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# Immunobiology of Bacterial CpG-DNA

Edited by H. Wagner

With 34 Figures and 12 Tables



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*Cover Illustration:* Schematic representation of immune cells stimulated by bacterial CpG-DNA. Antigen presenting cells (APCs) become directly stimulated to express co-stimulatory molecules and to synthesize cytokines and chemokines (transition to professional APCs). In addition haemopoietic growth factors become induced and haemopoiesis ensues. While CpG-DNA is mitogenic to B cells, upon B cell receptor (BCR) crosslinking CpG-DNA co-stimulates proliferating B cells. Susceptibility of NK cells to IL-12 becomes co-stimulated by CpG-DNA. T cells develop sensitivity to G-rich DNA upon T cell receptor (TCR) crosslinking.

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### Preface

When asked whether they were willing to contribute a chapter to this volume on the immunobiology of bacterial CpG-DNA, all colleagues approached who are working in this emerging new field responded favorably. Subsequently, and within the time required limit, they supplied fine chapters covering their respective areas of expertise.

Immune stimulatory CpG-motifs in bacterial DNA have recently attracted a great deal of interest throughout the academic community and the industry. For various reasons it is astounding that bacterial genomic DNA, so far considered only to be a genetic blueprint, may turn out to be a virulence factor, as is the case with endotoxin. If so, its role as a stimulus in septic shock needs to be evaluated. It also comes as a surprise that CpG-motifs within bacterial DNA target immature antigenpresenting dendritic cells (DCs) to transit to professional antigenpresenting cells (APCs). By virtue of this DC activating property, CpG-DNA may act as 'natural' adjuvants able to render proteinaceous antigens and T cell epitopes immunogenic. If so, the immunobiology of CpG-DNA explains its powerful adjuvanticity, which often surpasses the gold standard of complete Freund's adjuvants (CFA). Furthermore, immune recognition of CpG-motifs may represent an interesting example of the principle by which pattern recognition receptors in DCs detect subtle molecular characteristics that distinguish pathogens from our own cells. As a considerable body of information has become available – thanks to the continued efforts of much research – CpG-DNA-driven direct activation and costimulation of immune cells has become particularly useful to those who want to address details of the biological mechanisms and their relation to structure and function.

This volume attempted to cover as widely as possible current research on the immunobiology of pC-DNA. It encompasses the role of CpG-motifs in "naked DNA" used for vaccination protocols, its role in activating APCs in such a way to condition polarized Thl immune responses, and the signaling pathways VI Preface

involved for use as an adjuvant. Typically, in a newly developing research field, one leans back to wait for the first results from human clinical trials to learn whether the data obtained in animal model systems can be translated to humans.

The response and deep interest I experienced while presenting our experimental results during visits to many institutes, in different countries, on several continents encouraged me to undertake the task of editing the present volume. I wish to thank all the contributors, the editors of *Current Topics in Microbiology and Immunology* and, in particular, Springer-Verlag for their help in getting this volume to press.

Munich, November 1999

HERMANN WAGNER

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## Mechanism of Action of CpG DNA

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#### **1** Introduction

In recent years, the position of the innate immune system in regulating nearly all immune responses has become well established. A central tenet in understanding the function of the innate immune system is the concept that it is triggered by pattern recognition receptors (PRRs), which bind microbial structures that are not present in host tissues. Examples of PRRs that have become well accepted are the lipopolysaccharide (LPS) receptors, such as CD14, and mannose-binding protein.

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Likewise, complement proteins have a primitive ability to bind and be activated by microbial structures. Viral RNAs are thought to adopt certain structural conformations that are bound by PRRs and lead to the activation of the double-stranded-RNA-dependent protein kinase PKR (KUMAR et al. 1997). This mechanism is thought to be responsible for immune activation by the polynucleotide (rI, rC) (TALMADGE et al. 1985; WILTROUT et al. 1985). Evidence that bacterial DNA (bDNA) may also possess some structural feature that activates innate immune defenses was first provided by Tokunaga et al. who reported its surprising antitumor activity and ability to activate natural killer (NK) cells (TOKUNAGA et al. 1984; YAMAMOTO et al. 1988). Further studies by these investigators led to the proposal that this immune-stimulatory activity of bDNA resided in certain self-complimentary, palindromic sequences that contained CpG dinucleotides (KURAMOTO et al. 1992). Interestingly, methylation of the CpGs was initially thought to have no influence on the immune stimulatory activities of DNA (KURAMOTO et al. 1992). Independent studies by Pisetsky and colleagues showed that bDNA can also activate B-cell proliferation and that these proliferative activities can also be seen with polynucleotides, such as dG, dC (MESSINA et al. 1991, 1993), which also activate NK cells (TOKUNAGA et al. 1988).

Independent of these studies, several different groups of investigators using antisense oligonucleotides, which usually possessed nuclease-resistant phosphorothioate backbones, began to report the occurrence of immune stimulatory effects, some of which were initially thought to be due to an antisense mechanism of action but others of which were clearly due to some highly sequence-specific but nonantisense mechanism of action (KRIEG et al. 1989; HATZFELD et al. 1991; TANAKA et al. 1992; BRANDA et al. 1993; MCINTYRE et al. 1993; MOJCIK et al. 1993; PISET-SKY and REICH 1993; FISCHER et al. 1994; KOIKE et al. 1995). Initially, there seemed to be no common sequence element or motif among all of these different immunostimulatory effects. However, after several years of investigating the immunostimulatory effects of different oligonucleotides and DNAs, we finally came to the realization that a rather simple sequence motif based on an unmethylated CpG dinucleotide could, in particular base contexts, link together all of the reported immunostimulatory activities (Fig. 1; KRIEG et al. 1995; KRIEG 1998).

It makes tremendous teleological sense for the immune system to have evolved an innate defense triggered by these unmethylated CpG motifs. While CpGs are unmethylated and usually fairly abundant in bDNA, they are methylated and highly suppressed (only about one-quarter as frequent as would be predicted if base utilization were random) in vertebrate DNA (BIRD 1987). Moreover, the base context of CpG dinucleotides in the human genome is not random; in our genome, CpGs are most frequently preceded by a C or followed by a G, which is an unfavorable context for immune stimulation and which can even have neutralizing effects, blocking immune stimulation by optimal stimulatory CpG motifs (Fig. 1; KRIEG et al. 1998b).

It does not seem particularly logical that the recognition of CpG DNA evolved as a general defense strategy against bacteria, since most of these normally live outside our cells and their DNA would, therefore, be shielded from detection. The

CpG motifs can be species-specific: ODN #1826 – 5' TCCATGA<u>CG</u>TTCCTGA<u>CG</u>TT 3' (activates mouse but not human cells) ODN #2006 – 5' T<u>CGTCG</u>TTTTGT<u>CG</u>TTTGT<u>CG</u>TT 3' (activates human and mouse cells)

Fig. 1. Formula for a CpG motif. Not all DNA-containing, unmethylated CpG dinucleotides are immune stimulatory. If the CpG is preceded by a C or followed by a G, or if the CpGs are arranged in direct repeats, then they may have inhibitory rather than stimulatory effects on leukocyte activation by CpG DNA (KRIEG et al. 1998b). In general, mouse cells respond to a rather wide variety of CpG motifs, but human cells are optimally stimulated by a narrower subset. A particularly strong mouse motif is TGACGTT, while a particularly strong human motif is TGTCGTT. The figure shows examples of two phosphorothioate oligodeoxynucleotide (ODN) sequences that have immune stimulatory effects in mice, only one of which is stimulatory to human cells. Not all CpG motifs have the same range of immune activities - some are best at activating natural killer (NK) cells, with relatively little adjuvant effect, while others are potent adjuvants for antibody responses but have relatively little NK-stimulating activity. The exact range of immune effects that a given ODN will have depends on the CpG motif, the number and spacing of motifs in the ODN, the presence of other sequence motifs (such as poly-G sequences) and, very importantly, the backbone of the ODN. ODNs with nuclease-resistant phosphorothioate backbones can have dramatically increased potency if they have optimal or near-optimal CpG motifs but can have essentially no CpG activity if the motif is suboptimal, even though the same sequence may be stimulatory if made with a phosphodiester backbone (BALLAS et al. 1996; KRIEG et al. 1996). In general, phosphorothioate ODNs have lower potentials for inducing NK activation than do chimeric ODNs (BALLAS et al. 1996). The most potent backbones for stimulating B cells are, in order of potency, phosphorodithioate, phosphorothioate, phosphodiester, methylphosphonate, and 2'-O-methylribonucleotide (ZHAO et al. 1996a; KRIEG et al. 1996). Modification of the 5 position on the cytosine ring can also affect the level of immune stimulation of an ODN (Boggs et al. 1997)

CD14 pathway and detection of endotoxins is probably a far more important defense against gram-negative extracellular bacteria. Instead, we propose that the CpG-DNA innate defense has evolved and is specialized for protection against pathogens that replicate inside host cells, such as intracellular bacteria, viruses, and retroviruses. This would predict that the CpG pathway should trigger appropriate protective defenses, which would be primarily T-helper 1 cells (Th1) although, of course, this does not exclude production of antibodies. Indeed, CpG DNA appears to be the most potent Th1-like adjuvant described (CHU et al. 1997; LIPFORD et al. 1997b; ROMAN et al. 1997; WEINER et al. 1997; DAVIS et al. 1998; H. Davis, personal communication). In a recent comparison of the immune responses generated by 19 different vaccine adjuvants against 3 different antigens, CpG DNA gave by far the strongest Th1-like T-cell responses (P. Livingston, personal communication).

