **1'** for two NO molecules adsorbed on two neighboring O_{5c} anion sites at the terrace of BaO (*left, side view; right, top view*). *Color codes: yellow spheres* for N, *red spheres* for O, *large white spheres* for TIPs. Geometric details are given in Table S1 in supporting information (SI)



Table 1 The adsorption energies ΔH_{ad} (kcal/mol)^a for one and two NO (the triplet state) adsorbed on the surface anion sites (O_s) of BaO^b

Adsorption geometry	Terrace	Edge	Corner
One NO adsorption (1)	-28.4	-38.8	-46.1
Two NO adsorption (1')	-58.1	-66.9	-70.2

a The adsorption energies are calculated according to $\Delta H_{ad} = H_{tot}^{clus + n*NO} - H_{tot}^{clus} - n*H_{tot}^{NO}$, where n = 1, 2

 $^{\rm b}$ The corresponding adsorption configurations are illustrated in Fig. 2

although a similar stability on the O_{4c} sites was obtained, their adsorption energies on the O5c terrace sites were predicted to be less stable by ~10 kcal/mol than ours. The discrepancy is probably due to the difference in cluster size and the allowed surface relaxation. For adsorption on the terrace sites, besides the adsorbed NO molecule, the fourcenter ions of the (BaO)₁₆ cluster have been included in the present geometry optimization, while in the previous work [38], only the central O atom of the (BaO)₉ cluster was free to relax. The contribution to the adsorption energy associated with the surface relaxation can be significant. In our case, if only the central oxide anion was involved in the geometry optimization, the adsorption energy was found to decrease by ~4 kcal/mol. On the other hand, the effect of surface relaxation is usually well considered in the slab model calculations. For instance, the binding energies for NO adsorption on the BaO(100) surface were reported to be about -34 kcal/mol at a coverage of 0.25 [36, 37].

The frequency analysis shows that the N–O stretch band appears at 1,317 cm⁻¹ on the BaO terrace sites (Table 3), which may be compared to, for example, the observed frequency (1,300 cm⁻¹) for NO exposure on the BaO/Y₂O₃ catalyst [35]. With the decreasing of the coordination number of the adsorption sites, the N–O stretch band experiences a red shift to 1,241 cm⁻¹ on the O_{4c} edge sites, and to 1,096 cm⁻¹ on the O_{3c} corner sites, respectively. Therefore, we are inclined to associate the NO₂^{2–} species with the broad band at 1,206 cm⁻¹ that was observed by SedImair et al. [15] in their IR experiments, although they tentatively assigned this band to the bridged bidentate nitrites.

3.2 Two NO adsorption

A key theoretical concept in understanding the interaction between NO_x and the basic metal oxides is a pairwise, cooperative adsorption mechanism [36]:

$$M^{2+} - NO_x + NO_x - O^{2-} \longrightarrow M^{2+} - (NO_x)^{\delta-} / (NO_x)^{\delta+} - O^{2-}$$
(9)

where NO_x molecules, adsorbed on metal cation sites, and oxygen anion sites, respectively, accept and donate electrons. Such a unique ability of NO_x is a consequence of their high electron affinity and low ionization energy. The electron transfer between two NO_x adsorbates in close proximity through the substrate results in a substantial increase of the adsorption energies for the pair in comparison with two individual adsorbates in isolation. This effect Fig. 3 Optimized structures for NO dimer adsorbed on an O_{5c} anion site at the terrace of BaO, forming *cis*-N₂O₃^{2–} (2) and *trans*-N₂O₃^{2–} (2') (*left, side view; right, top view*). Color codes: yellow spheres for N, red spheres for O, large white spheres for Ba and small gray spheres for TIPs. Geometric details are given in Table S1 in supporting information (SI)



has been observed for the (NO, NO₂), (NO₂, NO₂), (NO₂, NO₃) pairs, but not for the (NO, NO) pair [36].

Adsorption of two NO molecules on the BaO surfaces is studied in the present work, where both NO molecules are found to adsorb on the O anion sites. The adsorption configuration on two adjacent O_{5c} sites is shown in Fig. 2 (1'), while geometries for some other configurations may be found in Table S1 in SI. As shown in Table 1, the adsorption energy per molecule on a terrace site is -29.05 kcal/ mol, no matter the electronic state is set to either a triplet or an open-shell singlet. This energy actually compares well with the adsorption energy of the singly adsorbed molecule, indicating that there is little enhanced pairwise cooperative effect. The same is true if two NO molecules are adsorbed on a pair of O_{5c} and O_{4c} sites. Interestingly, if adsorption occurs on a pair of (O_{5c}, O_{3c}) sites, a repulsion of 3.8 kcal/ mol is encountered (see Table 1).

The coverage dependence of NO adsorption on the BaO(100) surface was reported with the slab model calculations [37]. The results showed that the binding strength of NO on the anion sites almost linearly decreased with the increasing coverage from 34.6 kcal/mol at $\theta = 0.125$ –23.1 kcal/mol at $\theta = 0.5$. Therefore, as indicated by the present cluster model calculations and the previous slab model calculations [37], some other adsorption configurations containing NO dimer may be more favorable at higher NO coverage.

As described above, the isolated NO binds strongly on the O_s sites to form a NO_2^{2-} species even on the terrace

sites of BaO. As both the NO_2^{2-} species and the free NO molecule carry one unpaired electron on the N atom, respectively, a singlet $N_2O_3^{2-}$ species would be formed by coupling the two radical-like groups (see Fig. 3):

$$NO_{2(s)}^{2-} + NO \longrightarrow N_2O_{3(s)}^{2-}$$
(10)

The $N_2O_3^{2-}$ complex can also be regarded as the adsorption of a NO dimer (N₂O₂) on the surface O anion site. These have been studied before on MgO [41], as well as BaO [39]. The configurations of the $N_2O_3^{2-}$ surface complexes can be assigned to $cis-N_2O_3^{2-}$ and trans- $N_2O_3^{2-}$ with respect to *cis*- N_2O_2 and *trans*- N_2O_2 , respectively. The corresponding geometrical parameters are listed in Table S2 in SI. The N-N bond length is dramatically reduced from 1.96 Å for cis-N₂O₂ in the gas phase [42] to 1.29–1.32 Å, depending on the orientation, while the N–O bond length is drastically elongated from 1.16 Å in the gas phase [42] to 1.30–1.34 Å. Mulliken charge analysis shows that the adsorbed NO dimer on the terrace anion sites almost obtains one electron (-0.84 for the *cis* form and -0.75 for the *trans* form), which is increased to -1.0 and -1.3 for the low-coordination anion sites on the edges and corners, respectively (see Table S2 in SI). Such an electron transfer from the surface anion sites to the NO dimer reinforces the bond strength of N-N, while weakens the N-O bond simultaneously, which then triggers the direct dissociation channel from $N_2O_3^{2-}$ into N_2O as described in the next section.

Table 2 The adsorption energies ΔH_{ad} (kcal/mol)^a for a NO dimer adsorbed on the surface anion sites (O_s) of BaO^b

Adsorption geometry	Terrace	Edge	Corner
$cis-N_2O_3^{2-}(2)$	-41.4	-58.3	-89.7
$trans-N_2O_3^{2-}(2')$	-41.0	-68.5	-89.9

 a The adsorption energies are calculated according to $\Delta H_{ad}=H_{tot}^{clus+(NO)_2}-H_{tot}^{clus}-2*H_{tot}^{NO}$

 $^{\rm b}$ The corresponding adsorption configurations are illustrated in Fig. 3

The adsorption energies of the NO dimer on the terrace as well as edge and corner anion sites are summarized in Table 2. On the terrace sites, the stability of the *cis*- and *trans*-N₂O₃²⁻ is similar with an exothermicity of ~41 kcal/ mol with respect to two free NO molecules. On the lowcoordination anion sites, the N₂O₃²⁻ species are calculated to be more stable. On the edge sites, formations of the *cis*- and *trans*-N₂O₃²⁻ species are exothermic by 58.3 and 68.5 kcal/mol, respectively, while on the corner sites, both *cis*- and *trans*-N₂O₃²⁻ show a large stability with the adsorption energy of about -90 kcal/mol.

The formation of the stable $N_2O_3^{2-}$ species represents the N–N bond formation which should be identified by the IR spectra. Previously, the bands around 1,310 and 1,375 cm⁻¹ as observed by Prinetto et al. [13] and the peak around 1,380 cm⁻¹ as observed by Sedlmair et al. [15] were associated with the N–N stretching in the $N_2O_2^{2-}$ species. This is fairly reasonable if one compares them with the N–N stretching in the $Na_2N_2O_2$ salts, which appear around 1,314 and 1,419 cm⁻¹ [80]. However, as listed in Table 3, our calculated N–N stretching frequencies are also around 1,354 cm⁻¹ for *cis*-N₂O₃²⁻ and 1,413 cm⁻¹ for *trans*-N₂O₃²⁻, even though the corresponding peak appears at much low region of 1,120 cm⁻¹ in the Na₂N₂O₃ salts [80]. Hence, a band of the same nature may appear at different position on surfaces and in compounds. The present calculations offer a new insight into the assignment of the N–N stretching bands, which may indicate the formation of the N₂O₃²⁻ species on the surfaces.

3.3 N₂O₃²⁻ dissociation

The proposed mechanisms for NO decomposition via $N_2O_3^{2-}$ species are presented in Scheme 1. It is initialized by the formation of species **1** (NO_2^{2-}) via the adsorption of an NO molecule on the oxygen anion sites, then followed by the coupling of the second gas-phase NO to **1** to produce **2** (*cis*- $N_2O_3^{2-}$) or **2'** (*trans*- $N_2O_3^{2-}$), depending on the orientation. The dissociation of species **2** (**2'**) occurs through the cleavage of one of the N–O bonds, leading to the formation of species **3**(**3'**), which corresponds to a $N_2O_2^{2-}$ species plus a peroxide species of O_2^{2-} on the surface. A rearrangement of $N_2O_2^{2-}$ generates the lattice oxygen anion with an electrostatically adsorbed N_2O as in **4**(**4'**). The final step is envisioned as the desorption of N_2O , leaving the peroxide on the surfaces.

The transition states and intermediates in the dissociation processes on the terrace sites are presented in Figs. 4 and 5, respectively. The corresponding enthalpy changes, defined in Scheme 1, are summarized in Table 4, which are depicted in Fig. 6. The detailed geometric parameters are given in Tables S3 and S4 in SI.

Table 3 Calculated vibrationalfrequencies (cm^{-1}) for the		v ₁ (N–N stretch)	v ₂ (N–O stretch)
adsorption species on the BaO	Calc.		
terrace sites	NO adsorption (1)	-	1,317
	Two NO adsorption (1')	_	1,313 (symmetry), 1,309 (asymmetry)
	$cis-N_2O_3^{2-}(2)$	1,354	1,410
	<i>trans</i> - $N_2O_3^{2-}(2')$	1,413	1,282
The experimental IR spectra for free molecules ^a ions in	<i>trans</i> - $N_2O_2^{2-}(3)$	1,279	1,417
compounds ^a and the NO/BaO/	$cis-N_2O_2^{2-}(3')$	1,380	1,241
Al_2O_3 system ^b are also listed for	adsorbed N ₂ O (4)	1,884	1,118
comparison	adsorbed N ₂ O (4')	2,378	1,315
^a Data from Ref. [63] for	Expt.		
compounds	NO ^a	_	1,876
^b From IR spectra for NO/BaO/	N_2O^a	2,224	1,285
Al_2O_3 at room temperature [13]	NO_2^{-a}	-	1,260 (asymmetry), 1,330 (sym.)
^c Assigned to the surface	NO_3^{-a}	-	1,380 (asymmetry), 1,050 (sym.)
$N_2O_2^{2^-}$ species [13]	$N_2O_3^{2-a}$	1,120, 1,100	1,400, 1,380 (asymmetry NO ₂), 1,280 (sym. NO ₂), 980, 970
^d Assigned to the surface NO ₂ ⁻	cis-N ₂ O ₂ ^{2-a}	1,314	1,057 (sym.), 857 (asymmetry)
species [13]	trans- $N_2O_2^{2-a}$	1,419	1,120 (sym.), 1,030 (asymmetry)
^e Assigned to the surface NO_3^- species [13]	NO/BaO/Al ₂ O ₃	1375 ^c , 1310 ^c	1,050–950 ^c , 1,220–1,180 ^d , 1,600 ^d , 1470 ^e



Scheme 1 The possible reaction pathways of NO decomposition on BaO (100) surfaces

We first look at the dissociation of $cis-N_2O_3^{2-}$ (2) on the terrace sites of the BaO surfaces. A transition state (TS1 in Fig. 4) is located, which clearly indicates the cleavage of a N-O bond and the formation of an O_s-O bond. The asformed **3** is lower lying by 12.5 kcal/mol on the terrace with respect to 2. Notably in 3, a surface peroxide species (O_2^{2-}) can be envisioned. This is characterized by an O_s -O bond length of 1.523 Å as shown in Table S4 in SI, which approaches to the bond length of a standard peroxo species $(1.49 \pm 0.02 \text{ Å}, [81])$. Meanwhile, a relatively short O_s-N bond with the distance of 1.46 Å is maintained, which is accompanied by a N-N bond of 1.30 Å (See Table S4). The latter compares well with the N-N distance in the standard $N_2O_2^{2-}$ species (1.27 Å [80]). We thus assign this entity as a surface $N_2O_2^{2-}$ species. It has to be noted that the transformation from the $N_2O_3^{2-}$ (2) species to the $N_2O_2^{2-}$ species in 3 on the terrace sites needs to surmount an intrinsic reaction barrier of 52.8 kcal/mol. Hence, it is necessary to increase the reaction temperature to facilitate the $N_2O_2^{2-}$ formation.

Thereafter, **3** experiences a geometry relaxation and leads to the formation of **4** by overcoming a reaction barrier of 25.0 kcal/mol on the terrace sites via **TS2**. The geometrical features of **TS2** (see Fig. 4 and Table S3 in SI) clearly indicate the N₂O formation with the N–N bond length of 1.18 Å and the N–O length of 1.28 Å, which is accompanied by a strongly elongated O_s–N bond (2.11 Å). In **4** on the terrace, the as-formed N₂O is found to be adsorbed on the surface anion site via the middle N atom (see Fig. 5). There is still a sizable charge transfer from O_s to N₂O (0.57 as listed in Table S4 in SI), leading to a NNO which is bent by 138°, instead of adopting a linear NNO structure as in free N₂O. We find that there is an energy cost of 10.2 kcal/mol from **3** to **4**.

Now, we look at the dissociation of *trans*- $N_2O_3^{2-}$ (2') on the terrace site of the BaO surfaces. As shown in Fig. 6, the dissociation route is basically the same as that of the *cis*- $N_2O_3^{2-}$ species. The whole energetics is more favorable, as compared to the dissociation of *cis*- $N_2O_3^{2-}$ (2), due to a more favorable electrostatic interaction between the transconfigurations of the ad-species and the surfaces. Indeed,

TS1' is 5.7 kcal/mol lower lying than **TS1**, while **TS2'** is 17.3 kcal/mol more stable than **TS2**. The geometry features of N₂O in **4'** are shown in Fig. 5 and Table S4 in SI. The adsorbed N₂O is nearly linear (177°), which is nearly charge neutral. Hence, one may consider the N₂O₂²⁻ species in **3(3')** as a chemisorbed N₂O on the surface, the N₂O moiety in **4**, in particular **4'**, shall be considered as a physisorbed species.

The energy profiles show that the dissociation of $N_2O_3^{2-}$ on the terrace sites is characterized by a large barrier encountered at **TS1(TS1')** for the formation of $N_2O_2^{2-}$, followed by a small barrier at **TS2(TS2')** which eventually leads to the N_2O formation. Hence, if the first barrier can be overcome, e.g., with the help of other components in the catalyst at high temperature, it is possible that NO undergoes direct dissociation over BaO as it was observed in the BaO/MgO [31] or BaO/Y₂O₃ [35] system. On the other hand, we conclude that NO initially adsorbs on BaO as NO_2^{2-} and $N_2O_3^{2-}$ at low temperature in no presence of O_2 and Pt.

We have also examined the dissociation of $N_2O_3^{2-}$ on the low-coordination sites of the BaO surfaces. The corresponding enthalpy changes are also summarized in Table 4 and depicted in Fig. 6. The detailed geometric parameters may be found in SI. Even though the intrinsic barriers to transform $N_2O_3^{2-}$ (2 or 2') to $N_2O_2^{2-}$ (3 or 3') remain to be high (54.8-62.1 kcal/mol), the strong basicity of the low-coordination sites is found to significantly enhance the stabilities of the $N_2O_3^{2-}$ species, which actually brings the transition states (TS1 or TS1') lower (by -3.5 to -30.0 kcal/mol) than the entrance level (BaO + 2 NO), rendering these dissociation processes having no apparent barriers. Therefore, the $N_2 O_3^{2-}$ dissociation on the low-coordination surface sites is predicted to be easier than that on the terrace sites. In addition, the increased basicity from terrace to edge to corner sites has significantly increased the amount of the change transfer from 0.57 to 0.90 to 1.34 from the surfaces to the N₂O moiety (see Table S4 in SI), which stabilizes the bent N₂O in 4. In fact, both TS2 and TS2' are found to connect with 4 on the low-coordination sites.

Fig. 4 Optimized transition state (TS) structures on the terrace sites of BaO (left, side view; right, top view): TS1 and TS1' correspond to the dissociation of $cis-N_2O_3^{2-}$ and trans- $N_2O_3^{2-}$, respectively, to form *cis*- $N_2O_2^{2-}$ and *trans*- $N_2O_2^{2-}$; TS2 and TS2' correspond to the dissociation of $cis-N_2O_2^{2-}$ and *trans*- $N_2O_2^{2-}$, respectively, to form physisorbed N₂O. Color codes: yellow spheres for N, red spheres for O, large white spheres for Ba and small gray spheres for TIPs. Geometric details may be found in supporting information (SI)



The calculated N–N–O asymmetric stretching mode in **4** appears at 1,884 cm⁻¹ on the terrace sites, 1,756 cm⁻¹ on the edge sites, and 1,570 cm⁻¹ on the corner sites, which

are comparable to the experimental band observed around $1,780 \text{ cm}^{-1}$ for NO adsorbed on BaO/Y₂O₃ [35]. On the other hand, the calculated N–N–O asymmetric vibration

Fig. 5 Optimized structures on the terrace sites of BaO (*left, side view; right, top view*): **3** and **3'** correspond to *trans*-N₂O₂^{2–} and *cis*-N₂O₂^{2–}, respectively, plus a surface O₂^{2–} species. **4** and **4'** correspond to a physically adsorbed N₂O plus a surface O₂^{2–} species. *Color codes: yellow spheres* for N, *red spheres* for O, *large white spheres* for Ba and *small gray spheres* for TIPs. Geometric details may be found in supporting information (SI)















Table 4 Reaction enthalpychanges and intrinsic activationbarriers (kcal/mol) for reactionpaths defined in Scheme 1

	ΔH_1	ΔH_2	H_1^{\ddagger}	ΔH_3	H_2^{\ddagger}	ΔH_4	ΔH_5	$\Delta H_2{}^\prime$	$H_{I}^{\pm \prime}$	ΔH_{3}^{\prime}	$H_2^{\pm\prime}$	$\Delta H_4{}^\prime$	$\Delta H_5{}^\prime$
Terrace	-28.4	-13.0	52.8	-12.5	25.0	10.2	-0.6	-12.6	47.7	-14.8	7.7	6.1	5.5
Edge	-38.8	-19.5	54.8	-2.7	28.1	2.0	16.6	-29.7	58.3	-5.7	22.9	-	5.6
Corner	-46.1	-43.6	62.1	9.1	45.1	5.7	33.8	-43.8	59.9	1.7	36.7	-	5.2

4

20.0

0.0

-20.0

-40.0

-60.0

20.0

0.0

-20.0

-40.0

-60.0

-80.0

0.0

-20.0

-40.0

-60.0

-80.0

46.1

Corner

H at 298K(kcal/mol)

H at 298 K(kcal/mol)

H at 298 K (kcal/mol)

Fig. 6 Energy profiles for NO adsorption and transformations on the **a** terrace **b** edge and **c** corner sites of BaO surfaces. The reaction schemes are defined in Scheme 1. The profiles bifurcate at the formation of *cis*- and *trans*-N₂O₃²⁻ (**2** and **2'**, respectively) species



-30.0

-80.6

-88.2

frequency in 4' is around 2,377 cm⁻¹ on the terrace. This is in good agreement with the experimentally observed band at 2,350 cm⁻¹ for the NO/BaO/Y₂O₃ system [35] and is also close to that in the free N₂O [81]. All in all, our calculations support that the peaks around 1,780 and 2,350 cm⁻¹ are from N₂O, in accordance with the assignments reported by Ishihara et al. [35] for the NO/BaO/Y₂O₃ systems.

It is particularly interesting to compare the calculated N–N stretching spectra in $N_2O_2{}^{2-}$ with those observed

in experiments for the NSR systems [13, 15]. In fact, the calculated bands are around 1,279 and 1,380 cm⁻¹, which compare fairly well with the bands centered at 1,310 and 1,375 cm⁻¹ as observed by Prinetto et al. [13] and 1,380 cm⁻¹ as observed by SedImair et al. [15], which were assigned to the $N_2O_2^{2-}$ species. Hence, our calculations support that the surface $N_2O_2^{2-}$ will also contribute to the strong peaks observed around 1,310–1,380 cm⁻¹ if the barriers from **TS1** (**TS1'**) can be effectively overcome.

-35.5

TS2 -51.5

-74.9

-89.7

-89.9

 $BaO_2 + N_2O$

41.4

(c)

4 Concluding remarks

It is commonly reported that conversion of NO to NO_2 over noble metals is required for NO_x storage, while NO reduction also occurs over noble metals [6–10]. Due to the complexity of the NSR system, however, the understanding of the single role of BaO is still incomplete. Some reports suggested that BaO is inactive toward NO [10, 23], while some others showed that NO trapping is already significant on BaO alone [11, 12]. The present work presents a theoretic study on NO adsorption and reaction on BaO, trying to clarify this issue.

The XPS experiments by Schmitz and Baird have highlighted the importance of BaO [12]. Considering that NO_x in exhaust gas is primarily NO and that the surface area of the oxides far exceeds that of the dispersed noble metals, they proposed a unique mechanism that the initial trapping is dominated by the molecular adsorption of NO on BaO to form a nitrite-like (NO_2^-) ad-species [12]. Our calculations support this view of molecular adsorption of NO on BaO, although we are inclined to favor the picture that NO adsorbs on the surface anion sites to form NO_2^{2-} , which is in agreement with the results from the EPR study of NO on the lowcoordination sites on MgO and CaO surfaces [78, 79].

A key theoretical concept in understanding the interaction between NO_x and the basic metal oxides is a pairwise, cooperative adsorption mechanism [36], which is shown not to be effective for the (NO, NO) pair on BaO. At high NO pressure, dimerization is favored, in particular, on the low-coordination anion sites to form N₂O₃²⁻. Although the surface N₂O₂²⁻ species is a commonly assumed key intermediate in the NSR system, our calculations suggest that it has to surmount an intrinsic barrier of 50 kcal/mol for its formation. Notably, the high basicity of the low-coordination anion sites on BaO will bring the corresponding transition states below the entrance level, which shall facilitate the N₂O₂²⁻ formation.

Our calculations show that if this barrier can be overcome, the transformation from the surface $N_2O_2^{2-}$ species to N_2O is facile. Indeed, the formation of N_2O has been observed in some NSR and related systems [10, 25, 29, 33, 35]. It is well known that N_2O can be further reduced to N_2 on the metal oxide surfaces [82], which has been characterized with embedded cluster models for BaO(100) [83]. Hence, it is envisioned that BaO not only plays a role in NO storage, but also affects in a way the NO reduction.

The net reaction studied in the present work may be summarized as

$$2 \operatorname{NO} + \operatorname{O}_{(s)}^{2-} \longrightarrow \operatorname{N}_2\operatorname{O} + \operatorname{O}_{2(s)}^{2-}$$
(11)

It would be anticipated that the surface peroxide species may, in turn, serve as an active oxidant for further NO storage to form NO_2^- and NO_3^- [13, 15, 21, 31, 40], which deserves a detailed investigation.

Some calculated vibrational frequencies of the intermediates are summarized in Table 3, which shall be helpful to assign the bands observed in the experiments. In Table 3, we have also listed the experimental results for the corresponding free molecules (NO and N₂O) and ions (NO₂⁻, NO_3^- , $N_2O_3^{2-}$ and $N_2O_2^{2-}$) in compounds [80]. These are the reference numbers with which the IR spectra for surface species in the NSR systems are compared [84]. One has to note that coordination on surfaces will change the peak positions dramatically and the bands from various $N_x O_x^{\delta}$ species overlap. All of these have made the assignment of the corresponding vibrational spectra complicated and controversial. Indeed, the surface NO_2^{2-} and $N_2O_3^{2-}$ species provide an alternative explanation of the experimentally observed IR spectra which were previously assigned to the nitrite/nitrate and hyponitrite species [13, 15].

Further investigation is necessary to see how different reactant gases adsorb, react and even compete against each other for a specific active site, and how the interplay between different components in a real NSR catalyst will change the energy profiles or even change the reaction mechanisms.

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REGULAR ARTICLE

Microwave and infrared spectra of CO– $(pH_2)_2$, CO– $(oD_2)_2$, and mixed CO– pH_2 –He trimers

Xiao-Long Zhang · Hui Li · Robert J. Le Roy · Pierre-Nicholas Roy

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Abstract The microwave and infrared spectra of CO-(pH₂)₂, CO-(oD₂)₂, and CO-pH₂-He trimers are predicted by performing exact bound state calculations on the global potential energy surfaces defined as the sum of accurately known two-body pH₂-CO or oD₂-CO (in Li et al. J Chem Phys 139:164315, 2013), pH₂-pH₂ or oD₂-oD₂ (in Patkowski et al. J Chem Phys 129:094304, 2008), and pH₂-He pair potentials. A total of four transitions have been reported to date, three in the infrared region, and one in the microwave region, which are in good agreement with our theoretical predictions. Based on selection rules, new transitions for $J \leq 3$ have been predicted, and the corresponding transition intensities at different temperatures are also calculated. These predictions will serve as a guide for new experiments. The weak and tentatively assigned transitions are verified by our calculations. Three-body effects and the quality of the potential are discussed. A reduced-dimension

Dedicated to Professor Guosen Yan and published as part of the special collection of articles celebrating his 85th birthday.

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P.-N. Roy e-mail: pnroy@uwaterloo.ca treatment of the pH_2 or oD_2 rotation has been employed by applying the hindered-rotor averaging technique of Li et al. (J Chem Phys 133:104305, 2010). A technique for displaying the three-dimensional pH_2 or oD_2 density in the bodyfixed frame is used and shows that in the ground state, the two pH_2 or two oD_2 molecules are localized, while the He's are delocalized.

 $\label{eq:constraint} \begin{array}{ll} \mbox{Keywords} & \mbox{Microwave spectra} \cdot \mbox{Infrared spectra} \cdot \\ \mbox{CO-}(pH_2)_2 \mbox{ trimers} \cdot \mbox{CO-}(oD_2)_2 \mbox{ trimers} \cdot \mbox{CO-}pH_2\mbox{-He} \\ \mbox{trimers} \cdot \mbox{Reduced-dimension treatment} \cdot \mbox{Exact bound state} \\ \mbox{calculations} \cdot \mbox{Three-body effects} \end{array}$

1 Introduction

At a very low temperature (<2 K), high-resolution infrared or microwave spectra of a single chromophore molecule doped in helium droplets or clusters have provided a unique opportunity to study quantum solvation and microscopic superfluidity [1-20]. Para-hydrogen (pH₂) or ortho-deuterium (oD₂), like ⁴He atoms, are bosons. Similarly, a chromophore molecule embedded in pH_2 or oD_2 clusters can also be considered as a possible route to investigate quantum solvation and the superfluidity of pH₂ or oD₂ [21-30]. Recently, combining experimental measurements and theoretical simulations, superfluid response for pH₂ to dopant rotation has been first elucidated in CO2-(pH₂)_N clusters [26]. However, due to localization effects, the superfluid fraction of larger doped hydrogen clusters becomes suppressed. Carbon monoxide (CO) is a gentler probe molecule with much weaker and less anisotropic interaction with pH₂ and a relatively large rotational constant. These features lead to a delocalized distribution of pH₂ molecules with respect to CO. Persistent molecular superfluid response to dopant rotation in $CO-(pH_2)_N$ clusters has been observed, and this conclusion is supported by both microwave spectroscopy and the theoretical analysis [31].

An understanding of the experimental spectra at the microscopic scale requires complementary computer simulation studies based on accurate intermolecular potential energy functions. Very recently, two reliable ab initio analytical potential energy surfaces (PESs) for H2-CO dimer have been reported, referred to hereafter V_{12} [32, 33] and V_{MLRO} [34], respectively. Both PESs explicitly account for the intra-molecular C-O stretch coordinate and allow to describe the infrared spectrum in the region of the fundamental stretching vibration of C-O around 2,143.2712 cm⁻¹. The predicted infrared spectra calculated from these two PESs have already been shown to agree extremely well with the experimental spectra of the pH₂and oH2-CO dimers [32-34]. Studies of CO-(pH2)2 trimer are important for testing the accuracy of pair-wise additive approach by sum of two-body potentials to simulate systems with tens of pH₂ molecules, looking for three-body effects, and for testing the validity of quantum Monte Carlo methods for molecular clusters, since comparisons with exact quantum results are possible.

Although a large number of infrared and microwave transitions were recorded and assigned for CO–He and CO–pH₂ dimers [33, 35–37], only a few transitions for CO–(pH₂)₂, CO–(oD₂)₂, and CO–pH₂–He trimers were reported, and some weak or tentatively assigned transitions remain to be verified or reassigned [22, 31]. Therefore, reliable theoretical predictions based on exact quantum calculations will provide valuable guidance in confirming or reassigning the measured infrared spectra. Previous work on exact quantum calculations of infrared and microwave spectra for trimer complexes containing one CO, CO₂, N₂O, and OCS molecule attached to two He atoms have been reported, by Wang and Carrington [38–41] and by Li et al. [42], which yield excellent agreement with experiment

observations. Exact rovibrational spectra calculations for the trimer containing a single chromophore molecule doped with two pH₂ molecules, such as CO_2 -(pH₂)₂, were first reported by Li et al. [27] by applying the hindered-rotor averaging technique to treat pH₂ as a point-like particle and consequently reduce the dimension of the potential enemy surface. To date, however, exact quantum calculations for the trimer containing a chromophore molecule doped with two oD_2 or with mixed pH_2 and He have not been reported. Further more, exact quantum calculations for CO-(pH₂)_N clusters have only been limited to CO-pH₂ dimer. In the present paper, we extend such exact quantum calculations to the cases of two pH₂, or two oD₂, or mixed pH₂ and He, attached to one CO dopant molecule. The results are used to help assign or confirm the experimental spectra of CO- $(pH_2)_2$, CO– $(oD_2)_2$, and CO– pH_2 –He.

2 Computational methods

2.1 Geometry and reduced-dimension treatment

There are six atoms in CO-(pH₂)₂ system, and it takes a 12-dimensional (12D) PES to describe the interaction within the system. If we treat linear CO and the two H_2 as rigid molecules, taking only the intermolecular coordinates into consideration, there are still nine intermolecular degrees of freedom. The geometry of this complex in which CO is rigidly linear, and can be described naturally using the Jacobi coordinates $(q_1, q_2, \vartheta_1, \vartheta_2, \vartheta_3, \vartheta_4, \varphi, \varphi_1,$ φ_2) is shown on the left panel of Fig. 1, where q_1 and q_2 are the distances from the center of mass of CO to the centers of mass of the two H₂ molecules, ϑ_1 and ϑ_2 are the angles between \mathbf{q}_1 or \mathbf{q}_2 and a vector pointing from atom O to atom C, ϑ_3 and ϑ_4 are the angles between $\mathbf{q_1}$ and a vector pointing from atom H_2 to atom H_1 , and from q_2 to a vector pointing from atom H_4 to atom H_3 , φ is the dihedral angle between two planes defined by vector \mathbf{q}_1 and \mathbf{q}_2 with the

Fig. 1 Jacobi coordinates for the CO– $(H_2)_2$ complex: left side, the geometry describe with nine dimensions of intermolecular coordinates; right side, the geometry described with five dimensions of intermolecular coordinates and with the two para-hydrogen treated as spherical particles



axis of the CO molecule, φ_1 and φ_2 are the azimuthal angles for rotation of H₁-H₂ and H₃-H₄ about vectors **q**₁ or **q**₂, respectively.

It is still difficult to solve the nine-dimensional intermolecular Schrodinger equation. However, because the nuclear spins are anti-parallel for para-hydrogen, the rotational angular momentum, J, must be even. We know that the ground-state rotational wave function of an isolated ground-state (J = 0) H₂ molecule is precisely spherically symmetric, and the energy spacing to its first-excited (J = 2) level is relatively large (354.3735 cm⁻¹). It is therefore desirable to treat pH_2 as a spherical particle. Following this approximation, the nine-dimensional intermolecular coordinates for a CO- $(pH_2)_2$ complex can be reduced to five coordinates $(q_1, q_2, \vartheta_1, \vartheta_2, \varphi)$ shown on the right panel of Fig. 1, greatly reducing the computational effort. However, it should be noted that pH₂ is not truly spherical when interacting with other species. A simple spherical average over the orientations of the H₂ moiety does not provide an accurate description [43].

The rotational constant for $pH_2(59.322 \text{ cm}^{-1})$ is more than 31 times larger than that of CO in its ground $(1.9225288 \text{ cm}^{-1})$ or excited state $(1.9050256 \text{ cm}^{-1})$ [44]. It therefore seems reasonable to perform a Born-Oppenheimer-type separation of the fast rotational motion of pH₂ from the relative slower CO rotation. Recently, Li, Roy, and Le Roy introduced a much better reduced-dimension description which they call an "adiabatic-hindered-rotor (AHR)" method. This approximation yields an effective pH₂-{linear molecule} interactions that are an order of magnitude more accurate than those obtained using a simple spherical-average approximation. We expect that with such an AHR approximation, the CO-(pH₂)₂ trimer can be accurately described with the left over five degrees of freedom shown in the right panel of Fig. 1. Based on our previous tests for CO-oD₂ dimer, as shown in Table 2 of Ref. [43], we found the AHR treatment of oD_2 is still feasible and reliable, and the differences between $\{2D-4D\}$ are smaller than 0.092 cm⁻¹ for vibrational levels, which are about six times better than those obtained by spherical treatment. Therefore, similar treatment has been applied for the other two trimers of $CO-(oD_2)_2$ and $CO-pH_2$ -He.

The rovibrational energy levels of CO– $(pH_2)_2$, CO– $(oD_2)_2$, and CO– pH_2 –He were calculated using the radial discrete variable representation (DVR) and parity-adapted angular finite basis representation (FBR) methods described in Section II. The potential we used for each trimer is a sum of three pair potentials. The CO– pH_2 , CO– oD_2 , and oD_2 – oD_2 potentials were generated by an "adiabatic-hindered-rotor" average treatment of the H₂ or D₂ rotation using dimer potentials taken from Refs. [32, 34, 45], while pH_2 – pH_2 potentials were generated by a "spherical" average treatment of the H₂ rotation using dimer potential taken

from Ref. [45]. For CO-pH₂-He and CO-(He)₂ complexes, the CO-pH₂, CO-He, and He-He potentials of Refs. [34, 46, 47] were used. The rotational constants B_{CO} required for these calculations were fixed at the experimental values of 1.9225288 and 1.9050256 cm^{-1} for CO in their ground $(v_3 = 0)$ and excited $(v_3 = 1)$ states [44], respectively. In our calculations, the masses were set at 1.00782503207 u for H, 2.0141017778 u for D, 4.00260324 u for ⁴He, 15.994914635 u for ¹⁶O, and 12. u for ¹²C. An 40-point sine DVR grid range from 3.0 to 30.0 bohr was used for the radial r_1 or r_2 stretching coordinate. For the angular basis, we use $l_{\text{max}} = m_{\text{max}} = 25$; 30 Gauss–Legendre quadrature points were used for the integration over θ_1 or θ_2 , while 64 equally spaced points in the range $[0, 2\pi]$ were used for the integration over ϕ . It is useful to apply a potential ceiling value to reduce the spectral range and accelerate the convergence of the Lanczos calculation [48]. In our calculations, we use a ceiling of $1,000 \text{ cm}^{-1}$ and have confirmed that low-lying energy levels change by less than 0.001 cm^{-1} when the ceiling is raised to $10,000 \text{ cm}^{-1}$.

2.2 Hamiltonian and basis functions

Following our previous work for $CO_2-(pH_2)_2$ trimer [27], in Radau coordinate, we use the Hamiltonian and basis functions based mainly on the work of Wang et al. [39] and on fundamental ideas presented by Mladenovic [49]. In recent years, this approach have been successfully used to treat N₂O–(He)₂, CO–(He)₂, CO₂–(He)₂, OCS–(He)₂, and CO₂–(pH₂)₂ complexes. The method is briefly reviewed here, and the reader is referred to Refs. [39] and [27] for further details. With the AHR approximation, the rovibrational Hamiltonian of the CO–(pH₂)₂ complex in the CO molecule-fixed frame has the form (in a.u.) [39, 49–51],

$$\hat{H} = \hat{T}_{\rm str} + \hat{T}_{\rm diag} + \hat{T}_{\rm off} + \hat{T}_{\rm Cor} + \bar{V}(r_1, r_2, \theta_1, \theta_2, \phi)$$
(1)

in which

$$\hat{T}_{\rm str} = -\frac{1}{2m_{\rm H_2}} \frac{\partial^2}{\partial r_1^2} - \frac{1}{2m_{\rm H_2}} \frac{\partial^2}{\partial r_2^2}$$
(2)

$$\hat{T}_{diag} = -\left(\frac{1}{2m_{\rm H_2}r_1^2} + B_{\rm CO}\right) \left[\frac{\partial^2}{\partial\theta_1^2} + \cot\theta_1 \frac{\partial}{\partial\theta_1} - \frac{1}{\sin^2\theta_1} (\hat{J}_z - \hat{I}_{2z})^2\right] \\ + \left(\frac{1}{2m_{\rm H_2}r_2^2} + B_{\rm CO}\right) \hat{l}_2^2 + B_{\rm CO}[\hat{J}^2 - 2(\hat{J}_z - \hat{I}_{2z})^2 - 2\hat{J}_z\hat{I}_{2z}]$$
(3)

$$B_{\rm CO}[\hat{l}_{2+}\hat{a}_1^- + \hat{l}_{2-}\hat{a}_1^+] \tag{4}$$

$$\hat{T}_{\rm Cor} = -B_{\rm CO}[\hat{J}_-\hat{a}_1^+ + \hat{J}_+\hat{a}_1^- + \hat{J}_-\hat{l}_{2+} + \hat{J}_+\hat{l}_{2-}]$$
(5)

 $\hat{T}_{\text{off}} =$

where

$$\hat{J}_{\pm} = \hat{J}_x \pm i \hat{J}_y, \quad \hat{l}_{2\pm} = \hat{l}_{2x} \pm i \hat{l}_{2y}$$
 (6)

$$\hat{a}_1^{\pm} = \pm \frac{\partial}{\partial \theta_1} - \cot \theta_1 (\hat{J}_z - \hat{l}_{2z}) \tag{7}$$

The details of the coordinate system are shown in Fig. 2. r_1 and r_2 are the lengths of Radau (or orthogonalized satellite) vectors \mathbf{r}_1 and \mathbf{r}_2 , which are linear combinations of the Jacobi vectors \mathbf{q}_1 and \mathbf{q}_2 running from the center of mass of CO to the centers of the pH₂ molecules [49]. The polyspherical angles $(\theta_1, \theta_2, \phi)$ are determined by the three vectors (\mathbf{r}_0 , \mathbf{r}_1 , \mathbf{r}_2). θ_1 is the angle between the dopant axis \mathbf{r}_0 and $\mathbf{r_1}$, θ_2 and ϕ are the polar angles that specify the orientation of \mathbf{r}_2 with respect to the CO molecule-fixed frame. m_{H2} is the mass of the pH₂ molecule and B_{CO} is the rotation constant of CO. The operators J_x , J_y , and J_z are the components of the total angular momentum operator J in the body-fixed frame, the z axis of the body-fixed frame lies along the Jacobi radial vector \mathbf{r}_0 , and the x axis is in the plane that contains \mathbf{r}_0 and one pH₂ molecule. Here, $\overline{V}(r_1, r_2, r_3)$ θ_1, θ_2, ϕ) is the total potential energy function which is represented as a sum of two two-dimensional AHR pH2-CO potentials [34, 43] plus the one-dimensional AHR pH_2-pH_2 intermolecular potential [45]. The above Hamiltonian contains full vibration-rotation coupling.

A sine DVR grid [52] is used for the r_1 and r_2 degrees of freedom, while parity-adapted rovibrational basis functions are used for the angular part. A complete product basis function is

$$u_{l_1, l_2, m_2, K}^{J, M, P} = \frac{1}{\sqrt{2(1 + \delta_{m_2, 0} \delta_{K, 0})}} [|l_1, l_2, m_2, K; J, M\rangle + (-1)^{J+P} |l_1, l_2, -m_2, -K; J, M\rangle]$$
(8)

where K > 0, and P = 0 and 1 correspond to even and odd parities, respectively. If K = 0, then it is necessary to apply the constraint $m_2 \ge 0$ to get rid of the redundant basis. The combination $m_2 = K = 0$ and $(-1)^{J+P} = -1$ is not allowed. The parity-adapted basis makes it possible to calculate even and odd parity levels separately. Within each parity block, we use the symmetry-adapted Lanczos algorithm (SAL) [53, 54] to compute states that are symmetric (A) and antisymmetric (B) with respect to exchange of the two pH₂ or oD₂ in trimers of CO–(pH₂)₂ or CO–(oD₂)₂, while no exchange symmetry applied for CO–pH₂–He Trimer.

3 Results and discussion

3.1 Features of the five-dimensional potential energy surface

Figure 3 shows how the well depth of our vibrationally averaged five-dimensional V_{MLRO} ground-state PES for



Fig. 2 M is the center of mass of CO, and B is the canonical point for the Radau vectors. \mathbf{q}_1 and \mathbf{q}_2 are Jacobi vectors. \mathbf{r}_1 and \mathbf{r}_2 are Radau vectors. ϕ is a dihedral angle between \mathbf{r}_1 and \mathbf{r}_2 around \mathbf{r}_0 . $\theta_1(\theta_2)$ are angles between \mathbf{r}_0 and $\mathbf{r}_1(\mathbf{r}_2)$



Fig. 3 Minimum energy on our vibrationally averaged five-dimensional PES for $CO(v = 0)-(pH_2)_2$ trimer as a function of angles φ and ϑ_1 , for optimized values of ϑ_2 , q_1 , and q_2

 $CO(\upsilon = 0) - (pH_2)_2$ system depends on φ and ϑ_1 , when ϑ_2, q_1, q_2 are optimized to minimize the energy for each (φ, ϑ_1) . As seen there, the global minima with well depth of -126.06 cm^{-1} occur at the geometries ($\vartheta_1 = 95.6^\circ$, $\vartheta_2 = 95.6^{\circ}$) with $q_1 = q_2 = 3.57$ Å and $\varphi = 58.0^{\circ}$ (or by symmetry of $\varphi = 302.0^{\circ}$). Between these two identical global minima, there exists a saddle point with energy -101.90 cm^{-1} located at $\vartheta_1 = \vartheta_2 = 95.3^\circ$ with $q_1 = q_2 = 3.57$ Å and $\varphi = 180.0^\circ$, which corresponds to a coplanar geometry with the two pH₂ molecules on opposites of the CO molecule. In addition, Fig. 3 also shows local minimum with energy of -77.59 cm⁻¹ occurs at the coplanar geometry ($\vartheta_1 = 94.7^\circ$, $\vartheta_2 = 180.0^\circ$) with $q_1 = 3.56$ Å, $q_2 = 7.01$ Å, and $\varphi = 0.0^\circ$. Parameters characterizing the various stationary configurations and energies for the CO– $(pH_2)_2$ trimer are summarized in Table 1, where

	Global minimum	Saddle point	Local minimum
CO-(pH ₂₎₂ (V _{MLRQ})	{3.57, 3.57, 95.6, 95.6, 58.0, -126.06}	{3.57, 3.57, 95.3, 95.3, 180.0, -101.90}	{3.56, 7.01, 94.7, 180.0, 0.0, -77.59}
CO-(pH ₂) ₂ (V ₁₂)	$\{3.57, 3.57, 94.7, 94.7, 58.0, -126.22\}$	$\{3.57, 3.57, 94.7, 94.7, 180.0, -102.07\}$	$\{3.56, 7.01, 94.0, 180.0, 0.0, -77.74\}$
CO-(oD ₂) ₂ (V _{MLRQ})	$\{3.57, 3.57, 95.2, 95.2, 58.0, -126.91\}$	$\{3.56, 3.56, 94.6, 94.6, 180.0, -102.76\}$	$\{3.56, 7.01, 96.0, 180.0, 0.0, -78.07\}$
CO-(He) ₂ (V _{MLRQ})	$\{3.39, 3.39, 120.8, 120.8, 51.9, -53.47\}$	$\{3.39, 3.39, 120.8, 120.8, 180.0, -45.90\}$	$\{3.37, 6.31, 118.3, 0.0, 0.0, -32.22\}$

Table 1 Properties of stationary points of the $CO(v = 0)-(pH_2)_2$ potential energy surface, and comparisons with those results for $CO(v = 0)-(oD_2)_2$ and $CO(v = 0)-(He)_2$ surfaces

All entries are given as $\{q_1[Å], q_2[Å], \vartheta_1^\circ, \vartheta_2^\circ, \varphi^\circ, \Delta V[\text{cm}^{-1}]\}$

they are consistent with corresponding values obtained on V_{12} PES, and compared with properties of the analogous $CO-(oD_2)_2$ and $CO-(He)_2$ species. As shown in Table 1, all the stationary points (the global minimum, saddle point, and local minimum) for $CO-(pH_2)_2$ are more than twice as deep as those of $CO-(He)_2$. This will be mostly due to the polarizability of pH₂ (5.414 au) [55] is four times larger than that for He (1.383 au) [56]. Comparing CO– $(pH_2)_2$ and $CO_{-}(oD_2)_2$ yields the differences at the global minimum, saddle point, and local minimum are 0.85, 0.86, and 0.48 cm⁻¹, respectively. Those differences originate from the fact that $B_{D2}/B_{H2} \approx 0.5$, which means the solution of the two-dimensional angular rotational equation for oD_2 systems will have substantially larger contributions from basis functions corresponding to L > 0 [43], where L is the angular momentum quantum number of H₂ rotation. Figure 4 shows the minimum energy paths between equivalent global minima as a function of φ , for optimized values of $\vartheta_1, \vartheta_2, q_1 \text{ and } q_2.$

As clearly shown in Fig. 4(b), the comparison between $CO-(pH_2)_2$ and $CO-(He)_2$ shows that the anisotropy with respect to the dihedral torsional motion of the two pH_2 molecules in $CO-(pH_2)_2$ (48.47 cm⁻¹) is more than twice as strong as that for the analogous motion in the $CO-(He)_2$ (21.25 cm⁻¹) complex. It also shows that the barrier between two adjacent minima of $CO-(pH_2)_2$ (24.16 cm⁻¹) is more than three times higher than that for $CO-(He)_2$ (7.57 cm⁻¹). As would be expected, these barrier heights are approximately equal to the well depths for pH_2-pH_2 (24.71 cm⁻¹) [45] and He–He (7.65 cm⁻¹) [47] dimers, respectively. These differences in PESs will still be reflected in the different spectra of the Trimers.

3.2 Rovibrational energy levels and band origin shifts

The calculated intermolecular rovibrational energy levels for the ground ($\upsilon = 0$) states of CO–(pH₂)₂, CO–(oD₂)₂, and CO–pH₂–He trimers with J = 0, 1, 2, and 3 are listed in Table 2, which are relative to the corresponding zero point energies of -42.907, -59.326, and -26.671 cm⁻¹, respectively. Those for excited ($\upsilon = 1$) (not shown here) are very



Fig. 4 Minimum energy path on our vibrationally averaged fivedimensional PES for CO(v = 0)– $(pH_2)_2$ trimer as functions of angle φ for optimized values of ϑ_1 , ϑ_2 , and q_1 , q_2

similar and listed in Table S1 of the supplementary material. The rovibrational energy levels are labeled by quantum numbers $(J, P, n_{J,P})$. The only good quantum numbers are the overall angular momentum J and parity P = e or f. The label $n_{J,P}$ numbers consecutive states for each (J,P). As shown in Table 2, symmetry (A) and antisymmetry (B) are used to label the symmetries with respect to permutation of the two pH₂ or oD₂ molecules, (A, e) and (B, e) for even parity and (A, f) and (B, f) for odd parity. Because of the zero nuclear spin of the ⁴He, pH₂, oD₂, therefore for CO– (pH₂)₂, CO–(He)₂ and CO–(oD₂)₂ trimers, only (A, e) and (A, f) states are physically allowed. However, for CO–pH₂– He trimer, without exchange symmetry between pH₂ and He, only even and odd parity blocks are used to sort the rovibrational energy levels.

In order to assign the torsional motion states for CO– $(pH_2)_2$ trimer, it is very useful to yield a one-dimensional representation of the partial wave functions along the

Theor Chem Acc (2014) 133:1568

Table 2 Calculated $J = 0, 1,$	$CO(v = 0) - (pH_2)_2$				$CO(v = 0) - (oD_2)_2$				$CO(v = 0)-pH_2-He$			
(in cm^{-1}) for CO($v = 0$)–	(A,e)	(B,e)	(A,f)	(B,f)	(A,e)	(B,e)	(A,f)	(B,f)	e	f		
$(pH_2)_2$, $CO(v = 0)-(oD_2)_2$, and $CO(v = 0)-pH_2$ -He	J = 0											
complexes, which are relative	0.000 (0)	5.477	14.782	3.407 (1)	0.000 (0)	4.852	12.839	2.728 (1)	0.000 (0)	4.065 (1)		
to the corresponding zero point $2007 - 50326$	5.340 (2)	12.753	22.294	10.947 (3)	7.299 (2)	13.559	22.401	11.516 (3)	2.580 (2)	8.378 (3)		
and -26.671 cm^{-1} , respectively	6.951	14.491	24.140	14.932	7.915	15.141	23.522	14.116	4.991	11.280		
	11.901 (4)	18.603	27.663	19.647 (5)	11.140 (4)	18.436	27.299	16.553 (5)	6.651	11.407		
	J = 1											
	3.597	1.352	0.736	1.136	3.001	0.773	0.399	0.692	1.197	0.517		
	7.662	3.943	3.438	5.973	6.038	3.103	3.005	5.219	4.058	0.985		
	11.532	7.972	7.398	6.078	11.856	8.093	5.977	7.956	4.362	3.091		
	14.271	8.708	7.745	8.125	13.123	9.437	7.752	8.283	5.354	3.897		
	J = 2											
	2.083	2.426	4.126	2.945	1.181	1.410	2.508	1.650	1.494	2.319		
	4.074	4.758	4.865	4.857	2.521	3.910	3.740	3.830	1.778	3.235		
	4.558	7.337	8.609	5.168	3.754	5.958	6.839	3.941	3.080	4.848		
	7.307	8.967	9.926	10.283	6.659	8.591	9.742	8.589	4.197	5.198		
	J = 3											
	6.326	5.199	4.022	4.234	3.707	2.947	2.317	2.476	3.969	2.792		
	6.776	6.713	6.193	6.436	4.832	4.896	3.770	4.986	4.501	3.097		
The torsional motion	7.754	6.974	6.443	7.612	5.785	5.104	4.858	5.264	6.028	4.150		
assignments are indicated in	10.860	8.148	7.705	9.314	8.049	5.270	5.790	7.075	6.309	5.538		

ass parentheses as v_t

dihedral angle ϕ . In terms of the basis functions and Radau coordinate system, the wavefunction for the *n*'th energy level for a given set of $\{J, M, P\}$ quantum numbers is,

$$\begin{split} \Psi_{n}^{J,M,P}(r_{1},r_{2},\theta_{1},\theta_{2},\phi;\alpha,\beta,\gamma) \\ &= \sum_{\alpha_{1},\alpha_{2},l_{1},l_{2},m_{2},K} \langle \theta_{1},\theta_{2},\phi;\alpha,\beta,\gamma \mid l_{1},l_{2},m_{2};J,K,M,P\rangle\langle r_{1}\mid\alpha_{1}\rangle\langle r_{2}\mid\alpha_{2}\rangle \\ &\times \left\langle \alpha_{1},\alpha_{2},l_{1},l_{2},m_{2};J,K,M,P\mid\Psi_{n}^{J,M,P}\right\rangle, \end{split}$$

$$(9)$$

in which $\langle r_i | \alpha_i \rangle$ is a localized DVR basis function for particle j, and the coefficient $\langle \alpha_1, \alpha_2, l_1, l_2, m_2; J, K, M, P \mid \Psi_n^{J,M,P} \rangle$ is an eigenvector obtained via Lanczos diagonalization. The one-dimensional wave function along the dihedral angle ϕ , $\Psi_n^{J,M,P}$ is defined as

$$\psi^{J,M,P,n}(\phi) = \int dr_1 dr_2 \sin \theta_1 d\theta_1 \sin \theta_2 d\theta_2 \Psi_n^{J,M,P}$$

$$\times (r_1, r_2, \theta_1, \theta_2, \phi; \alpha, \beta, \gamma)$$
(10)

The above integral can be evaluated on the same grid as the potential integral. The torsional motion states for CO- $(pH_2)_2$ trimer are assigned and indicated in parentheses as (v_t) in Table 2, using the partial wave functions, which are vielded by integrating over all degrees of freedom except the dihedral angle ϕ . We make the assignment based on the nodal structure of the partial wave function shown in Fig. 5.

Table 3 presents the calculated zero point energies (ZPEs) and band origin shifts for CO containing dimer or trimer complexes formed from CO(v) with one or two pH₂, oD₂, He, or mixed one pH₂ and one He. ZPEs are relative to the energies at the dissociation limits for corresponding dimers or trimers. All the band origin shifts are calculated by $\Delta v_0 = E_{\nu=1}^{\text{ZPE}} - E_{\nu=0}^{\text{ZPE}}$. As shown in Table 3, the band origin shift for CO-(pH₂)₂ trimer predicted by our five-dimensional reduced-dimension V_{MLRO} PES is -0.369 cm⁻¹, which is almost twice the value of -0.179 cm^{-1} for COpH₂ dimer and slightly lower than the estimated experimental value of -0.355 cm⁻¹ [31], which was indirectly determined by combining the microwave and the a-type infrared transitions, assuming that the rotational frequencies are the same in the ground and first-excited vibrational states. For $CO-(oD_2)_2$ or $CO-(He)_2$ trimer, the predicted band origin shift of -0.420 or -0.049 cm⁻¹, is approximately equal to twice of -0.206 or -0.024 cm⁻¹ for CO-oD₂ or CO-He dimer.

For CO-pH2-He trimer, two CO-He potentials of CBS + Corr [57] and V333 [46] were used for comparison. The CBS + Corr surface by Peterson and McBane was a three-dimensional ab initio PES that explicitly incorporates dependence on the C-O stretching, obtained with state-ofthe-art ab initio and extrapolation methods. The V333 surface [46] by Chuaqui et al. [57] was a two-dimensional PES with CO fixed at its equilibrium geometry, obtained by fitting

Fig. 5 Partial wave functions for the torsional motion v_t of CO–(pH₂)₂ complex as a function of the azimuthal angle ϕ , compared with those of CO– (oD₂)₂, CO–(He)₂, CO–pH₂–He complexes



a potential to observed transitions near the fundamental band of CO. As we expected, the calculated band origin shift for CO–pH₂–He trimer is -0.205 cm^{-1} on CBS + Corr potential for CO–He, which is approximately equal to the sum of the band origin shifts of -0.179 and -0.024 cm^{-1} for CO– pH₂ and CO–He dimers, respectively. This indicates that the density for pH₂ or He in CO–pH₂–He trimer has very similar distribution as that in their dimers. Due to V333 potential for CO–He without taking into account the CO vibration, the predicted band origin shift on this surface for CO–pH₂–He trimer is -0.019 cm^{-1} different from that on CBS + Corr potential, which is very close to the calculated shift value of -0.024 cm^{-1} for CO–He dimer.

3.3 Predicted microwave and infrared transitions

For CO–(pH₂)₂, the predicted microwave (MW) and infrared (IR) transition frequencies (cm⁻¹) from our AHR 5D V_{MLRQ} potential are listed in Tables 4 and 5, respectively, and compared with those calculated on V_{12} potential and experimental results. The lower and upper states involved in the transitions are numbered with the set of parameters (*J*, *P*, *n*_{*J*,*P*}). For infrared transitions, to compare our computed energy transitions directly with experimental results, we calculate IR rovibrational transitions from

$$\upsilon = \upsilon_0(\text{CO}) + E_{\nu=1}^{\text{upper}} - E_{\nu=0}^{\text{lower}}$$
(11)

Table 3 The zero point energies (ZPEs) and band origin shifts for the complexes formed from stretch CO(v) with one or two pH₂, oD₂, or mixed one pH2 and one He

Complexes	$E_{\nu=0}^{\text{ZPE}}$	$E_{\nu=1}^{\text{ZPE}}$	Δv_0 (calc.)	Δv_0 (obs.)
$\overline{CO(\upsilon)-pH_2}$	-19.371	-19.550	-0.179	-0.179^{a}
CO(v)–(pH ₂₎₂	-42.907	-43.276	-0.369	-0.355^{b}
$CO(v)-oD_2$	-25.630	-25.836	-0.206	-0.202^{c}
$CO(\upsilon)$ – $(oD_2)_2$	-59.326	-59.746	-0.420	
CO(v)-He ^f	-6.431	-6.455	-0.024	-0.025^{d}
CO(υ)–(He) ₂ ^f	-13.130	-13.179	-0.049	-0.049^{e}
CO(v)-pH ₂ -He ^g	-26.671	-26.857	-0.186	
$CO(\upsilon)$ -pH ₂ -He ^f	-26.459	-26.664	-0.205	

ZPEs are relative to the energies at the dissociation limits for corresponding dimers or trimers. $\Delta v_0 = E_{\nu=1}^{ZPE} - E_{\nu=0}^{ZPE}$ is the resulting band origin shifts. All energies are in cm⁻¹

- ^a Reference [35]. ^b Reference [31]. ^c Reference [58]
- ^d Reference [36]. ^e Reference [38]
- ^f This calculation is with the CBS + Corr potential [57]
- ^g This calculation is with the V333 potential [46]

where, $v_0(CO) = 2,143.2712 \text{ cm}^{-1}$ is the experimental fundamental vibrational transition frequency of a free CO molecule.

As shown in column 2 of Tables 4 or 5, the transition energies yielded by our VMLRO PES are seen to agree very well with the experimental values shown in column 3 of Tables 4 or 5. The differences seen in column 5 are very small both for microwave and infrared transitions, yielding a root-mean-square (rms) discrepancy of only 0.010 cm^{-1} for all 4 observed transitions (1 for MW and 3 for IR), and also consistent with those obtained on the V_{12} potential

as shown in column 5 with the same rms discrepancy for those 4 experimental transitions. To predict more possible microwave and infrared transitions for CO-(pH₂)₂ trimer, the line strength and relative intensity at temperatures 0.6 K (effective experimental rotational temperature) are also computed from the wave functions. Note that the line strength does not depend on the temperature. Subject to the selection rules $\Delta J = 0, \pm 1, \Delta P = e \leftrightarrow f$, and with line strength larger than $10^{-3}(\mu_{CO}^2)$, 64 possible allowed transitions for MW and 129 for IR with $J \leq 3$ and $n \leq 4$ are predicted. To select most possible microwave and infrared transitions for $CO_{(pH_2)_2}$ trimer, all the relative intensities at temperature 0.6 K are calculated, and expressed relative to (1, f, 2)-(0, e, 1) transition, whose intensity is set to 1. With the restriction of the relative intensity larger than 10^{-3} , only 15 strongest transitions for MW and 18 for IR are kept and listed in Tables 4 and 5, respectively. The predicted 129 line strengths and 18 relative intensities for CO- $(pH_2)_2$ trimer at temperature of 0.6 K are clearly illustrated in the uppermost segment of Fig. 6.

For CO-(oD₂)₂ and CO-pH₂-He, the predicted infrared transition frequencies expressed relative to the band origin 2,143.2712 cm⁻¹ of a free CO from our AHR 5D V_{MLRO} potential are given in Tables 6 and 7, respectively. The lower and upper states involved in the transitions are also numbered with the set of parameters (J, P, n_{IP}) . Only the transitions between A+ and A- states are physically allowed for $CO_{-}(oD_{2})_{2}$, but this restriction does not apply to CO–pH₂–He. Subject to overall selection rules $\Delta J = 0$, ± 1 , $\Delta P = e \leftrightarrow f$, with line strength larger than 10^{-3} (μ_{CO}^2) , $J \leq 3$ and $n \leq 4$, 149 line strengths and 32 most possible allowed IR transitions at temperature of 0.6 K for

Table 4 Predicted microwavefrequencies (cm^{-1}) , linestrengths, and relativeinteractive of	Transitions	Diff. (Cal	lc. – Obs.)	Line	Intensity		
	Upper-lower	V _{MLRQ}	Obs.	V ₁₂	V _{MLRQ}	strength	(0.6 K)
$0.6 \text{ K for CO}_{(\text{pH}_2)_2} \text{ complex}$	(1, f, 1)–(0, e, 1)	0.736	0.741	0.002	-0.005	0.061	0.047
from vibrationally averaged	(2, f, 2)–(2, e, 1)	2.783				1.853	0.010
five-dimensional V_{12} or V_{12} pr	(1, e, 1)–(1, f, 1)	2.862				0.957	0.126
experimental results	(1, f, 2)–(0, e, 1)	3.438				1.294	1.000
1	(2, e, 3)–(1, f, 1)	3.822				0.679	0.090
	(3, f, 2)–(2, e, 1)	4.110				0.521	0.003
	(1, f, 3)–(2, e, 1)	5.315				0.225	0.001
	(3, f, 4)–(2, e, 1)	5.622				0.162	0.001
	(0, e, 3)–(1, f, 1)	6.951				0.205	0.158
	(2, f, 3)–(2, e, 1)	6.526				0.312	0.002
	(1, e, 2)–(1, f, 1)	6.926				0.079	0.011
The lower and upper states	(1, f, 3)–(0, e, 1)	7.236				0.105	0.082
involved in the transitions are numbered with the set of parameters $(J, P, n_{J,P})$. The line strength values are multiplied by 100	(1, f, 4)–(0, e, 1)	7.398				0.326	0.252
	(2, f, 4)–(2, e, 1)	7.844				0.489	0.003
	(1, e, 3)–(1, f, 1)	10.796				0.141	0.019

by 100

Table 5 Predicted infraredtransition frequencies (cm^{-1})	Transitions			Diff. (ca	lc. – obs.)	Line	Intensity
line strengths, and relative	Upper-lower	V _{MLRQ}	Obs.	V ₁₂	V _{MLRQ}	Strength	(0.6 K)
$0.6 \text{ K for CO}_{(\text{pH}_2)_2} \text{ complex}$	(0, e,1)–(1, f, 1)	2,142.167				0.062	0.008
from vibrationally averaged	(1, f, 1)–(0, e, 1)	2,143.638	2,143.657	0.007	0.019	0.001	0.051
five-dimensional V_{12} or V_{MLRQ} PES, comparison with experimental results	(2, e, 1)–(1, f, 1)	2,144.250				0.051	0.007
	(2, f, 1)–(2, e, 1)	2,144.945				0.100	0.001
	(2, f, 2)–(2, e, 1)	2,145.671				1.848	0.010
	(1, e, 1)–(1, f, 1)	2,145.747				0.959	0.127
	(1, f, 2)–(0, e, 1)	2,146.324	2,146.330	0.015	0.006	1.293	1.000
	(2, e, 3)–(1, f, 1)	2,146.713	2,146.712	0.011	-0.001	0.673	0.089
	(3, f, 2)–(2, e, 1)	2,147.586				0.514	0.003
	(1, f, 3)–(2, e, 1)	2,148.201				0.224	0.001
	(3, f, 4)–(2, e, 1)	2,148:893				0:161	0.001
	(0, e, 3)–(1, f, 1)	2,149.108				0.204	0.027
	(2, f, 3)–(2, e, 1)	2,149.414				0.311	0.002
	(1, e, 2)–(1, f, 1)	2,149.809				0.079	0.010
The lower and upper states involved in the transitions	(1, f, 3)–(0, e, 1)	2,150.283				0.103	0.079
are numbered with the set of	(1, f, 4)–(0, e, 1)	2,150.637				0.330	0.256
parameters $(J, P, n_{J,P})$. The line	(2, f, 4)–(2, e, 1)	2,150.733				0.498	0.003
strength values are multiplied by 10	(1, e, 3)–(1, f, 1)	2,153.693				0.141	0.019

 $CO_{-}(oD_2)_2$, and 118 line strengths and 37 most possible IR transitions for CO-pH2-He are predicted, and shown in the middle and lowermost segments of Fig. 6, respectively. Although MW or IR transitions for $CO_{-}(oD_{2})_{2}$ have not been reported yet, one transition marked "m" in Fig. 7 of Ref. [22] was tentatively assigned as b-type R(0) of the mixed ternary cluster CO-pH2-He. Based on the calculated line position and intensity, this line may correspond to the (1, f, 4)–(0, e, 1) transition. We are confident for this assignment, not only because the calculated transition at 2,146.958 cm⁻¹ are in good agreement with the experimental position at 2,146.93 cm⁻¹, but also the relative intensity for (1, f, 4)-(0, e, 1) at temperature 0.6 K is the strongest among all predicted transitions for CO-pH₂-He complex.

3.4 The wave functions and "solvent" density distributions

Examining the wave function of some of the vibrational states with v = 0 and J = 0 helps us to understand the nature about the motions of the pH₂, oD₂, and He in their complexes. We are most interested in the ϕ dependence of the wave function because this coordinate plays an important role in assigning and understanding many vibrational states. Figure 5 shows the reduced wave functions with (A,e) symmetry and (B,f) symmetry for $v_t = 0, 1, 2, 3,$ 4, and 5 levels of CO-(pH₂)₂, obtained by integrating the wave function over the coordinates of $r_1,\,r_2,\,\theta_1$ and θ_2 at each value of ϕ , and compared with those of CO–(oD₂)₂, $CO-(He)_2$, and $CO-pH_2$ -He trimers. As shown in Fig. 5,

the wave functions for the $v_t = 1, 3$, and 5 levels shown on the right side have strikingly regular nodal structures along the ϕ coordinate with exact zero at $\phi = 0$ or $\phi = 180^{\circ}$, which implies that there is little coupling between ϕ with the other coordinates. While the wave functions for $v_t = 0$, 2, and 4 levels shown on the left side have no strict characteristics as torsional states, which is probably due to the fact that there is strong coupling between ϕ with the other coordinates that distort their character. For CO-(He)2 and CO-pH₂-He trimers, there exist strong coupling between many (A,e) or (A,f) states, and it is hard to assign those states as torsional motions; thus, we only presented the lowest possible torsion states with $v_t < 4$. In Fig. 5, it can be clearly seen that the wave functions for the $v_t = 0, 2, and 4$ levels are quite different between the $CO-(pH_2)_2$ and CO-(He)₂ complexes, but very similar for CO-(pH₂)₂ and CO-(oD₂)₂, and for CO-(He)₂ and CO-pH₂-He complexes. In particular, for the $v_t = 0$ and 1 levels, the highly localized distribution for pH_2 or oD_2 in CO- $(pH_2)_2$ or CO- $(oD_2)_2$ complexes contrasts sharply with that for He in CO-(He)₂ or CO-pH2-He complexes. In addition to the expected excluded volumes near $\phi = 0$ and $\phi = 360$, for the ground state of CO- $(pH_2)_2$ or CO- $(oD_2)_2$, there are local maxima at $\phi = 58$ and $\phi = 302^{\circ}$ corresponding to the pH₂ or oD₂ particles lying at global minimum positions, while the He distribution in CO-(He)2 or CO-pH2-He is delocalized, even with a lot of distribution at $\phi = 180^{\circ}$.

Figure 7 shows the three-dimensional density for the two pH₂ molecules (upper), two He atoms (middle), or mixed
Fig. 6 Calculated infrared spectra of CO-(pH₂)₂ (upper panel), CO-(oD₂)₂ (middle panel), and CO-pH2-He (lower panel) for line strength (upward pointing lines) and relative intensity at 0.6 K (downward pointing lines), respectively. The intensities are expressed relative to (1, f, 2)-(0, e, 1) transition of CO-(pH₂)₂ or $CO-(oD_2)_2$, and (1, f, 4)-(0, e, f, 4)1) transition of CO-pH₂-He complexes, respectively, whose intensity is equal 1. All the line strength values are multiplied by 10. The relative intensity of $CO_{-}(pH_2)_2$ are also multiplied by 100



one pH₂ and one He (bottom) in the body-fixed frame, as calculated for the ground state of CO–(pH₂)₂, CO–(He)₂, or CO–pH₂–He using the procedure described in Ref. [27]. It is very useful to develop some intuitive feeling regarding the nature of these species. Iso-surfaces are used to represent the three-dimensional densities, and the z axis is defined to lie on the CO molecular axis. One significant feature of these representations is the fact that the two pH₂ molecules or two He molecules appear to have very different density distributions. It is clear that the density distribution of the first pH₂ or He particle in CO–(pH₂)₂ or CO–(He)₂ trimer is represented as a at disk and somewhat delocalized in the yz plane, while that of the second pH₂ in CO–(pH₂)₂ trimer is highly localized, with two density maxima at regions

associated with the pH₂–pH₂ potential minimum. In contrast, the density distribution of the second helium atom in CO–(He)₂ trimer (middle panel) is highly delocalized relative to the first, being distributed on an incomplete ring wrapped around the CO molecule. For CO–pH₂–He trimer, the density for pH₂ as particle-1 is located in the *yz* plane as reference, while the density for He as particle-2 is relative to pH₂. As shown in the lowest panel of Fig. 7, the more localized He distribution in CO–pH₂–He than in CO–(He)₂ is related to the stronger interaction (deeper well) between the pH₂–He interaction than the He–He one. For all three cases, the density for particle-2 is excluded from the region near particle-1 by the short-range repulsive wall of the pH₂– pH₂, He–He, or pH₂–He potential.

Table 6 Predicted infrared transition frequencies (cm⁻¹), line strengths, and relative intensities at temperature of 0.6 K for CO– $(oD_2)_2$ complex from vibrationally averaged five-dimensional V_{MLRQ} PES

Table 7	Predicted	infrared	transitic	on freque	ncies	(cm^{-1})), li	ne
strengths	s, and rela	tive inten	sities at to	emperature	e of 0.	6 K fo	or Co	-C
pH ₂ –He	complex	from vib	rationally	averaged	five-d	imensi	onal	V
MLRO PE	S							

Transitions		Line	Intensity
Upper-lower	Calc.	Strength	(0.6 K)
(1, f, 2)–(0, e, 1)	2,145.849	1.050	1.000
(1, e, 1)–(1, f, 1)	2,145.445	1.111	0.406
(2, e, 3)–(1, f, 1)	2,146.199	0.723	0.264
(1, f, 4)–(0, e, 1)	2,150.597	0.270	0.257
(1, f, 3)–(0, e, 1)	2,148.813	0.175	0.167
(2, f, 2)–(2, e, 1)	2,145.404	1.726	0.097
(1, e, 3)–(1, f, 1)	2,154.309	0.241	0.088
(0,e, 2)–(1, f, 1)	2,149.747	0.186	0.068
(1, e, 2)–(1, f, 1)	2,148.475	0.175	0.064
(1, f, 1)–(0, e, 1)	2,143.251	0.064	0.061
(2, e, 4)–(1, f, 1)	2,149.096	0.093	0.034
(3, f, 3)–(2, e, 1)	2,146.523	0.462	0.026
(0, e, 1)–(1, f, 1)	2,142.452	0.061	0.022
(2, e, 1)–(1, f, 1)	2,143.633	0.060	0.022
(2, f, 4)–(2, e, 1)	2,151.396	0.258	0.014
(1, f, 4)–(2, e, 1)	2,149.416	0.179	0.010
(2, f, 3)–(2, e, 1)	2,148.495	0.160	0.009
(3, f, 4)–(2, e, 1)	2,147.451	0.141	0.008
(3, e, 2)–(3, f, 1)	2,145.361	1.616	0.006
(1, f, 2)–(2, e, 1)	2,144.668	0.093	0.005
(1, f, 3)–(2, e, 1)	2,147.632	0.086	0.005
(1, f, 1)–(2, e, 1)	2,142.070	0.057	0.003
(3, f, 4)–(2, e, 2)	2,146.111	0.452	0.001
(3, e, 3)–(2, f, 1)	2,146.119	0.423	0.001
(1, f, 1)–(1, e, 1)	2,140.250	1.105	0.001
(0, e, 1)–(1, f, 2)	2,139.846	1.049	0.001
(1, f, 2)–(2, e, 2)	2,143.328	0.313	0.001
(1, e, 3)–(2, f, 1)	2,152.200	0.287	0.001
(1, e, 2)–(2, f, 1)	2,146.366	0.271	0.001
(2, f, 2)–(2, e, 2)	2,144.063	0.263	0.001
(2, e, 4)–(2, f, 1)	2,146.988	0.239	0.001
(1, f, 3)–(2, e, 2)	2,146.292	0.243	0.001

The lower and upper states involved in the transitions are numbered with the set of parameters $(J, P, n_{J,P})$. The line strength values are multiplied by 10

Figure 8 shows the "Solvent" pH₂(upper panels), He (middle panels), and pH₂–He (lower panels) densities in the first three torsional levels of CO–(pH₂)₂, CO–(He)₂, and CO–pH₂–He, respectively. For the $v_t = 0$ ground states, the density distribution patterns of particle-2 across three complexes are quite different, while for $v_t = 1$, 2, and 3 excited states, the density distribution patterns are similar among the three species, because at these levels energy patterns are determined by the nodal structures instead of