If the hypothesis that innate immune recognition of CpG motifs contributes to defense against infection is correct, then it would be expected that viruses and retroviruses may have evolved ways to overcome this defense. Indeed, genomic analysis of small DNA viruses and retroviruses shows that they have remarkably

repressed levels of CpG dinucleotides which, in some cases, are <10% of the expected level and which appear to represent strong evolutionary selection (SHPAER and MULLINS 1990; KARLIN et al. 1994; KRIEG 1996). Adenoviruses appear to have developed a different way of overcoming the CpG defenses by dramatically skewing the flanking bases of their CpG motifs so that the immune-stimulatory motifs are under-represented by approximately 30:1 compared with the neutralizing motifs (KRIEG et al. 1998b).

The purpose of this chapter is to review what is known about the molecular and cellular mechanisms of action of immune-stimulatory CpG DNA. This chapter will focus on studies of B cells and macrophages since comparatively little has yet been reported on other cell types.

#### 2 Molecular Mechanisms of Action of CpG DNA

#### 2.1 Cell-Surface Receptors and CpG DNA

It has long been appreciated that cell membranes have a variety of DNA-binding proteins on their surfaces (LERNER et al. 1971; AGGRARWAL et al. 1975; LOKE et al. 1989; YAKUBOV et al. 1989; BENNETT 1993). To date, none of these membrane DNA-binding proteins has actually been shown to mediate DNA uptake into cells, with the exception of a protein expressed on kidney epithelial cells that has been shown to function as a voltage-gated channel (HANSS et al. 1998a). However, oligodeoxynucleotide (ODN) uptake at low concentrations (below 1 $\mu$ M) appears consistent with receptor-mediated endocytosis while, at higher concentrations, fluid-phase endocytosis appears to be more important (BELTINGER et al. 1995).

Of course, a cell-surface CpG receptor could potentially transduce a signal into the cell without requiring uptake of the DNA. However, there is no evidence of any CpG-sequence specificity in the cell-membrane binding of DNA. Yamamoto et al. found that there was no difference in the spleen cell binding of <sup>32</sup>P-labeled ODN, which had stimulatory or non-stimulatory palindromes (YAMAMOTO et al. 1994). Likewise, we found no difference in the binding of CpG or non-CpG ODN to B cells or macrophages (KRIEG et al. 1995).

Nonetheless, human B cells were reported to be stimulated by immobilized ODN, implying activation through a cell-surface receptor (LIANG et al. 1996). In these experiments, the ODNs were synthesized with a lysine at the 5' end, coupled to cyanogen bromide-activated sepharose 4B beads, and then cultured with B cells for 4 days before performing <sup>3</sup>H-thymidine-incorporation assays. Aside from concerns over whether the ODNs actually remain immobilized under these culture conditions, Manzel and MacFarlane have recently found that ODNs coupled to sepharose beads in this manner are in fact taken up by cells during incubation in tissue culture, a phenomenon that could be responsible for the immune-stimulatory effects (MANZEL and MACFARLANE 1999). Under these same culture conditions,

CpG ODNs were not stimulatory if they were linked to latex or magnetic or gold beads, suggesting that cell uptake may indeed be required. This conclusion is in accord with results of our own studies, in which CpG ODNs were immobilized on Teflon fibers or avidin-coated plates, in which case stimulation was not seen using short-term-stimulation assays after 4h (KRIEG et al. 1995). The conclusion that ODN stimulation does not involve cell-surface signaling is further supported by the finding that lipofection of ODN into spleen cells enhances their immune-stimulatory effects (YAMAMOTO et al. 1994). Furthermore, while unmodified ODN usually must be at least 18 bases long to induce NK activity (YAMAMOTO et al. 1994), this length requirement is reduced to just 6 bases if the ODN are instead transfected into the cells (SONEHARA et al. 1996).

ODNs containing G-rich sequences, such as three or four Gs in a row ("poly-G ODNs"), have higher binding to cell membranes and have also been reported to have increased binding to the scavenger receptor on macrophages (HuGHES et al. 1994; KIMURA et al. 1994). The cell-surface integrin CD11B/CD18 (MAC-1) has also been reported to bind ODN in a non-sequence-specific fashion and mediate uptake and the generation of reactive oxygen species (ROSs; BENIMETSKAYA et al. 1997). However, mice genetically deficient in either one of these molecules appear to have essentially normal ODN uptake into cells (S. Crooke, personal communication). Furthermore, cells that do not express either molecule, such as B cells, are strongly activated by CpG DNA, indicating that neither receptor is required.

#### 2.2 Cellular Uptake and Intracellular Localization of CpG ODN

It appears that all cell types are capable of taking up DNA spontaneously in culture, although certain cell types have higher rates of uptake than others (KRIEG et al. 1991; ZHAO et al. 1996b; HARTMANN et al. 1998). It is important that investigators performing such experiments be aware of an artifact arising from the fact that apoptotic and dead cells have increased ODN uptake (ZHAO et al. 1993; GIACHETTI and CHIN 1996). In principle, mechanisms that could mediate ODN uptake include receptor-mediated endocytosis, adsorptive endocytosis, pinocytosis, potocytosis, and phagocytosis. It is clear that ODNs do not simply diffuse across cell membranes; cellular uptake is energy- and temperature-dependent and saturable (BELTINGER et al. 1995; KRIEG 1995). It is possible that several of these mechanisms contribute to ODN uptake. However, potocytosis seems unlikely to play a major role in ODN uptake because, although potocytosis has been reported to be dependent upon protein kinase C and cell-surface glycosyl phosphatidyl inositol-linked proteins (SMART et al. 1994), ODN uptake is not (A.M. Krieg, unpublished data). Following binding of ODNs, they are initially located in endosomes, which then become progressively acidified (BENNETT et al. 1985; TONKINSON and STEIN 1994; YI et al. 1998a). Most of the DNA remains in the endosomes but, based on studies with confocal microscopy, a small fraction appears to reach the nucleus (KRIEG et al. 1993; ZHAO et al. 1996b; ZHAO et al. 1993). The proportion of DNA that enters the nucleus can be increased by con-

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jugation of the ODN to compounds, such as cholesterol, or by transfection of the ODN into cells using cationic lipids (KRIEG et al. 1993; WAGNER et al. 1993; BENNETT 1998). If the DNA is directly micro-injected into the cytoplasm of a cell, it is localized to the nucleus within a few seconds (CHIN et al. 1990; LEONETTI et al. 1991). With the exception noted above for ODNs containing poly-G sequences, cellular uptake and intracellular localization of ODNs are non-sequence-specific (HANSS et al. 1998b; A.M. Krieg, unpublished data).

#### 2.3 Requirement for Endosomal Acidification/Maturation for CpG-Induced Immune Stimulation

Since DNA is internalized into cells via endocytosis into acidified vesicles, it was of interest to determine whether drugs that interfere with endosomal acidification and maturation alter the immune-stimulatory effects of CpG DNA. Indeed, we recently reported that monensin, chloroquine, and bafilomycin A, which interfere with endosomal acidification, completely block leukocyte activation by CpG DNA but not by other stimulatory agents, including phorbol 12-myristate 13-acetate, antiimmunoglobulin M (IgM), anti-CD40, or LPS (Y1 et al. 1998a). Other compounds related to chloroquine, such as quinacrine, are even more potent inhibitors of immune activation by CpG DNA (MAcFARLANE and MANZEL 1998). These compounds do not bind ODN nor do they prevent ODN uptake by cells (MAC-FARLANE and MANZEL 1998). However, it remains possible that these agents alter the intracellular localization of ODN. The ability of chloroquine and bafilomycin A to specifically inhibit CpG-induced activation of macrophages and dendritic cells (DCs) has been reported by Häcker et al. (HÄCKER et al. 1998). At present, the mechanism of these inhibitory effects is unclear. One interpretation is that endosomal acidification may be essential for the release of ODN from the endosomes into the cytoplasm, where they may exert their stimulatory effect. However, it appears that quinacrine and chloroquine successfully block immune stimulation by CpG ODN at concentrations that do not interfere with endosomal acidification (D. MacFarlane, personal communication). Alternatively, perhaps some other aspect of endosomal function is disrupted by these compounds and specifically blocks the effects of CpG DNA but not other lymphocyte mitogens that do not function through endosomes. Further studies will clearly be required to resolve this question. The inhibitory effects of chloroquine on CpG-induced activation are already apparent after 5min, indicating that the inhibitory effect occurs at a very early stage in the activation pathway (YI et al. 1998a).

#### 2.4 What is the Intracellular CpG Receptor?

Given the high degree of sequence specificity in the effects of CpG DNA and the fact that activation requires cell uptake and is not mediated through a cell-surface receptor, it appears clear that there must be some intracellular molecule that spe-

cifically interacts with CpG DNA to transduce the stimulatory signal. In order to identify proteins that may have CpG-specific patterns of binding to ODN, we performed electrophoretic mobility shift assays using ODN with stimulatory CpG motifs, methylated CpG motifs, or no motif as probes. In these preliminary studies, we identified factors in cytoplasmic and nuclear extracts of B-cell and monocytic cell lines; these factors specifically bind unmethylated CpG motifs (R. Tuetken, B. Noll, W. Shen, A.M. Krieg, unpublished data). Efforts to purify the binding proteins are currently underway. Identification of these proteins will be extremely important in understanding the CpG signaling pathways.

#### 2.5 Production of Intracellular Reactive Oxygen Species Following Stimulation by CpG DNA

Many lymphocyte mitogens alter the redox balance of the cell by inducing the generation of ROSs, such as superoxide, hydrogen peroxide, or lipid peroxides (HOTHERSALL et al. 1997; JONESON and BAR-SAGI 1998). Regulation of the redox balance of leukocytes is extremely important, because the binding of several transcription factors to DNA can be altered depending on the oxidative state of the cell (SEN and PACKER 1996; COTGREAVE and GERGES 1998). In lymphocytes, the generation of ROSs has been proposed to be important in mediating stimulation or apoptosis through the B-cell antigen receptor or CD40 (FANG et al. 1995; FANG et al. 1997; LEE and KORETZKY 1997). Expression of excessive amounts of intracellular ROS has been linked to apoptosis. The concentration, type, and intracellular localization of ROS are all likely to be important in determining their biologic effects.

To investigate whether leukocyte activation by CpG DNA may involve the generation of ROS, we stimulated cells in the presence of the ROS-sensitive dye dihydrorhodamine-123. These studies showed that leukocyte stimulation with CpG DNA, but not with non-CpG DNA, was associated with the rapid intracellular production of ROS, which was specifically blocked by chloroquine (Y1 et al. 1996c, 1998a). Multiple cellular enzymatic pathways are capable of producing ROSs, including xanthine oxidase, reduced nicotinamide adenine dinucleotide phosphate oxidase, cyclooxygenase, cytochrome p450 enzymes, and the mitochondria. Inhibition of xanthine oxidase with allopurinol has no effect on the ROS induction by CpG DNA, nor has inhibition of cyclooxygenase with indomethacin in our preliminary studies. Studies are currently underway to identify the enzymatic source of these ROSs and to specifically identify which ROSs are produced.

#### 2.6 Activation of Mitogen-Activated Protein Kinases by CpG DNA

Mitogen-activated protein kinases (MAPKs) are important mediators of many cellular activation responses. The three main MAPK pathways are the extracellular receptor kinase or ERK pathway, the p38 MAPK pathway, and the c-Jun NH<sub>2</sub>-

terminal kinase or JNK pathway. The p38 and JNK pathways are activated in both B cells and macrophages as early as 7min following activation with CpG DNA (HÄCKER et al. 1998; YI and KRIEG 1998b). These pathways are also activated in human primary B cells exposed to human activating CpG DNA motifs but not by non-CpG DNA (G. Hartmann and A.M. Krieg, unpublished data). More recently, we identified the upstream kinases MAPK kinase kinase 3 (MKK3), MKK4, and MKK6 as being activated very rapidly after CpG-DNA treatment of the murine B-cell line WEHI-231 (A.-K. Yi and A.M. Krieg, unpublished data). CpG DNA does not appreciably induce ERK-1 or ERK-2 activity in B-cell lines (YI and KRIEG 1998b) but does cause some activation of this pathway in macrophages (K.J. Stacey and D.A. Hume, personal communication).

#### 2.7 Activation of Transcription Factors by CpG DNA

The p38 and MAPK pathways have as their downstream targets several transcription factors, including activating transcription factor 2 and activator protein 1 (GUPTA and SIBER 1995; BIRELAND and MONROE 1997; DEFRANCO 1997). Therefore, it may not be surprising that these transcription factors become phosphorylated and show increased DNA binding and enhanced transcriptional activity in cells treated with CpG DNA (HÄCKER et al. 1998; YI and KRIEG 1998b).

Inhibition of p38 kinase with the specific inhibitor SB203580 inhibits CpGinduced cytokine production by macrophages and B cells, indicating the requirement for this kinase activity in mediating CpG effects (HÄCKER et al. 1998; YI and KRIEG 1998b). Nuclear factor  $\kappa B$  (NF $\kappa B$ ) has a critical role in mediating many inflammatory responses and is rapidly upregulated in leukocyte responses to many mitogens (KISTLER et al. 1998; SHA 1998). The first report that ODN could induce NF $\kappa$ B activation was an unexpected observation in an antisense experiment in which cells treated with a sense oligonucleotide to the p65 subunit of NF $\kappa$ B showed enhanced NFkB activity (MCINTYRE et al. 1993). This sense ODN contained a CpG motif that was subsequently found to be responsible for the observed upregulation of NFkB activity (A.M. Krieg, R. Narayanan, unpublished data). Following the identification of CpG DNA, the first report linking this to the activation of NFkB was from Stacey et al. who showed that macrophages treated with CpG DNA have enhanced transcriptional activity from the human immunodeficiency virus long, terminal repeat; this activity is regulated by  $NF\kappa B$ (STACEY et al. 1996). Supershift studies showed that this NFKB activation includes the p50 and p65 subunits but not p52 (Sparwasser et al. 1997b). NF $\kappa$ B activation was also seen following CpG-DNA treatment of B cells and was associated with the degradation of IkB $\alpha$  and IkB $\beta$  (Y1 et al. 1996b; Y1 and KRIEG 1998a). CpGinduced NF $\kappa$ B activation in B cells was associated with protection against the induction of apoptosis (YI and KRIEG 1998a). Similar CpG-specific induction of transcription-factor binding is seen in human primary B cells (G. Hartmann and A.M. Krieg, unpublished data). CpG DNA also induces increased levels of mRNA for other transcriptional factors, including c-myc, ets-2, and CCAAT enhancer

binding protein- $\beta$  and - $\Delta$  (Sweet et al. 1998; Y<sub>I</sub> and K<sub>RIEG</sub> 1998a; K.J. Stacey and D.A. Hume, personal communication).

#### 2.8 Induction of Gene Expression by CpG DNA

The cell-signaling pathways that are induced by CpG DNA lead to enhanced nuclear transcription within 15–30min (YI et al. 1996b, 1998b). Although the expression of multiple early-response genes, proto-oncogenes, and cytokine genes is induced by CpG DNA, the effects show a particular pattern that is suggestive of a Th1-like immune response. CpG DNA is a particularly strong inducer of the expression of interleukin 12 (IL-12), especially as compared to other agents, such as LPS, which induce IL-12 expression from macrophages or DCs more weakly (CHACE et al. 1997; LIPFORD et al. 1997b; JAKOB et al. 1998; SPARWASSER et al. 1998). There is no detectable induction of expression of the Th2-like cytokines, IL-4, or IL-5 (KLINMAN et al. 1996) although B cells are induced to express IL-10, which appears to be partially responsible for reducing the level of IL-12 secretion that is induced by CpG DNA (REDFORD et al. 1998).

The expression of tumor necrosis factor (TNF)-a is induced very rapidly following CpG stimulation and appears to be responsible for the induction of the systemic inflammatory-response syndrome in mice that have been rendered sensitive to the effects of TNF (HALPERN et al. 1996; SPARWASSER et al. 1997a,b). Other cytokines whose expression is induced by CpG DNA include IL-6, IL-1β, IL-1RA, macrophage inflammatory protein 1β, monocyte chemoattractant protein 1, interferon (IFN)- $\alpha/\beta$ , IL-18, and IFN- $\gamma$  (Y<sub>I</sub> et al. 1996c; ROMAN et al. 1997; Schwartz et al. 1997; Zhao et al. 1997; Sun et al. 1998; Yamamoto et al. 1988). Aside from these studies in mouse cells, more recently we have also found CpGspecific induction of IL-6 and TNF in human monocytes (HARTMANN et al. 1999), and IL-12 secretion from purified human DCs (G. Hartmann and A.M. Krieg, unpublished data). In addition, human peripheral blood mononuclear cells (PBMCs) cultured with CpG DNA show induction of expression of IFN- $\gamma$  (A.M. Krieg, unpublished data; YAMAMOTO et al. 1994b). Aside from the induction of these genes, CpG DNA also induces several cell-cycle-regulating proto-oncogenes and anti-apoptosis genes including, most notably, c-myc and bcl-xL (YI et al. 1996b, 1998b).