Upper-lowerCalc.Strength (0.6 K) $(1, f, 4)-(0, e, 1)$ $2,146.958^a$ 1.066 1.000 $(1, e, 2)-(1, f, 1)$ $2,146.599$ 0.898 0.244 $(1, f, 1)-(0, e, 1)$ $2,143.601$ 0.149 0.140 $(1, e, 3)-(1, f, 1)$ $2,146.908$ 0.374 0.102 $(0, e, 3)-(1, f, 1)$ $2,147.539$ 0.285 0.077 $(1, e, 3)-(1, f, 2)$ $2,146.440$ 0.627 0.055 $(1, e, 2)-(1, f, 2)$ $2,146.131$ 0.540 0.048 $(0, e, 1)-(1, f, 1)$ $2,142.569$ 0.148 0.040 $(0, f, 1)-(1, e, 1)$ $2,145.929$ 0.715 0.038 $(2, e, 1)-(1, f, 1)$ $2,144.061$ 0.130 0.035 $(2, e, 4)-(1, f, 1)$ $2,144.729$ 0.174 0.017 $(1, e, 1)-(1, f, 2)$ $2,143.299$ 0.186 0.016 $(2, f, 4)-(1, e, 1)$ $2,146.770$ 0.474 0.012 $(2, f, 4)-(1, e, 1)$ $2,146.710$ 0.229 0.012 $(1, f, 2)-(1, e, 1)$ $2,142.876$ 0.186 0.010 $(2, f, 3)-(2, e, 1)$ $2,146.129$ 0.712 0.009 $(3, f, 4)-(2, e, 1)$ $2,144.1209$ 0.122 0.006 $(2, e, 1)-(1, e, 1)$ $2,144.209$ 0.122 0.006 $(2, e, 1)-(1, e, 2)$ $2,$	Transitions		Line	Intensity
$(1, f, 4)-(0, e, 1)$ $2, 146.958^a$ 1.066 1.000 $(1, e, 2)-(1, f, 1)$ $2, 143.601$ 0.149 0.140 $(1, e, 3)-(1, f, 1)$ $2, 143.601$ 0.149 0.140 $(1, e, 3)-(1, f, 1)$ $2, 146.908$ 0.374 0.102 $(0, e, 3)-(1, f, 1)$ $2, 146.908$ 0.374 0.102 $(0, e, 3)-(1, f, 2)$ $2, 146.440$ 0.627 0.055 $(1, e, 2)-(1, f, 2)$ $2, 146.131$ 0.540 0.048 $(0, e, 1)-(1, f, 1)$ $2, 142.569$ 0.148 0.040 $(0, f, 1)-(1, e, 1)$ $2, 145.929$ 0.715 0.038 $(2, e, 1)-(1, f, 1)$ $2, 144.061$ 0.130 0.035 $(2, e, 4)-(1, f, 1)$ $2, 144.729$ 0.194 0.017 $(1, e, 1)-(1, f, 2)$ $2, 148.729$ 0.194 0.017 $(1, e, 1)-(1, f, 2)$ $2, 148.729$ 0.186 0.016 $(2, f, 4)-(2, e, 1)$ $2, 146.700$ 0.474 0.012 $(2, f, 3)-(1, e, 1)$ $2, 146.700$ 0.474 0.012 $(2, f, 3)-(1, e, 1)$ $2, 146.700$ 0.712 0.009 $(3, f, 4)-(2, e, 1)$ $2, 146.129$ 0.712 0.009 $(3, f, 4)-(2, e, 1)$ $2, 144.593$ 0.060 0.005 $(2, e, 2)-(1, f, 2)$ $2, 144.881$ 0.079 0.007 $(2, f, 1)-(1, e, 1)$ $2, 144.209$ 0.122 0.006 $(2, e, 1)-(1, e, 2)$ $2, 144.382$ 0.072 0.003 $(3, f, 4)-(2, e, 2)$ $2, 146.829$ 0.187 0.003 $(2$	Upper-lower	Calc.	Strength	(0.6 K)
(1, e, 2)-(1, f, 1) $2, 146.599$ 0.898 0.244 $(1, f, 1)-(0, e, 1)$ $2, 143.601$ 0.149 0.140 $(1, e, 3)-(1, f, 1)$ $2, 146.908$ 0.374 0.102 $(0, e, 3)-(1, f, 1)$ $2, 147.539$ 0.285 0.077 $(1, e, 3)-(1, f, 2)$ $2, 146.440$ 0.627 0.055 $(1, e, 2)-(1, f, 2)$ $2, 146.131$ 0.540 0.048 $(0, e, 1)-(1, f, 1)$ $2, 142.569$ 0.148 0.040 $(0, f, 1)-(1, e, 1)$ $2, 145.929$ 0.715 0.038 $(2, e, 1)-(1, f, 1)$ $2, 144.761$ 0.130 0.035 $(2, e, 4)-(1, f, 1)$ $2, 146.761$ 0.112 0.030 $(2, f, 3)-(2, e, 1)$ $2, 146.761$ 0.112 0.030 $(2, f, 4)-(1, e, 1)$ $2, 147.067$ 0.271 0.014 $(2, f, 4)-(1, e, 1)$ $2, 147.067$ 0.271 0.014 $(2, f, 4)-(2, e, 1)$ $2, 146.770$ 0.474 0.012 $(2, f, 3)-(1, e, 1)$ $2, 146.710$ 0.229 0.012 $(1, f, 2)-(1, e, 1)$ $2, 143.881$ 0.079 0.007 $(2, f, 3)-(2, e, 2)$ $2, 146.129$ 0.712 0.006 $(2, e, 1)-(1, f, 2)$ $2, 143.881$ 0.079 0.007 $(2, f, 4)-(2, e, 1)$ $2, 142.876$ 0.187 0.003 $(2, e, 1)-(1, f, 2)$ $2, 144.820$ 0.72 0.004 $(1, f, 1)-(2, e, 1)$ $2, 144.209$ 0.122 0.006 $(2, e, 1)-(1, e, 1)$ $2, 144.382$ 0.072 0.003 $(3, f$	(1, f, 4)–(0, e, 1)	2,146.958 ^a	1.066	1.000
(1, f, 1)-(0, e, 1) $2, 143.601$ 0.149 0.140 $(1, e, 3)-(1, f, 1)$ $2, 146.908$ 0.374 0.102 $(0, e, 3)-(1, f, 1)$ $2, 147.539$ 0.285 0.077 $(1, e, 3)-(1, f, 2)$ $2, 146.440$ 0.627 0.055 $(1, e, 2)-(1, f, 2)$ $2, 146.131$ 0.540 0.048 $(0, e, 1)-(1, f, 1)$ $2, 142.569$ 0.148 0.040 $(0, f, 1)-(1, e, 1)$ $2, 145.929$ 0.715 0.038 $(2, e, 1)-(1, f, 1)$ $2, 146.761$ 0.112 0.030 $(2, f, 3)-(2, e, 1)$ $2, 146.761$ 0.112 0.030 $(2, f, 3)-(2, e, 1)$ $2, 148.729$ 0.194 0.017 $(1, e, 1)-(1, f, 2)$ $2, 143.299$ 0.186 0.016 $(2, f, 4)-(1, e, 1)$ $2, 147.067$ 0.271 0.014 $(2, f, 4)-(1, e, 1)$ $2, 146.770$ 0.474 0.012 $(2, f, 3)-(1, e, 1)$ $2, 146.710$ 0.229 0.012 $(1, f, 2)-(1, e, 1)$ $2, 146.710$ 0.229 0.012 $(1, f, 2)-(1, e, 1)$ $2, 142.876$ 0.186 0.010 $(2, f, 3)-(2, e, 2)$ $2, 146.129$ 0.712 0.009 $(3, f, 4)-(2, e, 1)$ $2, 142.108$ 0.127 0.003 $(2, e, 1)-(1, f, 2)$ $2, 143.593$ 0.060 0.005 $(2, f, 1)-(1, e, 1)$ $2, 142.108$ 0.127 0.003 $(2, e, 1)-(1, f, 2)$ $2, 144.691$ 0.066 0.002 $(3, f, 4)-(2, e, 2)$ $2, 146.829$ 0.187 0.001 $(1,$	(1, e, 2)–(1, f, 1)	2,146.599	0.898	0.244
(1, e, 3)-(1, f, 1) $2,146.908$ 0.374 0.102 $(0, e, 3)-(1, f, 1)$ $2,147.539$ 0.285 0.077 $(1, e, 3)-(1, f, 2)$ $2,146.440$ 0.627 0.055 $(1, e, 2)-(1, f, 2)$ $2,146.131$ 0.540 0.048 $(0, e, 1)-(1, f, 1)$ $2,142.569$ 0.148 0.040 $(0, f, 1)-(1, e, 1)$ $2,145.929$ 0.715 0.038 $(2, e, 1)-(1, f, 1)$ $2,144.061$ 0.130 0.035 $(2, e, 4)-(1, f, 1)$ $2,146.761$ 0.112 0.030 $(2, f, 3)-(2, e, 1)$ $2,146.761$ 0.112 0.030 $(2, f, 3)-(2, e, 1)$ $2,148.729$ 0.194 0.017 $(1, e, 1)-(1, f, 2)$ $2,143.299$ 0.186 0.016 $(2, f, 4)-(1, e, 1)$ $2,147.067$ 0.271 0.014 $(2, f, 4)-(2, e, 1)$ $2,146.710$ 0.229 0.012 $(1, f, 2)-(1, e, 1)$ $2,142.876$ 0.186 0.010 $(2, f, 3)-(2, e, 2)$ $2,146.129$ 0.712 0.009 $(3, f, 4)-(2, e, 1)$ $2,147.113$ 0.303 0.008 $(2, e, 2)-(1, f, 2)$ $2,143.881$ 0.079 0.007 $(2, f, 1)-(1, e, 1)$ $2,144.209$ 0.122 0.006 $(2, e, 1)-(1, f, 2)$ $2,144.829$ 0.187 0.002 $(3, f, 4)-(2, e, 1)$ $2,144.209$ 0.127 0.003 $(0, f, 3)-(1, e, 1)$ $2,144.691$ 0.066 0.002 $(2, f, 1)-(2, e, 1)$ $2,144.627$ 1.150 0.001 $(1, f, 2)-(2, e, 1)$ <	(1, f, 1)–(0, e, 1)	2,143.601	0.149	0.140
(0, e, 3)-(1, f, 1) $2,147.539$ 0.285 0.077 $(1, e, 3)-(1, f, 2)$ $2,146.440$ 0.627 0.055 $(1, e, 2)-(1, f, 2)$ $2,146.131$ 0.540 0.048 $(0, e, 1)-(1, f, 1)$ $2,142.569$ 0.148 0.040 $(0, f, 1)-(1, e, 1)$ $2,145.929$ 0.715 0.038 $(2, e, 1)-(1, f, 1)$ $2,144.061$ 0.130 0.035 $(2, e, 4)-(1, f, 1)$ $2,146.761$ 0.112 0.030 $(2, f, 3)-(2, e, 1)$ $2,148.729$ 0.194 0.017 $(1, e, 1)-(1, f, 2)$ $2,143.299$ 0.186 0.016 $(2, f, 4)-(1, e, 1)$ $2,147.067$ 0.271 0.014 $(2, f, 4)-(2, e, 1)$ $2,146.710$ 0.229 0.012 $(1, f, 2)-(1, e, 1)$ $2,147.067$ 0.271 0.009 $(3, f, 4)-(2, e, 1)$ $2,147.113$ 0.303 0.008 $(2, e, 2)-(1, f, 2)$ $2,143.881$ 0.079 0.007 $(2, f, 1)-(1, e, 1)$ $2,144.209$ 0.122 0.006 $(2, e, 1)-(1, f, 2)$ $2,143.593$ 0.060 0.005 $(2, f, 4)-(2, e, 2)$ $2,146.486$ 0.297 0.004 $(1, f, 1)-(2, e, 1)$ $2,142.108$ 0.127 0.003 $(0, f, 3)-(1, e, 1)$ $2,142.108$ 0.127 0.003 $(0, f, 3)-(1, e, 1)$ $2,144.691$ 0.066 0.002 $(2, f, 1)-(2, e, 1)$ $2,144.691$ 0.066 0.002 $(3, f, 2)-(2, e, 1)$ $2,144.691$ 0.066 0.002 $(3, f, 2)-(2, e, 1)$ <	(1, e, 3)–(1, f, 1)	2,146.908	0.374	0.102
(1, e, 3)-(1, f, 2) $2,146.440$ 0.627 0.055 $(1, e, 2)-(1, f, 2)$ $2,146.131$ 0.540 0.048 $(0, e, 1)-(1, f, 1)$ $2,142.569$ 0.148 0.040 $(0, f, 1)-(1, e, 1)$ $2,145.929$ 0.715 0.038 $(2, e, 1)-(1, f, 1)$ $2,144.061$ 0.130 0.035 $(2, e, 4)-(1, f, 1)$ $2,146.761$ 0.112 0.030 $(2, f, 3)-(2, e, 1)$ $2,146.413$ 0.781 0.020 $(0, e, 4)-(1, f, 2)$ $2,148.729$ 0.194 0.017 $(1, e, 1)-(1, f, 2)$ $2,143.299$ 0.186 0.016 $(2, f, 4)-(2, e, 1)$ $2,146.770$ 0.474 0.012 $(2, f, 3)-(1, e, 1)$ $2,146.770$ 0.474 0.012 $(2, f, 3)-(1, e, 1)$ $2,146.710$ 0.229 0.012 $(1, f, 2)-(1, e, 1)$ $2,146.710$ 0.229 0.012 $(1, f, 2)-(1, e, 1)$ $2,144.129$ 0.712 0.009 $(3, f, 4)-(2, e, 1)$ $2,147.113$ 0.303 0.008 $(2, e, 2)-(1, f, 2)$ $2,143.593$ 0.660 0.005 $(2, f, 4)-(2, e, 2)$ $2,146.486$ 0.297 0.004 $(1, f, 1)-(2, e, 1)$ $2,142.108$ 0.127 0.003 $(0, f, 3)-(1, e, 1)$ $2,142.108$ 0.127 0.003 $(0, f, 3)-(1, e, 1)$ $2,144.691$ 0.066 0.002 $(2, f, 1)-(2, e, 2)$ $2,144.691$ 0.066 0.002 $(3, f, 1)-(2, e, 1)$ $2,142.295$ 0.078 0.001 $(3, e, 3)-(3, f, 1)$ <	(0, e, 3)–(1, f, 1)	2,147.539	0.285	0.077
(1, e, 2)-(1, f, 2) $2, 146.131$ 0.540 0.048 $(0, e, 1)-(1, f, 1)$ $2, 142.569$ 0.148 0.040 $(0, f, 1)-(1, e, 1)$ $2, 145.929$ 0.715 0.038 $(2, e, 1)-(1, f, 1)$ $2, 144.061$ 0.130 0.035 $(2, e, 4)-(1, f, 1)$ $2, 146.761$ 0.112 0.030 $(2, f, 3)-(2, e, 1)$ $2, 146.761$ 0.112 0.030 $(2, f, 3)-(2, e, 1)$ $2, 148.729$ 0.194 0.017 $(1, e, 1)-(1, f, 2)$ $2, 143.299$ 0.186 0.016 $(2, f, 4)-(1, e, 1)$ $2, 147.067$ 0.271 0.014 $(2, f, 4)-(2, e, 1)$ $2, 146.710$ 0.229 0.012 $(2, f, 3)-(1, e, 1)$ $2, 146.710$ 0.229 0.012 $(1, f, 2)-(1, e, 1)$ $2, 142.876$ 0.186 0.010 $(2, f, 3)-(2, e, 2)$ $2, 146.129$ 0.712 0.009 $(3, f, 4)-(2, e, 1)$ $2, 147.113$ 0.303 0.008 $(2, e, 2)-(1, f, 2)$ $2, 143.593$ 0.060 0.005 $(2, f, 1)-(1, e, 1)$ $2, 143.593$ 0.060 0.005 $(2, f, 4)-(2, e, 2)$ $2, 146.486$ 0.297 0.004 $(1, f, 1)-(2, e, 1)$ $2, 142.108$ 0.127 0.003 $(0, f, 3)-(1, e, 1)$ $2, 153.142$ 0.062 0.003 $(3, f, 4)-(2, e, 2)$ $2, 146.829$ 0.187 0.002 $(3, f, 1)-(2, e, 1)$ $2, 142.579$ 0.060 0.002 $(3, f, 2)-(2, e, 1)$ $2, 142.97$ 0.057 0.001 $(3, e$	(1, e, 3)–(1, f, 2)	2,146.440	0.627	0.055
(0, e, 1)-(1, f, 1) $2,142.569$ 0.148 0.040 $(0, f, 1)-(1, e, 1)$ $2,145.929$ 0.715 0.038 $(2, e, 1)-(1, f, 1)$ $2,144.061$ 0.130 0.035 $(2, e, 4)-(1, f, 1)$ $2,144.061$ 0.112 0.030 $(2, f, 3)-(2, e, 1)$ $2,146.131$ 0.781 0.020 $(0, e, 4)-(1, f, 2)$ $2,148.729$ 0.194 0.017 $(1, e, 1)-(1, f, 2)$ $2,143.299$ 0.186 0.016 $(2, f, 4)-(1, e, 1)$ $2,147.067$ 0.271 0.014 $(2, f, 4)-(2, e, 1)$ $2,146.770$ 0.474 0.012 $(2, f, 3)-(1, e, 1)$ $2,146.710$ 0.229 0.012 $(1, f, 2)-(1, e, 1)$ $2,146.710$ 0.229 0.012 $(1, f, 2)-(1, e, 1)$ $2,146.129$ 0.712 0.009 $(3, f, 4)-(2, e, 1)$ $2,146.129$ 0.712 0.009 $(3, f, 4)-(2, e, 1)$ $2,143.881$ 0.079 0.007 $(2, f, 1)-(1, e, 1)$ $2,144.209$ 0.122 0.006 $(2, e, 1)-(1, f, 2)$ $2,143.593$ 0.060 0.005 $(2, f, 4)-(2, e, 2)$ $2,146.486$ 0.297 0.004 $(1, f, 1)-(2, e, 1)$ $2,142.108$ 0.127 0.003 $(0, f, 3)-(1, e, 1)$ $2,144.382$ 0.072 0.002 $(3, f, 2)-(2, e, 1)$ $2,144.691$ 0.066 0.002 $(3, f, 2)-(2, e, 1)$ $2,144.691$ 0.066 0.002 $(1, f, 2)-(2, e, 1)$ $2,142.279$ 0.078 0.001 $(3, e, 3)-(3, f, 1)$ <	(1, e, 2)–(1, f, 2)	2,146.131	0.540	0.048
(0, f, 1)-(1, e, 1) $2,145.929$ 0.715 0.038 $(2, e, 1)-(1, f, 1)$ $2,144.061$ 0.130 0.035 $(2, e, 4)-(1, f, 1)$ $2,146.761$ 0.112 0.030 $(2, f, 3)-(2, e, 1)$ $2,146.761$ 0.112 0.020 $(0, e, 4)-(1, f, 2)$ $2,148.729$ 0.194 0.017 $(1, e, 1)-(1, f, 2)$ $2,143.299$ 0.186 0.016 $(2, f, 4)-(1, e, 1)$ $2,147.067$ 0.271 0.014 $(2, f, 4)-(2, e, 1)$ $2,146.770$ 0.474 0.012 $(2, f, 3)-(1, e, 1)$ $2,146.710$ 0.229 0.012 $(1, f, 2)-(1, e, 1)$ $2,146.710$ 0.229 0.010 $(2, f, 3)-(2, e, 2)$ $2,146.129$ 0.712 0.009 $(3, f, 4)-(2, e, 1)$ $2,147.113$ 0.303 0.008 $(2, e, 2)-(1, f, 2)$ $2,143.881$ 0.079 0.007 $(2, f, 1)-(1, e, 1)$ $2,144.209$ 0.122 0.006 $(2, e, 1)-(1, f, 2)$ $2,143.593$ 0.060 0.005 $(2, f, 4)-(2, e, 2)$ $2,146.486$ 0.297 0.004 $(1, f, 1)-(2, e, 1)$ $2,142.108$ 0.127 0.003 $(0, f, 3)-(1, e, 1)$ $2,153.142$ 0.062 0.003 $(3, f, 4)-(2, e, 1)$ $2,144.691$ 0.066 0.002 $(3, f, 2)-(2, e, 1)$ $2,144.691$ 0.066 0.002 $(3, f, 2)-(2, e, 1)$ $2,144.691$ 0.066 0.002 $(2, f, 1)-(2, e, 1)$ $2,142.295$ 0.078 0.001 $(3, e, 3)-(3, f, 1)$ <	(0, e, 1)–(1, f, 1)	2,142.569	0.148	0.040
(2, e, 1)-(1, f, 1) $2,144.061$ 0.130 0.035 $(2, e, 4)-(1, f, 1)$ $2,146.761$ 0.112 0.030 $(2, f, 3)-(2, e, 1)$ $2,146.413$ 0.781 0.020 $(0, e, 4)-(1, f, 2)$ $2,148.729$ 0.194 0.017 $(1, e, 1)-(1, f, 2)$ $2,143.299$ 0.186 0.016 $(2, f, 4)-(1, e, 1)$ $2,147.067$ 0.271 0.014 $(2, f, 4)-(2, e, 1)$ $2,146.770$ 0.474 0.012 $(2, f, 3)-(1, e, 1)$ $2,146.710$ 0.229 0.012 $(1, f, 2)-(1, e, 1)$ $2,142.876$ 0.186 0.010 $(2, f, 3)-(2, e, 2)$ $2,146.129$ 0.712 0.009 $(3, f, 4)-(2, e, 1)$ $2,147.113$ 0.303 0.008 $(2, e, 2)-(1, f, 2)$ $2,143.881$ 0.079 0.007 $(2, f, 1)-(1, e, 1)$ $2,142.108$ 0.127 0.003 $(2, e, 1)-(1, f, 2)$ $2,146.486$ 0.297 0.004 $(1, f, 1)-(2, e, 1)$ $2,142.108$ 0.127 0.003 $(0, f, 3)-(1, e, 1)$ $2,144.822$ 0.072 0.002 $(3, f, 4)-(2, e, 2)$ $2,146.829$ 0.187 0.002 $(3, f, 2)-(2, e, 1)$ $2,144.691$ 0.066 0.002 $(1, f, 2)-(2, e, 1)$ $2,142.579$ 0.060 0.002 $(2, f, 1)-(2, e, 2)$ $2,144.097$ 1.150 0.001 $(3, e, 3)-(3, f, 1)$ $2,146.770$ 0.268 0.001 $(3, f, 1)-(2, e, 2)$ $2,144.098$ 0.065 0.001 $(3, f, 1)-(2, e, 2)$ <	(0, f, 1)–(1, e, 1)	2,145.929	0.715	0.038
(2, e, 4)-(1, f, 1) $2,146.761$ 0.112 0.030 $(2, f, 3)-(2, e, 1)$ $2,146.413$ 0.781 0.020 $(0, e, 4)-(1, f, 2)$ $2,148.729$ 0.194 0.017 $(1, e, 1)-(1, f, 2)$ $2,143.299$ 0.186 0.016 $(2, f, 4)-(1, e, 1)$ $2,147.067$ 0.271 0.014 $(2, f, 4)-(2, e, 1)$ $2,146.770$ 0.474 0.012 $(2, f, 3)-(1, e, 1)$ $2,146.710$ 0.229 0.012 $(1, f, 2)-(1, e, 1)$ $2,142.876$ 0.186 0.010 $(2, f, 3)-(2, e, 2)$ $2,146.129$ 0.712 0.009 $(3, f, 4)-(2, e, 1)$ $2,147.113$ 0.303 0.008 $(2, e, 2)-(1, f, 2)$ $2,143.881$ 0.079 0.007 $(2, f, 1)-(1, e, 1)$ $2,142.109$ 0.122 0.006 $(2, e, 1)-(1, f, 2)$ $2,143.593$ 0.660 0.005 $(2, f, 4)-(2, e, 2)$ $2,146.486$ 0.297 0.004 $(1, f, 1)-(2, e, 1)$ $2,142.108$ 0.127 0.003 $(0, f, 3)-(1, e, 1)$ $2,153.142$ 0.062 0.003 $(3, f, 4)-(2, e, 2)$ $2,146.829$ 0.187 0.002 $(3, f, 2)-(2, e, 1)$ $2,144.691$ 0.066 0.002 $(1, f, 2)-(2, e, 1)$ $2,144.691$ 0.066 0.002 $(2, f, 1)-(2, e, 2)$ $2,144.097$ 1.150 0.001 $(3, e, 3)-(3, f, 1)$ $2,146.770$ 0.268 0.001 $(3, f, 1)-(2, e, 2)$ $2,144.407$ 0.051 0.001	(2, e, 1)–(1, f, 1)	2,144.061	0.130	0.035
(2, f, 3)-(2, e, 1) $2,146.413$ 0.781 0.020 $(0, e, 4)-(1, f, 2)$ $2,148.729$ 0.194 0.017 $(1, e, 1)-(1, f, 2)$ $2,143.299$ 0.186 0.016 $(2, f, 4)-(1, e, 1)$ $2,147.067$ 0.271 0.014 $(2, f, 4)-(2, e, 1)$ $2,146.770$ 0.474 0.012 $(2, f, 3)-(1, e, 1)$ $2,146.710$ 0.229 0.012 $(1, f, 2)-(1, e, 1)$ $2,142.876$ 0.186 0.010 $(2, f, 3)-(2, e, 2)$ $2,146.129$ 0.712 0.009 $(3, f, 4)-(2, e, 1)$ $2,147.113$ 0.303 0.008 $(2, e, 2)-(1, f, 2)$ $2,143.881$ 0.079 0.007 $(2, f, 1)-(1, e, 1)$ $2,144.209$ 0.122 0.006 $(2, e, 1)-(1, f, 2)$ $2,143.593$ 0.060 0.005 $(2, f, 4)-(2, e, 2)$ $2,146.486$ 0.297 0.004 $(1, f, 1)-(2, e, 1)$ $2,142.108$ 0.127 0.003 $(0, f, 3)-(1, e, 1)$ $2,153.142$ 0.062 0.003 $(3, f, 4)-(2, e, 2)$ $2,144.829$ 0.187 0.002 $(3, f, 2)-(2, e, 1)$ $2,142.579$ 0.060 0.002 $(1, f, 2)-(2, e, 1)$ $2,142.97$ 1.150 0.001 $(1, f, 2)-(2, e, 2)$ $2,144.091$ 0.065 0.001 $(3, e, 3)-(3, f, 1)$ $2,142.295$ 0.078 0.001 $(3, f, 1)-(2, e, 2)$ $2,144.098$ 0.065 0.001 $(2, f, 1)-(2, e, 2)$ $2,144.098$ 0.065 0.001	(2, e, 4)–(1, f, 1)	2,146.761	0.112	0.030
(0, e, 4)-(1, f, 2) $2,148.729$ 0.194 0.017 $(1, e, 1)-(1, f, 2)$ $2,143.299$ 0.186 0.016 $(2, f, 4)-(1, e, 1)$ $2,147.067$ 0.271 0.014 $(2, f, 4)-(2, e, 1)$ $2,146.770$ 0.474 0.012 $(2, f, 3)-(1, e, 1)$ $2,146.710$ 0.229 0.012 $(1, f, 2)-(1, e, 1)$ $2,142.876$ 0.186 0.010 $(2, f, 3)-(2, e, 2)$ $2,146.129$ 0.712 0.009 $(3, f, 4)-(2, e, 1)$ $2,147.113$ 0.303 0.008 $(2, e, 2)-(1, f, 2)$ $2,143.881$ 0.079 0.007 $(2, f, 1)-(1, e, 1)$ $2,144.209$ 0.122 0.006 $(2, e, 1)-(1, f, 2)$ $2,143.593$ 0.060 0.005 $(2, f, 4)-(2, e, 2)$ $2,146.486$ 0.297 0.004 $(1, f, 1)-(2, e, 1)$ $2,142.108$ 0.127 0.003 $(0, f, 3)-(1, e, 1)$ $2,153.142$ 0.062 0.003 $(3, f, 4)-(2, e, 2)$ $2,144.829$ 0.187 0.002 $(3, f, 2)-(2, e, 1)$ $2,144.382$ 0.072 0.002 $(3, f, 2)-(2, e, 1)$ $2,142.579$ 0.060 0.002 $(1, f, 2)-(2, e, 1)$ $2,142.295$ 0.078 0.001 $(3, e, 3)-(3, f, 1)$ $2,142.295$ 0.078 0.001 $(3, e, 3)-(2, f, 1)$ $2,144.098$ 0.065 0.001 $(2, f, 1)-(2, e, 2)$ $2,144.098$ 0.065 0.001 $(3, f, 2)-(2, e, 2)$ $2,144.098$ 0.063 0.001	(2, f, 3)–(2, e, 1)	2,146.413	0.781	0.020
(1, e, 1)-(1, f, 2) $2,143.299$ 0.186 0.016 $(2, f, 4)-(1, e, 1)$ $2,147.067$ 0.271 0.014 $(2, f, 4)-(2, e, 1)$ $2,146.770$ 0.474 0.012 $(2, f, 3)-(1, e, 1)$ $2,146.710$ 0.229 0.012 $(1, f, 2)-(1, e, 1)$ $2,142.876$ 0.186 0.010 $(2, f, 3)-(2, e, 2)$ $2,146.129$ 0.712 0.009 $(3, f, 4)-(2, e, 1)$ $2,147.113$ 0.303 0.008 $(2, e, 2)-(1, f, 2)$ $2,143.881$ 0.079 0.007 $(2, f, 1)-(1, e, 1)$ $2,144.209$ 0.122 0.006 $(2, e, 1)-(1, f, 2)$ $2,143.593$ 0.060 0.005 $(2, f, 4)-(2, e, 2)$ $2,146.486$ 0.297 0.004 $(1, f, 1)-(2, e, 1)$ $2,142.108$ 0.127 0.003 $(0, f, 3)-(1, e, 1)$ $2,153.142$ 0.062 0.003 $(3, f, 4)-(2, e, 2)$ $2,146.829$ 0.187 0.002 $(3, f, 4)-(2, e, 1)$ $2,144.382$ 0.072 0.002 $(3, f, 2)-(2, e, 1)$ $2,144.691$ 0.066 0.002 $(1, f, 2)-(2, e, 1)$ $2,142.295$ 0.078 0.001 $(1, f, 2)-(2, e, 2)$ $2,144.098$ 0.065 0.001 $(3, e, 3)-(3, f, 1)$ $2,142.295$ 0.078 0.001 $(3, f, 1)-(2, e, 2)$ $2,144.098$ 0.063 0.001 $(3, f, 2)-(2, e, 2)$ $2,144.098$ 0.065 0.001 $(3, f, 2)-(2, e, 2)$ $2,144.07$ 0.051 0.001	(0, e, 4)–(1, f, 2)	2,148.729	0.194	0.017
(2, f, 4)-(1, e, 1) $2,147.067$ 0.271 0.014 $(2, f, 4)-(2, e, 1)$ $2,146.770$ 0.474 0.012 $(2, f, 3)-(1, e, 1)$ $2,146.710$ 0.229 0.012 $(1, f, 2)-(1, e, 1)$ $2,142.876$ 0.186 0.010 $(2, f, 3)-(2, e, 2)$ $2,146.129$ 0.712 0.009 $(3, f, 4)-(2, e, 1)$ $2,147.113$ 0.303 0.008 $(2, e, 2)-(1, f, 2)$ $2,143.881$ 0.079 0.007 $(2, f, 1)-(1, e, 1)$ $2,144.209$ 0.122 0.006 $(2, e, 1)-(1, f, 2)$ $2,143.593$ 0.060 0.005 $(2, f, 4)-(2, e, 2)$ $2,146.486$ 0.297 0.004 $(1, f, 1)-(2, e, 1)$ $2,142.108$ 0.127 0.003 $(0, f, 3)-(1, e, 1)$ $2,153.142$ 0.062 0.003 $(3, f, 4)-(2, e, 2)$ $2,144.829$ 0.187 0.002 $(3, f, 4)-(2, e, 1)$ $2,144.391$ 0.066 0.002 $(3, f, 2)-(2, e, 1)$ $2,142.579$ 0.060 0.002 $(1, f, 2)-(2, e, 1)$ $2,142.295$ 0.078 0.001 $(3, e, 3)-(3, f, 1)$ $2,142.295$ 0.078 0.001 $(3, e, 3)-(2, f, 1)$ $2,144.098$ 0.065 0.001 $(3, f, 1)-(2, e, 2)$ $2,144.098$ 0.063 0.001 $(3, f, 2)-(2, e, 2)$ $2,144.407$ 0.051 0.001	(1, e, 1)–(1, f, 2)	2,143.299	0.186	0.016
(2, f, 4)-(2, e, 1) $2,146.770$ 0.474 0.012 $(2, f, 3)-(1, e, 1)$ $2,146.710$ 0.229 0.012 $(1, f, 2)-(1, e, 1)$ $2,142.876$ 0.186 0.010 $(2, f, 3)-(2, e, 2)$ $2,146.129$ 0.712 0.009 $(3, f, 4)-(2, e, 1)$ $2,147.113$ 0.303 0.008 $(2, e, 2)-(1, f, 2)$ $2,143.881$ 0.079 0.007 $(2, f, 1)-(1, e, 1)$ $2,144.209$ 0.122 0.006 $(2, e, 1)-(1, f, 2)$ $2,143.593$ 0.060 0.005 $(2, f, 4)-(2, e, 2)$ $2,146.486$ 0.297 0.004 $(1, f, 1)-(2, e, 1)$ $2,142.108$ 0.127 0.003 $(0, f, 3)-(1, e, 1)$ $2,153.142$ 0.062 0.003 $(3, f, 4)-(2, e, 2)$ $2,144.829$ 0.187 0.002 $(3, f, 1)-(2, e, 1)$ $2,144.579$ 0.066 0.002 $(1, f, 2)-(2, e, 1)$ $2,144.691$ 0.066 0.002 $(2, f, 1)-(2, e, 1)$ $2,142.295$ 0.078 0.001 $(3, e, 3)-(3, f, 1)$ $2,142.295$ 0.078 0.001 $(3, e, 3)-(2, f, 1)$ $2,144.098$ 0.065 0.001 $(2, f, 1)-(2, e, 2)$ $2,144.098$ 0.065 0.001 $(3, f, 2)-(2, e, 2)$ $2,144.098$ 0.063 0.001	(2, f, 4)–(1, e, 1)	2,147.067	0.271	0.014
(2, f, 3)-(1, e, 1) $2,146.710$ 0.229 0.012 $(1, f, 2)-(1, e, 1)$ $2,142.876$ 0.186 0.010 $(2, f, 3)-(2, e, 2)$ $2,146.129$ 0.712 0.009 $(3, f, 4)-(2, e, 1)$ $2,147.113$ 0.303 0.008 $(2, e, 2)-(1, f, 2)$ $2,143.881$ 0.079 0.007 $(2, f, 1)-(1, e, 1)$ $2,144.209$ 0.122 0.006 $(2, e, 1)-(1, f, 2)$ $2,143.593$ 0.060 0.005 $(2, f, 4)-(2, e, 2)$ $2,146.486$ 0.297 0.004 $(1, f, 1)-(2, e, 1)$ $2,142.108$ 0.127 0.003 $(0, f, 3)-(1, e, 1)$ $2,153.142$ 0.062 0.003 $(3, f, 4)-(2, e, 2)$ $2,146.829$ 0.187 0.002 $(3, f, 1)-(2, e, 1)$ $2,144.691$ 0.066 0.002 $(1, f, 2)-(2, e, 1)$ $2,142.579$ 0.060 0.002 $(2, f, 1)-(2, e, 1)$ $2,142.957$ 0.078 0.001 $(3, e, 3)-(3, f, 1)$ $2,142.295$ 0.078 0.001 $(3, e, 3)-(2, f, 1)$ $2,144.098$ 0.065 0.001 $(3, f, 1)-(2, e, 2)$ $2,144.098$ 0.063 0.001 $(3, f, 2)-(2, e, 2)$ $2,144.407$ 0.051 0.001	(2, f, 4)–(2, e, 1)	2,146.770	0.474	0.012
(1, f, 2)-(1, e, 1) $2,142.876$ 0.186 0.010 $(2, f, 3)-(2, e, 2)$ $2,146.129$ 0.712 0.009 $(3, f, 4)-(2, e, 1)$ $2,147.113$ 0.303 0.008 $(2, e, 2)-(1, f, 2)$ $2,143.881$ 0.079 0.007 $(2, f, 1)-(1, e, 1)$ $2,144.209$ 0.122 0.006 $(2, e, 1)-(1, f, 2)$ $2,143.593$ 0.060 0.005 $(2, e, 1)-(1, f, 2)$ $2,144.209$ 0.122 0.004 $(1, f, 1)-(2, e, 2)$ $2,146.486$ 0.297 0.004 $(1, f, 1)-(2, e, 1)$ $2,142.108$ 0.127 0.003 $(0, f, 3)-(1, e, 1)$ $2,153.142$ 0.062 0.003 $(3, f, 4)-(2, e, 2)$ $2,144.829$ 0.187 0.002 $(3, f, 4)-(2, e, 1)$ $2,144.382$ 0.072 0.002 $(3, f, 2)-(2, e, 1)$ $2,144.579$ 0.066 0.002 $(1, f, 2)-(2, e, 1)$ $2,142.295$ 0.078 0.001 $(1, f, 2)-(2, e, 2)$ $2,144.098$ 0.065 0.001 $(3, e, 3)-(2, f, 1)$ $2,144.098$ 0.065 0.001 $(3, f, 1)-(2, e, 2)$ $2,144.407$ 0.051 0.001	(2, f, 3)–(1, e, 1)	2,146.710	0.229	0.012
(2, f, 3)-(2, e, 2) $2, 146.129$ 0.712 0.009 $(3, f, 4)-(2, e, 1)$ $2, 147.113$ 0.303 0.008 $(2, e, 2)-(1, f, 2)$ $2, 143.881$ 0.079 0.007 $(2, f, 1)-(1, e, 1)$ $2, 144.209$ 0.122 0.006 $(2, e, 1)-(1, f, 2)$ $2, 143.593$ 0.060 0.005 $(2, e, 1)-(1, f, 2)$ $2, 143.593$ 0.060 0.005 $(2, f, 4)-(2, e, 2)$ $2, 146.486$ 0.297 0.004 $(1, f, 1)-(2, e, 1)$ $2, 142.108$ 0.127 0.003 $(0, f, 3)-(1, e, 1)$ $2, 153.142$ 0.062 0.003 $(3, f, 4)-(2, e, 2)$ $2, 146.829$ 0.187 0.002 $(3, f, 4)-(2, e, 1)$ $2, 144.382$ 0.072 0.002 $(3, f, 2)-(2, e, 1)$ $2, 144.691$ 0.066 0.002 $(1, f, 2)-(2, e, 1)$ $2, 144.691$ 0.066 0.002 $(2, f, 1)-(2, e, 1)$ $2, 142.295$ 0.078 0.001 $(3, e, 3)-(3, f, 1)$ $2, 142.295$ 0.078 0.001 $(3, e, 3)-(2, f, 1)$ $2, 144.098$ 0.065 0.001 $(3, f, 1)-(2, e, 2)$ $2, 144.098$ 0.063 0.001 $(2, f, 1)-(2, e, 2)$ $2, 144.407$ 0.051 0.001	(1, f, 2)–(1, e, 1)	2,142.876	0.186	0.010
(3, f, 4)-(2, e, 1) $2, 147.113$ 0.303 0.008 $(2, e, 2)-(1, f, 2)$ $2, 143.881$ 0.079 0.007 $(2, f, 1)-(1, e, 1)$ $2, 144.209$ 0.122 0.006 $(2, e, 1)-(1, f, 2)$ $2, 143.593$ 0.060 0.005 $(2, e, 1)-(1, f, 2)$ $2, 143.593$ 0.060 0.005 $(2, e, 1)-(1, f, 2)$ $2, 143.593$ 0.060 0.005 $(2, f, 4)-(2, e, 2)$ $2, 146.486$ 0.297 0.004 $(1, f, 1)-(2, e, 1)$ $2, 142.108$ 0.127 0.003 $(0, f, 3)-(1, e, 1)$ $2, 153.142$ 0.062 0.003 $(3, f, 4)-(2, e, 2)$ $2, 146.829$ 0.187 0.002 $(3, f, 1)-(2, e, 1)$ $2, 144.382$ 0.072 0.002 $(3, f, 2)-(2, e, 1)$ $2, 144.691$ 0.066 0.002 $(1, f, 2)-(2, e, 1)$ $2, 143.912$ 0.057 0.001 $(3, e, 3)-(3, f, 1)$ $2, 142.295$ 0.078 0.001 $(3, e, 3)-(2, f, 1)$ $2, 144.098$ 0.065 0.001 $(3, f, 1)-(2, e, 2)$ $2, 144.098$ 0.063 0.001 $(3, f, 2)-(2, e, 2)$ $2, 144.407$ 0.051 0.001	(2, f, 3)–(2, e, 2)	2,146.129	0.712	0.009
(2, e, 2)-(1, f, 2) $2,143.881$ 0.079 0.007 $(2, f, 1)-(1, e, 1)$ $2,144.209$ 0.122 0.006 $(2, e, 1)-(1, f, 2)$ $2,143.593$ 0.060 0.005 $(2, e, 1)-(1, f, 2)$ $2,143.593$ 0.060 0.005 $(2, f, 4)-(2, e, 2)$ $2,146.486$ 0.297 0.004 $(1, f, 1)-(2, e, 1)$ $2,142.108$ 0.127 0.003 $(0, f, 3)-(1, e, 1)$ $2,153.142$ 0.062 0.003 $(3, f, 4)-(2, e, 2)$ $2,146.829$ 0.187 0.002 $(3, f, 1)-(2, e, 1)$ $2,144.382$ 0.072 0.002 $(3, f, 2)-(2, e, 1)$ $2,144.691$ 0.066 0.002 $(1, f, 2)-(2, e, 1)$ $2,142.579$ 0.060 0.002 $(2, f, 1)-(2, e, 1)$ $2,143.912$ 0.057 0.001 $(3, e, 3)-(3, f, 1)$ $2,142.295$ 0.078 0.001 $(3, e, 3)-(2, f, 1)$ $2,144.098$ 0.065 0.001 $(3, f, 1)-(2, e, 2)$ $2,144.407$ 0.051 0.001	(3, f, 4)–(2, e, 1)	2,147.113	0.303	0.008
(2, f, 1)-(1, e, 1) $2,144.209$ 0.122 0.006 $(2, e, 1)-(1, f, 2)$ $2,143.593$ 0.060 0.005 $(2, f, 4)-(2, e, 2)$ $2,146.486$ 0.297 0.004 $(1, f, 1)-(2, e, 1)$ $2,142.108$ 0.127 0.003 $(0, f, 3)-(1, e, 1)$ $2,153.142$ 0.062 0.003 $(3, f, 4)-(2, e, 2)$ $2,146.829$ 0.187 0.002 $(3, f, 1)-(2, e, 1)$ $2,144.382$ 0.072 0.002 $(3, f, 2)-(2, e, 1)$ $2,144.691$ 0.066 0.002 $(1, f, 2)-(2, e, 1)$ $2,142.579$ 0.060 0.002 $(2, f, 1)-(2, e, 1)$ $2,142.977$ 1.150 0.001 $(3, e, 3)-(3, f, 1)$ $2,146.297$ 1.150 0.001 $(3, e, 3)-(2, f, 1)$ $2,146.770$ 0.268 0.001 $(3, f, 1)-(2, e, 2)$ $2,144.098$ 0.065 0.001 $(3, f, 1)-(2, e, 2)$ $2,144.407$ 0.051 0.001	(2, e, 2)–(1, f, 2)	2,143.881	0.079	0.007
(2, e, 1)-(1, f, 2) $2,143.593$ 0.060 0.005 $(2, f, 4)-(2, e, 2)$ $2,146.486$ 0.297 0.004 $(1, f, 1)-(2, e, 1)$ $2,142.108$ 0.127 0.003 $(0, f, 3)-(1, e, 1)$ $2,153.142$ 0.062 0.003 $(3, f, 4)-(2, e, 2)$ $2,146.829$ 0.187 0.002 $(3, f, 1)-(2, e, 1)$ $2,144.382$ 0.072 0.002 $(3, f, 2)-(2, e, 1)$ $2,144.691$ 0.066 0.002 $(1, f, 2)-(2, e, 1)$ $2,142.579$ 0.060 0.002 $(2, f, 1)-(2, e, 1)$ $2,143.912$ 0.057 0.001 $(3, e, 3)-(3, f, 1)$ $2,142.295$ 0.078 0.001 $(3, e, 3)-(2, f, 1)$ $2,144.098$ 0.065 0.001 $(3, f, 1)-(2, e, 2)$ $2,144.407$ 0.051 0.001	(2, f, 1)–(1, e, 1)	2,144.209	0.122	0.006
(2, f, 4)-(2, e, 2) $2,146.486$ 0.297 0.004 $(1, f, 1)-(2, e, 1)$ $2,142.108$ 0.127 0.003 $(0, f, 3)-(1, e, 1)$ $2,153.142$ 0.062 0.003 $(3, f, 4)-(2, e, 2)$ $2,146.829$ 0.187 0.002 $(3, f, 1)-(2, e, 1)$ $2,144.382$ 0.072 0.002 $(3, f, 2)-(2, e, 1)$ $2,144.691$ 0.066 0.002 $(1, f, 2)-(2, e, 1)$ $2,142.579$ 0.060 0.002 $(2, f, 1)-(2, e, 1)$ $2,143.912$ 0.057 0.001 $(3, e, 3)-(3, f, 1)$ $2,142.295$ 0.078 0.001 $(3, e, 3)-(2, f, 1)$ $2,144.098$ 0.065 0.001 $(3, f, 1)-(2, e, 2)$ $2,143.628$ 0.063 0.001 $(3, f, 2)-(2, e, 2)$ $2,144.407$ 0.051 0.001	(2, e, 1)–(1, f, 2)	2,143.593	0.060	0.005
(1, f, 1)-(2, e, 1) $2,142.108$ 0.127 0.003 $(0, f, 3)-(1, e, 1)$ $2,153.142$ 0.062 0.003 $(3, f, 4)-(2, e, 2)$ $2,146.829$ 0.187 0.002 $(3, f, 1)-(2, e, 1)$ $2,144.382$ 0.072 0.002 $(3, f, 2)-(2, e, 1)$ $2,144.691$ 0.066 0.002 $(1, f, 2)-(2, e, 1)$ $2,142.579$ 0.060 0.002 $(2, f, 1)-(2, e, 1)$ $2,143.912$ 0.057 0.001 $(3, e, 3)-(3, f, 1)$ $2,142.295$ 0.078 0.001 $(3, e, 3)-(2, f, 1)$ $2,144.098$ 0.065 0.001 $(3, f, 1)-(2, e, 2)$ $2,143.628$ 0.063 0.001 $(3, f, 2)-(2, e, 2)$ $2,144.407$ 0.051 0.001	(2, f, 4)–(2, e, 2)	2,146.486	0.297	0.004
(0, f, 3)-(1, e, 1) $2,153.142$ 0.062 0.003 $(3, f, 4)-(2, e, 2)$ $2,146.829$ 0.187 0.002 $(3, f, 1)-(2, e, 1)$ $2,144.382$ 0.072 0.002 $(3, f, 2)-(2, e, 1)$ $2,144.691$ 0.066 0.002 $(1, f, 2)-(2, e, 1)$ $2,142.579$ 0.060 0.002 $(2, f, 1)-(2, e, 1)$ $2,142.912$ 0.057 0.001 $(3, e, 3)-(3, f, 1)$ $2,146.297$ 1.150 0.001 $(1, f, 2)-(2, e, 2)$ $2,142.295$ 0.078 0.001 $(3, e, 3)-(2, f, 1)$ $2,146.770$ 0.268 0.001 $(3, f, 1)-(2, e, 2)$ $2,144.098$ 0.065 0.001 $(2, f, 1)-(2, e, 2)$ $2,143.628$ 0.063 0.001 $(3, f, 2)-(2, e, 2)$ $2,144.407$ 0.051 0.001	(1, f, 1)–(2, e, 1)	2,142.108	0.127	0.003
(3, f, 4)-(2, e, 2) $2,146.829$ 0.187 0.002 $(3, f, 1)-(2, e, 1)$ $2,144.382$ 0.072 0.002 $(3, f, 2)-(2, e, 1)$ $2,144.691$ 0.066 0.002 $(1, f, 2)-(2, e, 1)$ $2,142.579$ 0.060 0.002 $(2, f, 1)-(2, e, 1)$ $2,142.979$ 0.057 0.001 $(3, e, 3)-(3, f, 1)$ $2,146.297$ 1.150 0.001 $(1, f, 2)-(2, e, 2)$ $2,142.295$ 0.078 0.001 $(3, e, 3)-(2, f, 1)$ $2,146.770$ 0.268 0.001 $(3, f, 1)-(2, e, 2)$ $2,144.098$ 0.065 0.001 $(2, f, 1)-(2, e, 2)$ $2,143.628$ 0.063 0.001 $(3, f, 2)-(2, e, 2)$ $2,144.407$ 0.051 0.001	(0, f, 3)–(1, e, 1)	2,153.142	0.062	0.003
(3, f, 1)-(2, e, 1) $2,144.382$ 0.072 0.002 $(3, f, 2)-(2, e, 1)$ $2,144.691$ 0.066 0.002 $(1, f, 2)-(2, e, 1)$ $2,142.579$ 0.060 0.002 $(2, f, 1)-(2, e, 1)$ $2,143.912$ 0.057 0.001 $(3, e, 3)-(3, f, 1)$ $2,146.297$ 1.150 0.001 $(1, f, 2)-(2, e, 2)$ $2,142.295$ 0.078 0.001 $(3, e, 3)-(2, f, 1)$ $2,146.770$ 0.268 0.001 $(3, f, 1)-(2, e, 2)$ $2,144.098$ 0.065 0.001 $(2, f, 1)-(2, e, 2)$ $2,143.628$ 0.063 0.001 $(3, f, 2)-(2, e, 2)$ $2,144.407$ 0.051 0.001	(3, f, 4)–(2, e, 2)	2,146.829	0.187	0.002
(3, f, 2)-(2, e, 1) $2,144.691$ 0.066 0.002 $(1, f, 2)-(2, e, 1)$ $2,142.579$ 0.060 0.002 $(2, f, 1)-(2, e, 1)$ $2,143.912$ 0.057 0.001 $(3, e, 3)-(3, f, 1)$ $2,146.297$ 1.150 0.001 $(1, f, 2)-(2, e, 2)$ $2,142.295$ 0.078 0.001 $(3, e, 3)-(2, f, 1)$ $2,146.770$ 0.268 0.001 $(3, f, 1)-(2, e, 2)$ $2,144.098$ 0.065 0.001 $(2, f, 1)-(2, e, 2)$ $2,143.628$ 0.063 0.001 $(3, f, 2)-(2, e, 2)$ $2,144.407$ 0.051 0.001	(3, f, 1)–(2, e, 1)	2,144.382	0.072	0.002
(1, f, 2)- $(2, e, 1)$ $2,142.579$ 0.060 0.002 $(2, f, 1)$ - $(2, e, 1)$ $2,143.912$ 0.057 0.001 $(3, e, 3)$ - $(3, f, 1)$ $2,146.297$ 1.150 0.001 $(1, f, 2)$ - $(2, e, 2)$ $2,142.295$ 0.078 0.001 $(3, e, 3)$ - $(2, f, 1)$ $2,146.770$ 0.268 0.001 $(3, f, 1)$ - $(2, e, 2)$ $2,144.098$ 0.065 0.001 $(2, f, 1)$ - $(2, e, 2)$ $2,143.628$ 0.063 0.001 $(3, f, 2)$ - $(2, e, 2)$ $2,144.407$ 0.051 0.001	(3, f, 2)–(2, e, 1)	2,144.691	0.066	0.002
(2, f, 1)-(2, e, 1) $2,143.912$ 0.057 0.001 $(3, e, 3)-(3, f, 1)$ $2,146.297$ 1.150 0.001 $(1, f, 2)-(2, e, 2)$ $2,142.295$ 0.078 0.001 $(3, e, 3)-(2, f, 1)$ $2,146.770$ 0.268 0.001 $(3, f, 1)-(2, e, 2)$ $2,144.098$ 0.065 0.001 $(2, f, 1)-(2, e, 2)$ $2,143.628$ 0.063 0.001 $(3, f, 2)-(2, e, 2)$ $2,144.407$ 0.051 0.001	(1, f, 2)–(2, e, 1)	2,142.579	0.060	0.002
(3, e, 3)-(3, f, 1) $2,146.297$ 1.150 0.001 $(1, f, 2)-(2, e, 2)$ $2,142.295$ 0.078 0.001 $(3, e, 3)-(2, f, 1)$ $2,146.770$ 0.268 0.001 $(3, f, 1)-(2, e, 2)$ $2,144.098$ 0.065 0.001 $(2, f, 1)-(2, e, 2)$ $2,143.628$ 0.063 0.001 $(3, f, 2)-(2, e, 2)$ $2,144.407$ 0.051 0.001	(2, f, 1)–(2, e, 1)	2,143.912	0.057	0.001
(1, f, 2)-(2, e, 2) $2,142.295$ 0.078 0.001 $(3, e, 3)-(2, f, 1)$ $2,146.770$ 0.268 0.001 $(3, f, 1)-(2, e, 2)$ $2,144.098$ 0.065 0.001 $(2, f, 1)-(2, e, 2)$ $2,143.628$ 0.063 0.001 $(3, f, 2)-(2, e, 2)$ $2,144.407$ 0.051 0.001	(3, e, 3)–(3, f, 1)	2,146.297	1.150	0.001
(3, e, 3)-(2, f, 1) $2,146.770$ 0.268 0.001 $(3, f, 1)-(2, e, 2)$ $2,144.098$ 0.065 0.001 $(2, f, 1)-(2, e, 2)$ $2,143.628$ 0.063 0.001 $(3, f, 2)-(2, e, 2)$ $2,144.407$ 0.051 0.001	(1, f, 2)–(2, e, 2)	2,142.295	0.078	0.001
(3, f, 1)-(2, e, 2) $2,144.098$ 0.065 0.001 $(2, f, 1)-(2, e, 2)$ $2,143.628$ 0.063 0.001 $(3, f, 2)-(2, e, 2)$ $2,144.407$ 0.051 0.001	(3, e, 3)–(2, f, 1)	2,146.770	0.268	0.001
(2, f, 1)- $(2, e, 2)$ $2,143.628$ 0.063 0.001 $(3, f, 2)$ - $(2, e, 2)$ $2.144.407$ 0.051 0.001	(3, f, 1)–(2, e, 2)	2,144.098	0.065	0.001
(3, f, 2)–(2, e, 2) 2, 144,407 0,051 0,001	(2, f, 1)–(2, e, 2)	2,143.628	0.063	0.001
	(3, f, 2)–(2, e, 2)	2,144.407	0.051	0.001