#### **3** Cellular Mechanisms of Action of CpG DNA

#### 3.1 B-Cell Activation by CpG DNA

All of the above pathways described as mediating CpG effects are inducible in primary B cells or B-cell lines following treatment with CpG DNA. CpG DNA is

the most potent single B-cell mitogen that has been described and is capable of driving more than 95% of B cells into the cell cycle (KRIEG et al. 1995). In contrast to some other B-cell mitogens, such as 8-substituted guanines, CpG DNA appears to be equally stimulatory to both resting and activated B-cell subsets (GOODMAN 1991; KRIEG 1998). B cells enter the G1 phase of the cell cycle within a few hours and progress through the cell cycle under appropriate conditions. B-cell stimulation results in the secretion of IL-6 and IL-10 within a few hours (Y1 et al. 1996c; REDFORD et al. 1998). In contrast to B-cell activation by LPS, which is suppressed by the addition of IFN- $\gamma$ , B-cell activation by CpG DNA synergizes with IFN- $\gamma$  to promote stronger responses (Y1 et al. 1996a). CpG-activated B cells are induced to secrete IgM in an IL-6-dependent fashion (Y1 et al. 1996c).

Under certain well-described conditions, B cells will undergo apoptosis. Two experimental situations in which this has been investigated include the spontaneous apoptosis of primary B cells (ILLERA et al. 1993; MOWER et al. 1994) and the induction of apoptosis in the B-cell line WEHI-231, in which crosslinking of the B-cell receptor leads to programmed cell death (BOYD and SCHRADER 1981; FISCHER et al. 1994). The regulation of apoptosis in the WEHI-231 system is thought to involve the activity of NFkB and the expression of c-myc and bcl-x<sub>L</sub>, which are thought to be protective (FISCHER et al. 1994; GOTTSCHALK et al. 1994; FANG et al. 1995; LEE et al. 1995; MERINO et al. 1995). CpG treatment is a very strong rescuer of B cells from apoptosis in these experimental models. In the case of the WEHI-231 model, CpG DNA can be added as late as 8h after the apoptosisinducing signal of ligating-surface IgM without losing its anti-apoptotic activity (YI et al. 1996b; YI and KRIEG 1998a). This rescuing activity of CpG DNA appears to require the induction of NFkB, since protection is lost if cells are treated with inhibitors of NFkB activation (YI and KRIEG 1998a,b).

In addition to these changes, B cells activated by CpG DNA show increased expression of surface class-II major histocompatibility (MHC) molecules and the co-stimulatory molecules B7-1 and B7-2 (KRIEG et al. 1995; DAVIS et al. 1998). This suggests the possibility that CpG may enhance the antigen-presenting function of B cells directly. No other cell type is required for these activities of CpG DNA on B cells.

The responses of human B cells to CpG DNA generally appear to be very similar to those of murine cells, with the exception that the optimal motifs for stimulation are different (Fig. 1). ODNs with good human CpG motifs are very strong B-cell mitogens, but there is also a relatively non-sequence-specific stimulatory effect of the phosphorothioate backbone; this effect becomes evident at higher ODN concentrations of around 10 $\mu$ M (LIANG et al. 1996). Seemingly paradoxically, *Escherichia coli* DNA does not induce <sup>3</sup>H-thymidine incorporation by human B cells (LIANG et al. 1996) but does trigger B cells to proliferate, as detected by the CFSE (carboxyfluorescein succinimidyl ester) assay or by the expression of cell-surface CD86 (G. Hartmann and A. Krieg, manuscript in preparation). These seemingly contradictory results may be understood if we consider that the intracellular degradation of bacterial DNA (which, unlike phosphorothioate, is highly susceptible to degradation) releases free thymidine that effectively competes with

radioactive thymidine in the proliferation assay, thereby giving a false negative result (MATSON and KRIEG 1992). To avoid this effect in experimental assays using non-nuclease-resistant ODN, it is important to design appropriate readouts that will not be affected. In our experimental systems, phosphodiester-backbone CpG DNA shows strong and reproducible stimulatory activities on human cells that are CpG-sequence-specific even at high concentrations. In fact, many CpG motifs that will not activate human cells when present in an ODN with a phosphorothioate backbone are stimulatory when used in a phosphodiester or chimeric phosphorothioate/phosphodiester backbone (BALLAS et al. 1996; G. Hartmann and A.M. Krieg, manuscript in preparation).

#### 3.2 T-Cell Activation by CpG DNA

The effects of CpG DNA on T cells are much less clear. An initial report using partially purified T-cell populations (about 85% pure) reported the direct stimulation of cytokine secretion by CpG DNA in T cells (KLINMAN et al. 1996). However, subsequent studies by several labs have shown that, in fact, there is no direct activation of T cells by CpG DNA (LIPFORD et al. 1997b; SUN et al. 1998b). However, by inducing production of type-I IFNs from adherent cells, CpG DNA activates T cells to express CD69 and B7-2, although their proliferative responses to ligation of the T-cell receptor are reduced (SUN et al. 1998b). This functional inhibition resulted from the production of type-I IFN by antigen-presenting cells (SUN et al. 1998a). However, highly purified T cells that are stimulated through the T-cell receptor show synergistic proliferative responses to CpG DNA, indicating a mechanism through which CpG DNA could promote antigen-specific T-cell responses (BENDIGS et al. 1999). Further studies will be required to clarify the significance of these stimulatory and inhibitory activities of CpG DNA in vivo.

#### 3.3 NK-Cell Activation by CpG DNA

Early studies by Tokunaga and colleagues demonstrated that bDNA is a strong stimulus for NK lytic activity and for IFN- $\gamma$  production (Tokunaga et al. 1984, YAMAMOTO et al. 1992). However, these effects of CpG DNA are not observed with highly purified NK cells (BALLAS et al. 1996; Cowdery et al. 1996). Instead, these studies demonstrated that NK cell activation by CpG DNA was blocked by neutralizing antibodies to IL-12, TNF- $\alpha$ , and/or type-I IFNs. Since these cytokines are produced by adhering cells, this raised the possibility that the effect of CpG DNA on NK cells may be indirect. It is now clear that this is partially true, but it also appears that NK cells incubated with IL-12 together with CpG DNA are much more strongly stimulated than those incubated with an equivalent amount of IL-12 alone (Cowdery et al. 1996). Thus, as with T cells, NK cells appear to be costimulated by the combination of CpG DNA and specific activating cytokines. Interestingly, there is no such co-stimulatory activity for IL-2, which is otherwise an extremely potent activator of NK cells (Z. Ballas, G. Weiner, J. Cowdery, A.M. Krieg).

#### 3.4 Monocytes, Macrophages and DC Activation by CpG DNA

These antigen-presenting cell types appear to be directly stimulated by CpG DNA in the mouse to produce a variety of cytokines. A common theme that has emerged from studies of the cytokine responses of antigen-presenting cells (APCs) to CpG DNA is the prominence of Th1-like responses. For example, skin-derived DCs are stimulated by LPS to produce large amounts of TNF- $\alpha$  and relatively low levels of IL-12 but are stimulated by CpG DNA to produce very large amounts of IL-12 with relatively low amounts of TNF- $\alpha$  (JAKOB et al. 1998). APCs are also induced to produce IL-1 and IL-6 and to express increased levels of cell-surface class-II MHC and the co-stimulatory molecules B7-1 and B7-2 (STACEY et al. 1996; LIPFORD et al. 1997a; SPARWASSER et al. 1997a,b). These effects appear to be directly mediated, since no accessory cells are required.

Some differences are observed in CpG-induced activation of human cells. The cytokine response of human monocytes to CpG DNA is delayed compared to that of LPS. As measured by intracellular cytokine staining, LPS-activated monocytes produce high levels of TNF- $\alpha$  and IL-6 within 4h; CpG-activated monocytes have no detectable cytokine response at this time point (HARTMANN et al. 1999). Instead, the human monocyte cytokine response to CpG DNA becomes detectable at approximately 12–18h and is lower in magnitude than that which can be induced by LPS (HARTMANN et al. 1999). Human DCs show much faster kinetics in their responses to CpG DNA (Hartmann).