The lower and upper states involved in the transitions are numbered with the set of parameters $(J, P, n_{J,P})$. The line strength values are multiplied by 10

^a The observed value is 2,146.93 cm⁻¹

inter-particle interactions and can be explained by a simple one-dimensional particle-in-a-box model along the ϕ coordinate [27, 39].



Fig. 7 Three-dimensional representations of the pH₂ density for the ground state CO–(pH₂)₂ trimer (*upper panel*) in the body-fixed frame, compared with the helium atom density for a ground-state CO–(He)₂ trimer (*middle panel*) and the pH₂–He density of a ground-state CO–pH₂–He trimer (*lower panel*). The position of the first pH₂ or He defines the location of the yz plane, while the density of the second pH2 or He particle is shown in the xy plane. These results were obtained with the values of the Gaussian standard deviations defining the densities set at $\sigma_x = \sigma_z = 0.25$ Å and $\sigma_y = 0.1$ Å for particle-1 and at $\sigma_x = \sigma_x = \sigma_z = 0.25$ Å for particle-2

4 Conclusions

Predicted rovibrational energy levels, microwave, and infrared spectra for CO– $(pH_2)_2$, CO– $(oD_2)_2$, and CO– pH_2 – He complexes have been obtained from the vibrationally averaged five-dimensional PESs as a sum of accurate CO– pH_2 , CO– oD_2 , or CO–He and pH_2 – pH_2 , oD_2 – oD_2 , or pH_2 –He pair potentials. For CO– $(pH_2)_2$ complex, it is remarkable that the predictions are in good agreement with the experimental measurements of the microwave and infrared transitions, showing that the spectrum can be reliably predicted by this additive approach. Small remaining differences between experiment and theory may in part be due to our neglect of three-body contributions to the interaction energy. For CO– $(oD_2)_2$ trimer, the infrared spectrum



Fig. 8 Three-dimensional representations of the "solvent" (pH₂, He and pH₂–He) density for CO–(pH₂)₂ (*upper panels*), CO–(He)₂ (*mid-dle panels*), and CO–pH₂–He (*lower panels*) trimers in their four lowest torsion vibrational levels

has been predicted for the first time. For CO-pH₂-He, one observed IR transition tentatively assigned as b-type R(0)was confirmed based on the calculated line position and intensity. Although many of the transitions for those three complexes have not been observed, our predictions will be a good guidance for future experimental research. The calculated band origin shift from vibrationally averaged five-dimensional PESs associated with the fundamental transition of CO is -0.369 cm^{-1} for CO-(pH₂)₂, which is also in good agreement with the experimental value of -0.355 cm⁻¹. The band origin shifts for CO-(oD₂)₂ and CO-pH₂-He are also predicted for the first time. In any case, the result indicates that a reduced-dimension treatment of pH2 or oD2 rotation based on a "adiabatic-hindered-rotor" average over its relative orientations can yield accurate spectra predictions. Three-dimensional representation of the pH₂, oD₂, or He density in the body-fixed frame is very useful to develop some intuitive feeling regarding the nature of these species, and it clearly shows the density distribution of the second pH₂, oD₂, or He relative to the first one. The pH₂ and oD₂ distributions in the vibrational ground state of $CO-(pH_2)_2$ and $CO-(oD_2)_2$ are highly localized, with two maxima, while the He distribution in CO-(He)₂ or CO-pH₂-He is delocalized, with one maximum lying in an incomplete ring wrapped around the CO molecule axis at the O end.

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REGULAR ARTICLE

Numerical solution of solvent reorganization energy and its application in electron transfer reaction

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Abstract According to our recent studies on the nonequilibrium solvation, the solvent reorganization energy λ_s is found to be the cost of maintaining the residual polarization, which equilibrates with the constraining extra electric field. In this work, a matrix form of λ_s has been formulated based our new analytical expression of the solvent reorganization energy. By means of the integral equation formulation-polarizable continuum model (IEF-PCM), a new numerical algorithm for λ_s has been implemented as a subroutine coupled with the Q-Chem package. Then, we have performed a comparison of numerical results with analytical solution obtained by two-sphere model for λ_s in selfexchange electron transfer (ET) reaction of He-He⁺ system. The numerical results and analytical solution coincide as the distance between the donor and the acceptor. In order to compare with our pervious numerical algorithm with dielectric polarizable continuum model, self-exchange ET reactions between tetracyanoethylene, tetrathiafulvalene, and their corresponding ionic radicals in acetonitrile have been studied. Overall, the solvent reorganization energy calculated by IEF-PCM is more reasonable since IEF-PCM gives a better self-energy of surface element.

Dedicated to Professor Guosen Yan and published as part of the special collection of articles celebrating his 85th birthday.

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1 Introduction

Through thermal fluctuation [1] or light irradiation [2], an electron transfer (ET) from a donor chemical species to the acceptor is a fundamental step in all oxidation–reduction reactions that impacts a variety of chemical and biochemical processes. The studies about ET in photosynthesis, respiration, and various enzyme-catalyzed redox reactions have made many significant progresses.

ET reactions are also remarkable for their simple theoretical concept. The most widespread theory for ET was developed by Marcus and Hush [3, 4], which provides a helpful physical-chemical reference framework. With the harmonic approximation, two parabolas along with a reaction coordinate can be used to represent the energy curves of reaction and product states as showed in Fig. 1. In the diabatic representation, two curves intersect, and crossing point (q_c) corresponds to the transition state. With the assumption of both parabolas with same curvature, the activation energy ΔG^+ is given by

$$\Delta G^{+} = \frac{(\lambda + \Delta G^{0})^{2}}{4\lambda} \tag{1}$$

where ΔG^0 is the reaction heat, in other words, the driving force, and λ is called the reorganization energy including contributions from both the solute and the solvents. An ET is the simplest chemical reaction because it does not involve any bond-breaking or bond-forming. For a self-exchange ET, the activation energy is only related to reorganization energy, i.e., $\Delta G^+ = \lambda/4$. λ is usually separated into two



Fig. 1 Potential energy curves of ET reaction

parts: the inner reorganization energy λ_m corresponding to the energy needed to distort the bonds and angles in the reactant, while the solvent reorganization energy λ_s arising from the polarization of surrounding solvent. For λ_m , many theoretical computational methods, such as the most common Nelsen's four-point methods [5] and its correction forms [6], have been established. Another widely used method is based on the diabatic state through constrained density functional theory (CDFT) [7]. For evaluation of λ_s , the key problem is to correctly deal with the scheme of nonequilibrium solute–solvent interaction. Therefore, how to set up a rational expression of nonequilibrium solvation energy is a big challenge in theory [8, 9].

By introducing the constrained equilibrium approach in continuum model, our group proposed a new way to the nonequilibrium solvation model and obtained the expression for electrostatic solvation energy in the nonequilibrium polarization case [10–12]. This expression has been successfully applied to the evaluation of vertical ionization energy of hydrated electron making use of approximation of spherical cavity and point charge [10], solvatochromic shift of absorption spectra with solvation model of sphere cavity and point dipole moment [11], and λ_s of ET reactions by two-sphere model [12] and numerical solution [13]. In present work, we re-derive a matrix expression of λ_s for the gain of the popularization potential and the self-energy of surface element. IEF-PCM has been applied for the implementation of numerical solution of λ_s .

2 Theory

As shown in Scheme 1, for an ET, or a photo absorption/ emission process in the solvent, the system firstly establishes an initial equilibrium state 1 [ρ_1, φ_1^{eq}], where ρ_1 and φ_1^{eq} stand for solute charge distribution and solvent polarization potential (total electric potential excluding the vacuum potential due to solute charge), respectively. After a fast change of solute charge from ρ_1 to ρ_2 , the dynamic polarization of the solvents adjusts itself quickly to equilibrate with ρ_2 , while the inertial polarization keeps fixed. The system reaches the nonequilibrium polarization state 2 $[\rho_2, \varphi_2^{\text{non}}]$, where φ_2^{non} is the nonequilibrium polarization potential. After enough long time, the inertial polarization finally reaches to the one which equilibrates to the solute charge ρ_2 , and the system relaxes to the equilibrium state 2 $[\rho_2, \varphi_2^{eq}]$, with φ_2^{eq} being the equilibrium polarization potential corresponding to ρ_2 .

The rational description of λ_s requires a proper account of thermodynamical nonequilibrium effect, and how to obtain the energy for nonequilibrium state theoretically is in general a difficult task [9-14]. Usually, classical thermodynamics cannot be directly adopted to handle the nonequilibrium polarization of dielectric media, but the constrained equilibrium approach proposed by Leontovich [15] provides in some cases a way to the gain of the free energy of a thermodynamically nonequilibrium state. This approach is based on the following three assumptions [10, 11, 16–18]. Firstly, by imposing suitable external conservative forces, a nonequilibrium state of an isothermal system without flow can be mapped to a constrained equilibrium state, which has the same internal variables as those in the nonequilibrium state. Secondly, the differences in state function between the constrained equilibrium and any other equilibrium state can be calculated simply by means of classical thermodynamics. Thirdly, the external forces can be removed suddenly without friction from the constrained equilibrium system so as to recover the true nonequilibrium situation. Recently, our group applied this thermodynamical principle to construct the constrained equilibrium states $[\rho_2 + \rho_{ex}, \varphi_2^{non}]$ where ρ_{ex} represents the external electric charge. In this way, λ_s can be obtained as below [10–12]:

$$\lambda_{\rm s} = -\frac{1}{2} \int\limits_{V} \rho_{\rm ex} \left(\varphi_2^{\rm non} - \varphi_2^{\rm eq}\right) \mathrm{d}V \tag{2}$$

The detailed derivation for Eq. (2) can be found in the pervious papers [16–18]. An alternative form of Eq. (2) can be derived by means of Gauss formula,

Scheme 1 .

$$[\rho_1, \varphi_1^{\text{eq}}] \xrightarrow{\text{fast}} [\rho_2, \varphi_2^{\text{non}}] \xrightarrow{-\lambda} [\rho_2, \varphi_2^{\text{eq}}] \xrightarrow{+\rho_{\text{ex}}, \text{quasistatic}} [\rho_2 + \rho_{\text{ex}}, \varphi_2^{\text{non}}]$$

$$\lambda_{s} = -\frac{1}{2} \oint_{S} V_{\text{ex}}(\sigma_{2}^{\text{non}} - \sigma_{2}^{\text{eq}}) dS$$
$$= -\frac{1}{2} \oint_{S} (\Delta \sigma^{\text{dyn}} - \Delta \sigma^{\text{eq}}) V_{\text{ex}} dS$$
(3)

where V_{ex} is electric potential arising from ρ_{ex} , σ_2^{non} and σ_2^{eq} are the nonequilibrium and equilibrium polarization distributions on the surface, respectively. $\Delta \sigma^{\text{dyn}}$ is surface polarization charge density generated by the change of the solute charge $\Delta \rho$ ($\Delta \rho = \rho_2 - \rho_1$) in the medium with a dielectric constant of ε_{op} , while $\Delta \sigma^{\text{eq}}$ is that in the medium with a dielectric constant of ε_{s} . Equation (3) can be easily changed into matrix form, i.e.,

$$\lambda_{\rm s} = -\frac{1}{2} V_{\rm ex}^{\rm T} (\Delta q^{\rm dyn} - \Delta q^{\rm eq}) \tag{4}$$

here, V_{ex}^{T} is transposed matrix of the column matrix V_{ex} for electrostatic potentials produced by ρ_{ex} . Δq^{dyn} and Δq^{eq} are the matrix forms of $\Delta \sigma^{\text{dyn}}$ and $\Delta \sigma^{\text{eq}}$, respectively. In order to obtain the elements of Δq^{dyn} , Δq^{eq} , and V_{ex} , we can solve the linear matrix equation of IEF-PCM [19, 20], i.e.,

$$D_{q} = -V \tag{5}$$

where D is the interaction square matrix collecting elements related to geometry of the cavity and the dielectric constants $\varepsilon(\varepsilon_s \text{ or } \varepsilon_{op})$. The column vector q collects the polarization surface charges [21]. According to the equilibrium polarization, we have

$$D_{\varepsilon_s} q_2^{\text{eq}} = -V_{2c} \tag{6}$$

$$D_{\varepsilon_{\rm s}} \Delta q^{\rm eq} = -\Delta V_{\rm c} \tag{7}$$

$$D_{\varepsilon_{\rm op}} \Delta q^{\rm dyn} = -\Delta V_{\rm c} \tag{8}$$

$$D_{\varepsilon_{s}}(\Delta q^{dyn} - \Delta q^{eq}) = \Delta V_{ex}$$
(9)

Then, a new matrix form for the solvent reorganization energy is given by

$$\lambda_{s} = \frac{1}{2} \left[(D_{\varepsilon_{s}}^{-1} - D_{\varepsilon_{op}}^{-1}) \times \Delta V_{c} \right]^{T} \times D_{\varepsilon_{s}} \times \left[(D_{\varepsilon_{s}}^{-1} - D_{\varepsilon_{op}}^{-1}) \times \Delta V_{c} \right]$$
(10)

Equation (10) tells that the solvent reorganization energy from our new theory depends on the interaction matrix D and the solute potential change ΔV_c . The detailed derivation of Eq. (10) can be found in our recently submitted paper [22]. This equation can easily be obtained by IEF-PCM. More interestingly, differing from D-PCM, the selfenergy of a given surface element is not involved in the above expressions. It means that IEF-PCM is more suitable for our new model. In this work, we have developed a subroutine coupled with the Q-Chem package [23] to numerically calculate the values of λ_s .

3 Results and discussion

The self-exchange ET reaction in helium and its cation radical (He–He⁺) system is here taken as a simple test example. The motivation of choosing this system is to check our numerical algorithm by comparing the numerical values with the results of analytical two-sphere model. It is well known that the diatomic system is an ideal system for us to apply two-sphere model; thus, we can take this ET as a benchmark example. In our previous work [18], we obtained the expression of analytic solution for λ_s with two-sphere model as follow:

$$\lambda_{\rm s} = \frac{\Delta q^2}{2} \left(\frac{1}{r_{\rm D}} + \frac{1}{r_{\rm A}} - \frac{2}{\rm d} \right) \frac{(\varepsilon_{\rm op}^{-1} - \varepsilon_{\rm s}^{-1})^2}{(1 - \varepsilon_{\rm s}^{-1})}$$
(11)

where Δq is defined as the amount of point charge transferred from sphere D to A as showed in Fig. 2. r_A and r_D are radii of the acceptor and donor sphere, respectively, while *d* is the distance between the two spherical centers. ε_{op} and ε_s are the optical and static dielectric constants. According to our pervious study [12], this equation gives a satisfying estimation of λ_s for pyrene-fumaronitrile and ferrous-ferric systems compared with available experimental values. Here, we use Eq. (11) to calculate λ_s of He–He⁺. When the two helium atoms are far from each other, the cavity consists of two separated spheres. When *d* is large enough, the values of λ_s by numerical solution ought to approach to the results by two-sphere model. So, this example can be considered as the criteria to detect whether the numerical program is correct or not.

For the numerical algorism, the well-known Generating Polyhedra (GEPOL) procedure [24] has been applies, with the Waals radii being scaled by a factor of 1.2 in the construction of the cavity. The van der Waals radius of helium



Fig. 2 Two-sphere model



Fig. 3 Distance dependency of the solvent reorganization energy

is 1.2 Å. Thus, the cavity radius equals to 1.44 Å. Each sphere is divided into 960 tesserae in order to guarantee the computation precision. We take water as solvent, for which the static and the optical dielectric constants are 78.5 and 1.778, respectively. When the distance of two helium atoms is less than 2.88 Å, the solvent cavity consists of two intersected spheres. Increasing the distance to 4.0 Å, the cavity is divided into two independent spheres, and each has 960 tesserae.

The values of λ_s with the increasing of the separation distance by both two-sphere model and numerical solution are depicted in Fig. 3. It can be seen that the numerical solution and two-sphere model give consistent results when the distance is larger than about 4.5 Å. At the critical value of 2.88 Å of separation distance, i.e., $d = r_A + r_D$, where the two cavities just contact with each other, the value of λ_s is 33.12 kcal/mol by two-sphere model and 36.47 kcal/mol by the numerical solution. If we take the numerical solution as accurate, it will be found that two-sphere model underestimates λ_s by about 10 %. This indicates that two-sphere approximation behaves very well when the two spheres are far from each other. However, the results of two-sphere model become worse when the two spheres intersect (d < 2.88 Å). Therefore, we should pay attention to the limitation of two-sphere model in applying it to calculate λ_s of ET reactions.

Recently, CDFT was used to construct the diabatic states of ET [7, 20, 21, 23]. We studied the self-exchange ET reactions between TCNE, TTF, and their corresponding ionic radicals in acetonitrile. The structures TCNE and TTF are given in Fig. 4. We use the same method of CDFT as pervious paper [13] to optimize the geometries of the species and construct the diabetic states of those two ET systems. The values of the inner reorganization energy λ_m were reported in Ref. [13] where one can find the detailed description of the geometry optimization and the calculation of λ_m . Here, we recalculate λ_s for those two systems by means of the above IEF-PCM-based matrix expression of numerical algorithm. With explicitly constraining one moiety in the complex to have one charge more than the other moiety, the geometries have been optimized by changing the separation distance (R_{AD}) from 2.5 to 4.0 Å by CDFT with B3LYP/6-31 + g(d) method. The minimum energy structures are located at $R_{AD} = 3.35$ Å for TCNE/TCNE⁻ and $R_{AD} = 3.42$ Å for TTF/TTF⁺.

At the fixed donor–acceptor distances, CDFT method has been adopted to perform the calculation. The charge densities for the complex systems before and after the ET have been obtained [13]. With these charge distributions, the solvent reorganization energy has been calculated with Q-Chem according to Eq. (10), by introducing the Q-Chem output files to a separated subroutine that carries out the cavity construction and the calculation of polarized charges in the framework of IEFPCM. The obtained results for λ_s , with the experimental values available [25–27], are listed in Table 1. Besides, other theoretical data from the smoothed multisphere model (SMSM) [26] have been listed as well.

A quick glance at the SMSM results given in Table 1 leads us to a wrong impression that the traditional theory gives the rational reorganization energy. However, if we revisit the theoretical treatments by other authors, we will find there are some inconsistent treatments in selecting the cavity sizes comparing with the well-established

Fig. 4 Structures of **a** TCNE/TCNE⁻ and **b** TTF/TTF⁺ complexes



Table 1 Comparison of the theoretically calculated reorganization energies (cm⁻¹) with the experiment values in acetonitrile for ET systems TCNE/TCNE⁻ and TTF/TTF⁺

System	λ_{s}	λ _m	λ
TCNE/TCNE-	5008 ^a 5350 ^b	2265 ^a 2300 ^b	7273 ^a 7620 ^b 7250 ^c
TTF/TTF ⁺	3623 ^a 4520 ^b	2274 ^a 2270 ^b	$5897^a6820^b5810^c$

 $^a\,$ This work. λ_m is from our previous work based on D-PCM [13]

^b Values based on the SMSM [26]

^c Experiment data obtained from the intervalence absorption, Refs. [25–27]

equilibrium solvation. It is well-known that in the treatment of DPCM for equilibrium solvation, the solute cavity is constructed by taking the atomic sphere with van der Waals radii being scaled by a factor of 1.2. Such a cavity size usually exhibits reasonable equilibrium solvation energies. However, in the calculation of nonequilibrium solvation energy, authors in general need to use much larger solute cavities so as to obtain the solvent reorganization energy comparable to the experimental measurements, since the same cavity size as that used for equilibrium solvation energy calculation definitely leads to a much larger λ_s by the previous theory. Here, we take the following examples for our arguments. As for experimental values, the energy of the diagnostic NIR band in the Class II mixed-valence TCNE/TCNE⁻ and TTF/TTF⁺ complex provides the direct measurement of the reorganization energy for ET as $\lambda = \nu_{IV}$ with ν_{IV} being the energy of the NIR band maximum [26, 28]. In the theoretical aspect, Kochi et al. applied the SMSM for the calculation of solvent reorganization. In the construction of the cavity, a value of ~2 Å for $r_{\rm p}$, the radius of an effective sphere was taken for the solvent acetonitrile, and hence, the sphere radius takes a value of van der Waals radius plus $r_{\rm p}$ in constructing the solute cavity [26]. We see that the H atomic sphere will take a radius of 3.2 Å, while in DPCM treatment for equilibrium salvation, this value is only 1.44 Å. So, in the SMSM treatment for nonequilibrium solvation, a much larger solute cavity is spanned compared with DPCM cavity for equilibrium solvation calculation. In fact, if one uses the widely accepted cavity size spanned by the spheres with 1.2 times of atomic van der Waals radii, it is expected that the SMSM will produces the doubled solvent reorganization energies, rather than those as listed in Table 1 by means of SMSM.

Apparently, the calibration to the experimental results in solvent reorganization energy calculation by enlarging the solute cavity is illogical, and it covers up the original mistakes in the traditional theories of nonequilibrium solvation and will cause the confusions in the sphere radius choice in solvent effect evaluation by continuum model. Moreover, these confused treatments for solute cavity size are also harmful to the confidence for the applications of continuum model, since people sometimes incorrectly attribute the ill results of solvent reorganization energy by Marcus model to the inaccuracies of continuum model [29].

4 Conclusion

In this work, a numerical algorithm for λ_s has been introduced. After the development of a subroutine coupled with the Q-Chem package based on IEF-PCM algorithm, the matrix form of the solvent reorganization energy has been calculated. In order to prove its rationality, the selfexchange ET in He-He⁺ system was investigated using both numerical solution and two-sphere model. Actually, He–He⁺ system is just a positively charged diatom system. The choice of He atom here is just for the purpose of the gain of the atomic radius. For the point charge approximation, the calculation in fact has nothing to do with other properties of the atom selected, except the atom radius. From Fig. 2, one can see that the numerical and analytical treatments give consistent results when the distance between two helium atoms is larger than 4.5 Å. However, when the two spheres intersect (d < 2.88 Å), the results of two-sphere model become worse. This fact reminds us to be careful with the limitation of two-sphere model in applying to calculate λ_s of ET. Finally, self-exchange ET reactions between TCNE, TTF, and their corresponding ionic radicals in acetonitrile have been investigated. The obtained reorganization energies λ of 7,273 cm⁻¹ for TCNE and $5,897 \text{ cm}^{-1}$ for TTF are in good agreement with available experimental results of 7,250 and 5,810 cm^{-1} , respectively.

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REGULAR ARTICLE

Adsorption of water molecules on sodium chloride trimer

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Abstract The hydration of NaCl has been widely studied and believed to be important for understanding the mechanisms of salt dissolution in water and the formation of ice nucleus, cloud, and atmospheric aerosols. However, understanding on the *poly*-NaCl ion pair interacting with water is very limited. Here, we investigated the adsorption of water molecules on (NaCl)₃, using both theoretical calculations and anion photoelectron spectroscopy measurements. The calculated vertical detachment energies and the experimental ones agree well with each other. Furthermore, we found that, for neutral (NaCl)₃(H₂O)_n (n = 2-7) clusters, the water-doped cuboid and structures formed by adding water molecules on the Na–Cl edges of the cuboid are energetically favored; water molecules preferentially bind to the

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Y. Q. Gao Biodynamic Optical Imaging Center (BIOPIC), Peking University, Beijing 100871, China Na–Cl edge if the NaCl ion pair has larger partial charges than others. We also found the anionic structures are more various compared with neutral ones, and the Na⁺ and Cl⁻ ions are hydrated more easily in the anionic clusters than in the corresponding neutrals.

1 Introduction

Salt ions can strongly affect the water/air surface tension and the solubility of proteins [1–5]. They are also related to the formation and reactivity of sea salt aerosols [6–8]. Thus, one challenge is to understand the ion–water and water–water interactions at the molecular level and to obtain the detailed solvation mechanisms of ions and/or ion pairs in water [9–11]. Molecular clusters are considered to play important roles in providing the connection between the properties of isolated molecules/atoms and those of the condensed phase systems. In particular, due to their significance in many fields, small clusters of sodium chloride and water molecules NaCl(H₂O)_n have called attention from both experimental [11–16] and theoretical [17–33] groups.

In contrast to the large number of studies on the clusters formed by a single sodium chloride ion pair and water molecules, i.e., $\text{NaCl}(\text{H}_2\text{O})_n$, studies on $(\text{NaCl})_x$ (with x > 1) are rare. There do exist several studies on sodium chloride clusters [34–37]. For example, Sunil and Jordan [34] performed theoretical calculations on negative $(\text{NaCl})_{2-4}$ clusters and discussed the properties of the excess electron binding. This topic was revisited by Anusiewicz et al. [35] and they argued that there are two types of anions in this system, dipole-bound anions and solvated-electron species based on their high-level calculations. Iwona et al. [36] reported results on (NaCl)₁₋₆ calculations in which various isomers with different bonding forms are presented. Ayuela et al. [37] also theoretically predicted the structures of $(\text{NaCl})_n (n \le 19)$ clusters, using the ab initio perturbed-ion model. It is noted that these two studies [36, 37] actually gave two similar minimal structures of (NaCl)₃. However, these studies do not include water molecules in the clusters, and thus, the detailed interactions of (NaCl)_r with water remain unexplored. Motivated by the potential importance of $(NaCl)_3(H_2O)_n$ clusters in the formation of doped crystals [38], atmospheric processes at the ocean surface [39, 40], and salt dissolution processes [41], we herein investigated the initial hydration of (NaCl)₃ by using theoretical calculations and photoelectron spectroscopy experiments, for clusters $(NaCl)_3(H_2O)_n$ (n = 0-6). The calculations were performed for the clusters both in the anionic and neutral states.

Photoelectron spectroscopy in combination with quantum calculations is a very mature tool to study clusters [42]. However, computational studies on the clusters with relatively large size are hindered by the outstanding difficulty in covering the large configuration space of the clusters. There exist many local minima on the potential energy surfaces; thus, it is computationally challenging to locate all the stable isomers. In the current study, to overcome this difficulty, we used our well-established integrated tempering sampling (ITS) molecular dynamics [43] to generate structures of low configuration energies. ITS allows efficient search of structures by sampling over a broad energy range and thus permits fast identification of low-energy structures and transition states. The low-energy structures obtained from the ITS calculations are then optimized using both classical force field and DFT functional. This procedure was proven effective and efficient in structure search for $(H_2O)_n$, $LiI(H_2O)_n$, $CsI(H_2O)_n$ [44], and $NaCl(H_2O)_n$ [45] with *n* being up to 20 in our previous studies. We note here that other global minimization methods besides ITS can be efficient and effective for searching the low-energy structures. However, through ITS simulation, one can readily obtain the various thermodynamic properties in addition to stable structures as well [44].

The rest of this account is organized as follows: In Sect. 2, we described the experimental details and theoretical methodologies; experimental spectra and computational results of both anions and neutrals and discussions are arranged in Sect. 3; and Sect. 4 is the concluding remarks.

2 Experimental details and theoretical methodologies

In this part, we will first describe the details of the anion photoelectron spectroscopy experiments. Then, a brief introduction of the basic principles of the integrated tempering sampling (ITS) method will be discussed. The theoretical details, including the classical MD (and ITS) simulations and the DFT calculations, are described in the last part.

2.1 Photoelectron spectroscopy

The experiments were conducted on a homebuilt apparatus consisting of a time-of-flight (TOF) mass spectrometer and a magnetic-bottle photoelectron spectrometer, which has been described previously [46]. Briefly, the $(NaCl)_{3}^{-}(H_{2}O)_{n}$ clusters were produced in a laser vaporization source, by laser ablating a rotating and translating NaCl disk target with the second harmonic (532 nm) light pulses from a Nd:YAG laser, while helium carrier gas with ~4 atm backing pressure seeded with water vapor was allowed to expand through a pulsed valve to generate and cool the hydrated $(NaCl)_3^-$ clusters. The cluster anions were mass-analyzed by the TOF mass spectrometer. The $(NaCl)_{3}^{-}(H_{2}O)_{n}$ (n = 0-6) clusters were each mass-selected and decelerated before being photodetached by the 532 nm (2.331 eV) photons from another Nd:YAG laser. The photodetached electrons were energy-analyzed by the magnetic-bottle photoelectron spectrometer. The instrumental resolution was approximately 40 meV for electrons with 1 eV kinetic energy.

2.2 Integrated tempering sampling (ITS) method

Briefly, in the ITS simulation, a modified potential energy function is obtained from the original potential by making use of an energy distribution function with the form of a summation of Boltzmann factors over a series of temperatures:

$$P(U) = \sum_{k=1}^{N} n_k e^{-\beta_k U} \tag{1}$$

where $\beta = 1/k_B T$, k_B is the Boltzmann constant, T the temperature, and U the potential energy of the system.

The sampled potential energy range by such a method is largely expanded, to both low and high potential energies, when compared to standard MD simulations. In the current study, we used 300 discrete temperatures even spaced in the range of 100–250 K. In the ITS simulation, the system is governed with a biased potential function in the form of Eq. (1). An equilibrium simulation with ITS yields a biased distribution function as a function of the potential energy of the system,

$$\rho'(r) = \frac{e^{-\beta U'(r)}}{Q'} \tag{2}$$

where $Q' = \int e^{-\beta' U(r)} dr$. In ITS, since U'(r) is a known function of U(r), the distribution function $\rho(r)$ for the system with the unbiased potential can be recovered back from $\rho'(r)$ as:

$$\rho(\mathbf{r}) = \frac{e^{-\beta U(r)}}{Q} = \rho'(\mathbf{r}) \frac{e^{-\beta [U(r) - U'(r)]} Q'}{Q}$$
(3)

where $Q = \int e^{-\beta U(r)} dr$. More details can be found in our previous publications [43, 47].

2.3 Simulation details

We used SANDER module of AMBER 9.0 program package [48] to perform MD simulations of $(NaCl)_3(H_2O)_n$ (n = 0-7), in which the ions 08 parameters [49] and single-point-charge/extended (SPC/E) water model [50] were utilized. The ion-water cross terms were calculated with the Lorentz Berthelot combination rules. These models are popular and well established in AMBER program package, and they give convincing bulk solution properties comparing to experimental data [49, 50]. One may expect more reliable thermodynamic properties in the classical simulations when more accurate models, such as polarizable force field, are utilized to these systems, but our earlier studies did show that the classical force field used in this study provides reasonable description of the cluster structures compared with the quantum calculations [44]. The initial configurations used for MD were generated randomly and followed by an energy minimization. To avoid the molecules to escape from the clusters to the vacuum, a spherical potential is applied to the center of mass of every atom. The restricting force of the added potential has the form of:

$$F(R) = \frac{\varepsilon}{1 + e^{-2\alpha\varepsilon|(r-r_0)|}}$$
(4)

where *r* is the spatial vector of an atom, ε is the maximal restrict force [here, we set = 5 kcal/(mol Å)], r_0 is the radius of sphere (here, we set $r_0 = 7$ Å), and is the flexibility constant of the sphere shell (we set = 5). In this way, the molecules are confined in a spherical space with a radius of 7 Å.

We collected a 100-ns-long ITS simulation trajectory for each system of $(NaCl)_3(H_2O)_n$. Hundreds of different low-energy configurations were chosen from simulated trajectory as the initial structures for structure optimization using classical force field. These structures then converged to several dozens of local minimal structures, and then, re-optimizations by DFT calculations were performed. The DFT calculations were performed using GAUSSIAN 09 program package [51]. The structures of $(NaCl)_3(H_2O)_n$ were fully optimized with DFT employing the dispersion-corrected hybrid functional ω B97XD [52]. The Pople's all-electron basis set 6-311++G(d,p) was used for all the atoms. The ω B97XD functional has been shown to perform extraordinarily well on describing dispersion interaction [53], which is the main consideration for such weak-interaction cluster systems. As the energetics and geometries of one electron-bound anion clusters will be evaluated, we add the diffuse functions to all atoms. After these structures were fully optimized, harmonic vibrational frequencies were calculated to confirm that they are indeed local minima and to obtain the zeropoint energy corrections and the Gibbs free energies. Zeropoint vibrational energy corrections were incorporated to the sum of the electron energies.

3 Results and discussions

3.1 The experimental and theoretical vertical detachment energies

The photoelectron spectra of $(NaCl)_3(H_2O)_n$ (n = 0-6)clusters recorded at 532 nm are shown in Fig. 1. In general, all the spectral features of the clusters are broad, and their maxima are about 0.99, 1.04, 1.09, 1.11, 1.03, 1.05, and 1.07 eV for n = 0-6, respectively. Specifically, for $(NaCl)_{3}^{-}(H_{2}O)$, there is an additional small feature centered at about 1.51 eV. Figures 2 and 3 present the typical low-energy structures for each cluster size along with their relative energies and calculated VDEs (the geometries and energetics are discussed below). The calculated VDEs of the lowest energy structures of $(NaCl)_{3}^{-}(H_{2}O)_{n}$ (n = 0, n)1, and 3-6) are in good accord with the experimental values. While for $(NaCl)_{3}^{-}(H_{2}O)_{2}$, the theoretical VDE of the lowest energy isomer, which is a distorted cuboid structure, is 0.54 eV and much lower than the experimental value of 1.09 eV. The VDE value is also 0.54 eV when calculated at the MP2/aug-cc-pvtz level of theory. These results at both ω B97XD and MP2 levels suggest that the structure 2A⁻ is unlikely the structure observed in the experiments. By examining the Gibbs free energies of these structures, it is found that structure 2C⁻ has the lowest Gibbs free energy and its theoretical VDE is 1.05 eV, consistent with the experimental measurement of 1.09 eV very well. For $(NaCl)_{3}^{-}(H_2O)_n$ (n = 0, 1, and 3-6) clusters, we also checked their Gibbs free energies and found that the lowest electronic energy structures also have the lowest Gibbs free energies. These results suggest that the photoelectron spectra may relate to the structures of relative large stability in Gibbs free energies. The good performance of $\omega B97XD/6$ -311++G(d,p) on the $(NaCl)_{3}^{-}(H_{2}O)_{n}$ clusters shown above provides the credence of the following discussions on the corresponding neutral clusters.



Fig. 1 Photoelectron spectra of the $(NaCl)_{3}^{-}(H_{2}O)_{n}$ (n = 0-6) clusters taken with 532 nm photons

3.2 Structures of $(NaCl)_3^-(H_2O)_n$

3.2.1 Typical minimal energy structures of $(NaCl)_3^-$

The typical minimal energy structures of $(NaCl)_3^-$ are displayed in Fig. 2. As discussed later, the structures of salt– water molecules being adsorbed on $(NaCl)_3^-$ mostly evolve from these pure salt structures. The global minimum of the anion $(0A^-$ in Fig. 2) is a "A-shaped" structure, similar to that reported in the previous studies [34–37]. The VDE of this structure is calculated to be 1.045 eV, in good agreement with the experimental measured value of 0.99 eV. Structure 0B⁻ is higher in energy by only 0.001 eV than the structure 0A⁻. To the best of our knowledge, this "fishshaped" C_{2V}-symmetric structure (structure 0B⁻) has not been reported so far for either anionic or neutral forms. The kite-shaped 0C⁻ and a little distorted kite-shaped 0F⁻, which are optimized from the initial configuration of by changing the positions of Na⁺ and Cl⁻ of structure 0C⁻, are higher in energy than 0A⁻ by 0.170 and 0.519 eV, respectively. The linear 0D⁻ and Y-shaped 0E⁻ structures are much less stable than others identified here. In summary, most of the structures found in the current study are very similar to those reported previously. Therefore, we followed Anusiewicz et al. [35] for the naming of these structures. Besides, the current study also identifies a new low-energy "fish-shaped" structure (0B⁻), which has not been reported before. Furthermore, the excess electron substantially increases the complexity of the potential energy surface, which will be shown by the richness of the anionic structures in comparison with the corresponding neutral ones.

3.2.2 Typical minimal energy structures of $(NaCl)_3^-(H_2O)_n$ (n = 1-6)

Figure 3 shows the typical minimal energy structures of $(NaCl)_{3}^{-}(H_2O)_n$ (n = 1-6). For n = 1, five of the six lowest energy structure are formed through adsorbing one water molecule to the " Λ -shaped" structure of $(NaCl)_{3}^{-}$ (see $0A^{-}$ in Fig. 2). In structure $1A^{-}$, the O atom of the water interacts with two Na atoms, and the two H atoms of the water are dangled. Due to the excess electron, it can be seen that the two dangling H atoms forming two O-H…e⁻ bonds (see Fig. S1 in Supporting Information), which have been suggested to be strongly favored in the liquid phase of methanol [54].

For n = 2, the global minimum structure $2A^-$ is a slightly distorted cuboid. As mentioned above, the calculated VDE of this configuration is 0.54 eV. Yang et al. [55] have previously found that the neutral $(\text{NaCl})_4$ forms a particularly stable cuboid and should have relatively low electron affinity. The other "A-shaped"-based structures are formed by re-orientation of the water molecule to interact with Na or Cl ions in different ways for both n = 1 and 2. In addition to these structures, the $1D^-$ and $2C^-$ structures are derived from the newly identified "fish-shaped" structure $(0B^- \text{ in Fig. 2})$.

For $(NaCl)_{3}^{-}(H_{2}O)_{n}$ (n = 3-6) clusters, the excess electron causes a significant perturbation to the neutral structures. It can be seen that all the local minimal structures are derived from the distorted cuboid $(NaCl)_{3}^{-}(H_{2}O)_{2}$ ($4A^{-}$ and $6A^{-}$) or much more distorted structures ($3A^{-}$ and $5A^{-}$). For these clusters with larger number of water molecules, the cuboid-based structures no longer dominate. At the same time, the Na⁺ and Cl⁻ ions are much more hydrated in the anions (such as $3A^{-}$, $3C^{-}$, and $3D^{-}$) than in the corresponding neutrals, which can be seen later. It therefore appears that the excess electron causes enhanced hydration of Na and Cl ions. For the rest of the large number of

Fig. 2 Typical minimal energy structures of $(NaCl)_3^-$. The relative energies to the corresponding smallest values (*first line*) and the theoretical VDEs (*second line*) are presented. All the energies are in eV



Fig. 3 Typical minimal energy structures of $(\text{NaCl})_3^-(\text{H}_2\text{O})_n$ (n = 1-6). They are presented in the similar way as those in Fig. 2

anionic clusters not discussed in details here, we provided their Cartesian coordinates, corresponding relative energies, and theoretical VDEs in Supporting Information.

3.3 Structures of $(NaCl)_3(H_2O)_n$

3.3.1 Typical minimal energy structures of $(NaCl)_3(H_2O)_n$ (n = 0-2)

To understand salt-water interactions, it is certainly more interesting to study the neutral salt-water clusters. Figure 4 shows the typical minimal energy structures for $(NaCl)_3(H_2O)_n$ (n = 0-2) clusters. For $(NaCl)_3$, the previous studies [34-37] identified and discussed two minimal energy structures, i.e., the global minimum 0A, which is a 2-dimensional (2D) plane structure and the D_{3h}-symmetric 0B structure as shown in Fig. 4. In the current study, a third minimal energy structure ("fish-shaped" OC in Fig. 4) was also identified though it is relatively high in energy (0.174 eV higher than 0A). It is also noted here that the D_{3h} -symmetric structure is not found to be stable in the anionic form. The optimization from 0B in the anionic state leads the structure into the arrow-shaped 0D⁻ structure. These findings are consistent with previous studies [35] and could be a result of the high symmetry of 0B, whose dipole moment is zero, and thus, its binding of an excess electron is too weak to form a stable anion.

Among the $(NaCl)_3(H_2O)$ structures, 1A is resulted from adsorbing the water molecule on the edge of Na–Cl of the structure 0A of $(NaCl)_3$. Adsorption of one water leads to a structure transition from planar $(NaCl)_3$ ("arrow-shaped" 0A) to structure 1B, which is higher in energy than 1A by 0.034 eV. In structure 1B, $(NaCl)_3$ is actually very similar to the " Λ -shaped" structure, which is the global minimum of the anionic state (0A⁻ in Fig. 2). The other local minimum structures with much higher energies than the global minimum structure can be seen to arise from the cyclic (1C and 1D), "fish-shaped" (1E), or "arrow-shaped" (1F) (NaCl)₃ structures.

It is well known that the lowest energy structure of the $(NaCl)_4$ cluster is a T_d-symmetric cuboid [36, 37]. This study finds that the global minimum structure of $(NaCl)_3(H_2O)_2$ cluster is a distorted cuboid (2A in Fig. 4). In structure 2A, two water molecules occupy the sites of Na^+ and Cl^- of $(NaCl)_4$, respectively, with one water molecule interacting with the other two Na⁺ through its O atom, and another one connecting to two Cl⁻ via hydrogen bonds. One hydrogen bond also forms between the two water molecules, and one hydrogen atom of the Na-coordinated water is dangling. Other structures such as 2B, 2C, 2D, 2E, and 2F can all be regarded as the distorted forms of the 2A structure. In these structures, the positions and orientations of the two water molecules deviate from the cuboid 2A structure. 2G is a derivative of the planar arrow-shaped structure 0A and is much higher in energy than others. Different from (NaCl)₃(H₂O), most of the seven low-energy structures of the (NaCl)₃(H₂O)₂ cluster are non-planar. From the above analyses, we can see that the adsorption of one/two water molecule(s) or one electron causes a (NaCl)₃ structure transition from planar to non-planar.

3.3.2 Typical minimal energy structures of $(NaCl)_3(H_2O)_n$ (n = 3-7)

We show the typical low-energy structures of $(NaCl)_3(H_2O)_n$ (n = 3-7) clusters in Fig. 5. Interestingly,



Fig. 4 Typical minimal energy structures of $(NaCl)_3(H_2O)_n$ (n = 0-2). The relative energies to the corresponding smallest values are presented. All the energies are in eV



Fig. 5 Cuboid structure evolution in the clusters of $(NaCl)_3(H_2O)_n$ (n = 2-7). The relative energies to the corresponding smallest values are presented. All the energies are in eV

the cuboid formed by $(NaCl)_3(H_2O)_2$ is maintained in many low-energy structures. A chart of possible structure evolution is given in Fig. 5.

For $(NaCl)_3(H_2O)_3$, 3A and 3B are almost degenerate in energy with the latter being higher by only 0.001 eV. The water molecule that does not belong to the cuboid, as mentioned above, binds to the Na–Cl edges. By examining the Mülliken charge of the atoms in structure 2A, it is found that water molecules prefer to bind to the Na–Cl edges, for which the Na and Cl atoms have the largest sum of partial charges (SPCs) (see the Supporting Information for the Mülliken charges in Table S1). Interestingly, when the SPC values of the two atoms are both small, binding of the water molecule to a Na–Cl edge causes the separation of Na⁺ and Cl⁻ (see 3C in Fig. 5 and the Na–Cl distance in Table S2 in Supporting Information). Adsorption on the Na–Cl edge of medium SPC values results in two structures much higher in energy (3D and 3E in Fig. 5). In these two structures, Na⁺ and Cl⁻ remain in contact. The lowest energy structure of $(NaCl)_3(H_2O)_4$ cluster 4A is resulted from adsorbing the other two water molecules on the two Na–Cl edges, and it is noted that the SPCs of these two edges are the highest and second highest values in structure 3A. When adsorption of water is on the other Na–Cl edges (4B to 4G), the resulted structures are of higher energies. A penta-prism structure 4H, higher in energy than 4A by 0.074 eV, was also found for the $(NaCl)_3(H_2O)_4$ cluster. In this structure, the four water molecules form a four-member ring, and this ring further connects to the Λ -shaped-like $(NaCl)_3$ structure.

The lowest energy structure of $(NaCl)_3(H_2O)_5$ cluster 5A can be constructed from 4A with the newly added water molecule interacting with two Cl⁻ and one Na⁺ ions. The dipole of the water molecule is almost parallel to the plane composed of Cl–Na–Cl. The binding pattern of the water molecule in this cluster is similar to that on NaCl(001) surface [56]. The adsorbed water molecules, as mentioned above, preferentially interact with Na and Cl ions rather than other water molecules. In the latter case, less stable structures form, such as structures 5D and 5F in Fig. 5.

The low-energy structures of $(NaCl)_3(H_2O)_6$ also remain the basic cuboid form. The adsorbed water molecules on the NaCl ions can be seen in 6A, 6B, and 6C. Interestingly, a fused cuboidal structure 6D, which can be seen as two cuboid units sharing a face formed by $(NaCl)(H_2O)_2$ fragment, is also obtained. Although the water molecules form more water–water hydrogen bonds in 6D than in 6A, 6B, and 6C, it is still less stable than 6A by 0.023 eV.

Different from $(NaCl)_3(H_2O)_n$ (n = 2-6), the lowenergy structures of $(NaCl)_3(H_2O)_7$ are derivatives of the fused face-sharing cuboid 6D. The seventh water molecule can be adsorbed on the different Na–Cl edges (7A, 7B, 7C, 7D, and 7F). As can be predicted from the partial charge analysis of structure 6D, the most stable structure of $(NaCl)_3(H_2O)_7$ is formed by adding water molecule on the Na–Cl edge of the largest SPC value, which is the edge composed of "1Cl" and "5Na" (see the label in Table S1 in Supporting Information). Two structures 7E and 7G, maintaining only one cuboid consisting of $(NaCl)_3(H_2O)_2$ fragment, also appear as local minimal structures on the energy surface, and they are higher in energy than 7A by 0.056 and 0.069 eV, respectively.

By using enhanced sampling simulations followed by optimization using ω B97XD density functional, we also obtained a larger number of structures of the noncuboid nature, including the structures formed by adsorbing water molecule(s) on the several basic structures of (NaCl)₃ and other structures with Na⁺ and Cl⁻ being hydrated. These structures are not shown or discussed here, and the Cartesian coordinates and energetics of all the fully optimized cluster structures are provided in Supporting Information.



Fig. 6 The change of water binding energies (WBEs) versus the number of water molecules n for the most stable isomers of (NaCl)₃(H₂O)_n

3.4 Water binding energies of the lowest energy structures

To examine the stability of the cluster as a function of cluster size, we define the water binding energy (WBE) using Eq. (5):

$$\mathbf{E}\left[(\mathrm{NaCl})_{3}(\mathrm{H}_{2}\mathrm{O})_{n}\right] = \mathbf{E}\left[(\mathrm{NaCl})_{3}(\mathrm{H}_{2}\mathrm{O})_{n-1}\right] + \mathbf{E}(\mathrm{H}_{2}\mathrm{O}) - \mathrm{WBE}$$
(5)

In the above equation, $\mathbf{E}[(\text{NaCl})_3(\text{H}_2\text{O})_n]$ and $\mathbf{E}[(\text{NaCl})_3(\text{H}_2\text{O})_{n-1}]$ are the energy of the clusters with the number of water molecules being *n* and n - 1, respectively; $E(H_2O)$ is the energy of water monomer (zeropoint energy correction is included) and the value is -76.410660 atomic unit. WBE characterizes the absolute value of the energy decrease when a water molecule binds on the cluster. Figure 6 shows the WBEs as a function of the number of water molecules n in the lowest energy structures for both neutrals and anions. The binding energies for $(NaCl)_3(H_2O)_n$ (n = 1-7) are 0.644, 0.765, 0.610, 0.597, 0.565, 0.502, and 0.563 eV, respectively. The largest WBE of 0.765 eV is responsible for the second water molecule binding to structure 1A to form structure 2A. When *n* changes from 3 to 6, the WBE slightly decreases, which means that the further binding of water molecules to the clusters stabilizes the cluster less than the second water molecule does. When n changes from 6 to 7, a small increase of the WBE can also be found, which, as reflected from the structures changes, is due to that the fused face-sharing cuboid structure (7A) becomes the lowest energy structure in (NaCl)₃(H₂O)₇. The two maxima of the WBE in Fig. 6 are in accord with the first and second cuboid structures being the lowest energy ones when n = 2 and 7.

For the anions, the maximum WBE is 0.741 eV when n = 1, corresponding to the energy difference when the structure changes from $0A^-$ to $1A^-$. As can be seen in Figs. 2 and S1 and as mentioned earlier, the two dangling $O-H\cdots e^-$ bonds likely stabilize the cluster; the binding of the second water molecule results in the cuboidal $2A^-$ structure, but the WBE of 0.613 eV is smaller than when the first water is added. For n = 2 to 6, the WBEs are all smaller than the corresponding value for n = 1. A maximum of WBE occurs at n = 3 and a slight increase exists when n changes from 5 to 6. However, these sudden increases in WBE cannot be apparently indicated by the structural changes because, as mentioned in Sect. 3.2, the anionic structures are largely perturbed and Na⁺ and Cl⁻ are more hydrated than the corresponding neutral ones.

4 Concluding remarks

Water adsorption on (NaCl)₃ in both anionic and neutral states is investigated in this study by photoelectron spectroscopy experiments and DFT calculations. The possible observed structures in experiments are assigned according to the DFT results. The calculated vertical detachment energies (VDEs) and the experimental ones agree well with each other. The enhanced sampling method, ITS, was employed to ensure that the low-energy structures of the clusters are fully sampled. The efficiency of this approach has been shown in finding new minimal energy structures of (H₂O)₁₂ and (H₂O)₂₀ clusters [44], which were missed by other methods. The effectiveness of this method was again demonstrated in the present study. For instance, we identified a new "fish-shaped" C2V-symmetric structure in both anionic and neutral states, which has not been reported before [35–37, 57]. The anionic one is very competitive to (higher in energy by only 0.001 eV) the previously regarded global minimum structure in terms of our DFT calculations. In addition, by using dispersion-corrected ω B97XD functional and diffusive basis set, a more convincing ranking of these structures (by energy) was obtained.

For neutral $(NaCl)_3(H_2O)_n$ clusters, the water-doped cuboid and the structures derived from adding water molecules on the Na–Cl edges of the cuboid are energetically favored. We found that the most stable structure of $(NaCl)_3(H_2O)_2$ is a water-doped cuboid (2A in Fig. 5). This finding expands the class of dopant sodium chloride clusters: This study shows not only the metals [58] and hydroxide [59], but also water molecules can be the proper candidate for the doped NaCl clusters (crystals). The cuboid is so stable that the low-energy structures of $(NaCl)_3(H_2O)_n$ (n = 3-6) are those that all maintain the cuboid. The fused face-sharing cuboid structure appears when n is up to 6 though its energy

is slightly higher than the water-doped cuboid structure. Finally, the derivative structures from the fused face-sharing cuboid become dominant for the low-energy structures of $(NaCl)_{3}(H_{2}O)_{7}$. The lowest energy structures change from n = 1 to 2 and 6 to 7 are well reflected on the two maxima, which can be seen from the plot of the water binding energy (WBE) versus the cluster size (Fig. 6). Through this study, the pattern of water binding on (NaCl)₂ is established. From the structure evolution of the cuboid (Fig. 5), we can clearly find that the water molecule preferentially binds to the Na-Cl edge if the NaCl ion pair has larger partial charges than others. The fused face-sharing cuboid (6D) is the first structure in which the salt is fully covered by one layer of water molecules. The seventh water molecule again binds to the Na-Cl edge instead of other water molecules in the lowenergy structures of (NaCl)₃(H₂O)₇. Thus, water molecules will preferentially bind to the Na-Cl edges than interact with other water molecules.

Finally, we want to note that the anionic clusters are more distorted and have higher configuration diversity than the corresponding neutral ones. For example, for the waterfree (NaCl)₃ cluster, only three local minima were found for neutral form with relatively low energies, but for anionic clusters, seven low-energy stable structures were identified. For (NaCl)₃⁻(H₂O)_n (n = 1-6) clusters, although the majority of structures are similar to the corresponding neutral ones, the Na⁺ and Cl⁻ ions are much more hydrated in the former.

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REGULAR ARTICLE

Thermal decomposition of FOX-7 studied by ab initio molecular dynamics simulations

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Abstract Ab initio molecular dynamics simulations were conducted to study the thermal decomposition mechanisms of solid FOX-7. We found that the initial decomposition of FOX-7 has three main reaction routes. The C–NO₂ bond fission as reported previously is indeed the most common route. However, the inter- and intramolecular hydrogen transfers are also found to be valid, which is contradictory to previous energy minimization calculation results. The simulations agree well with known experimental data in terms of activation energy and reaction side products. A complete reaction network that describes how the main products, H₂O, CO₂ and N₂ form in FOX-7 decomposition, is constructed from the simulation trajectories.

Keywords FOX-7 \cdot Ab initio simulation \cdot Reaction mechanism \cdot Activation energy

1 Introduction

FOX-7 (1,1-diamino-2,2-dinitroethene) is a new energetic molecule superior to well-known RDX (1,3,5-trinitro-2-oxo-1,3,5-triazacyclo-hexane) and HMX (1,3,5,7-tetrani-tro-1,3,5,7-tetraazacyclo-octane) due to low sensitivity and easy synthesis procedure [1]. Although all of these molecules have similar molecular stoichiometry, their decomposition mechanisms are expected to be different because of

Dedicated to Professor Guosen Yan and published as part of the special collection of articles celebrating his 85th birthday.

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the differences in detonation performances. Understanding the decomposition mechanisms of FOX-7 is of great importance to design appropriate usage, transport and storage methods for this new energetic material [2, 3].

Considerable amount of experimental and theoretical work has been conducted with FOX-7 to study its reaction mechanisms [4-8]. A recent investigation using laser and mass spectroscopy techniques has been carried out to detect reaction products and transient intermediates [4]. Combining the measured data with quantum mechanical calculations, a reaction network was proposed [4]. Several calculations [5-8] were performed to study the decomposition mechanisms for FOX-7. The C-NO₂ and C-NH₂ bond dissociation energies were compared, and the former (293 kJ/ mol) was found to be more favorable than the later (466 kJ/ mol) [5]. The nitro-to-nitrite rearrangement was investigated, and an energy barrier of 247 kJ/mol was obtained [6]. The effects of charged and excited states on the decomposition mechanism of FOX-7 were studied [7]. The intramolecular and intermolecular hydrogen transfers were also investigated [8]. All of these calculations were based on characterizations of potential energy surfaces for proposed reaction pathways using a series of energy minimizations. We refer this type of calculations as *energy minimizations* in this manuscript.

Decompositions of energetic materials are usually characterized by complex reaction networks consisting of a large number of unit reactions and intermediates, which are correlated with the thermodynamic conditions such as temperature, pressure and concentration [9, 10]. Unlike reactions occurring in low-pressure gas phase or low-concentration solution, which can be usually described by a few unit reactions, reactions in high pressure or high concentration cannot be treated well by energy minimizations on preset reaction pathways because the *real* reaction pathways

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are too complicated to be postulated based on chemical intuitions. Although the energy minimizations can be used to accurately predict energies and structures of the reactants, products, intermediates and transition states, the calculations are not suitable for predicting new (unknown) reaction pathways, especially for those strongly depend on thermodynamic conditions. Generally speaking, it is highly challenging to obtain complete decomposition mechanisms of energetic materials.

Reactive molecular dynamics (RMD) simulation offers a promising way to study the complex chemical reactions without prior knowledge of reaction pathways and to include thermodynamic conditions into consideration. In RMD simulations, atoms are treated as classical particles moving on potential energy surfaces that represent the multi-atom interactions. The time trajectories of atoms are determined by numerical integration of the Newton's equation of motion. From the trajectories, chemical species including reactants, products and intermediates, as well as chemical reaction types that connect the species can be obtained. If the trajectories are ergodic, complex reaction mechanisms can be uncovered by the simulation. Two types of representation of the potential energy surfaces are used for RMD. One is to use empirical functional forms to describe the atom-atom interactions (called reactive force field or RFF); the other is to solve Schrödinger equation or Kohn-Sham equation instantly for each configuration generated in the simulation (ab initio molecular dynamics or AIMD). The empirical RFF method can be applied to systems containing up to millions of atoms and lasting up to microsecond time length, but it requires tedious and difficult parameterization. The AIMD method is computationally expensive so that it can only be applied to small systems and a short period of simulation time, but it is ab initio, unbiased by empirical inputs and suitable for discovering complex reaction pathways. The AIMD method has been used in studies of various chemical reactions, including the initial reactions of energetic materials such as CL-20 and nitromethane [3, 11, 12].

In this work, we carried out AIMD simulations using the Car–Parrinello scheme [13] to study the thermal decomposition of solid FOX-7. The purpose of these simulations was to discover unknown reaction pathways. By analyzing the simulation trajectories obtained at different temperatures thoroughly, we found important species and reaction types that have not been reported in previous theoretical and experimental studies. In addition, the reaction network that connects the major species by reaction types from the initial reactant (FOX-7) to the main end products is obtained. Due to exceptionally high expenses, the AIMD simulations were conducted with fairly small time and length scales, and consequently the statistical data reported

in this manuscript are associated with large uncertainties. Given the uncertainties, the data are still very valuable for us to describe the reactions mechanisms in a qualitative manner. The reaction pathways identified in this work provide a basis for further potential energy calculations, from which RFF can be developed to simulate decomposition reactions in much larger time and length scales.

2 Computational details

The constant number, volume and temperature (NVT) MD simulations were conducted using the Kohn–Sham DFT method with the plane wave and pseudopotential formulation as implemented in CPMD software [14]. The BLYP exchange–correlation functional and Troullier–Martins norm-conserving pseudopotentials [15] for core electrons were applied. A plane wave cutoff of 85 Ry was utilized. The mass for orbitals was set to 400 au. The Nosé–Hoover thermostats [16] were applied to control the temperature. A 0.05 fs time step was used for the integration of motion equations.

FOX-7 solid was represented by a super-cell with the periodic boundary condition. The super-cell consists of two unit cells, and the unit cell structure was taken from the Cambridge Structural Database (a-FOX-7 at 298 K) [17]. The initial configuration was equilibrated for 1 ps using NVT MD simulation at room temperature (298 K) to release any tensions between the reported structures and DFT predicted structures. No chemical reaction was observed by the end of the equilibration period. The relaxed structure was then used for four parallel simulations conducted at different simulations: 2,500, 3,000, 3,500 and 4,000 K, which are in the range of rapid laser detonation (2,500–5,000 K) [11]. The temperatures were assigned to the simulations with the same initial configuration so that the decomposition rates at different temperatures can be evaluated.

For each frame of the trajectories, chemical species were identified based on atom-atom distances. If the distance between two atoms is shorter than a threshold value, which is the sum of the covalent radius of two atoms plus a buffer size of 0.2 Å, the atoms are considered to be connected with a chemical bond. The covalent bond radii for C, H, O and N atoms are 0.77, 0.32, 0.73 and 0.75 Å, respectively. The species were uniquely labeled using the standard IUPAC International Chemical Identifier (InChI) code, version 1 [18]. By tracking the evolution of the identified species in the trajectories, chemical reaction types that connect species were tabulated. The analysis requires large amount of data processing, so we implemented a computer program based on relational database concept to facilitate data searching and sorting.

Fig. 1 Population distributions of reactant FOX-7 (*black*); intermediates NO₂ (*red*), NH₃ (*lawn green*), CO (*forest green*); and products H₂O (*blue*), N₂ (*magenta*), CO₂ (*yellow*) for simulation time, at (**a**) 2,500 K, (**b**) 3,000 K, (**c**) 3,500 K and (**d**) 4,000 K



3 Results and discussion

3.1 Overview of the reactions

The numbers of seven stable molecules (FOX-7, NH₃, NO_2 , CO, N_2 , H_2O and CO_2) with simulation time at different temperatures are plotted in Fig. 1. Due to the small size of these simulations, the fluctuations in the population curves are large, but general trends can be seen. The reactant FOX-7 population declines from initial value of eight, and the decline rate increases as the temperature increases. FOX-7 molecules decomposed completely near 2.5 ps at 2,500 K, 1.0 ps at 3,000 K, 0.5 ps at 3,500 K and less than 0.5 ps at 4,000 K. As soon as FOX-7 decomposition starts, NO₂ appears in the trajectory, as shown in Fig. 1a (2,500 K). NO₂ is an intermediate as its population reaches a maximum value of 5 and then decreases. The second compound appears in the trajectory is water (H₂O) which is one of the main products. The population of water fluctuates largely, indicating water acts as catalyst in part as well. Almost at the same time of water generation, two more intermediates, CO and NH₃, and another main product CO₂ are formed. The populations of intermediates CO and NH₃ are relatively low and their maximum values are about two. The CO₂ is not very stable as shown by the large fluctuations in its population curve. The last main product formed in the simulation is N_2 , which is a very inert compound with triple nitrogen-nitrogen bonds and its fluctuation in population curve is very small. Due to the small simulation length and time scales, the reactions are not completely finished at any temperature. However, the evolutions of species at 3,500 and 4,000 K as shown in Fig. 1 are similar. At the end of simulation, FOX-7 decomposition is finished and intermediates including NO₂, CO and NH₃ are converted to more stable products. In addition, main products N₂, H₂O and CO₂ are formed in the simulation system. In other words, the reaction processes at 3,500 and 4,000 K is close to completion.

The initial decomposition rates of FOX-7 are estimated from the population curves in Fig. 1. Assuming the initial decomposition of FOX-7 is a first-order reaction, we estimated the rate constants by fitting the population data of FOX-7 to:

$A(t) = A(0) \exp(-kt)$

where A(0) is the initial number density and A(t) is the number density at certain time. The fitting curves with the simulation data are shown in Fig. 2. The fitting is not quite accurate for the low-temperature (2,500 K) data, but fairly good for other temperatures. The rate constants in logarithm at different temperatures are depicted in Fig. 3. Apparently the rate constants cannot be fit by a straight line using the Arrhenius law:

$$\ln(k) = \ln(A) - \left[\frac{E_a}{R}\right] \frac{1}{T}$$

Fig. 2 Exponential fits of reaction rates at different temperatures, a 2,500 K, b 3,000 K, c 3,500 K, d 4,000 K. A(0) is the initial number and A(t) is the instant number of FOX-7. The *black line* is measured from the simulation trajectory, and the *red line* is exponential fit curve





Fig. 3 The decomposition rate constant versus reciprocal temperature. The *dots* are calculated rate constants. The *lines connecting two dots* are used to derive activation energies using the Arrhenius equation

The nonlinearity is presumably due to the large temperature spans (1,500 K) in which the frequency factor A is temperature dependent. Using two adjacent data to estimate the activation energies in different temperature ranges, we obtained 98 kJ/mol from 3,500 to 4,000 K, 142 kJ/mol from 3,000 to 3,500 K and 204 kJ/mol from 2,500 to 3,000 K. These values are in reasonable

 Table 1
 Molecular weight (MW) distribution of chemical species generated in the simulations at different temperatures

MW	2,500 K	3,000 K	3,500 K	4,000 K
1–10	1	1	1	1
11-20	5	7	11	10
21-50	22	51	65	59
100–199	42	155	300	327
101-200	148	219	300	359
201-300	56	43	42	88
301-400	8	8	6	7
401–500	2	0	0	0
Total	284	484	725	851

agreement with the experimental value of 242 kJ/mol, which was measured at the temperature range between 483 and 523 K [19].

3.2 Generated molecular species

A large number of molecular species were found in the trajectories of different temperatures. The numbers of molecular species found at different temperatures, grouped by molecular weight (MW), are listed in Table 1. The only species with MW less than 10 is hydrogen atom (either in

radical or proton form), and most species are in the range of 100–200 MWs. The total number of species appears to be dependent on temperature. The species number is 284 at 2,500 K, 484 at 3,000 K, 725 at 3,500 K and 851 at 4,000 K, respectively. Two factors contribute to this trend: The extents of reaction are different (see Fig. 1) and the entropy effect that favors smaller fragments. The first factor can be seen from the fact that the species numbers are similar for results at 3,500 and 4,000 K, since reactions are close to completion at these two temperatures. The second factor is evident from the MW distributions at different temperatures—the distribution shifts slightly to smaller species and the total number of species increases as the temperature increases.

The numbers of reaction types that connect the species are about twice larger than the numbers of species. Table 2 lists the number of reaction types at different temperatures, grouped by the number of occurrence. The numbers of reaction types are 571 at 2,500 K, 923 at 3,000 K, 1,360 at 3,500 K and 1,480 at 4,000 K. Again we see that the reaction type number at 3,500 K is close to that at 4,000 K. The majority of reaction types (~78 %) occurred only once and only a couple of them (<0.1 %) occurred more than 100 times. In addition, ~19 % of reaction types occurred from 2 to 10 times and a few (~3 %) occurred from 11 to 100 times. Many reactions are conjugated (paired) forward and backward reactions. If we count the conjugated reaction types only, the total numbers are reduced by less than a half to 318, 494, 784 and 925, respectively. Furthermore, we examined the reaction types and found that about a half of these reaction types are unimolecular decompositions, while the other half are bimolecular collisions. This ratio is about the same at different temperatures.

Tables 1 and 2 show the reactions are very complicated, and the number of species and reactions are much greater than that in typical energy minimization calculations. The large numbers of species and reaction types were caused by the non-equilibrium process in which new species were generated and eliminated constantly. For a complete trajectory, the numbers of species and reactions should be close to constant at a given temperature. For analysis of reaction mechanism, we need a complete trajectory from initial reactant to final products. Since the reactions are close to finish at 3,500 and 4,000 K, we take the data of 3,500 K for following analysis.