#### 4 Interaction of CpG DNA with other Cell Activation Pathways

## 4.1 Synergy versus Antagonism between CpG DNA and the B-Cell Antigen Receptor

A common biological theme in cellular responses to environmental stimuli is the interaction between different cell signaling pathways, which can synergize or interfere with one another. In the case of CpG DNA, its role as a "danger signal" that indicates the presence of infection makes it logical that the immune system should have evolved to enhance cellular activation in response to foreign antigens that are encountered in the presence of CpG DNA. Indeed, although high concentrations of CpG DNA can drive more than 95% of B cells into the cell cycle, at low concentrations of CpG DNA synergy with the B-cell receptor (BCR) is very prominent, resulting in at least a tenfold increase in the proliferation and Ig secretion of B cells that have also been activated through their antigen receptor

(KRIEG et al. 1995). Thus, although CpG DNA by itself is, of course, a nonantigen-specific signal, it preferentially promotes the induction of antigen-specific antibody responses. CpG DNA and the BCR also synergize for induction of B-cell IL-6 secretion (Y1 et al. 1996c). With the identification of the intracellular kinase pathways that are activated by CpG DNA, it has become possible to determine the molecular mechanism of the synergy at a more proximal level. CpG DNA and the BCR synergize for inducing MKK3, MKK4, MKK6, and JNK, but not ERK (Y1 and KRIEG 1998b; A.K. Yi and A.M. Krieg, unpublished data).

Although mature B cells are typically activated by crosslinking of the BCR, immature B cells may be induced to undergo apoptosis (Norvell et al. 1995). This process is thought to be important in maintaining immune tolerance to self antigens present in the bone marrow, which will prevent autoreactive B cells from maturing, thereby reducing the risk of autoimmunity. A B-cell line with an immature phenotype, WEHI-231, has been commonly used as a model of this activation-induced apoptosis, since crosslinking of the BCR causes WEHI-231 cells to undergo rapid growth arrest followed by apoptotic death (BOYD et al. 1981; SCOTT et al. 1985). Apoptotic death has been linked to changes in c-myc gene expression and to a loss of NFkB activity (McCormack et al. 1984; Fischer et al. 1994; Lee et al. 1995; SCHAUER et al. 1996; WU et al. 1996). These studies have shown that agents that restore NFkB activation, such as LPS and CD40L, can prevent apoptosis of WEHI-231 cells following BCR ligation. CpG DNA, which also induces c-myc expression and NFkB activation, also protects WEHI-231 cells from cell death induced by BCR ligation (YI and KRIEG 1998a; YI et al. 1996b). BCR ligation in WEHI-231 cells does not immediately lead to irreversible apoptosis, since CpG DNA can still provide substantial protection even when added 8h after BCR ligation (Y1 et al. 1996b). The protective effect of CpG DNA is associated with increased binding of the p50/c-Rel heterodimer of NFkB as compared to the p50 homodimer, which is dominant after BCR ligation alone (Y1 and KRIEG 1998a).

Given its ability to interfere with apoptosis in this model of tolerance induction, it might be expected that CpG DNA would interfere with the deletion of autoreactive cells in vivo and trigger autoimmune disease. Surprisingly, this is not the case; CpG DNA does not induce lupus in normal mice nor flare lupus in genetically lupus-prone mice. In fact, the cytokines induced by CpG DNA actually reduce disease severity (GILKESON et al. 1996; MOR et al. 1997; GILKESON et al. 1998)! Thus, it appears that additional mechanisms that are not blocked by CpG DNA must protect against autoimmunity in vivo. Nonetheless, studies in the WEHI-231 system show the complexity of the signaling pathways induced by CpG DNA and the BCR and the ways in which these pathways can antagonize as well as synergize.

#### 4.2 Interaction of CpG DNA with Other B-Cell Activation Pathways

Aside from the BCR, several other cell-surface molecules on B cells are known to regulate their activation. For example, B cells are stimulated by crosslinking of

CD40 or class-II MHC molecules, both of which can also synergize with activation through the BCR. Therefore, we also examined whether CpG DNA may synergize with these signaling pathways. These studies showed that CpG DNA synergized with B-cell activation through class-II MHC but not through CD40 (A.M. Krieg and G. Bishop, unpublished data).

B cells can also be affected by cytokines, such as IFN- $\gamma$ . The effects of IFN- $\gamma$  on B cells can be rather complex, with both positive and negative effects reported, depending on the experimental system (LEIBSON et al. 1984; COWDERY et al. 1992; CHACE et al. 1993). For example, IFN- $\gamma$  attenuates LPS-induced B-cell activation (CHACE et al. 1993). In contrast, IFN- $\gamma$  promotes B-cell activation by CpG DNA, as demonstrated by the fact that this activation is weaker in mice, which are genetically deficient in IFN- $\gamma$  (Y1 et al. 1996a). Since CpG DNA is, indirectly, a strong stimulus for NK cell production of IFN- $\gamma$ , this would tend to promote antigen-specific B-cell responses. CpG DNA synergizes with LPS for the activation of B-cell proliferation and Ig secretion (A.-K. Yi and A.M. Krieg, unpublished data).

#### 4.3 Interaction of CpG DNA with Monocyte Activation by LPS

LPS is a potent stimulus for monocyte cytokine production, which superficially shares many similarities with CpG DNA. Both agents are mitogenic for mouse B cells and induce high levels of monocyte TNF- $\alpha$ , IL-6, and other cytokines. However, differences between the biologic activities of these agents have been noted. For example, macrophages are induced to express inducible nitric oxide synthase by LPS, but CpG DNA does not exert this effect unless the macrophages have also been primed with IFN- $\gamma$  (STACEY et al. 1996, 1998). Conversely, RAW264 macrophage cells are stimulated to produce IL-12 by CpG DNA but not by LPS (ANITESCU et al. 1997). More marked differences have been observed in the responses of human cells to LPS and CpG. In human PBMCs treated with LPS, the monocytes show rapid induction of TNF- $\alpha$  and IL-6 production within a few hours, but the production of these cytokines in response to CpG DNA is delayed until about 18h (HARTMANN et al. 1999). In fact, this difference in the kinetics of cytokine secretion in response to LPS and CpG DNA can be used to detect the presence of LPS in synthetic ODN or bDNA samples.

Several investigators have reported synergy between CpG DNA and LPS for induction of cytokine production in vitro and in vivo and for induction of the systemic inflammatory-response syndrome in vivo. Thus, although CpG DNA by itself is quite non-toxic, it effectively primes for the Schwartzman reaction, as demonstrated by a very high mortality in mice given a sublethal dose of endotoxin 4h after a dose of CpG DNA (COWDERY et al. 1996). This mortality is associated with synergistic induction of TNF- $\alpha$ , which is dramatically enhanced by administration of both CpG DNA and LPS (COWDERY et al. 1996; HARTMANN et al. 1996; SPARWASSER et al. 1997a).

#### 5 Conclusion

Our understanding of the mechanism of action of CpG DNA has advanced remarkably far in the few years since the discovery that the immune system recognizes bDNA as "foreign" because of its content of unmethylated CpG motifs. The ability to detect CpG DNA lies both in cells of the innate immune system, such as monocytes and macrophages, but also in the acquired immune system, where it is a remarkably potent B-cell mitogen. As a result, CpG DNA not only triggers rapid innate immune responses that can provide a sort of non-specific innate protection against challenge with a broad range of infectious pathogens (KRIEG et al. 1998a; ZIMMERMAN et al. 1998; ELKINS et al. 1999), it also strongly enhances the generation of antigen-specific B-cell responses (KRIEG et al. 1995). In fact, CpG DNA has now emerged as an astonishingly potent and non-toxic vaccine adjuvant whose therapeutic effects are even more marked than those of the "gold standard" (complete Freund's adjuvant), but without the associated toxicity (CHU et al. 1997; LIPFORD et al. 1997b; ROMAN et al. 1997; WEINER et al. 1997; DAVIS et al. 1998; SUN et al. 1998a).

Several of the molecular signaling pathways through which the CpG DNA signal is transduced have been identified, as reviewed above. Perhaps the most important immediate question for understanding the molecular mechanism of CpG DNA is the identification of its receptor. It appears that this receptor is a cytoplasmic protein or belongs to a family of proteins, and it is hoped that, with its identification, the remaining pieces of the activation pathway will gradually be filled in.

Along with our understanding of the mechanism of action of CpG DNA has come increasing evidence of its potent therapeutic effects in a wide variety of animal models. As of this writing, CpG will imminently be entering human clinical trials, which will be the final test for whether this fascinating immune modulator can be used successfully to improve human health.