The reactant, transient intermediates and products can be identified by counting activity of each species in the trajectory. For each species, its activity is defined by two parameters: the number of reactions acting as reactant (R) and the number of reactions acting as product (P). If R > P, there is net gain of this species and it is a product. If R < P, there is a net loss and it is a reactant. If R = P, the species is an intermediate. There are much more intermediates than

 Table 2
 Numbers of reaction types found in the simulations at different temperatures, grouped by the numbers of occurrence

Count	2,500 K	3,000 K	3,500 K	4,000 K
1	415	716	1,059	1,196
2-10	137	181	267	258
11-100	17	25	32	24
>100	2	1	2	2
Total	571 (318)	923 (494)	1,360 (784)	1,480 (925)

Most of the reaction types are conjugated (forward and backward) reactions. The total numbers of conjugated reaction types are in the parenthesis

 Table 3
 The occurrence numbers of reactant and product molecules in the 3,500 K simulation

Formula	Molecular name	R	Р	Net
$\overline{C_2H_4N_4O_4}$	2,2-dinitro-1,1-ethenediamine	66	58	-8
$C_2H_2N_4O_2$	(See InChi code a)	1	2	1
C ₅ H ₃ N ₅ O ₃	(See InChI code b)	1	2	1
HO	Hydroxyl radical	252	253	1
H ₃ N	Ammonia	60	61	1
CHNO	Isocyanic acid	50	51	1
HNO_2	Nitrous acid	39	40	1
C_2N_2O	Cyanic isocyanate	23	24	1
CHN	Hydrocyanic acid	13	14	1
CH ₂ O ₃	Carbonic acid	11	12	1
N ₂ O	Dinitrogen monoxide	6	7	1
CH ₂ NO	Imidoformate	6	7	1
Н	Hydrogen radical	921	923	2
CO ₂	Carbon dioxide	32	35	3
NO	Nitric oxide	89	93	4
N ₂	Nitrogen	10	15	5
H ₂ O	Water	307	314	7

The occurrence numbers are listed for the species acting as reactant (R) and product (P) in reactions found in the simulation trajectory. The net number of gain or loss, calculated as the difference between R and P, indicates the species is a reactant (loss) or product (gain)

(a) InChI = $1S/C_2H_2N_4O_2/c3-1-2-4-6(8)5-7/h2,4H,$ (b) InChI = $1S/C_5H_3N_5O_3/c6-2-9-4(11)10-3(7)1-8-5(12)13/h1,10,12H$

products and reactant in the system. Table 3 lists the activity of products and reactant at 3,500 K. There was only one reactant ($C_2H_4N_4O_4$) FOX-7, which acted as a reactant for 66 times and as a product for 58 times in the simulation time, leading to the net loss of 8, as expected. The data also indicate that the initial decomposition of FOX-7 was a reversible reaction; it reversed for 58 times in the simulation time period. As shown in the table, all of the species were participants in reversible reactions and the difference is how active they were. Some were involved in reactions more frequently than others, indicating some molecules are easier to be activated.

Table 4 The occurrence number (Occur.) distribution of transient intermediates acting as reactants (R), products (P) and net gain (Net) in 3,500 K simulation

R	Р	Net	Occur.
1	1	0	406
2	2	0	117
3	3	0	56
4	4	0	27
5	5	0	20
6	6	0	12
7	7	0	8
8	8	0	7
9	9	0	8
10	10	0	3
>10	>10	0	44
Total			708

There was a total of 16 products at the end of 3,500 K simulation trajectory, ranked differently by the net populations in the last column. In addition to the main products $H_2O(7)$, $N_2(5)$ and $CO_2(3)$ (number in parenthesis denotes the population, and the same as follows), which can be derived from stoichiometry with assumption of complete reaction, we also obtained side products: NO(4), H(2), CH₂NO(1), N₂O(1), CH₂O₃(1), HCN(1), C₂N₂O(1), HONO(1), CHNO(1), NH₃(1), HO(1), C₂H₂N₄O₂(1) and C₅H₃N₅O₃(1). These side products overlap with those reported by laser and mass spectroscopic measurements [4], but cannot be predicted by energy minimization calculations [6].

The existence of H by the end of the reaction is not surprising because of the very high temperature (3,500 K). As shown in Table 3, H was very active; it acted 921 times as reactant and 923 times as product. Therefore, there is a good chance to see a couple of H left by the end of the simulation. The reaction types and products were also determined by the high pressure in the simulation system. Due to the fixed volume, the pressure in the simulation box was estimated to be in the order of 10^{3} - 10^{4} atm. The existence of two relatively large products, O=N-N(O)=N-CH₂-CN and HO-C(O)-N=CH-C(N)-NH-C(O)-N=C=N, in the list is due to the high pressure. Relevant to real explosion, the simulation condition represents a small region of the material and a very short time in which the reaction has started but the expansion has not yet. Once the expansion is allowed, the temperature and pressure will decrease, and the products will be different since the large fragments will decompose into smaller ones.

Subtracting one reactant and 16 products (Table 3) from the total number of 725 (Table 1), we obtained 708

 Table 5
 Average life time (ALT) of products and reactant in 3,500 K simulation

Formula	Molecular species	R	Р	Net	ALT (fs)
$C_2H_4N_4O_4$	2,2-dinitro-1,1-ethenedi- amine	66	58	-8	106
N_2	Nitrogen	10	15	5	1,151
NO	Nitric oxide	89	93	4	912
N ₂ O	Dinitrogen monoxide	6	7	1	560
CO_2	Carbon dioxide	32	35	3	542
C_2N_2O	Cyanic isocyanate	23	24	1	297
H ₂ O	Water	307	314	7	240
CHN	Hydrocyanic acid	13	14	1	208
H ₃ N	Ammonia	60	61	1	122
CHNO	Isocyanic acid	50	51	1	115
CH ₂ O ₃	Carboxylic acid	11	12	1	21
CH ₂ NO	Imidoformate	6	7	1	10
HNO ₂	Nitrous acid	39	40	1	29
$C_2H_2N_4O_2$	(See InChI code a)	1	2	1	7
$C_5H_3N_5O_3$	(See InChI code b)	1	2	1	7
Н	Hydrogen radical	921	923	2	111
OH	Hydroxyl radical	252	253	1	57

(a) InChI = $1S/C_2H_2N_4O_2/c^{3-1-2-4-6}(8)^{5-7/h^2,4H}$; (b) InChI = $1S/C_5H_3N_5O_3/c^{6-2-9-4}(11)^{10-3}(7)^{1-8-5}(12)^{13/h1,10,12H}$

transient intermediates. Table 4 shows the activity distribution of these intermediates. Most (57 %) intermediates acted as reactant and product only once. The number of intermediates with higher activity parameters is smaller. Only 44 (6.7 %) acted more than 10 times. However, it should be pointed out that the intermediate activity does not necessarily reflect its importance. In fact, some critical intermediates appeared only a couple of times in the reaction trajectories.

The average life time (ALT) of a species indicates its stability. For each species, its life time was calculated by subtracting its appearing time from its disappearing time and the ALT was calculated by averaging over multiple existences. Table 5 lists the ALT for products and reactant in femtosecond (fs). It is interesting to note that the ALT of the reactant FOX-7 is only 106 fs, which is similar to that of the hydrogen atom (ion) 111 fs, indicating FOX-7 is highly unstable at 3,500 K. The most stable one is N₂ (1,151 fs), as expected, other stable compounds are nitrogen oxides and carbon dioxide. An interesting point to make is that water (H₂O, 240 fs), hydrogen cyanide (HCN 208 fs) and ammonia (NH₃, 122 fs) are fairly active molecules at high temperature like 3,500 K.

In Table 6, we list the most stable intermediates with ALT longer than 102 fs. Many of these intermediates exhibit ALT longer than the products listed in Table 5. Similar to the discussion of activity parameters, the

Table 6Average life time (ALT) of intermediates in 3,500 K simulation

Formula	Molecular species		Р	Net	ALT (fs)
NO ₃	Nitric acid	14	14	0	613
СО	Carbon monoxide	5	5	0	369
H_4N	Ammonium	34	34	0	349
CH_2N_2	Cyanamide	45	45	0	300
C ₃ HN ₄ O	(a)	6	6	0	256
C ₃ N ₃ O	(b)	14	14	0	197
NO ₂	Nitrogen dioxide	68	68	0	191
C_2N_2	Oxalonitrile	2	2	0	186
C ₂ HNO ₂	(Hydroxyimino)ethenone	2	2	0	158
C ₂ HN ₂ O	Cyano(nitroso)methanide	3	3	0	143
CH_4N	Aminomethyl	9	9	0	137
CN	Cyanide	33	33	0	136
$CH_{3}N_{3}(c)$	(c)	9	9	0	133
$CH_4N_2O(d)$	(d)	24	24	0	133
$CH_3N_2O_3(e)$	(e)	11	11	0	129
$CH_3N_3O_2(f)$	(f)	5	5	0	129
$CH_2NO_2(g)$	(g)	36	36	0	125
$CH_3NO_2(h)$	(h)	42	42	0	124
CN ₂ O	Oxocyanamide	12	12	0	123
$C_2H_3N_2O_2(i)$	(i)	4	4	0	121
$C_2H_3N_2(j)$	(j)	12	12	0	116
$C_{3}H_{2}N_{2}O_{2}\left(k ight)$	(k)	3	3	0	115
CHN	Nitriliomethanide	8	8	0	111
HNO	Nitroxyl	27	27	0	110
$C_3H_3N_3O_2$	(1)	7	7	0	109
C ₂ HO	Ethynyloxidanyl	10	10	0	106
$C_2H_4N_2O$	Diaminoethenone	6	6	0	102

(a) $InChI = 1S/C_3HN_4O/c4-1-6-3(8)7-2-5/h6H$, (b) $InChI = 1S/C_3N_3O/c4-1-3(7)6-2-5$, (c) $InChI = 1S/CH_3N_3/c2-1-4-3/h2-4H$, (d) $InChI = 1S/CH_4N_2O/c2-1(3)4/h2-3H_2$, (e) $InChI = 1S/CH_3N_2O_3/c4-2-1-3(5)6/h1,5-6H$, (f) $InChI = 1S/CH_3N_3O_2/c2-1(5)3-4-6/h3H,2H_2$, (g) $InChI = 1S/CH_2NO_2/c2-1(3)4/h2-3H$, (h) $InChI = 1S/C_2H_3N_2O_2/c3-1(4)2(5)6/h3H,4H_2$, (i) $InChI = 1S/C_2H_3N_2/c3-1-2-4/h3H,1H_2$, (j) $InChI = 1S/C_3H_2N_2O_2/c4-1-2-5-3(6)7/h2,6H$, (k) $InChI = 1S/C_3H_3N_3O_2/c4-2-5-3(7)1-6-8/h5H,4H_2$, (l) $InChI = 1S/C_3H_3N_3O_2/c4-2-5-3(7)1-6-8/h5H,4H_2$

higher ALT does not equal to more importance; in fact, some critically important intermediates exhibit very short time. We list the intermediates with long ALT for the purpose that these species may be detected using experimental techniques. The transient intermediate species H, CN, OH, NO, C and N were detected before [4], but species like NO₃, NH₄⁺ and CH₂N₂ which are fairly stable based on our simulation data have not been reported by experimental work yet. We should make a note that one of the intermediates, NO₂, was treated as a product by the measurement [4].

3.3 Reaction network and mechanisms

The large amount of data generated in the simulation is very valuable for understanding the FOX-7 reaction mechanisms at extreme condition. However, in order to draw a clear picture on the reaction network, we should focus on specific purposes. We used a reverse mapping strategy to identify the reaction pathways from initial reactant, FOX-7 to the most common products N_2 , CO₂ and H₂O. Starting from the products, we used the atom sequence numbers (a unique number assigned to each atom) to find out all species that contain the same atoms in the trajectory. Then, we looked up the reaction list in a backward manner and found out reactions that connect those species. By doing so, we built up a simplified reaction network that describes how the most common products are formed, as shown in Fig. 4.

Reaction networks have been proposed previously based on experimental and quantum chemistry data [4, 6]. In those published papers, water is considered to be formed between OH and H, and CO₂ is formed by carbon oxidization step by step with reaction intermediates such as C=CO(NO₂), C=C=O, C-CH=O and CO₂. The network reported in this work is more comprehensive and unbiased; it gives a complete description of the formation of the three common products. We obtained the formations of H₂O and CO₂ in similar routes as previously suggested. However, the formation of N₂ has not been described before because it occurred in later stage and is much more complicated than those of H₂O and CO₂.

With the reaction network established, we outline the pathways as follows. In the following schemes (1-11), each species is labeled with a code and ALT for convenience in discussion. First of all, we identified three types of initial decompositions for reactant FOX-7. They are (1) C–NO₂ bond fission, (2) intermolecular hydrogen transfer and (3) intramolecular hydrogen transfer.

$$\begin{array}{c} H_2N \\ H_2N \\ H_2N \\ (X1, 106 \text{ fs}) \\ (X2, 7 \text{ fs}) \\ (X3, 191 \text{ fs}) \end{array} \xrightarrow{H_2N \\ H_2N \\ (X3, 191 \text{ fs}) \\ (X3, 191 \text{ fs}) \end{array}$$
(1)







Fig. 4 Reaction map from reactant FOX-7 to main products H₂O, CO₂ and N₂

The occurrence times of the three initial decomposition types are 4, 3 and 1, respectively (for 8 FOX-7 molecules). Due to small sizes and short simulation times, the occurrence times are associated with large uncertainties. However, analysis shows that all of the three types of initial reactions were observed in simulations at different temperatures. The occurrence percentage of the three initial decomposition types are 45, 33 and 22 % at 2,500 K, 33, 50 and 17 % at 3,000 K, 50, 13 and 17 % at 3,500 K, and 50, 33 and 17 % at 4,000 K. The most popular route is through the C-NO₂ bond fission, as indicated in previous energy minimization calculations [5]. This reaction generates an intermediate, 2-nitroethene-1,1-diamine (X2, 7 fs) and NO₂ (X3, 191 fs), which explains why NO₂ is the first intermediate generated in the reactions as we discussed above (Fig. 1). We did not observe nitro-to-nitrite rearrangement as initial reaction in the trajectory. The important finding here is that both intramolecular hydrogen transfer (Scheme 2) and intermolecular hydrogen transfer (Scheme 3) were found to be feasible, contradictory to the previous energy minimization calculation results [8]. This shows that the significant differences between energy minimization calculations and dynamic simulations. With many atoms, the thermal fluctuation is a driving force for the reactions, but it is not feasible by energy minimization calculations. It should be noted that this study is aimed to discover unknown reaction pathways that are beyond what can be postulated based on chemical intuition and to prepare reaction information for developing reactive force field, which can be used to get quantitative prediction of reaction mechanisms, kinetics with enhanced sampling techniques, large simulation size and time scale.

As a matter of fact, the hydrogen transfer reactions (1) and (2) are very important as they produced two highly active intermediates (X4, 22 fs) and (X6, 14 fs), in which the C–N bonds are activated and will be broken rapidly to generate nitrosyl hydroxide (X7, 29 fs) and another intermediate. From (X6, 14 fs), it produces:

$$\begin{array}{c} H_{2}N \\ H_{2}N \\ H_{2}N \\ H_{2}N \\ O \\ (X6, 14 \text{ fs}) \end{array} \xrightarrow{NO_{2}} H_{2}N \\ H_{2}N \\ (X2, 7 \text{ fs}) \\ (X7, 29 \text{ fs}) \end{array}$$
(4)

In addition, nitrous acid, HONO (X7, 29 fs), can be generated from NO_2 (X3, 191 fs) by accepting a proton from other intermediate. Nitrous acid is reactive (29 fs), and it rapidly decomposes to NO and OH radicals:

HO-NO
$$\longrightarrow$$
 OH + NO
(X7, 29 fs) (X8, 57 fs) (X9, 912 fs) (5)

OH radical reacts with proton to produce one of the final products, H_2O . NO is relative stable, but it can react with other species as shown in scheme (9) below.

The intermediate 2-nitroethene-1,1-diamine (X2, 7 fs) is very reactive and may react with a proton to form NH_3 and intermediate (X10, 42 fs):

In addition, (X2, 7 fs) undergoes C–NO₂ isomerization through a three-member-ring transition state (X12, 8 fs), in which the first C–O bond starts to form. (X12, 8 fs) is then converted to the intermediate (X13, 5 fs), in which a carbonyl (C=O) bond is formed. Afterward, (X13, 5 fs) loses a NO group to generate a relatively stable intermediate (X14, 99 fs).



(X14, 99 fs) is oxidized by NO₂ to form the intermediate (X15, 69 fs), which undergoes the C–C bond fission to produce the intermediate (X16, 6 fs) and another final product (CO₂, 541 fs). (X16, 6 fs) is very reactive and quickly releases two hydrogen atoms to form carbodiimide (X17, 68 fs).



Carbodiimide (X17, 68 fs) is a reactive molecule widely used in the direct conjugation of carboxylic acid to primary amine for organic synthesis. In the simulated trajectory, carbodiimide (X17) reacts with NO to generate the first N–N bond:

HN=C=NH + NO
$$\longrightarrow$$
 NO (9)
(X17, 68 fs) (X9, 912 fs) (X19, 20 fs)

The intermediate (X19, 20 fs) reacts with a reducing agent denoted by R to form the intermediate (X20, 25 fs), which generates the final products N_2 (X21, 1,150 fs) and HCN (X22, 207 fs) by internal hydrogen transfer and C–N bond fission.

$$N \equiv C - NH \xrightarrow{R} N \equiv C - N \equiv NH \longrightarrow N \equiv N + H - C \equiv N$$
(X19, 20 fs)
(X20, 25 fs)
(X21, 1150 fs) (X22, 207 fs)
(10)

(X19, 20 fs) has another route in which the C–N bond fission occurs before the reduction. The C–N bond fission and internal hydrogen transfer generates HCN (X22, 207 fs) and N₂O (X23, 560 fs), which is then reduced to form N₂ (X21, 1,150 fs).



4 Conclusion

In order to obtain the thermal decomposition mechanism of solid FOX-7 at high temperature, we carried out NVT ensemble AIMD simulations at 2,500, 3,000, 3,500 and 4,000 K. The simulated reactions progress in different extents as the reaction rate is directly related with temperature. The simulation results show that reactions at the end of 3 ps simulation are close to completion at 3,500 and 4,000 K while they are still in the middle way at 2,500 and 3,000 K. Taking data obtained at different temperatures, we obtained a whole picture of the decomposition process for FOX-7 under extreme condition.

The simulation data demonstrates that the reactions in high-pressure gas phase are very complex. There are hundreds of unit reactions and about the same number of transient intermediates. Unlike reactions in dilute solution or low-pressure gas phase in which reaction pathways consist few unit reactions, the complex reaction pathways in high-pressure gas phase are too complicated to be made solely based on chemical intuitions. It is necessary to use molecular dynamics method to study this type of reactions. Similar problems exist for reactions in high-concentration solutions, such as the condensation reactions in polymers and zeolite syntheses.

One of the main findings of this work is about the initial decomposition of FOX-7. The C–NO₂ bond fission mechanism [5] is found to be the most common route. However, the inter- and intramolecular hydrogen transfers which have been excluded by previous energy minimization calculations [8] are found to be plausible. The most popular initial decomposition is the C–NO₂ bond fission process, followed by the intramolecular hydrogen transfer, and the least popular one is the intermolecular hydrogen transfer. The nitro-to-nitrite rearrangement suggested by previous energy minimization calculation [6] is not observed in the simulated data. These findings demonstrate that the reactions in ideal gases are dramatically different from those in highly condensed phases.

The simulation results agree well with known experimental data [4]. Using the multiple temperature data, the activation energy of initial decomposition was estimated and the result is in agreement with the experimental value [19]. The molecules identified in the simulation process agree with experimental observation. It is important to note that, in addition to the main products, which can be deduced from stoichiometry, we found more than a dozen of side products. Among those side products, HCN, NO, NH₃, H₂O, HCN, HONO, NO₂ are identified in the experiment [4], which is pretty encouraging result. The quantitative agreement has not been achieved because the simulations condition is still different from the real experimental considering the system size and reaction time scale.

In order to capture reaction details, we collected trajectory at an interval of 0.5 fs, which is short enough to catch very fast reactions. The data set was too large to be analyzed by hand. With computer software, we carried out statistical analysis of the data and tracked out how the main products are generated. For the first time, we were able to map out a complete reaction network for H_2O , CO_2 and N_2 formations in the thermal decomposition of FOX-7. Although the network may not be the same for all products, the investigation method is generally applicable. One can use this method to track back any product to find out how it is produced, and then possibly how to promote or demote its production. This would be very useful for targeted synthesize or for environment protection.

There is still large room for future improvement. The small simulation size, short simulation time and high

simulation temperature are obviously problems to be addressed in future work. One of the possibilities is to develop RFF based on AIMD simulation result obtained from this study. If RFF represents the reaction potential energy surface well, MD simulations with RFF can be carried out with much larger (in 2–3 orders of magnitude higher) size and time scales than what can be done using AIMD. The quantitative results with smaller fluctuation can be anticipated from large-scale MD simulation with RFF and high-performance supercomputer.

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REGULAR ARTICLE

Hemibond complexes between H₂S and free radicals (F, Cl, Br, and OH)

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Abstract The interaction of hydrogen sulfide (H_2S) with F, Cl, Br, and OH is investigated using ab initio methods to identify the two-center three-electron hemibond responsible for their complexation. The binding energies are found to be stronger than those in the analogous water complexes, but follow the same trend of increasing strength: F > Cl > Br > OH. The radicals are located nearly perpendicular to the H_2S plane forming an angle of about 90°. Analysis of molecular orbitals and natural bond orbitals are carried out to understand the energetics, structures, and bonding characteristics of these hemibonded complexes.

Keywords Hemibond · Radical · Ab initio

1 Introduction

Hydrogen sulfide (H_2S) is present in many atmospheric, interstellar, and industrial environments. In Earth's atmosphere, for example, H_2S released from biota, volcanoes, and industrial processes may contribute to climate changes [1]. This molecule is also known to react with free radicals such as OH in atmosphere [2] and in combustion [3]. Such reactions typically result in the abstraction of a hydrogen from H_2S by the radical, and these processes are activated with a barrier. Very recently, we have systematically studied similar reactions involving H_2O and free radicals such

Dedicated to Professor Guosen Yan and published as part of the special collection of articles celebrating his 85th birthday.

B. Alday · R. Johnson · J. Li · H. Guo (⊠) Department of Chemistry and Chemical Biology, University of New Mexico, Albuquerque, NM 87131, USA e-mail: hguo@unm.edu as F [4–8] and Cl [9] and found that there often exists a pre-reaction complex between the two reactants. The existence of such complexes has also been reported in other ab initio studies [10–13]. Interestingly, the F…H₂O complex has been shown to aid the F + H₂O reaction by guiding the trajectories to the transition state, resulting in a significant enhancement of the reactivity, particularly at low collision energies [14]. More interestingly, it was demonstrated that such complexes are not van der Waals or electrostatic in nature, but due to covalent interactions based on a twocenter three-electron bond [15].

Such a two-center three-electron bond was first discussed by Pauling [16]. Due to the fact that one electron is in the antibonding orbital, such a covalent bond is usually quite weak with a relatively long bond length and hence called a hemibond [17]. Hemibond bonding has been found in many branches of chemistry [18], particularly between molecules with lone pairs of electrons and a free radical with an unpaired electron [19-31]. As mentioned above, such hemibond complexes are not only of a fundamental interest, but might also play an important role in molecule-radical reactions. Like oxygen-containing molecules, sulfur compounds may also provide lone electron pairs on S, thus allowing the formation of hemibonds with free radicals. In the literature, however, most such complexes investigated before are charged [17, 19, 21, 22, 24–26], which complicates the bonding with electrostatic interactions. To further understand the sulfur-containing hemibond complexes, we report here an extensive highlevel ab initio and density functional theory (DFT) study of the complexes formed by H_2S and several free radicals. The major aim is to understand the similarities and differences between the H₂O…X and H₂S…X complexes with X = F, Cl, Br, and OH by examining the structural, energetic, and electronic aspects of these interesting systems.

For the $H_2S\cdots OH$ complex, we also investigate the competing hydrogen bond complex.

2 Theory

Most of the calculations were carried out at an F12 version of the unrestricted coupled cluster method with singles, doubles, and perturbative triples based on restricted open-shell Hartree-Fock reference wave functions (ROHF-UCCSD(T)-F12a) [32, 33]. The 1s, 2s, and 2p core electrons in S were frozen, similar to the work done in the analogous water systems [15]. The F12a method converges well to the complete basis set (CBS) limit with relatively small basis sets; so, the augmented correlation-consistent polarized valence triple zeta (aug-cc-pVTZ or AVTZ) basis set [34] was used. The minimum energy structures of both the complex and monomers were obtained by optimization at the UCCSD(T)-F12a level. The binding energy $(E_{\rm b})$ of the complexes was computed directly from the difference of the complex and separated monomers. This direct approach is justified as the basis set superposition error (BSSE) was examined for the F...H2S complex and was found to be quite small (less than 0.02 cm^{-1}). The calculations were performed using MOLPRO 2010.1 [35].

Additional calculations have been carried out using the spin-restricted open-shell density functional theory (RO-DFT) with the modified Perdew–Wang 1-parameter



Fig. 1 Geometries and binding energies for the $H_2S \cdots X$ (X = F, Cl, Br, and OH) complexes obtained at the UCCSD(T)-F12a/AVTZ level of theory

functional for kinetics (MPW1K) suggested by Truhlar and coworkers [36], which has been shown to yield comparable energetics for the F····H₂O complex with the UCCSD(T)/ cc-pV5Z level of theory [11]. The same AVTZ basis set was used. The DFT calculations are used as a diagnostic tool to understand the bonding nature. Molecular orbitals within the DFT treatment were also used in the natural bond orbital (NBO) analysis [37], which allows the description of the bonding and lone pair orbitals within the intuitive Lewis structure framework. These molecular orbital calculations were carried out using Gaussian 09 [38].

3 Results and discussion

All $H_2S \cdots X$ complexes were found to have the C_s symmetry, with the mirror plane bisecting the H-S-H bond. The relative position of the two moieties is defined by θ (the angle between the X-S line and H₂S plane) and R (the distance between X (O in the case of OH) and S) as indicated in Fig. 1. For the H_2S ...OH complex, an additional bond length for the OH and the S-O-H angle are needed. A summary of the geometries of the complexes formed between H₂S and X = F, Cl, Br, and OH is given in Table 1. For comparison, the equilibrium geometries of the isolated H₂S and OH are also included for comparison. In addition, the optimized DFT geometries are also given for comparison. Interestingly, the intramolecular H-S and O-H bonds are only slightly elongated in the complex, and the bonding angle remains roughly same as the H₂S and OH monomers. This observation suggests that the complexation has an insignificant effect on the H_2S monomer as well as OH in the $H_2S\cdots OH$ complex.

It is interesting to note that the X–S bond lengths are quite long, suggesting a relatively weak interaction characterizing the hemibonds. The long bond lengths are consistent with previous theoretical results for hemibond complexes involving sulfur [17, 19, 21, 22, 24–26]. For example, the S–O distance in the H₂S···OH complex is 2.62 Å, which can be

Table 1 Geometric parameters of H_2S and its complexes with various free radicals obtained at the UCCSD(T)-F12a/AVTZ level and at the MPW1K/AVTZ level (in parentheses)

	R _{SH1}	R _{SH2}	R	R _{OH3}	$\theta_{\rm H1SH2}$	θ	ϕ
H ₂ S (Expt.) [40]	1.3356	1.3356			92.12		
H ₂ S	1.325 (1.336)	1.325 (1.336)			92.9 (92.7)		
OH				0.970 (0.963)			
$H_2S\cdots F$	1.338 (1.335)	1.338 (1.335)	1.948 (1.993)		94.4 (94.1)	87.3 (87.3)	
H ₂ S…Cl	1.340 (1.336)	1.340 (1.336)	2.580 (2.591)		92.8 (93.1)	87.8 (87.8)	
H ₂ S…Br	1.340 (1.336)	1.340 (1.336)	2.802 (2.788)		92.6 (93.0)	87.9 (87.9)	
H ₂ S…OH	1.339 (1.335)	1.339 (1.335)	2.621 (2.825)	0.969 (0.961)	92.6 (92.8)	68.0 (72.3)	87.6 (81.6)
H ₂ S…HO	(1.343)	(1.343)	(3.569)	(0.979)	(92.7)	(158.8)	(16.6)

The intermolecular geometric parameters are defined in Fig. 1. Bond length in angstroms and angle in degrees



Fig. 2 Comparison of the binding energies and structures $H_2S\cdots X$ and $H_2O\cdots X$ (X = F, Cl, Br, and OH) complexes obtained at the UCCSD(T)-F12a/AVTZ level of theory

 Table 2
 Comparison of binding energies obtained at the UCCSD(T)-F12a/AVTZ level and at the MPW1K/AVTZ level (in parentheses)

Complex	<i>E</i> _b (kcal/mol) hemibond	$E_{\rm b}$ (kcal/mol) hydrogen bond
H ₂ S…F	-21.44 (-16.87)	_
H_2S ····Cl	-9.76 (-11.59)	_
H ₂ S…Br	-6.96 (-8.82)	_
H ₂ S…OH	-2.71 (-2.07)	-2.66* (-1.11)

* Single point CCSD(T)-F12a result from DFT geometry

compared with that in the NCS⁻···OH complex (2.179 Å) [26]. (The shorter distance in the latter complex is presumably due to the additional charge–dipole interaction.) In addition, the free radical typically bonds above (or equivalently below) the molecular plane of H₂S, which is analogous to the hemibond complexes involving H₂O [15]. However, the angle θ is somewhat different for the two types of complexes. While the angle θ is larger than 90° in the H₂O···X (X = Cl, Br) complexes, the corresponding angle in the H₂S···X complexes is near 90°. Similar right-angled hemibond complexes have been reported before between sulfur-containing molecules and free radicals [17, 19, 21, 22, 24–26].

As shown in Fig. 2 and Table 2, binding energies of the complexes are in general much stronger than that in the corresponding water complexes. Among them, the H₂S…F complex is the strongest, with a large binding energy of 21.44 kcal/mol. The binding energy decreases in the order $F \gg Cl > Br > OH$. These binding energies are consistent with those found previously for hemibond complexes involving sulfur [19, 22, 24–26].

In Table 3, the harmonic vibrational frequencies of these complexes are listed along with those of the monomers.

Fable 3 Harmon	vic vibrational frequen	rcies (cm^{-1}) for monon	ners and complexes at t	he UCCSD(T)-F1	2a/AVTZ level an	d at the MPW1K/	AVTZ level (in par	entheses)	
	HSH bend	S-H symmtrical. stretch	S-H asymmtrical. stretch	S-X stretch	H ₂ S–X wagging	H ₂ S–X bending	S-O twisting	OH wagging	O-H stretch
OH expt. [40]									3,737.76
H ₂ S expt. [40]	1,183	2,626	2,615						
H_2S	1,212.3 (1,231.3)	2,719.4 (2,782.5)	2,735.3 (2,796.3)						
H_2SF	$1,181.1\ (1,208.6)$	2,715.0 (2,787.4)	2,740.6 (2,804.0)	435.9 (347.4)	552.8 (461.6)	609.7 (539.3)			
H ₂ S…Cl	1,205.1 (1,229.9)	2,714.31 (2,756.5)	2,730.8 (2,769.3)	237.6 (242.0)	366.0 (384.9)	396.6 (426.8)			
H ₂ S…Br	1,205.5 (1,227.4)	2,711.08 (2,783.1)	2,726.1 (2,797.1)	164.7 (175.0)	306.4 (321.1)	332.5 (346.8)			
H ₂ S…OH	$1,206.2\ (1,232.0)$	2,739.9 (2,765.5)	2,756.9 (2,777.8)	78.0 (132.5)	149.0 (257.8)	251.8 (316.9)	284.6 (397.2)	504.9 (606.1)	3,765.6 (3,787.6)

Fig. 3 Molecular orbital energy level diagram for $H_2S \cdots X$ (X = F, Cl, Br, OH) complexes with corresponding MO wavefunctions obtained at the MPW1K/aug-cc-pVTZ level of theory



Again, the complexation changes the monomer frequencies very little. The vibrational frequencies between the monomers in the complex are typically low, consistent with the weak interactions in the hemibonded complexes. Similar conclusions can be formed from the DFT frequencies also listed in the Table 3.

To understand these structural and energetic trends, we resort to the molecular orbital (MO) analysis using the DFT method. In Fig. 3, MO energy level diagrams for the H₂S…X complexes are displayed. The two major monomer MO wavefunctions that contribute to the hemibond MO wavefunctions in the complexes are displayed with isosurface plots. The resulting electronic structures clearly indicate the formation of a two-center three-electron hemibond. It is also clear that the magnitude of the binding energy can be attributed to the energy difference between the highest occupied molecular orbital (HOMO) of S (i.e., a lone pair orbital) and the singly occupied molecular orbital (SOMO) of the radical. Since the hemibond strength is controlled by the stabilization energy of the bonding orbital between the two, the smaller this HOMO-SOMO gap, the greater the stabilization of the resulting bonding MO, the stronger the bond. The energy difference is the smallest for the H₂S…F complex, resulting in the strongest binding energy. The energy gap increases in the order F < Cl < Br < OH. The larger binding energies in the H₂S…X complexes than those in the H₂O···X complexes can be largely attributed to the higher energy of the H_2S HOMO (-0.312 au) than that of H₂O (-0.384 au) [15].

The geometry of the complex can also be understood in terms of the hemibond as well. For the $H_2O\cdots X$ complexes, it was demonstrated [15] that the geometry of the complexes is largely dictated by the overlap between the radical SOMO



Fig. 4 Images of the lone pair orbitals from NBO and corresponding MOs of $\mathrm{H_2S}$

and molecular HOMO. In H₂S, there are two lone pair nonbonding MOs in S. One is perpendicular to the molecular plane while the other is within the molecular plane, as shown in Fig. 4 by both the MO and NBO wavefunctions. They are the HOMO (E = -0.312 a.u.) and HOMO-1 (E = -0.419 a.u.), respectively. As discussed above, the radical SOMO interacts primarily with the H₂S HOMO, which is perpendicular to the H₂S plane. As a result, the complexes are all formed with an angle θ close to 90°.

For the H_2S ...OH complex, there is another configuration in which the H of the OH moiety points toward S. This isomer of the complex is apparently due to the dipole– dipole or hydrogen bond interaction. Although this hydrogen-bonded complex was identified by DFT optimization, we were unable to converge the CCSD(T)-F12 calculation for this complex, presumably due to flatness of the potential energy surface near the minimum. The DFT geometric parameters of this complex are also included in Table 1. Its binding energy is -1.11 kcal/mol at the DFT level. The single point CCSD(T)-F12 calculation at the DFT geometry gives a binding energy of -2.66 kcal/mol. These results suggest that the hemibond complex is slightly more stable than the hydrogen bond complex. There have been extensive discussions in the literature on the relative stability of the similar isomers in the complexes formed between H₂O and OH, which also have comparable binding energies [29, 31].

Finally, we note in Tables 1 and 2 that DFT values are qualitatively similar to those obtained at the UCCSD(T)-F12/AVTZ level, although there are quantitative differences. This is in further support of the covalent nature of these complexes as DFT is considered incapable of handling dispersion and other weak interactions. This observation is consistent with the recent work that DFT is capable of describing hemibonds [31, 39], although such a description is not expected to be quantitative. This is important as it lends credence to DFT based simulations in hemibond systems.

4 Conclusions

In summary, our high-level ab initio calculations predict that H_2S forms a strong hemibond with a free radical such as F, Cl, Br, and OH. These hemibond complexes have much stronger binding energies than those involving H_2O , thanks to good energy match between the SOMO of the radical and lone pair HOMO of H_2S , which is perpendicular to the H_2S molecular plane. The complexes formed between H_2S and the radicals assume a C_s geometry where the radical is above (or below) the molecular plane immediately above the S atom, which can again be attributed to the SOMO–HOMO interaction. These interesting properties of the hemibond complexes will have an important impact on our future investigations of the reactions between H_2S and free radicals.

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REGULAR ARTICLE

DFT and TD-DFT studies on osmacycle dyes with tunable photoelectronic properties for solar cells

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Abstract B3LYP and CAM-B3LYP functionals have been used to determine structures, electronic and optical properties of osmium-bridged tricyclic aromatic compounds. Calculations show that the optical properties and charge separation features of these osmacycle derivatives can be well modified by incorporating different π -bridge groups. In particular, the newly designed osmacycle dyes 5 and 6 by embedding thiophene and thienothiophene bridge units to osmium polycyclic aromatic system show very strong and broad adsorptions in the whole visible region and excellent charge separation in the first excited state of $(\pi\pi^*)$ from the hightest occupied molecular orbital to the lowest unoccupied molecular orbital excitation. Furthermore, the predicted relatively high light harvesting efficiency and large driving force for electron injection suggest that they are quite promising for design of high-performance dye-sensitized solar cells.

Keywords Metal-bridged tricyclic aromatic systems · Density functional calculations · Osmacycle dyes · Structural modification · Absorption spectra · Electron injection

Dedicated to Professor Guosen Yan and published as part of the special collection of articles celebrating his 85th birthday.

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1 Introduction

Solar power plays an important role in turning energy systems toward sustainability. Among the photovoltaic devices, the dye-sensitized solar cell (DSSC) is the only solar device that mimics the photosynthesis process in a green leaf. Owing to its simplicity of manufacturing process, low cost, and relatively high energy conversion efficiency, DSSCs become the promising one of renewable energy devices [1, 2]. In general, DSSC is composed of five components, including conductive mechanical support, semiconductor film, sensitizer, electrolyte containing I^-/I_3^- redox couple and counter electrode with a trioxide reduction catalyst [3–6]. As one of the crucial parts in DSSCs, the photosensitizer to harvest sunlight has been viewed as a breakthrough for performance improvement of DSSCs.

Since O'Regan and Grätzel reported the first successful DSSC based on the mechanism of a fast regenerative photoelectrochemical process in 1991, the design of high-performance sensitizers has attracted considerable attention, both experimentally and theoretically [3–20, 29, 41, 46–48], in order to enhance the power conversion efficiency (PCE) of DCCSs. High-efficient dyes commonly should fulfill some essential requirements, including the broad absorption ranging from the whole visible region to the near-infrared part, the novel anchoring groups with strong adsorption on the semiconductor surface, the adaptation of corresponding energy levels among the conduction edge of semiconductor, the excited-state and oxidized-state levels of dyes, and the redox potential of electrolyte, as well as high electrochemical and thermal stability.

At present, among widely used metal–organic and metal-free organic sensitizers for DSSCs [7], the metalfree organic dyes such as coumarin, indoline, cyanine, triphenylamine, and porphyrin have attracted considerable interest for renewable energy devices because of their significant advantages in terms of synthesis procedures, diversiform structures, and low costs [8–12]. However, the PCE for the metal–organic dyes has been recently climbed to 12.3 % [13, 14]. Among the numerous metal–organic dyes, ruthenium dyes, such as Ru bipyridine complex (red dye) [15] and Ru terpyridine complex (black dye) [16], have been extensively investigated [17, 18], and at the same time, the Os terpyridine complex was also examined for comparison. As a consequence, the osmium compounds exhibit extremely broad UV–Vis absorption, perfectly reversible electrochemistry, and prominent photochemical stability, although they suffer from lower V_{oc} or J_{sc} values under different conditions [19, 20].

Very recently, a series of novel metal-bridged polycyclic aromatic compounds have been successfully synthesized, and they show outstanding stability and unusual optical property [21–23]. To have an insight into their photoelectronic properties and possible applications in DSSCs, herein extensive density functional calculations have been performed. Based on calculations, electronic and structural properties, as well as structural modification effects on charge separation and light harvesting ability of the osmium metallacycle dyes, have been discussed.

2 Computational details

2.1 Theoretical background

It is well known that the performance of DSSCs can be characterized by the overall power conversion efficiency (η) and the monochromatic incident photo-to-electron power conversion efficiency (IPCE). The power conversion efficiency is generally defined as the ratio of the maximum solar power, P_{max} , and the incident solar power on the cell, P_{inc} , and it can be obtained by the following equation [5, 24]:

$$\eta = \frac{P_{\max}}{P_{inc}} = \frac{I_{\max} \cdot V_{\max}}{P_{inc}} = \frac{J_{\max} \cdot V_{\max} \cdot S}{P_{inc}}$$
$$= \frac{FF \cdot J_{sc} \cdot V_{oc} \cdot S}{P_{inc}} = \frac{FF \cdot J_{sc} \cdot V_{oc}}{I_{s}}$$
(1)

where $J_{\rm sc}$ is the short-circuit current density, $V_{\rm oc}$ is the open-circuit photovoltage, FF is the fill factor of the cell, $I_{\rm s}$ is the intensity of the incident light, and $J_{\rm max}$ and $V_{\rm max}$, respectively, are the photocurrent density and photovoltage under the maximum solar power. The fill factor ranging from 0 to 1 is the ration of the maximum power ($P_{\rm max}$) and the product of the open-circuit voltage ($V_{\rm oc}$) and the short-circuit current ($I_{\rm sc}$), exhibiting the extent to which the power is lost due to ohmic resistance and overpotential. According to Eq. 1, manipulation of $J_{\rm sc}$ and $V_{\rm oc}$ is an effective way to enhancement of η .

The photocurrent density, J_{sc} , is defined by integrating IPCE(λ) over the solar spectral density, $I_s(\lambda)$:

$$J_{\rm sc} = e \int \mathrm{IPCE}(\lambda) I_{\rm s}(\lambda) \mathrm{d}\lambda \tag{2}$$

$$IPCE(\lambda) = \frac{n_e}{n_p} = LHE(\lambda) \times \Phi_{inject} \times \eta_{collect} \times \eta_{reg}$$
(3)

where LHE(λ) is the light harvesting efficiency at a given wavelength, Φ_{inject} is the electron injection efficiency, $\eta_{collect}$ is the charge collection efficiency at the conducting glass substrate, and η_{reg} is the dye regeneration efficiency, and therein LHE(λ) and Φ_{inject} affect prominently on IPCE(λ). LHE(λ) can be expressed as:

$$LHE(\lambda) = 1 - 10^{-f} \tag{4}$$

where A(f) is the absorption (oscillator strength) of the dye at the maximum wavelength [25], and the oscillator strength associated with the λ_{\max}^{ICT} can be written as:

$$f = \frac{2}{3} \lambda_{\text{max}}^{\text{ICT}} |\vec{\mu}_0 - \text{ICT}|^2 \tag{5}$$

Hence, *f* is determined by the maximum wavelength $\lambda_{\text{max}}^{\text{ICT}}$ (ICT: intramolecular charge transfer) and the dipolar transition moment $|\vec{\mu}_0 - \text{ICT}|$ associated with the electronic excitation [26, 27].

The electron injection efficiency Φ_{inject} is in direct proportion to the driving force (ΔG_{inject}) of the electron injection from the photoinduced excited states of sensitizer to semiconductor surface, and it can be expressed as (in eV) [28]:

$$\Delta G_{\rm inject} = E_{\rm ox}^{\rm dye*} - E_{\rm CB}^{\rm TiO_2} \tag{6}$$

where $E_{\text{ox}}^{\text{dye*}}$ is the oxidation potential of the dye in the excited state, and $E_{\text{CB}}^{\text{TiO}_2}$ is the bottom of conduction band of titanium dioxide whose experimental value is considered as -4.0 eV [29]. As for $E_{\text{ox}}^{\text{dye*}}$, it was assumed that the electron injection occurs from the unrelaxed excited state because the primary electron transfer event in DSSCs is much faster than the vibrational relaxation of the photoexcited dyes [28], and $E_{\text{ox}}^{\text{dye*}}$ can be computed as:

$$E_{\rm ox}^{\rm dye*} = E_{\rm ox}^{\rm dye} - \lambda_{\rm max} \tag{7}$$

 λ_{max} is the vertical transition energy which is the approximation of the 0–0 transition energy E_{0-0} [28, 30], and $E_{\text{ox}}^{\text{dye}}$ is the redox potential of the ground state of the dye which can be estimated by the Nernst equation and can also be written in the energy scale as Eq. 8.

$$E_{\rm ox}^{\rm dye} = \frac{\Delta G_{\rm (aq)}}{nF} \tag{8}$$

The value of $\Delta G_{(aq)}$ is Gibbs free energy change due to the oxidation of the dye in solution, and it can be



Scheme 1 Thermodynamic cycle for calculation of redox potentials

calculated according to the Born–Haber cycle, as shown in Scheme 1, where *F* represents the Faraday constant of 96,485.3 C mol⁻¹, and *n* is the number of electrons involved in the redox couple [dye] $^{2+}/[dye]^+$ (here n = 1) [31, 32].

2.2 Computational methods

Here, theoretical calculations of geometries, frequencies, electronic structures, as well as electronic absorption spectra for the osmapentalyne derivatives have been performed by the B3LYP functional [33, 34] with the basis sets of LANL2DZ for Os and P atoms and the 6-31G(d) basis set for other atoms implemented in Gaussian09 package [35]. Polarization functions of P ($\zeta(d) = 0.340$), S($\zeta(d) = 0.421$), and Os (ζ (d) = 0.886) [36] are also added to the LANL2DZ basis set. The solvent effects of dichloromethane used in experiment have been considered using the SMD continuum solvation model [37]. Previous time-dependent density functional theory (TD-DFT) assessment suggested that the hybrid functionals of B3LYP, CAM-B3LYP, M06-2X, and PBE0 are suitable for the optical 0-0 transition of dyes in solution [30]. The coulomb-attenuating method CAM-B3LYP functional taking long-range correction into account has been proved to be appropriate for electronic excitations with character of charge transfer (CT), but the conventional exchange-correlation functional such as B3LYP is also practicable for some complexes [38-42]. Herein TD-DFT calculations with both CAM-B3LYP and B3LYP functionals were used for determination of the vertical excitation energies and corresponding oscillator strengths. In order to reduce computational costs, the triphenylphosphine ligand in the experimental complex is replaced by the phosphine (PH₃), and test calculations indicate that such simplification of ligands has only negligible influences on the structural and electronic properties of the relatively large metallacycles.

3 Results and discussion

3.1 Structural and optical properties of osmium polycyclic aromatic complexes

The geometry and absorption spectrum of the osmiumbridged tricyclic aromatic compound **1** were determined by B3LYP calculations. The optimized structure and selected bond lengths are shown in Fig. S1 and Table S1 in Supporting Information). As shown in Table S1, the B3LYPpredicted bond lengths of complex 1 with the 6-31G(d) basis set are similar with those obtained with the relatively large 6-311++G(d,p) basis set, and they are all consistent with the experimental values. Furthermore, the B3LYP/6-31G(d)-predicted strong absorptions at 350, 378, 421, and 549 nm can reasonably match the experimental bands at 353, 367, 407, and 557 nm, respectively, as shown in Fig. S2. Accordingly, the B3LYP functional with the 6-31G(d) basis set is practicable for this kind of osmium polycyclic aromatic systems.

Among the most recently synthesized osmium polycyclic aromatic systems [23], the complex with an oxygen heterocycle (2' in Fig. 1) was found to have strong and broad adsorptions in almost whole visible region, suggesting that it may be of high potential for the photosensitizer in solar cells, and this osmium tricyclic aromatics 2' served as a parent was considered for subsequently computational design of osmacycle dyes.

Here, DFT calculations show that the osmacycle complex 2 from substitution of PH_3 for PPh_3 groups in 2' has the singlet ground state, like 2'. As shown in Fig. 1 and Table 1, both complexes 2 and 2' have quite similar geometries, lending support to simplification of PPh₃ ligands in calculation. In particular, all predicted Os-C bond lengths in 2 (2.062-2.190 Å) as well as 2' (2.062–2.208 Å) are within the range of Os-C single- and double-bond lengths, in which Os1-C27 (2.190 Å), Os1-C7 (2.098 Å), and Os1-C12 (2.135 Å) bear more single-bond character, while Os1-C16 (2.062 Å) has partial character of double bond [43]. Their corresponding Wiberg bond indices of 2 are 0.761, 0.894, 0.779, and 0.912, respectively. The natural charge populations at atoms Os1, C7, C12, C16, and C27 are -1.337, -0.180, -0.115, 0.432, and -0.118, respectively. Therein the relatively large natural positive charges at C16 may stem from the adjoining oxygen with large electronegativity. The dihedral angles of C27-C16-C7-C10, C22-Os1-C13-C9 are 177.6° and 178.2°, respectively, and thus complex 2 has an approximately planar conformation.

The canonical molecular orbital nucleus-independent chemical shift (CMO-NICS) analyses [44] indicate that NICS(1) values at the center of A, B, and C rings (see Scheme S1 and Table S2) are -4.3, -6.1, and -9.0 ppm, respectively. Such negative NICS values reveal the presence of notable aromaticity in this metallacycle from extending π conjugation interaction, which may play an important role in stabilizing metal-bridged tricyclic aromatic complexes.

The hightest occupied molecular orbital (HOMO) and the lowest unoccupied molecular orbital (LUMO) are generally responsible for the strong electronic transition



Fig. 1 Optimized structures of osmium-bridged tricyclic aromatic compounds 2 and 2' in dichloromethane solution

Table 1 Selected predicted bond lengths (Å), bond angles (°), dihedral angels (°), charge populations, and Wiberg bond indexes (WBI) of **2** and **2'** (in parenthesis)

Bonds		Angles/dihedral angle	s	Charge	;	WBI	
Os1–C7	2.098 (2.107)	C7–Os1–C12	74.3 (73.3)	Os1	-1.337 (-1.160)	Os1–C7	0.8942 (0.9006)
Os1–C12	2.135 (2.126)	C12-Os1-C16	73.7 (73.5)	C7	-0.180 (-0.199)	Os1-C12	0.7791 (0.7740)
Os1-C16	2.062 (2.062)	C16-Os1-C27	74.1 (73.9)	C12	-0.115 (-0.131)	Os1-C16	0.9122 (0.9176)
Os1-C27	2.190 (2.208)	C27-C16-C7-C10	177.6 (179.7)	C16	0.432 (0.423)	Os1-C27	0.7614 (0.7427)
		C22-Os1-C13-C9	178.2 (178.0)	C27	-0.118 (-0.132)		

from the ground state to the first excited state. To understand the nature of absorption spectra, the electron density differences between the first excited state and the ground state and distribution of related molecular orbitals were discussed. The electron density diagrams and energies of selected frontier molecular orbitals of the complex 2 in dichloromethane solution are displayed in Fig. 2. As Fig. 2 shows, the energy levels of HOMO and LUMO are -5.20and -2.32 eV, respectively, indicating that the HOMO– LUMO excitation may occur in the visible region.

The orbital composition analysis shows that HOMO is mainly localized over the phenanthrene moiety, while LUMO is contributed by the metal-bridged blended five membered rings in **2**. As shown in Fig. 3b, the electron densities are significantly depleted at the phenanthrene group, whereas there are increases at the metal-bridged tricyclic moiety, resulting in partial charge transfers as the electronic transition occurs from the ground state to the first excited state.

Absorption spectra of complexes 2 and 2' in dichloromethane solution were investigated by both TD-B3LYP and TD-CAM-B3LYP calculations with the SMD solvation model, and experimental and theoretical absorption spectra are displayed in Fig. 3a. Experimentally, the dominant absorptions appear at 397 and 541 nm for



Fig. 2 Selected key frontier molecular orbitals and energy levels in the complex ${\bf 2}$

complex 2' [21–23], respectively. TD-B3LYP calculations indicate that the electronic absorptions at 396 and 529 nm have relatively large oscillator strengths of 0.11 and 0.13, respectively, showing good agreement with the experimental observations. We note that here TD-CAM-B3LYP calculations systematically overestimate the transition energies, as shown in previous studies [26]. However,



Fig. 3 a Electronic absorption spectra of 2 and 2' predicted by TD-B3LYP and TD-CAM-B3LYP in dichloromethane solution; b Electron density differences between the first excited state and the ground state of the osmacycle 2 (*Yellow* and *blue* refer to a decrease and an increase in electron density, respectively; isovalue: 0.0004 au.)

the TD-CAM-B3LYP-predicted spectra nicely match the experimental bands both in peak positions and relative intensities if a redshift of 0.57 eV is introduced to the TD-CAM-B3LYP spectrum (refer to Fig. S3). Hence, both B3LYP and CAM-B3LYP approaches have been applied to simulation of UV/Vis spectra in following calculations. As Fig. 3a shows, the absorption spectra of complexes 2 and 2' are similar and they all have broad absorptions in the visible region. Accordingly, the ligand-simplified complex 2 is also promising for design of high-performance dyes through introduction of various π -bridge-conjugated groups and anchoring groups.

3.2 Effects of structural modification on electronic and optical properties

To have an insight into effects of structural modification on charge separation and light harvesting ability, here four metal-bridged tricyclic aromatic derivatives in Scheme 2 have been considered, which are derived from the embedding of various π -bridge-conjugated groups to the complex 2, based on previous studies [41, 45–47]. Considering the direction of charge transfer, the steric effect of triphenylphosphine groups, and the features of frontier orbitals, the C10 site in 2 was served as the π -bridge junction. Predicted electronic and light harvesting properties related to the dye performance are collected in Table 2. The electron density differences between the first excited state and the ground state calculated by Multiwfn [48], and distributions of HOMO and LUMO orbitals in dyes 3–6 are presented in Fig. 4 and Fig. S4, respectively.

We note that the donor (D) moiety has dominant contribution to HOMO orbitals, while the π -bridge (B) and the acceptor (A) groups dominate the LUMO orbitals in dyes **3–6** here. The predicted electron density differences in Fig. 4 show that the electronic densities in the donor moiety decrease remarkably, while sharp increases appear in the bridge and acceptor moieties in the low-lying states from the S₀ \rightarrow S₁ transition in **3–6**. This suggests that introduction of the electron-deficient pyrimidine and electron-rich thiophene bridges can further facilitate charge separation during the electronic transition in the dyes **3–6**, compared to **2**.

As Fig. 5 shows, the osmacycle **3–6** have much stronger and broad absorptions in the visible region, compared to the complex **2**. In particular, the newly designed dyes **5** and **6** containing thiophene and thienothiophene units exhibit very strong absorptions in the whole visible region. These novel spectroscopic features are highly required for enhancement of the light harvesting ability of dyes and improvement of the performance of DSSCs.

The predicted transition energies, corresponding oscillator strengths, the light harvesting efficiency LHE(λ), and the driving force ΔG_{inject} are compiled into Table 2. The CAM-B3LYP functional is proved to be appropriate for the charge-transfer (CT) excitations, and it has been used to determine the vertical energies and the oscillator strengths within the TD-DFT framework. In Table 2, all strongest $S_0 \rightarrow S_1$ transitions are generally mainly contributed by the HOMO–LUMO excitation, and the first excited state is characterized by ${}^1(\pi\pi^*)$.

Here, all LUMO energy levels of these dyes are more positive than the bottom of conduction band of TiO_2



Table 2 HOMO and LUMO energies (eV), transition energies (λ in nm), corresponding oscillator strengths (f), the light harvesting efficiency (LHE), redox potentials (ΔE^{dye} in eV), and the driving force (ΔG_{inject} in eV) for osmacycle dyes by B3LYP and CAM-B3LYP

Dye	HOMO/LUMO	λ _{max}	Main configurations	f	LHE	$\Delta E^{\rm dye}$	ΔG_{inject}
B3LY	p						
1	-5.15/-2.49	549	$H \rightarrow L(0.70)$	0.33	0.53	4.96	-1.30
2	-5.20/-2.32	529	$H \rightarrow L(0.69)$	0.13	0.26	5.23	-1.11
2'	-5.48/-2.60	524	$H \rightarrow L(0.69)$	0.14	0.28	4.98	-1.39
3	-5.51/-3.02	595	$H \to L(0.69) H-1 \to L(0.12)$	0.37	0.57	5.31	-0.77
4	-5.48/-3.11	603	$H \to L(0.68) H \to L + 1(0.19)$	0.38	0.58	5.24	-0.81
5	-5.44/-2.99	590	$H \to L(0.66) H \to L + 1(0.24)$	0.77	0.83	5.29	-0.81
6	-5.43/-2.99	591	$H \to L(0.65) H \to L + 1(0.26)$	1.03	0.91	5.27	-0.82
CAM-	B3LYP						
1	-6.31/-1.32	431	$\mathrm{H} \rightarrow \mathrm{L}(0.58) \mathrm{H}\text{-}1 \rightarrow \mathrm{L}(0.32)$	0.53	0.70	5.11	-1.76
2	-6.44/-1.17	434	$\mathrm{H} \rightarrow \mathrm{L}(0.63) \ \mathrm{H}\text{-}1 \rightarrow \mathrm{L}(0.14)$	0.17	0.32	5.32	-1.54
2′	-6.70/-1.41	424	$H \to L(0.62) H \to L + 1(0.15)$	0.17	0.32	5.16	-1.76
3	-6.71/-1.83	448	$H \to L(0.55) H \to L + 1(0.32)$	0.45	0.65	5.68	-1.08
4	-6.68/-1.91	445	$H \to L + 1(0.50) H \to L(0.38)$	0.49	0.68	5.39	-1.40
5	-6.64/-1.83	451	$H \to L(0.43) H \to L + 1(0.44)$	1.00	0.90	5.38	-1.37
6	-6.62/-1.84	452	$\mathrm{H} \rightarrow \mathrm{L} + 1(0.44) \ \mathrm{H} \rightarrow \mathrm{L}(0.41)$	1.22	0.94	5.68	-1.06

Fig. 4 Electron density differences between the first excited state and the ground state of dyes 3–6 (*yellow* and *blue* refer to a decrease and an increase in electron density, respectively; isovalue: 0.00002 au)





Theor Chem Acc (2014) 133:1531





Fig. 5 Predicted absorption spectra of the new designed osmacycle dyes **2–6** in the dichloromethane solution by B3LYP

(-4.0 eV vs. vacuum) and HOMO energies are more negative than the potential of I^-/I_3^- redox couple (-4.8 eV vs. vacuum) [29]. Such nice adaptation among corresponding energy levels could ensure effective electron injection of the excited-state dyes and fast regeneration of the oxidized dyes. The dyes **5** and **6** have relatively larger LHE values among them, indicating that the introduction of thiophene and thienothiophene units can enhance the light absorption to a great extent. We note that there is a balance between λ_{max} and ΔG_{inject} under this circumstance which ΔE^{dye} keeps certain values. Hence, the value of ΔG_{inject} can maintain around -1.2 eV with a red-shifted absorption as expected [49]. These results suggest that the dyes **5** and **6** among this series of osmacycles are the most promising candidates for the high-performance photosensitizers.

4 Conclusion

Both B3LYP and CAM-B3LYP calculations on electronic and optical properties of the recently synthesized osmiumbridged tricyclic aromatic complex 2' show good agreement with the experimental observation, and the calculated CMO-NICS values reveal the strong aromaticity for its relatively high stability. Several new osmacycle dyes, derived from introduction of various π -bridge-conjugated groups based on the complex 2', have been designed and characterized theoretically. Calculations show that their electronic and optical properties can well be modulated by incorporating thiophene or thienothiophene bridge unit. The electron density differences between the first excited state and the ground state show much remarkable charge transfer features in the first excited states of dyes 3-6, compared to 2. In particular, the newly designed osmacycles 5 and 6 were predicted to have excellent charge separation and strong light harvesting ability, as well as large driving forces for electron injection, and they may be potential photosensitizers for high-performance DSSCs.

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REGULAR ARTICLE

First-principles investigations on the anisotropic charge transport in 4,4'-bis((*E*)-2-(naphthalen-2-yl)vinyl)-1,1'-biphenyl single crystal

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Abstract We applied the master equation method to investigate the anisotropic transport property of the 4,4'-bis((*E*)-2-(naphthalen-2-yl)vinyl)-1,1'-biphenyl molecular crystal based on first-principles calculation. It is found that the hole mobility has the largest value along the [100] direction, while electrons have the best transport property along the [010] direction. The anisotropic transport property was found to have close relationship with the charge transfer integral which is determined by the molecular stacking network in the crystals as well as the intermolecular frontier orbital overlap. In addition, the effect of the charge carrier density and the electronic field on the charge

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Hefei National Laboratory for Physical Sciences at the Microscale, University of Science and Technology of China, Hefei 230026, Anhui, China transport was also studied, and little effect was found except that the density is larger than 0.01 and the electronic field is increased to 1.0×10^6 V/cm. The kinetic Monte Carlo simulation method has also been used to study the anisotropic charge transport property, and consistent results were obtained as with the master equation method.

KeywordsOrganic electronics \cdot Carrier mobility \cdot Charge transfer \cdot Master equation \cdot Crystal anisotropy

1 Introduction

Organic molecular crystals, as one of the most important components in organic solar cells, organic field-effect transistors, and organic light-emitting diodes, have attracted much attention of both theoretical and experimental researchers, and many excellent molecular crystals have been designed and integrated into functional devices [1-3]. Since the charge transport is an essential process in all these devices, its mechanism has become an enthusiastically controversial topic regarding both electrical and optical measurements [4-6] and some general rules have been proposed [7, 8]. To better understand the transport mechanism, it is desirable to gain knowledge of the anisotropic transport property of organic molecular crystals which is closely related to the stacking configurations and the molecular structures. To date, the anisotropy of transport property has been investigated only in a few molecular crystals, such as rubrene (5,6,11,12-tetraphenyltetracene), CH4T(dicyclohexyl-α-quaterthiophene), pentacene, and tetracene, all of which are studied on the twodimensional (2-D) charge transport [9-14]. Recently, Tao group for the first time performed the measurement of the three-dimensional (3-D) charge transport in the 4,4'-bis((E)-2-(naphthalen-2-yl)vinyl)-1,1'-biphenyl (BNVBP) molecular crystal with two kinds of morphology and orientation and the highest hole mobility was found along the *a*-axis direction [15]. In this work, we apply the master equation (ME) method to investigate the anisotropic property of the charge carrier mobility based on the first-principles calculations. The calculated results are found in good agreement with the experiment. The influence of the electric field and the carrier density on mobility has also been studied. The 3-D transport property of the intrinsic diffusion mobility was also investigated based on the kinetic Monte Carlo (MC) simulations, and the similar anisotropic property was found. The theoretical investigation helps for better understanding of the transport mechanism and intrinsic property of the BNVBP molecular crystal.

2 Methodology

In this paper, both the drift mobility and the diffusion mobility are investigated. For the drift mobility, the incoherent motion of the carriers is described by the Pauli Master equation [16]:

$$\frac{\partial \rho_i}{\partial t} = -\sum_{j \neq i} \left[W_{ji} \rho_i (1 - \rho_j) - W_{ij} \rho_j (1 - \rho_i) \right] \tag{1}$$

Here, ρ_i is the charge carries population density of site *i* and the $(1 - \rho_j)$ represents the Coulomb penalty of site *j* due to the occupation of charge carriers. W_{ji} is defined as the charge carrier transition rate from site *i* to site *j*, and it can be calculated using the Marcus–Hush electron transfer theory at the high-temperature limit [17, 18]:

$$W_{ji} = \frac{V_{ji}^2}{\hbar} \sqrt{\frac{\pi}{K_B T \lambda}} \cdot \exp\left[-\frac{\left(\mp \Delta G_{ji} + \lambda\right)^2}{4\lambda K_B T}\right]$$
(2)

Here, V_{ji} is the transfer integral between site *i* and site *j*, and λ is the reorganization energy. ΔG_{ji} is the site energy difference between the site *i* and site *j*, and it is defined as

$$\Delta G_{ji} = E_j - E_i \tag{3}$$

where E_i is the free energy of site *i*. The sign of minus before the site energy difference in Eq. (2) is adopted for holes, and the positive sign is used for the electrons. When an electric field is applied, the site energy is modulated as

$$\Delta G_{ji} = E_j - E_i + |e|\vec{F} \cdot \vec{R}_{ji} \tag{4}$$

where \overline{F} is the electronic field and \overline{R}_{ji} is the displacement between site *i* and site *j*:

$$\overline{R}_{ji} = \overline{R}_j - \overline{R}_i. \tag{5}$$

As the measurement is performed at the steady state, the charge carrier population is assumed to be constant. The steady master equation is expressed as follows:

$$0 = -\sum_{j \neq i} \left[W_{ji} \rho_i (1 - \rho_j) - W_{ij} \rho_j (1 - \rho_i) \right]$$
(6)

The charge carrier population at the steady state can be calculated by the iterative procedure:

$$\rho_i = \frac{\sum_j W_{ij}\rho_j}{\sum_j W_{ji} + \sum_j \left(W_{ij} - W_{ji}\right)\rho_j}$$
(7)

Then, the charge carrier mobility can be calculated:

$$\mu = \frac{\sum_{i,j} W_{ji} \rho_i (1 - \rho_j) \widehat{F} \cdot \overline{R}_{ji}}{\rho_{\text{tot}} |F|}$$
(8)

Here, F is the unit vector of the electric field, and ρ_{tot} is the total charge carrier population in the investigated supercell. In this paper, the 5 × 5 × 5 supercell is large enough to give the consistent result and periodic boundary condition has also been applied. Based on these equations, we can also investigate the influence of the charge carrier density and the electric field.

When there is no electric field, the charge carriers can also diffuse in the molecular crystals, and the intrinsic charge mobility can be calculated using the Einstein equation

$$\mu = \frac{eD}{K_{\rm B}T} \tag{9}$$

where *D* is the diffusion coefficient, and it can obtained by performing the kinetic Monte Carlo simulation. In the simulation, the crystal structure is used and one charge is only allowed to transport to neighbor molecules with the probability as $P_{ki} = \frac{W_{ki}}{\sum_{j} W_{kj}}$. The sum runs over all the possible paths when one charge is located at site *k*. At each step, one random number *r* is generated. If the number *r* satisfies $\sum_{i=1}^{\alpha} P_{ki} < r \le \sum_{i=1}^{\alpha} P_{ki}$, then the charge will transport through the α path. For the detailed simulation method, one can refer to the Ref. [19]. By repeating the process for thousands times, the averaged value with respect to the time follows a linear relationship. The ratio is the diffusion coefficient according to

$$D = \lim_{t \to \infty} \frac{r^2}{2nt} \tag{10}$$

where *r* and *t* are the averaged displacement and total time span. Here, *n* represents the dimension of the charge transport in the crystal. In our calculation, the $10 \times 10 \times 3$ supercell is adopted. The time limit for each simulation was set as 0.01 ns and 2,000 trajectories were chosen to get the average value of the diffusion coefficient *D*.





In this work, the charge carrier transition rate is calculated using the Marcus rate equation where the transfer integral and the reorganization energy are two important parameters. Lots of methods have been developed to calculate the transfer integral [20–26], and here, the site energy correction method is adopted [23-25]. The reorganization energy is calculated using the adiabatic potential energy surface method (AP) [23]. The normal mode analysis method (NM) [27] has also been used for validation [28]. The calculation of the transfer integral is based on the electronic structure calculation using the PW91PW91/6-31g* method. To calculate the reorganization energy, the geometry of the molecule is first optimized and then, the frequency and electronic structure calculation are performed using the B3LYP/6-31g* method. All the quantum chemistry calculation is realized in the Gaussian 09 program [29].

3 Results and discussion

3.1 Geometric comparison

To calculate the reorganization energy, the BNVBP molecule (see Fig. 1a) was first optimized in the gas phase. Compared with the structure in the crystal, the most significant difference lies in the dihedral angle between the two phenyls at the center of the molecule (see Fig. 1b, c). In the crystal structure, the two phenyl rings in the center are at the same surface, so are the C=C double bonds. However, significant dihedral angle (about 34.68°) can be found between the two phenyl rings in the center of the molecule in gas phase. From Fig. 1b, c, we can also see that the

Table 1 Structure parameters of the BNVBP molecule in the crystaland gas phase. The numbers shown in the first column are the atomnumbers shown in Fig. 1a (unit: degree)

Angles	Crystal	Gas	Difference
47-7-45-39	-9.337	1.087	10.424
53-37-10-24	11.295	0.189	11.106
45-39-15-51	4.039	1.637	2.402
13-32-53-37	175.110	179.860	4.750
28-5-4-30	0.051	34.682	34.631
47-7-45	121.647	123.043	1.396
45-39-15	128.324	127.111	1.213
39-15-51	124.696	123.968	0.728
28-5-4	121.935	121.287	0.648
13-32-53	118.421	118.969	0.548
53-37-10	125.405	127.120	1.715
37-10-11	119.293	119.049	0.244

dihedral angles between the end naphthalene groups and the central part also differ with each other. In the crystal structure, the naphthalene groups at two ends outstand the central surface and the angles are as large as 10.42° and 11.10° , respectively; however, they keep the same surface with the phenyl rings connected by the C=C double bond at each side in gas phase. Other significant differences between the structures in crystal and gas phase are listed in Table 1. We can see that the differences mainly arise from the dihedral angles and the angles between bonds. The bond length involved in this molecule has little difference (no more than 0.1 Å) between the crystal and gas structures which is therefore not listed here.

Table 2 Reorganization energy of electrons and holes calculatedusing the AP method and NM method (unit: meV)

Method	Electron	Hole
AP	215	175
NM	215	175

3.2 Electronic structure

Based on the electronic structure calculations of the BNVBP molecule in crystal structure, the highest occupied molecular orbital (HOMO) and the lowest unoccupied molecular orbital (LUMO) are shown in Fig. 1d, e. Clearly, both the HOMO and LUMO are delocalized orbitals; thus, similar transfer integrals are expected both for holes and electrons. The calculated energy for HOMO is -5.14 eV which is close to the work function of gold (5.1 eV). The energy for LUMO is -1.81 eV which does not match with the metal electrode and is easy to be ionized in the air. Consequently, BNVBP should be used as a p-type semiconductor material.

3.3 Reorganization energy

To simulate the property of BNVBP molecules in crystals, the dihedral angle 28-5-4-30 is frozen during the calculation of the reorganization energy. Two kinds of methods have been employed to calculate the reorganization energy, namely the AP method and NM method. It is shown in Table 2 that the reorganization energy calculated with the NM method is fully consistent with the AP method. As illustrated in this table, holes have smaller reorganization energy than electrons and larger mobility is expected for holes. The comparison between the structures of the cation and the neutral state is shown in Fig. 1f. These two structures have very good overlap, and no significant difference can be found, which reflects small reorganization energy induced when one electron is withdrawn from the molecule. The similar phenomenon can also be found for the anion and neutral states (see Fig. 1g).

The reorganization energy of holes contributed by every vibration modes can also be obtained with the NM method, as illustrated in Fig. 2. It is found that the profiles for the neutral state and the cation state are quite similar. Most of the vibration modes have little contribution to the reorganization energy. The main contribution comes from the C=C stretching vibration modes in the biphenyl of the molecule and the C=C bonds. The stretching mode of the two naph-thalene groups in the molecule and the C–H in-plane bending modes have also significant contributions. The reorganization energy for anions contributed by vibration modes has also been calculated, and similar profile is obtained as that for holes.



Fig. 2 Reorganization energy contributed by vibration modes of neutral state (a) and cation state (b). The vibration profiles of vibration modes with significant contribution to the reorganization energy are shown at the *bottom* of the figure

3.4 Transfer integral

Another important factor that determines the charge carrier mobility is the intermolecular transfer integral which depends on both the molecule geometry and the stacking structure. The unit cell of the BNVBP molecular crystal is shown in Fig. 3a, where the crystal structure comes from Ref. [12]. One can see that molecule arranged in a herringbone pattern. The main transfer paths on the (001) and (010) surface are illustrated in Fig. 3b, c. The transfer integral of holes and the intermolecular distances are listed in Table 3. It can be seen that the transfer integrals along the six different paths on the (001) surface are much larger than those on the (010) surface. The large separation between the molecules along the paths on (010) surface makes the transfer integral very small. It has been widely accepted that larger transfer integral can be generated along π stacking directions. In this study, the intermolecular distance along the π stacking direction is larger than the other directions on the (001) surface; thus, smaller transfer integral is obtained. Correspondingly, the transfer integral of electrons is also shown in Table 3. One can see that



Fig. 3 a The unit cell of the BNVBP molecular crystal detected in experiment, **b** and **c** present the transfer paths on the (001) surface and the (010) surface

the transfer integrals along the paths on the (010) surface are quite small as well. The transfer integrals along the π stacking direction become much smaller than those along the other four paths on the (001) surface, which implies lower electron mobility along the π stacking direction.

3.5 Electric field and charge carrier density dependences of drift mobility

Based on the above calculations, one can find that the transfer integral is much smaller than the reorganization

energy for both holes and electrons, which implies that charge carriers transport by hopping between molecules. The hopping rate can be calculated according to the Marcus rate equation; consequently, the drift mobility can be obtained by solving the master equations. In the calculations of the drift mobility, two parameters need to be determined. One is the charge carrier density; the other is the magnitude of the electronic field. In the experiment, the applied voltage is 80 V and the length of the OFET is about 7.3 µm [15]. The electric field is about $F = V/L = 1.0 \times 10^5$ V/cm. The charge carrier density is determined by gate voltage, difficult to be measured in experiment. Nevertheless, the influence of the charge carrier density on the drift mobility can be simulated in our theoretical method (see Fig. 4a). Here, the electric field is assumed to be 1.0×10^5 V/cm, and the voltage is applied along the [100] direction. It is illustrated that there is no significant influence on the mobility when the charge carrier density is small. When the carrier density is larger than 0.01, the mobility decreases significantly with the increase in the carrier density. This is due to less available sites left for charge to transport in the molecular crystal; consequently, charge transport will be hindered.

The influence of the electric field on the drift mobility is quite complicated (see Fig. 4b). When the electronic field is in the range of 10^2 – 10^6 V/cm, the drift mobility has no significant change. When the electric field is larger than 1.0×10^6 V/cm, the drift mobility increases at first and then decreases. This happens when the inverted regime of Marcus theory is approached (i.e., at the semiclassical level, when $\Delta G_{ii} > \lambda$). In that regime, the increase in driving force will lead to a reduction in transfer rate and hence in mobility. It should be noted that the charge carrier density of 0.001 is adopted. Similar properties can be found for electrons. It is found that the drift mobility for electrons is about half of the values of holes. In addition, one can find that the turning point of the electronic field for electrons is at 5.04 \times 10⁶ V/cm, but at 2.01 \times 10⁶ V/cm for holes. From Eqs. (2) and (4), one can conclude that the difference

 Table 3 Transfer integrals of holes and electrons along different paths in molecular crystals

Transfer paths	Central distances (Å)	Transfer integral of holes (meV)	Transfer integral of electrons (meV)
P1	5.97	-17.41	9.26
P2	4.80	22.24	-30.87
P3	4.80	22.24	-30.87
P4	5.97	-17.41	9.26
Р5	4.80	19.91	-29.67
P6	4.80	19.91	-29.67
P7	26.95	0.012	0.014
P8	26.90	0.347	-0.427
Р9	26.95	0.015	0.038
P10	26.90	0.350	-0.421



Fig. 4 a Charge carrier density dependence of the hole mobility (*dashed line*) and electron mobility (*solid line*), b electronic field dependence of hole mobility and electron mobility

mainly arises from the deviation of the reorganization energy of electrons and holes.

3.6 Anisotropic property

In the study of the anisotropic property of the drift mobility, the electric field is fixed as 1.0×10^5 V/cm and the charge carrier density of 0.001 is adopted. The anisotropic properties of the holes as well as the electrons calculated with both ME and MC methods are shown in Fig. 5. The angles on the (001), (100), and (010) surface in Fig. 5 are defined as the angles between the electric field and the OA ([100]), OB ([010]), and OC ([001]) directions, respectively. It is clear that the mobility calculated with ME method has the same anisotropic property as the diffusion mobility calculated with the MC method for both holes and electrons. Considering the computational errors, the mobility calculated with both methods is in good agreement with each other. As illustrated in Fig. 5a, the holes have larger mobility along the [100] direction than that along the [010] direction. The hole mobility along the [001] direction is much smaller than that along the [010] direction (see Fig. 5b) and the [100] direction (see Fig. 5c). It is concluded that holes have the largest mobility along the [100] direction. This is understandable if we analyze the result of the transfer integrals. From Table 3 and Fig. 3b, we can see that not only the transfer integrals of holes along P1 and P4, but also that along P2, P3, and P4 and P5 will have large contribution to the charge transport along the [100] direction, although the transfer integral along P1 and P4 is smaller than that along the other four paths. However, charge transport along [010] direction is only determined by the transfer integral along P3, P4, P5, and P6. From Fig. 5d, we find that the electrons have the largest mobility along the [010] direction, which is different from that of holes. Similarly, the reason mainly arises from the difference of the transfer integral. One can see that the transfer integral of electrons along the π stacking direction ([100]) is much smaller compared with that of holes, while the transfer integrals along other paths are relatively large which will induce large electron mobility on the [010] direction. The electron mobility on the [001] direction is also the smaller than that along [100] direction (see Fig. 5f), which is in consistent with that of holes. Our theoretical investigation shows the close relationship between the anisotropic property of the mobility and the transfer integrals.