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## Oligodeoxyribonucleotides with 5'-ACGT-3' or 5'-TCGA-3' Sequence Induce Production of Interferons

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### **1** Introduction

More than 1 billion children in 182 countries throughout the world have been vaccinated with the Bacillus of Calmette and Guerin (BCG) against tuberculosis. BCG appears to be able to elicit protective cell-mediated immune responses in the absence of progressive disease. In addition, experimental uses of BCG and related mycobacteria have greatly contributed to the progress of modern immunology. Trials of cancer immunotherapy with BCG were carried out on a worldwide scale in the 1970s and contributed much information to the fields of basic and clinical immunology (BAST 1974). Evidence foreshadowing our finding of senseless immunostimulatory DNA was also obtained by the experimental use of BCG.

Until recently, DNA had only been considered as the blueprint of life and was thought to be immunologically uniform and essentially inert. However, recent advances in research on antisense DNA and gene vaccination changed this view; this research indicated that DNA can operate as a genetic-code-based intracellular

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signal. Our findings – that immunocytes of mice and humans can be stimulated in a senseless manner by DNA derived from BCG, resulting in interferon (IFN) production and tumor-growth inhibition (TOKUNAGA 1984; SHIMADA 1986; YAMAMOTO S 1988) – further transformed this conceptual framework (STEIN 1995; PISETSKY 1996; KRIEG 1996b). Using a variety of synthetic, single-stranded oligo-deoxyribonucleotides (ODN), we reported that hexamer palindromic sequences with 5'-ACGT-3' or 5'-TCGA-3' motifs are essential for inducing IFNs (KURA-MOTO 1992; TOKUNAGA 1992). We also reported that bacterial, viral, or invertebrate DNA (but not vertebrate or plant DNA) can induce IFNs, and suggested that this mechanism may contribute to the self-non-self discrimination for foreign DNA (YAMAMOTO S, YAMAMOTO T, SHIMADA S, et al. 1992).

Interest in this line of study was galvanized by the demonstration of murine B-cell activation by CpG motifs in bacterial DNA (KRIEG, YI, et al. 1995). Today, immunostimulation with DNA is one of the most active topics in the research fields of host-defense mechanisms, plasmid vaccination, etiology of certain diseases, and therapeutic application. In this review, we will describe how our BCG studies led to the discovery of immunostimulatory DNA.

#### 2 Antitumor Activity of DNA from BCG

In the 1970s, trials of cancer immunotherapy with BCG were carried out energetically and extensively. The mode of antitumor action of BCG was considered to be host mediated, and major effector cells were thought to be macrophages activated by lymphokines produced by BCG-sensitized lymphocytes. Intratumor administration of viable BCG was effective, but the trials were given up because the side reactions induced by viable BCG were recognized to be severe. Many investigators made efforts to isolate components of BCG or related bacteria possessing antitumor activity but not side effects. We found that BCG cytoplasm precipitated by streptomycin contained a substance strongly active against mouse and guinea pig syngeneic tumors. This fraction was purified, and a fraction possessing strong activity but few side effects was obtained. We named this fraction MY-1 but, in this review, we call it BCG-DNA.

The BCG-DNA was mostly composed of nucleic acids (98%); its protein content was only 0.15%, and the sugar content was 0.20%. The elution profile indicated that BCG-DNA was distributed over a wide range of molecular weights, and its elution peak corresponded to molecular weights of about  $2-4 \times 10^5$ kDa. The DNA was single stranded.

The antitumor activities of BCG-DNA were examined against various syngeneic tumors of mice or guinea pigs. Repeated intra-regional injections with BCG-DNA regressed or inhibited almost all of the various solid tumors. Repeated subcutaneous injections against leukemia were also effective (TOKUNAGA 1984; SHIMADA 1985). We were very hesitant to conclude that DNA itself is active against tumor, because few incidences of biologic activity of DNA had been reported before, although synthetic double-stranded RNA, such as polyI:C, and synthetic single-stranded poly(dG,dC) were known to induce IFN and activate both natural killer (NK) cells and macrophages (TOKUNAGA 1988). In the early days of IFN research, Isaacs and his co-workers had observed that foreign DNA induced IFN, but they later questioned their observation. We further treated BCG-DNA with either RNase or DNase, purified the products by shaking with chloroform/alcohol and eluting with Sephadex G-100, and obtained the fractions named RNase-digest or DNAse-digest, respectively (TOKUNAGA 1984). The RNase-digest contained mostly DNA (97.0%), and the DNase-digest contained 97.0% RNA and 1.5% DNA. Amazingly, RNase-digest produced IFN and showed antitumor activity, as did BCG-DNA, while DNase-digest lost its activity completely, suggesting strongly that the activity is a result of its DNA.

Direct in vitro cytotoxicity of DNA extracted from various cells had been reported by others, but BCG-DNA (1000µg/ml) showed no cytotoxicity in vitro for cells of 13 cell lines, suggesting that the action of BCG-DNA is host mediated (SHIMADA 1985). An intra-peritoneal (ip) injection of BCG-DNA (100µg) into mice rendered peritoneal cells cytotoxic (SHIMADA 1986; YAMAMOTO 1988). The effector activity was destroyed by anti-asialo-GM1 antiserum or carrageenan in vitro, but not by anti-Thy1.2 or carbonyl-iron uptake, suggesting that the effector cells are NK cells (SHIMADA 1986; YAMAMOTO 1988). Ip injection with DNase-digest of BCG-DNA did not bring such NK activation.

Intra-dermal (id) injection of BCG-DNA induced in situ infiltration of mononuclear cells (most of which were asialo-GM1-positive) and lasted for a week. The infiltrating cells were also positive for Ly-5 and partially positive for Thy1.2 but negative for Mac-1, Ia, mu-chain, Lyt-1, Lyt-2, L3T4, and Fc receptor II. They contained neither peroxidase nor nonspecific esterase, suggesting that the cells were NK. DNase destroyed all these effects of BCG-DNA (KURAMOTO 1989, 1992).

Peripheral blood lymphocytes (PBL) from healthy human donors and 20 cancer patients were assessed for NK activity after 24h of incubation with BCG-DNA, RNase-digest, or DNase-digest. BCG-DNA or RNase-digest (but not DNase-digest) augmented the NK activity of PBL from healthy donors and all cancer patients, although the degree of augmentation varied depending upon the PBL used (MASHIBA 1988).

When normal mouse spleen cells were incubated with BCG-DNA or RNase-digest (but not DNase-digest) for 20h, the cells produced both IFN and macrophage-activating factor (MAF). IFN was measured by observing inhibition of the cytopathic effects of vesicular stomatitis virus (VSV); MAF was measured by observing the extent to which normal peritoneal macrophages became cytotoxic towards tumor cells (YAMAMOTO 1988). The MAF activity was destroyed by anti-IFN- $\gamma$  antiserum, and the anti-virus activity was eliminated by anti-IFN- $\alpha/\beta$ .

# **3** Particular Base Sequences in BCG-DNA Required for Immuno-Stimulation

#### 3.1 Significance of Palindromic Sequences Having -CG- Motifs

We wondered whether or not the activity of BCG-DNA is base-sequence dependent. Based on the fact that the molecular sizes of the DNA molecules contained in BCG-DNA preparation were distributed over a broad range but peaked at 45 bases (TOKUNAGA 1984), we synthesized 13 kinds of 45-mer singlestranded ODNs having sequences randomly selected from the cDNA encoding BCG proteins and tested their ability to produce IFN- $\alpha$  and - $\gamma$  and augment NK cells (TOKUNAGA 1992). Amazingly, 6 of the 13 ODNs activated NK cells, while the others did not. IFN-inducing activities paralleled these results, suggesting that some stereochemical structures constructed by particular sequences are required for expression of the biologic activity (TOKUNAGA 1992). ODNs that were biologically active in vitro also inhibited tumor growth (KATAOKA 1992).

We further synthesized a variety of 30-mer single-stranded ODNs with sequences chosen from the cDNA encoding proteins of various species, including humans. ODNs having palindromic sequences of either GACGTC, AGCGCT, or AACGTT were active, whereas ODNs with ACCGGT were inactive. When a portion of the sequence of the inactive ODNs was substituted with GACGTC, AGCGCT, or AACGTT (but not ACCGGT), the ODN acquired biologic activity (YAMAMOTO S, YAMAMOTO T, KATAOKA T, et al. 1992).

The 30-mer ODNs active in mice were also active in humans. IFN activity was detected in the culture fluid of PBL after 8h, and the amount of IFN reached the maximum after 18h. IFN- $\alpha$  was predominantly produced, and small amounts of IFN- $\beta$  and IFN- $\gamma$  were also found (YAMAMOTO T, YAMAMOTO S, KATAOKA T, KOMURO K, et al. 1994).

In order to know the minimum and essential sequences responsible for the biological activity, the palindromic sequence (GACGTC) in an active 30-mer ODN, 5'-ACCGAT GACGTC GCCGGT GACGGC ACCACG-3', was replaced with each of the 63 theoretically possible 6-mer palindromic sequences, and they were tested for NK-enhancing ability (KURAMOTO 1992). The analogue 30-mer ODNs that enhanced NK activity more strongly than the original ODN were 8 ODNs with one of the following palindromic sequences: AACGTT, AGCGCT, ATCGAT, CGATCG, CGTACG, CGCGCG, GCGCGC, and TCGCGA. All included one or more 5'-CG-3' motif(s). The converse was not true; for instance, ACCGGT and CCCGGGG were impotent. Palindromes composed entirely of adenines (A) and thymines (T) and those with a sequence of either Pu (purine)-Pu-Pu-Pu (pyrimidine)-Py-Py or Py-Py-Pu-Pu-Pu were generally unfavorable for the activity. Since stacking between Py is less stable than that between Pu or Pu-Py, stable helical structures formed by the palindromes including the 5'-CG-3' sequence but without a Py-Py-Py sequence seemed to be favorable (KURAMOTO

1992). BALLAS et al. pointed out that palindromic sequences are not always necessary for NK activation (BALLAS 1986).

#### 3.2 Significance of Sequences Outside of Potent Hexamers

Sequences outside of hexamer palindromic sequences were also meaningful; studies were designed to get a comprehensive picture of the sequence requirements of ODNs. A comparison was made of the NK-augmenting activities of single-stranded 30-mer homo-oligomers, all of which possess one of the potent sequences AACGTT, CGATCG, GACGTC, or ATCGAT at the center position (KURAMOTO 1992). The homo-oligomers with no palindromic structures showed no activity. Oligo-G having a potent palindromic sequence showed the highest activity, but oligo-A and oligo-C having this sequence gave only marginal activity. No activity was seen in the oligo-G containing an impotent palindrome. These results indicate the independent and cooperative effects of palindromic and extra-palindromic sequences on the activity of ODN (KURAMOTO 1992).

The effects of the number and location of the palindromic sequences were also investigated. Among the 30-mer oligo-G nucleotides containing a different number of AACGTT palindromes, an ODN with one particular palindrome showed the strongest activity. The ODN with AACGTT at the 5'-end or 3'-end showed slightly stronger activity than the ODN with the sequence in the center, although the activity was influenced more drastically by the number of palindromes than by their location (KURAMOTO 1992).

It is unlikely that the cooperation between the palindromic and extra-palindromic sequences is due to secondary structures, such as bulges and hairpin loops possibly composed of these sequences, for several reasons. First, the thermostability of secondary structures was not correlated with the activity of ODNs, as judged by temperature-gradient gel electrophoresis. Second, there was no correlation between the activity and the preferred secondary structure of the ODN as predicted by the thermodynamic calculation. Third, the activity of a single-stranded ODN is comparable to that of its double-stranded counterpart, although the latter is less likely to form secondary structures than the former.

The effect of ODN base length on the ability of ODNs to induce IFN was also investigated. Ten kinds of 12- to 30-mer nucleotides having the AACGTT sequence were synthesized. Immunostimulatory activity was observed in ODNs of 18 bases or more in length and was proportional to the base length, with a maximum of 22– 30 bases. However, the ODNs 16 bases or less in length were not active even if they possessed an AACGTT sequence (YAMAMOTO T, YAMAMOTO S, KATAOKA T and TOKUNAGA T 1994a).

The NK-enhancing and IFN-producing activities of 30-mer single-stranded oligo-G having AACGTT (ODN-1) were inhibited by a homo-oligomer of G, dextran sulfate, and polyvinyl sulfate. ODN-1 inhibited acetyl-low-density-lipoprotein binding to the scavenger receptor, suggesting that the sequences outside of the potent hexamer may play a role in the binding of ODN to cell surfaces (KIMURA 1994).

#### 4 DNA from Bacteria, but not Vertebrates, Activates Immunocytes

In 1984, we noticed that DNA-rich fractions from various species of bacteria exhibited biological activities similar to those of BCG-DNA but DNA from calf thymus or salmon testis did not. Since these results were difficult to explain, we hesitated to publish them. We prepared DNA fractions from six bacterial species, calf thymus, and salmon testis, using exactly the same procedures used in the preparation of BCG-DNA and by other methods. The DNA fractions from all of the bacteria caused remarkable elevations of NK activity in mouse spleen cells and induced generation of both MAF and IFN in a concentration-dependent manner. However, the DNA fractions from calf thymus and salmon testis did not show any activity. The bacterial DNA fractions caused tumor inhibition, but the vertebrate DNA fraction did not. The activities of the bacterial DNA were completely abrogated by pre-treatment with DNase but not RNase (YAMAMOTO S, YAMAMOTO T, SHIMADA S, et al. 1992).

Further, we examined the increased number of DNA samples from a variety of animals and plants. As illustrated in Fig. 1, all of the DNA from four species of bacterium augmented NK activity and induced IFN, while the DNA from ten kinds of vertebrate, including three fish and five mammals, showed no such activities. The DNA from four species of plant were also inactive (YAMAMOTO S, YAMAMOTO T, SHIMADA S, et al. 1992).

To explain such differences of biological activities among DNAs from different sources, we set up the following six hypotheses: (i) the bacterial DNA fraction might contain toxic substances like lipopolysaccharide (LPS); (ii) the molecular size of the DNA contained in the DNA fractions might be different in bacteria and animals, a fact which might influence biologic activities; (iii) DNA from animals might be more sensitive than bacterial DNA to the DNase employed, which was derived from higher animals; if so, the bacterial DNA would persist longer in the cultures of mouse spleen cells; (iv) G + C ratios might influence the activities; (v) the majority of -CG- motifs are known to be methylated in vertebrates but not in bacteria, and the presence of methyl cytosine may obstruct the activity of -CG- motifs; (vi) the frequency of a particular base sequence essential for the biologic activity of the DNA may be different.

We examined all of the six possibilities. (i) The activities of the bacterial DNA fractions were not influenced by the presence of polymyxin B, an inhibitor of LPS; these activities were present even in spleen cells from LPS-insensitive C3H/HeJ mice, indicating that the activities were not ascribed to LPS contamination. (ii) The profiles of elution patterns obtained by agarose-gel electrophoresis were essentially the same in all of the DNA fractions from eight different sources. (iii) The absorbance of BCG-DNA and calf thymus DNA at 260nm before and after DNase treatment was the same. (iv) The (G+C) ratios of the bacterial DNAs used varied from 70.7% for *Streptomyces aureofaciens* to 27.1% for *Clostridium perfringens*,


Induction of NK and IFN activity of mouse spleen cells incubated with DNA samples from various sources

Fig. 1. BALB/c mouse spleen cells  $(1 \times 10^7 \text{cells/ml})$  were incubated with  $100\mu\text{g/ml}$  of each of the DNA samples for 20h and centrifuged. The cell fractions were used for natural killer (NK) assays, and the culture supernatants were used for interferon (IFN) assays. NK cell activity was measured by a 4-h <sup>51</sup>Cr-release assay using YAC-1 lymphoma cells as target cells. The level of IFN in the supernatant was measured in terms of its ability to inhibit the cytopathic effects of vesicular stomatitis virus

and those in calf and salmon DNAs are 50.2% and 40.2%, respectively. No correlation between the (G+C) ratio and the activity of DNA was found. (v) We previously reported that a synthetic, single-stranded poly(dG,dC) is active in vitro. Therefore, we synthesized poly(dG,dC) where all cytosines were substituted with methyl cytosines. The resulting poly (dG,methyl-dC) augmented NK activity and induced IFN as effectively as the original poly(dG,dC). Furthermore, we synthesized both a 30-mer nucleotide having 5'-AACGTT-3' and its analogue, in which C was replaced with methyl cytosine; both ODNs could induce IFN equally. Accordingly, we thought that the fifth hypothesis was incorrect. In 1994, we found that certain hexamers, acting alone, could induce IFN when they were encapsulated in liposomes (YAMAMOTO T, YAMAMOTO S, KATAOKA T and TOKUNAGA T 1994b). Using this liposome system, we recognized that methylation of the cytosine of AACGTT resulted in a significant decrease of the activity (SONEHARA 1996). We also found that, when *Escherichia coli* DNA was incubated with CpG methylase, the resulting DNA reduced its IFN-inducing activity with incubation time. Therefore, we now think that hypothesis (v) must be one of the reasons for the difference in activity between vertebrate and non-vertebrate DNAs. (vi) We tried to test the incidence of the nine potent palindromic sequences of cDNA reported in the GenBank DNA database. Among 17 kinds of living things registered in the database, we randomly chose one or more sequences, and the incidences of the potent palindromic sequences in all the hexamer palindromic sequences were expressed as the number per 1000 base pairs. The incidence of the potent palindromic sequences in all the DNAs from the vertebrates and plants was less than 1.0, while the incidence in most of the bacteria, viruses, and silkworms was larger than 1.0. There were some exceptions; the incidence, for instance, in the cDNA from  $M_{y-1}$ coplasma pneumoniae and Cl. perfringens was very low (0.2-0.4), but the activities of these DNAs were high. This may be due to the fact that our analysis was limited to only parts of the genomic DNA. In addition, particular 8-, 10- or 12-mer palindromic sequences show stronger immunostimulatory activities than certain hexamers (SONEHARA 1996). Bird stated that CpG in bulk vertebrate DNA occurs at about one-fifth of the expected frequency (BIRD 1986). We think that the different frequencies of potent sequences in DNA in vertebrate and invertebrate DNAs must be one reason for their different activities.