The mobility of holes and electrons calculated with both ME and MC methods along [100], [010], and [001] directions is listed in Table 4. The anisotropic property of the hole mobility is [100] > [010] > [001]; however, for electrons, it is [010] > [100] > [001]. It is clear that the electron mobility along [100] direction is about one half of the hole mobility, while the electron mobility along the [010] direction is almost the same as the hole mobility. It can also be seen that the hole mobility along the [100] direction is $0.532 \text{ cm}^2/\text{Vs}$, which is a little smaller than the experimental value (2.3 \pm 0.2 cm²/Vs) measured on the plateshaped single-crystal OFETs but almost the same as the value (0.40 \pm 0.25 cm²/Vs) measured on the rod-shaped single-crystal OFETs [15]. The calculated hole mobility along [100] direction is about one and half times as large as that along the [010] direction, which is in consistent with the experimental result. From the anisotropic property on the (001) surface, we can deduce that the hole mobility along [110] direction should be in the middle of the value along [100] direction and [010] direction. However,



Fig. 5 a-c Hole mobility on the (001), (100), and (010) surface, d-f electron mobility on the (001), (100), and the (010) surface

Table 4 Mobility of holes and electrons along different directions calculated with both ME and MC methods and the averaged diffusion mobility (unit: cm^2/Vs)

Methods Direction	ME	ME			MC			
	[100]	[010]	[001]	[100]	[010]	[001]	Average	
Hole	0.532	0.358	0.013	0.538	0.349	0.000	0.444	
Electron	0.337	0.450	0.020	0.330	0.458	0.000	0.394	

the experimental measurement shows that the hole mobility along [110] direction is smaller than that along the [010] direction. This discrepancy may be induced by many factors, such as the defect, traps, grain boundaries, and so on.

4 Conclusion

In summary, the anisotropic property of the hole mobility and electron mobility of the BNVBP molecular crystal were investigated with both ME and MC methods. The charge carrier mobility values calculated with two methods are consistent with each other. The anisotropic property of electron mobility on the (001) surface is different from that of hole mobility, due to difference in intermolecular frontier orbital overlap: HOMO for hole and LUMO for electron. The BNVBP molecular crystal has the largest hole mobility on the [100] direction, while the best transport direction for electrons is along the [010] direction. The relationship between the anisotropic property of mobility and transfer integral has been established. Our calculation confirms that the BNVBP molecular crystal has better transport property as p-type materials than the n-type materials. The study indicates that the charge carrier density and the electronic field have little effect on the drift mobility except that the density is larger than 0.01 and the electronic field is increased to 1.0×10^6 V/cm. It should be noted that the basic assumption here is the localized charge hopping model, which is in qualitative agreement with the experiment. More general considerations should include charge delocalization effect, quantum nuclear tunneling, as well as non-perturbative charge dynamics, which deserve further investigations [30].

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REGULAR ARTICLE

Iron carbonyl thioboronyls: effect of substitution of sulfur for oxygen in the viability of binuclear complexes toward dissociation reactions

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Abstract The recent discovery of boronyl complexes of the type $(R_3P)_2Pt(BO)X$ (R = cyclohexyl; X = halogen) makes of interest the chemistry of complexes of the related thioboronyl (BS) ligand. In this connection, the binuclear iron carbonyl complex Fe₂(BS)₂(CO)₈ is predicted by density functional theory to have a symmetrical unbridged structure similar to the valence isoelectronic $Mn_2(CO)_{10}$. Higher-energy unsymmetrical $(OC)_5Fe \rightarrow Fe(BS)_2(CO)_3$ structures are also found as well as a doubly bridged $Fe_2(BS)_2(CO)_6(\mu-CO)_2$ structure. The complex Fe₂(BS)₂(CO)₈ is predicted to be viable toward symmetrical dissociation into Fe(BS)(CO)₄ fragments. However, the unsymmetrical dissociation of $Fe_2(BS)_2(CO)_8$ into $Fe(CO)_5 + Fe(BS)_2(CO)_3$ is predicted to be exothermic by ~9 kcal mol⁻¹. The low-energy structures of the mononuclear Fe(BS)₂(CO)₃ include structures in which the

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Department of Chemistry and Center for Computational Quantum Chemistry, University of Georgia, Athens, GA 30602, USA e-mail: rbking@chem.uga.edu two BS ligands have coupled to form a B_2S_2 ligand through B–B bond formation.

Keywords Iron · Boron sulfides · Metal–metal bonding · Metal carbonyls · Density functional theory

1 Introduction

Metal carbonyl chemistry has evolved since the discovery of nickel carbonyl in 1890 into one of the most important and substantial subareas of coordination and organometallic chemistry [1]. However, coordination chemistry of the diatomic boron ligands BF, BO, and BS related to CO has only very recently begun to emerge to an extremely limited extent. Syntheses of metal complexes of these diatomic boron ligands have proven to be challenging because of the instability of the free ligands. Thus, the first well-characterized fluoroborylene (BF) metal complex $Cp_2Ru_2(CO)_4(\mu$ -BF) was synthesized only in 2009 by Vidović and Aldridge [2] and characterized structurally by X-ray crystallography. Similarly, the first boronyl (BO) metal complex $(Cy_3P)_2Pt(BO)Br$ (Cy = cyclohexyl) was synthesized and structurally characterized only in 2010 by Braunschweig et al. [3, 4]. Metal thioboronyl (BS) complexes remain unknown. Free BS is unstable under ambient conditions but can be detected spectroscopically in the gaseous mixture obtained by passing a microwave discharge through B_2S_3 vapor [5, 6].

The instability of the free BX (X = F, O, S) ligands under ambient conditions restricts the scope of viable methods for the syntheses of their metal complexes as compared with syntheses of similar metal carbonyls, which can use the very stable free CO gas as a chemical reagent, sometimes even at elevated pressures. For this reason, theoretical studies on metal BS complexes are important for predicting the potentially most productive areas of their coordination chemistry to explore by indirect synthetic methods using viable boron-sulfur compounds as sources of the BS ligand. In this connection, we have used density functional theory (DFT) to study mononuclear and binuclear metal carbonyl fluoroborylene (BF) and boronyl (BO) derivatives of manganese [7], iron [8, 9], and cobalt [10]. The theoretical work on the iron carbonyl boronyl $Fe_2(BO)_2(CO)_8$ led to the discovery of unbridged dimeric $(OB)(OC)_4$ Fe-Fe $(CO)_4(BO)$ structures analogous to the well-known unbridged experimental (OC)5Mn- $Mn(CO)_5$ structure [11, 12]. Such $Fe_2(BO)_2(CO)_8$ structures are predicted to be viable with respect to unsymmetrical dissociation into $Fe(CO)_5 + Fe(BO)_2(CO)_3$. We now report similar theoretical studies on the analogous Fe₂(BS)₂(CO)₈ system. Again, unbridged (SB)(OC)₄Fe- $Fe(CO)_4(BS)$ structures are predicted analogous to the $Mn_2(CO)_{10}$. However, such $Fe_2(BS)_2(CO)_8$ species are predicted to differ from their oxygen analogs by being disfavored energetically relative to unsymmetrical dissociation into $Fe(CO)_5 + Fe(BS)_2(CO)_3$.

2 Theoretical methods

Electron correlation effects were considered by employing DFT, which has evolved as a practical and effective computational tool, especially for organometallic compounds [13– 19]. Two DFT methods were used in this study, namely the B3LYP and BP86 methods. The popular B3LYP method combines the Becke three-parameter functional (B3) [20] with the Lee–Yang–Parr (LYP) generalized gradient correlation functional [21]. The BP86 method combines Becke's 1988 exchange functional (B) with Perdew's 1986 gradientcorrected correlation functional (P86) [22, 23]. In general, these two methods predict geometries and relative energies in reasonable agreement. However, the BP86 method has been found to predict vibrational frequencies closer to experimental results.

For consistency with our previous research, double- ζ plus polarization (DZP) basis sets were adopted in the present study. Thus, one set of pure spherical harmonic *d* functions with orbital exponents $\alpha_d(B) = 0.7$, $\alpha_d(C) = 0.75$, $\alpha_d(O) = 0.85$, and $\alpha_d(S) = 0.7$ for boron, carbon, oxygen, and sulfur, respectively, was added to the standard Huzinaga–Dunning contracted DZ sets, designated as (9s5p1d/4s2p1d) for boron, carbon, and oxygen and (11s7p1d/6s4p1d) for sulfur [24, 25]. The loosely contracted DZP basis set for iron is the Wachters primitive set augmented by two sets of p functions and one set of d functions, contracted following Hood et al. [26], designated as (14s11p6d/10s8p3d).

The geometries of all structures were fully optimized using the two DFT methods. Harmonic vibrational frequencies and the corresponding infrared intensities were evaluated analytically. All computations were performed with the Gaussian 03 program package [27]. The fine grid (75, 302) was the default for evaluating integrals numerically. The finer grid (120, 974) was used for more precise resolution of the imaginary vibrational frequencies [28]. The tight (10^{-8} hartree) designation was the default for the selfconsistent field (SCF) convergence. All of the predicted triplet structures in the present study were found to have negligible spin contamination, with S(S + 1) values close to the ideal value of 2.0. The energies reported in the text and figures are those obtained after zero-point vibrational correction.

A given $Fe_a(BS)_a(CO)_b$ structure is designated as **abAc** where **a** and **b** are the number of BS and CO groups, **c** orders the structures according to their relative energies, and **A** indicates whether the structure is a singlet (**S**), doublet (D), triplet (**T**), or quartet (Q). Thus, the lowest energy singlet structure of $Fe_2(BS)_2(CO)_8$ is designated **28S-1**.

3 Results and discussion

3.1 Binuclear Fe₂(BS)₂(CO)₈ structures

Eleven optimized singlet structures were found for $Fe_2(BS)_2(CO)_8$ (Figs. 1, 2; Table 1) within 25 kcal mol⁻¹ of the global minimum **28S-1**. The lowest energy triplet $Fe_2(BS)_2(CO)_8$ structures were found to lie more than 25 kcal mol⁻¹ in energy above the global minimum **28S-1**. Therefore, triplet $Fe_2(BS)_2(CO)_8$ structures are not considered in this paper. The singlet $Fe_2(BS)_2(CO)_8$ structures include nine unbridged structures, one singly bridged structure, and one doubly bridged structure.

3.1.1 Unbridged $Fe_2(BS)_2(CO)_8$ structures

Among the nine unbridged $Fe_2(BS)_2(CO)_8$ structures (Fig. 1; Table 1), five structures, namely **28S-1**, **28S-2**, **28S-3**, **28S-4**, and **28S-5**, all consist of two $Fe(BS)(CO)_4$ units linked by an iron–iron bond with no bridging CO or BS groups. Structures **28S-1**, **28S-2**, and **28S-3** have staggered configurations of the CO and BS groups, whereas **28S-4** and **28S-5** have eclipsed arrangements of the CO and BS groups. The other four unbridged $Fe_2(BS)_2(CO)_8$ structures **28S-6**, **28S-7**, **28S-8**, and **28S-11** all consist of one $Fe(BS)_2(CO)_3$ and one $Fe(CO)_5$ unit linked by an iron–iron bond (Fig. 1; Table 1). Structures **28S-6**, **28S-7**, and **28S-8** have staggered arrangements of the CO and BS groups, whereas **28S-11** has an eclipsed arrangement of the CO and BS groups. Their structural differences mainly lie in the



Fig. 1 Nine unbridged $Fe_2(BS)_2(CO)_8$ structures. In these figures, the *upper numbers* refer to distances (in Å) obtained by B3LYP and the *lower numbers* to those obtained by BP86



relative positions of the two terminal BS groups. Structures **28S-5** and **28S-11** are predicted by both the B3LYP and BP86 methods to have imaginary vibrational frequencies.

Following the corresponding normal modes leads from the eclipsed structures to the staggered structures, namely **28S-2** and **28S-8**, respectively.

Table 1 Zero-point energies (*E*, in hartrees), relative energies (ΔE , in kcal mol⁻¹), and the numbers of imaginary vibrational frequencies (Nimag) for the optimized Fe₂(BS)₂(CO)₈ structures

Species	B3LYP			BP86		
	$-E_{\rm ZPVE}$	ΔE_{ZPVE}	Nimag	$-E_{\rm ZPVE}$	ΔE_{ZPVE}	Nimag
28S-1 (C_s)	4,280.499435	0.0	None	4,281.046436	0.0	None
28S-2 (C_2)	4,280.498604	0.5	None	4,281.048253	-1.1	None
28S-3 (C_2)	4,280.495127	2.7	None	4,281.046199	0.2	None
28S-4 (D_{4h})	4,280.493995	3.4	None	4,281.037744	5.5	1 (37 <i>i</i>)
28S-5 (C _{2h})	4,280.490659	5.5	1 (27 <i>i</i>)	4,281.041927	2.8	1 (21 <i>i</i>)
28S-6 (C_s)	4,280.488789	6.7	None	4,281.036497	6.2	None
28S-7 (C_s)	4,280.483466	10.0	None	4,281.032484	8.8	None
28S-8 $(C_{2\nu})$	4,280.476025	14.7	None	4,281.019173	17.1	1 (19 <i>i</i>)
28S-9 $(C_{2\nu})$	4,280.473249	16.4	1 (78 <i>i</i>)	4,281.032683	8.6	None
28S-10 (C_1)	4,280.472733	16.8	None	4,281.037039	5.9	None
28S-11 $(C_{2\nu})$	4,280.467981	19.7	1 (25 <i>i</i>)	4,281.010348	22.7	2 (30 <i>i</i> , 24 <i>i</i>)

The Fe₂(BS)₂(CO)₈ structures **28S-1**, **28S-2**, and **28S-**3 have staggered arrangements of the CO and BS ligands very similar to that of the isoelectronic $Mn_2(CO)_{10}$ (Fig. 1; Table 1). The energies of these three structures lie within 2.7 kcal mol^{-1} , suggesting a fluxional system. Structure 28S-1 is composed of two Fe(BS)(CO)₄ units linked by an iron-iron single bond of length 2.885 Å (B3LYP) or 2.847 Å (BP86). This iron-iron bond distance is very close to the experimental Mn-Mn distances in Mn₂(CO)₁₀ of 2.92 Å [29] and 2.895 Å determined by X-ray crystallography [30]. Structure **28S-1** is a genuine minimum by both the B3LYP and BP86 methods. The axial Fe-B distance of 1.943 Å (B3LYP) or 1.937 Å (BP86) in 28S-1 is slightly shorter than the equatorial Fe-B distance of 1.967 Å (B3LYP) or 1.962 Å (BP86). This implies stronger Fe \rightarrow B back-bonding in the axial positions than in the equatorial position. This may be a consequence of the stronger trans effect of a strongly back-bonding CO group than that of an iron-iron bond.

The Fe₂(BS)₂(CO)₈ structures **28S-2** and **28S-1** differ only in the position of one of the terminal BS groups: equatorial for **28S-2** and axial for **28S-1**. The iron–iron bond distance of length 2.945 Å (B3LYP) or 2.895 Å (BP86) in **28S-2** is slightly longer than that in **28S-1** but still corresponds to a formal single bond. The B3LYP and the BP86 methods differ in the relative energy ordering of these structures. Thus, the B3LYP method predicts **28S-1** to lie 0.5 kcal mol⁻¹ in energy below **28S-2** (Fig. 1; Table 1). However, the BP86 method predicts **28S-2** to lie 1.1 kcal mol⁻¹ in energy below **28S-1**. Since these energy differences are small, these two structures may be considered to be nearly degenerate in energy.

The $Fe_2(BS)_2(CO)_8$ structure **28S-3** lies only 2.7 kcal mol⁻¹ (B3LYP) or 0.2 kcal mol⁻¹ (BP86) above **28S-1** (Fig. 1; Table 1). The iron–iron bond distance in **28S-3** of length 2.956 Å (B3LYP) or 2.911 Å (BP86)

is slightly longer than those in **28S-1** and **28S-2**, but still implies a formal single bond.

The two eclipsed $Fe_2(BS)_2(CO)_8$ structures 28S-4 and **28S-5** consist of two completely equivalent $Fe(BS)(CO)_4$ units linked by an iron-iron bond, but differing in only the location of the two terminal BS groups. Thus, the terminal BS groups are located in axial positions for 28S-4 and in equatorial positions for 28S-5 (Fig. 1; Table 1). Structure **28S-4** lying 3.4 kcal mol⁻¹ (B3LYP) or 5.5 kcal mol⁻¹ (BP86) above 28S-1 has a small imaginary vibrational frequency of $37i \text{ cm}^{-1}$ (BP86). The iron–iron bond distance of 2.897 Å (B3LYP) or 2.860 Å (BP86) in 28S-4 implies a formal single bond. The other eclipsed Fe₂(BS)₂(CO)₈ structure **28S-5**, lying 5.5 kcal mol^{-1} (B3LYP) or 2.8 kcal mol⁻¹ (BP86) in energy above **28S-1**, has a small imaginary vibrational frequency of $27i \text{ cm}^{-1}$ (B3LYP) or 21i cm⁻¹ (BP86). The iron–iron bond distance in **28S-5** of 3.065 Å (B3LYP) or 2.995 Å (BP86) is significantly longer than that in **28S-4**, but still implies a formal single bond. The axial Fe-B distance of 1.939 Å (B3LYP) or 1.935 Å (BP86) in 28S-4 is somewhat shorter than that in 28S-1.

The $Fe_2(BS)_2(CO)_8$ structures **28S-6**, **28S-7**, **28S-8**, and **28S-11** are different types of unbridged structures in which an $Fe(BS)_2(CO)_3$ unit is linked to an $Fe(CO)_5$ unit by an iron–iron bond (Fig. 1; Table 1). These unsymmetrical unbridged $Fe_2(BS)_2(CO)_8$ structures are significantly higher-energy structures than the symmetrical unbridged $Fe_2(BS)_2(CO)_8$ structures discussed above. Structures **28S-6**, **28S-7**, and **28S-8** have staggered arrangements of the ligands very similar to that of the isoelectronic $Mn_2(CO)_{10}$. These three structures have Fe–Fe distances of ~2.9–3.0 Å similar to $Mn_2(CO)_{10}$ implying formal single bonds. Their structural differences mainly lie in the relative positions of the two terminal BS groups.

The eclipsed $Fe_2(BS)_2(CO)_8$ structure **28S-11** is the highest energy of the eleven structures, lying 19.7 kcal mol⁻¹

(B3LYP) or 22.7 kcal mol⁻¹ (BP86) above **28S-1** (Fig. 1; Table 1). The Fe–Fe distance in **28S-11** of 3.293 Å (B3LYP) or 3.125 Å (BP86) is also the longest iron–iron distance in all eleven Fe₂(BS)₂(CO)₈ structures but implies a formal single bond. Structure **28S-11** has small imaginary vibrational frequencies of 25*i* cm⁻¹ (B3LYP) or 30*i* and 24*i* cm⁻¹ (BP86). The longer Fe–Fe distances in **28S-6**, **28S-7**, **28S-8**, and **28S-11** relative to **28S-5** can be considered as a donor–acceptor single bonds with the Fe(CO)₅ unit being the donor and the Fe(BS)₂(CO)₃ unit being the acceptor, i.e., (OC)₅Fe \rightarrow Fe(BS)₂(CO)₃.

3.1.2 Bridged Fe₂(BS)₂(CO)₈ structures

The $Fe_2(BS)_2(CO)_8$ structure **28S-9**, lying 16.4 kcal mol⁻¹ (B3LYP) or 8.6 kcal mol⁻¹ (BP86) in energy above **28S**-1, is predicted to have two bridging CO groups (Fig. 2; Table 1). Structure 28S-9 is a genuine minimum by BP86 but has a small imaginary vibrational frequency of 78i cm^{-1} by B3LYP. An analogous doubly bridged $Mn_2(CO)_{10}$ structure is predicted in a previous theoretical study [31] to lie 21.0 kcal mol⁻¹ (B3LYP) or 14.8 kcal mol⁻¹ (BP86) in energy above the corresponding unbridged global minimum. The $Fe_2(BS)_2(CO)_8$ structure **28S-10**, lying 16.8 kcal mol^{-1} (B3LYP) or 5.9 kcal mol^{-1} (BP86) in energy above 28S-1, has an unusual bridging µ-SBCO ligand formed by coupling the BS group with one of the CO groups and bridging the Fe₂ unit through the boron atom. Structure **28S-10** is a genuine minimum by both the B3LYP and BP86 methods. The Fe-Fe distances in both 28S-9 and 28S-10 of 2.732 Å and 2.800 Å (B3LYP) or 2.703 Å and 2.711 Å (BP86) are about 0.1 Å shorter than those in the unbridged structures, indicating the shortening of a formal iron-iron single bond by increasing the number of bridging groups.

3.2 Mononuclear Fe(BS)(CO)₄ and Fe(BS)₂(CO)₃ structures

3.2.1 The tetracarbonyl $Fe(BS)(CO)_4$

Both DFT methods (B3LYP and BP86) predict two doublet structures for $Fe(BS)(CO)_4$ with a 17-electron configuration for the central iron atom (Fig. 3; Table 2). These two structures are genuine minima without any imaginary vibrational frequencies by both B3LYP and BP86 methods. All of the quartet $Fe(BS)(CO)_4$ structures are predicted to have higher energies and thus are not discussed in this paper.

The global minimum $Fe(BS)(CO)_4$ structure **14D-1** is a C_s symmetry doublet trigonal bipyramidal structure with a linear BS group in an axial position (Fig. 3; Table 2). The other $Fe(BS)(CO)_4$ structure **14D-2**, lying 7.7 kcal mol⁻¹ (B3LYP) or 6.6 kcal mol⁻¹ (BP86) in energy above **14D-1**,



Fig. 3 Two optimized Fe(BS)(CO)₄ structures

Table 2 Zero-point energies (*E*, in hartrees) and relative energies $(\Delta E, \text{ in kcal mol}^{-1})$ for the optimized Fe(BS)(CO)₄ structures

Species	B3LYP		BP86		
	$-E_{\rm ZPVE}$	ΔE_{ZPVE}	$-E_{\rm ZPVE}$	ΔE_{ZPVE}	
14D-1 (C_s)	2,140.241313	0.0	2,140.507374	0.0	
14D-2 (C_{2v})	2,140.229054	7.7	2,140.496890	6.6	

Neither structure has any imaginary vibrational frequencies

is a doublet $C_{2\nu}$ square pyramid with a linear BS group in the apical position. Structure **14D-2** has a shorter Fe–B distance than that in **14D-1** by both methods. In previous related theoretical studies, the singlet Fe(CS)(CO)₄ [32] and Fe(BF)(CO)₄ [33] structures with trigonal bipyramidal rather square pyramid geometry and linear CS and BF groups in the axial positions are found to have lower energies than those with the CS and BF groups, respectively, in the equatorial positions. However, the lowest energy Fe(BO)(CO)₄ structure has the BO group located in a basal position of the FeC₄B square pyramid.

3.2.2 The tricarbonyl $Fe(BS)_2(CO)_3$

Both DFT methods (B3LYP and BP86) predict three singlet structures and three triplet structures for Fe(BS)₂(CO)₃ (Fig. 4; Table 3). The five structures 23S-1, 23S-2, 23T-1, 23T-2, and 23T-3 are genuine minima without any imaginary frequencies by both the B3LYP and the BP86 methods. Structures 23S-1, 23S-2, 23S-3, and 23T-3 have square pyramidal geometry. In 23T-1 and 23T-2, the two BS ligands have coupled to form a coordinated B_2S_2 unit. This B_2S_2 unit can be considered to be a two-electron donor to the iron atom, through either the BS bond in 23T-1 or the BB bond in 23T-2. The two BS ligands in the singlet structure 23S-2 are also weakly coupled with B–B distances of 2.206 Å (B3LYP) or 2.063 Å (BP86).



Fig. 4 Six optimized Fe(BS)₂(CO)₃ structures

Table 3 Zero-point energies (<i>E</i> , in hartrees), relative energies (ΔE , in kcal mol ⁻¹), and numbers of imaginary vibrational frequencies (Nimag) for the optimized Fe(BS) ₂ (CO) ₃ structures	Species	B3LYP			BP86		
		$-E_{\rm ZPVE}$	ΔE_{ZPVE}	Nimag	$-E_{\rm ZPVE}$	ΔE_{ZPVE}	Nimag
	23S-1 (C_s)	2,449.934503	0.0	None	2,450.221965	0.0	None
	23T-1 (C_s)	2,449.940103	-3.5	None	2,450.206853	9.5	None
structures	23T-2 (<i>C</i> _s)	2,449.934439	0.0	None	2,450.210802	7.0	None
	23S-2 (C_s)	2,449.930616	2.4	None	2,450.218032	2.5	None
	23S-3 (C_{2y})	2,449.921136	8.4	None	2,450.203551	11.6	1 (63 <i>i</i>)
	23T-3 $(C_{2\nu})$	2,449.913590	13.1	None	2,450.191690	19.0	None

The global minimum of $Fe(BS)_2(CO)_3$ is either a singlet or a triplet spin state structure, depending upon the theoretical method. The B3LYP method favors the high spin state, while the BP86 favors the low spin state in accord with observations by Reiher et al. [34, 35]. The global minimum predicted by B3LYP is a C_s triplet structure 23T-1, lying 3.5 kcal mol⁻¹ below the singlet structure **23S-1**. However, the BP86 method predicts a C_s singlet structure 23S-1 to be the global minimum, lying 9.5 kcal mol⁻¹ below **23T-1**.

3.3 Vibrational frequencies

The $\nu(CO)$ and $\nu(BS)$ frequencies for the optimized $Fe(BS)(CO)_n$ structures (n = 4, 3, 2, 1), $Fe(BS)_2(CO)_3$,

and $Fe_2(BS)_2(CO)_8$ (Tables 4, 5) were predicted by the BP86 method, which has been shown to be more reliable for ν (CO) frequencies than the B3LYP method when comparison with experimental data is available [36, 37]. The predicted terminal $\nu(CO)$ frequencies for the Fe(BS)(CO)_n (n = 4, 3, 2, 1), Fe(BS)₂(CO)₃, and Fe₂(BS)₂(CO)₈ structures range from 1,950 to 2,108 cm^{-1} , which is typical for terminal CO groups. Table 4 also compares the terminal $\nu(CO)$ frequencies of the Fe(BS)(CO)₄ global minimum 14D-1 (Fig. 3) with those of the corresponding Co(BO) $(CO)_4$, $Fe(BF)(CO)_4$, $Fe(CS)(CO)_4$, and $Fe(BO)(CO)_4$ structures predicted by using the same BP86 method. Our predicted $\nu(CO)$ frequencies for the Fe(BS)(CO)₄ structure lie within 25 cm⁻¹ of the corresponding ν (CO)

Table 4 ν (CO) and ν (BS) frequencies (in cm ⁻¹) of the	Structures	ν(CO)	v(BS)
mononuclear $Fe(BS)(CO)_n$	14D-1 (<i>C_s</i>)	1,999 (1,189), 2,006 (719), 2,012 (525), 2,063 (281)	1,191 (112)
(n = 4, 3, 2, 1) structures (BP86 method)	14D-2 (C_{2v})	1,993 (1,291), 2,001 (1,163), 2,011 (169), 2,070 (83)	1,216 (239)
incurou)	$Co(BO)(CO)_4 (C_{3v})$	2,026 (893), 2,026 (893), 2,039 (471), 2,095 (90)	-
	$Fe(BF)(CO)_4 (C_{2v})$	1,984 (1,310), 1,993 (1,011), 2,000 (293), 2,060 (113)	-
	$Fe(CS)(CO)_4(C_{2v})$	1,990 (1,047), 2,008 (331), 2,008 (1,239), 2,070 (191)	-
	$Fe(BO)(CO)_4(C_s)$	2,006 (1,216), 2,010 (708), 2,018 (516), 2,075 (96)	_
	13D-1 (C_{2v})	1,985 (1,420), 1,998 (833), 2,059 (168)	1,206 (199)
	13D-2 (C_s)	1,967 (1,006), 1,985 (713), 2,035 (557)	1,192 (129)
	13Q-1 (C_s)	1,979 (1,049), 1,979 (1,049), 2,024 (453)	1,141 (7)
	12Q-1 $(C_{2\nu})$	1,964 (1,651), 2,026 (163)	1,162 (11)
	$12Q-2(C_s)$	1,964 (1,196), 2,021 (655)	1,170 (52)
	12D-1 $(C_{2\nu})$	1,957 (1,568), 2,035 (30)	1,202 (164)
	12D-2 (C_s)	1,959 (815), 2,011 (759)	1,194 (167)
	11Q-1 $(C_{\infty \nu})$	2,000 (902)	1,200 (124)
	11D-1 (C_s)	1,950 (526)	1,195 (163)
	23S-1 (C_s)	2,009 (1,121), 2,020 (679), 2,071 (199)	1,197 (172), 1,226 (175)
	23T-1 (C_s)	2,006 (649), 2,009 (801), 2,054 (613)	1,151 (80), 1,191 (65)
	23T-2 (C_s)	2,016 (523), 2,017 (1,059), 2,071 (29)	1,168 (95), 1,189 (19)
	23S-2 (C_s)	1,971 (1,506), 1,995 (651), 2,037 (345)	1,076 (90), 1,422 (88)
	238-3 $(C_{2\nu})$	1,994 (1,290), 1,995 (572), 2,038 (431)	1,121 (179), 1,230 (70)
Infrared intensities in parentheses are in km mol ⁻¹	23T-3 $(C_{2\nu})$	1,996 (1,156), 2,009 (620), 2,049 (181)	1,124 (1), 1,129 (4)

Table 5 ν (CO) and ν (BS) frequencies (in cm⁻¹) of the Fe₂(BS)₂(CO)₈ structures (BP86 method)

Structures	ν(CO)	ν(BS)
Mn ₂ (CO) ₁₀	1,992, 2,025, 2,053 (gas phase)	_
Mn ₂ (CO) ₁₀	1,983, 2,014, 2,045 (n-hexane solution)	-
28S-1 (<i>C</i> _s)	1,989 (306), 2,004 (68), 2,009 (60), 2,028 (1,286), 2,028 (1,168), 2,049 (132), 2,049 (1,068), 2,099 (26)	1,218 (277), 1,232 (303)
28S-2 (<i>C</i> ₂)	1,997 (92), 2,011 (582), 2,015 (434), 2,024 (67), 2,033 (645), 2,035 (889), 2,055 (1,409), 2,090 (26)	1,219 (439), 1,221 (67)
28S-3 (<i>C</i> ₂)	1,996 (28), 2,002 (118), 2,017 (615), 2,022 (437), 2,022 (1,016), 2,028 (539), 2,052 (1,288), 2,087 (225)	1,221 (108), 1,229 (325)
28S-4 (<i>D</i> _{4h})	1,998 (0), 1,998 (0), 2,023 (0), 2,027 (1,788), 2,027 (1,788), 2,043 (0), 2,043 (463), 2,102 (0)	1,229 (708), 1,232 (0)
28S-5 (<i>C</i> _{2<i>h</i>})	1,989 (0), 2,006 (100), 2,018 (1,683), 2,023 (0), 2,026 (816), 2,026 (0), 2,052 (1,526), 2,084 (0)	1,221 (503), 1,223 (0)
28S-6 (<i>C_s</i>)	1,985 (379), 1,998 (178), 2,024 (437), 2,027 (1,033), 2,034 (825), 2,040 (662), 2,048 (298), 2,103 (231)	1,198 (210), 1,226 (292)
28S-7 (<i>C_s</i>)	1,981 (421), 1,990 (141), 2,021 (463), 2,034 (713), 2,040 (877), 2,040 (992), 2,060 (308), 2,104 (232)	1,161 (198), 1,201 (196)
28S-8 (<i>C</i> _{2v})	1,986 (525), 2,008 (194), 2,036 (305), 2,041 (734), 2,045 (1,159), 2,047 (768), 2,057 (0), 2,108 (184)	1,186 (485), 1,213 (13)
28S-9 (<i>C</i> _{2v})	1,864 (704), 1,890 (2), 1,998 (0), 2,019 (1,754), 2,024 (34), 2,034 (564), 2,053 (726), 2,082 (186)	1,231 (395), 1,235 (229)
28S-10 (<i>C</i> ₁)	1,986 (148), 1,993 (193), 2,008 (360), 2,011 (798), 2,031 (572), 2,042 (1,221), 2,055 (1,300), 2,084 (377)	780 (70), 1,123 (160)
28S-11 (<i>C</i> _{2<i>v</i>})	1,987 (355), 2,008 (148), 2,035 (203), 2,036 (1,366), 2,045 (759), 2,046 (720), 2,059 (123), 2,106 (250)	1,181 (418), 1,207 (19)

Infrared intensities in parentheses are in $\rm km\;mol^{-1}$

The bridging (CO) and (BS) frequencies are in bold face

Table 6 Dissociation energies (kcal mol^{-1}) for the symmetrical and unsymmetrical dissociations of $Fe_2(BS)_2(CO)_8$

	B3LYP	BP86
$Fe_2(BS)_2(CO)_8 (28S-1) \rightarrow 2 Fe(BS)(CO)_4 (14D-1)$	10.6	19.9
$\operatorname{Fe}_2(\operatorname{BS})_2(\operatorname{CO})_8 (\mathbf{28S-1}) \to \operatorname{Fe}(\operatorname{BS})_2(\operatorname{CO})_3$	-10.3	-8.2
$(23S-1) + Fe(CO)_5$		
$\operatorname{Fe}_2(\operatorname{BO})_2(\operatorname{CO})_8 \to \operatorname{Fe}(\operatorname{BO})_2(\operatorname{CO})_3 + \operatorname{Fe}(\operatorname{CO})_5$	25.6	21.7

All results reported here refer to the global minima of reactant and products

frequencies of Co(BO)(CO)₄, Fe(BF)(CO)₄, Fe(CS)(CO)₄, and Fe(BO)(CO)₄. The terminal ν (CO) frequencies (1,981– 2,108 cm⁻¹) for the staggered Fe₂(BS)₂(CO)₈ structures are also agree well with the experimental infrared ν (CO) frequencies in the isoelectronic Mn₂(CO)₁₀, namely 1,983, 2,014, and 2,045 cm⁻¹ in *n*-hexane solution and 1,992, 2,025, and 2,053 cm⁻¹ in the gas phase (Table 5).

The ν (CO) frequencies for the bridging carbonyl groups in the binuclear Fe₂(BS)₂(CO)₈ derivatives are predicted to range from 1,864 to 1,890 cm⁻¹. They are typically 100– 200 cm⁻¹ lower than the terminal ν (CO) frequencies in similar compounds relating to the lower C=O bond orders in carbonyl groups bonded to two metal atoms rather than a single metal atom.

The terminal $\nu(BS)$ frequencies for the Fe(BS)(CO)_n (n = 4, 3, 2, 1), Fe(BS)₂(CO)₃, and Fe₂(BS)₂(CO)₈ structures with uncoupled BS ligands are predicted to range from 1,121 to 1,235 cm⁻¹. The weak coupling of the terminal BS ligands in the Fe(BS)₂(CO)₃ structure **23S-2** leads to an abnormally low terminal $\nu(BS)$ frequency of 1,076 cm⁻¹ and an abnormally high $\nu(BS)$ frequency of 1,422 cm⁻¹. The $\nu(BS)$ frequency of 780 cm⁻¹ for the bridging BS group in **28S-10** is much lower than those for the terminal BS groups. This suggests a lower effective B–S bond order in a BS group bonded to two metal atoms relative to a BS group bonded to a single metal atom.

3.4 Symmetrical and unsymmetrical dissociation of Fe₂(BS)₂(CO)₈ into mononuclear fragments

The only possible dissociation pathway of the homoleptic $Mn_2(CO)_{10}$ into mononuclear fragments involves symmetrical dissociation into $\cdot Mn(CO)_5$ radicals. This is an endothermic process with calculated and experimental dissociation energies around 40 kcal mol⁻¹ [38]. The analogous symmetrical dissociation of the isoelectronic binuclear thioboronyl Fe₂(BS)₂(CO)₈, like that of Mn₂(CO)₁₀, is also an endothermic process, but significantly less endothermic, requiring 10.6 kcal mol⁻¹ (B3LYP) or 19.9 kcal mol⁻¹ (BP86) (Table 6). However, the predicted dissociation energy for the unsymmetrical dissociation of Fe₂(BS)₂(CO)₈ into the stable Fe(CO)₅ plus the 16-electron species Fe(BS)₂(CO)₃ is exothermic by -10.3 kcal mol⁻¹ (B3LYP) or -8.2 kcal mol⁻¹ (BP86) suggesting that Fe₂(BS)₂(CO)₈ is thermochemically disfavored. This contrasts with the oxygen analog Fe₂(BO)₂(CO)₈ for which the unsymmetrical dissociation into the stable Fe(CO)₅ and the 16-electron complex Fe(BO)₂(CO)₃ is highly endothermic at ~23 kcal mol⁻¹. If the kinetic barrier for the unsymmetrical dissociation of Fe₂(BS)₂(CO)₈ into Fe(CO)₅ + Fe(BS)₂(CO)₃ is sufficiently high, it might still be possible to synthesize Fe₂(BS)₂(CO)₈ by dimerization of Fe(BS)(CO)₄, since that is an exothermic process.

4 Summary

The lowest energy structures for the thioboronyl $Fe_2(BS)_2(CO)_8$ are symmetrical unbridged structures having two $Fe(BS)(CO)_4$ units linked solely by an Fe–Fe bond. Such structures are analogous to the well-known unbridged structure of the valence isoelectronic $Mn_2(CO)_{10}$. Unsymmetrical unbridged structures consisting of an $Fe(CO)_5$ unit bonded to an $Fe(BS)_2(CO)_3$ unit by an Fe \rightarrow Fe dative bond lie at least ~6 kcal mol⁻¹ in energy above the symmetrical unbridged structures. The lowest energy bridged $Fe_2(BS)_2(CO)_6(\mu$ -CO)₂ structure lies ~12.5 kcal mol⁻¹ in energy above the lowest energy isomeric unbridged structure. A higher-energy isomeric $Fe_2(BS)(CO)_7(\mu$ -SBCO) structure is also found in which a BS and CO ligand have coupled to form a bridging SBCO ligand.

The structures of the mononuclear derivatives Fe(BS) $(CO)_n$ (n = 4, 3, 2, 1) and Fe(BS)₂(CO)₃ were also investigated in order to evaluate the thermochemistry of the dissociation of Fe₂(BS)₂(CO)₈ into mononuclear fragments. The low-energy Fe(BS)₂(CO)₃ structures include triplet structures in which the two BS ligands have coupled through B–B bond formation to form a B₂S₂ ligand. The binuclear derivative Fe₂(BS)₂(CO)₈ is found to be viable with respect to symmetrical dissociation into two Fe(BS)(CO)₄ units. However, unsymmetrical dissociation of Fe₂(BS)₂(CO)₈ is an exothermic process by ~9 kcal mol⁻¹, suggesting that Fe₂(BS)₂(CO)₈ is thermochemically unstable.

5 Supporting information

Tables S1 to S30: theoretical Cartesian coordinates (in Å) for the Fe(BS)(CO)_{*n*} (n = 4, 3, 2, 1), Fe(BS)₂(CO)₃, and Fe₂(BS)₂(CO)₈ structures using the B3LYP/DZP and BP86/DZP methods; Tables S31 to S60: Theoretical harmonic vibrational frequencies (in cm⁻¹) for the Fe(BS) (CO)_{*n*} (n = 4, 3, 2, 1), Fe(BS)₂(CO)₃, and Fe₂(BS)₂(CO)₈

structures using the B3LYP/DZP and BP86/DZP methods; Figure S1 and Table S61: $Fe(BS)(CO)_3$ structures; Figure S2 and Table S62: $Fe(BS)(CO)_2$ structures; Figure S3 and Table S63: Fe(BS)(CO) structures; Complete Gaussian reference (Ref. [27]).

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