### **5** Immunostimulatory Sequences

As described, we reported first that hexamer palindromic sequences having -CGmotifs are essential for IFN production and NK augmentation, with some exceptional cases. Krieg et al. used the term "CpG motif" for the basic structure of B-cell stimulation (KRIEG 1995); this motif consists of the CpG dinucleotide flanked by two 5' purines (preferably a GpA dinucleotides) and two 3' pyrimidines (preferably a TpT). Ballas et al. (BALLAS 1996) examined the necessity of palindromic sequences for NK activation and observed that the presence of a hexamer palindrome is irrelevant for NK activation; for instance, GTCGTT and GACGTT are active, and GACGTC is inactive. They concluded that unmethylated CpG motifs, but not palindromic sequences, are definitely required, and two flanking bases at the 5' and 3' ends showed stringent requirements. Boggs et al. tested various sequences of ODNs and confirmed that the CpG motif was stimulatory for NK cells only in specific sequence contexts (BoGGS 1997). Monteith et al. tested the ability of various ODN sequences to induce splenomegaly and B-cell stimulation; a 5'-AACGTT-3' palindrome was the most effective (MONTEITH 1997).

Since we found previously that a hexamer palindromic ODN (5'-AACGTT-3'), acting alone, induced IFN from mouse spleen cells when added with cationic liposomes (YAMAMOTO 1994), 32 kinds of hexamer palindromic ODN were synthesized and tested for their IFN-inducing ability in the presence of liposomes (SONEHARA 1996). The sequence-activity analysis revealed a conspicuous dependency of the IFN-inducing activity on the presence of CG motifs, and the sequences showing the highest activity were able to express as NACGTN and NTCGAN,

where N represents any complementary pair of bases. Moderate activity was observed in ACGCGT and TCGCGA. The activity of CGNNCG sequences was quite low. These results agree (with some exceptions) with previous results obtained with the use of the 64 theoretically possible kinds of hexamer palindromic sequences contained in a 30-mer ODN (SONEHARA 1996). An exception was that CGATCG showed only marginal activity, although it was among the most potent sequences in the previous experiment. We think that the results obtained with liposomeencapsulated ODN definitely reveal the sequence-activity relationship of IFN-inducing ODN, because possible effects of the flanking sequence, which is important for efficient uptake of ODN (KIMURA 1994), can be excluded, and a possible influence of secondary and tertiary structures on the activity is minimal for short, hexameric ODNs. In addition, the base-substitution study is free of the complication in which elimination of a palindrome gives rise to another palindrome. To obtain more precise information, each of the bases of AACGTT except CG were replaced with a distinct base and encapsulated in a liposome. Original palindromic sequences showed the highest activity, and the activity was markedly reduced by any single base substitution except substitution of G for the first A (that is, GA-CGTT; SONEHARA 1996). A high activity of GACGTT was also reported in other laboratories (BALLAS 1996; BOGGS 1997). We noticed, however, that some incomplete palindromic sequences, such as AACGTT and CACGTT, showed reduced but significant activities (SONEHARA 1996). Nevertheless, we found that one 30-mer ODN (C13, which encodes a regulator protein of IFN-ß and has the sequence 5'-GGGCAT CGGTC GAAGTG AAAGTG AAAGTGA-3') showed extremely strong activity. This ODN contains no hexamer palindromic sequence but does contain TCGA.

It is also true that the formula, Pu-Pu-CG-Py-Py, proposed by Krieg et al. (KRIEG 1996b) and Klinman et al. (KLINMAN 1996), which has been used widely by many investigators, has many exceptions. At present, therefore, the general term "immunostimulatory sequence (ISS) of DNA" (SATO 1996; ROMANN 1997; TIGHE 1998) or "CpG dinucleotides in particular base contexts" (SATO 1996; BOGGS 1997; YI and TUETKEN 1998) may be appropriate. Therefore, we will use "ISS DNA" in this review.

After the observation that ISS DNA induces mouse B cells to proliferate and secrete immunoglobulin (KRIEG 1995), it was reported that the same ODNs induced the rapid and coordinated secretion of interleukin 6 (IL-6), IL-12, and IFN- $\gamma$  but not IL-2, IL-3, IL-4, IL-5, or IL-10 (KLINMAN 1996). It was also reported that bacterial DNAs stimulate macrophages/monocytes to secrete IL-12 and tumor necrosis factor  $\alpha$  (TNF- $\alpha$ ), resulting in IFN- $\gamma$  production from NK cells (HALPERN 1996), and that IFN- $\gamma$  promotes IL-6 and immunoglobulin M (IgM) secretion responses to ISS-DNA (YI and CHASE 1996). Bacterial DNA induces macrophage IL-12 production which, in turn, stimulates NK cell IFN- $\gamma$  production (CHASE 1997). IL-10 inhibits IL-12 secretion induced by bacterial DNA (ANITESCCU 1997). Nitric oxide production from macrophages in response to bacterial DNA is absolutely dependent on IFN- $\gamma$  priming (SWEET 1998). ISS DNAs trigger maturation and activation of mouse dendritic cells (SPARWASSER 1998).

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Interestingly, in the field of plasmid DNA immunization, preferential induction of T-helper 1 (Th1) response and inhibition of specific IgE formation by plasmid DNAs were reported (RAz 1996). Sato et al. pointed out that the immunogenicity of plasmid DNA required short ISSs that contain a CpG in a particular base context (SATO 1996). Roman et al. showed that particular ODNs potently stimulate immune responses to administered antigens; they suppress IgE synthesis but promote IgG and IFN- $\gamma$  production and initiate IFN- $\gamma$ , IFN- $\alpha$ , IFN- $\beta$ , IL-12, and IL-18 production, resulting in fostering of Th1 responses and enhancement of cell-mediated immunity (ROMAN 1997).

Intracellualr mechanisms of immunostimulatory DNA are still unclear. We showed that scavenger receptors of mouse macrophages are the binding sites on cell surfaces in the particular case of G homo-oligomers having –AACGTT– (KIMURA 1994). In addition, we observed that bone-marrow cells from scavenger-receptor knockout mice could be stimulated with ISS ODN. However, it was reported that Mac-1 protein is an ODN-binding protein on polymorphnuclear cells, although the base sequence of ODN is unclear (BENIMETSAYA 1997).

It seems clear that ISS ODNs enter into cells and react with the cytoplasmic and nuclear apparatus. In our experiments, a high level of IFN was produced by a hexamer ODN alone when entrapped in liposome (YAMAMOTO T, YAMAMOTO S, KATAOKA T, TOKUNAGA T 1994b), suggesting that the ODN worked inside the cells. The necessity of cellular uptake of the DNA was proven in B cells (KRIEG 1995) and macrophages (YI and KRIEG 1998).

In the early days of our research, we speculated that a potent, palindromic sequence may bind to a certain negative transcriptional factor like IkB, or a suppressive factor released from the IFN regulatory gene, resulting in activation of the IFN gene (YAMAMOTO S, YAMAMOTO T, KATAOKA T, et al. 1992). Although many transcriptional factors, such as *c-myc* and AP-1, were known to recognize short palindromic sequences, no transcriptional factor regulating IFN-gene expression was known to recognize active ODN sequences. Then we proposed an alternative candidate: the cyclic adenosine monophosphate (cAMP) response-element (CRE)-binding protein, which recognizes the CRE (GACGTC; SONEHARA 1996). Our preliminary data showed that IFN induction by ISS ODN is inhibited by forskolin, a cAMP agonist (SONEHARA 1996).

Yi et al. reported that ISS-DNA-induced IL-6 production from B cells and monocytes is mediated through the generation of reactive oxygen species (ROSs) (YI, TUETKEN et al. 1998). The ROS burst is linked to the activation of nuclear factor  $\kappa B$  (NF $\kappa B$ ), which induces leukocyte gene transcription and cytokine secretion. It was also stated that ISS DNAs taken up by macrophages activate a signaling cascade leading to activation of NF $\kappa B$  (YI and KRIEG 1998). The inhibitors of NF $\kappa B$ , I $\kappa B$ - $\alpha$  and I $\kappa B$ - $\beta$ , are degraded rapidly in cells treated with ISS ODN.

It was found that ISS ODNs inhibit apoptosis of B cells (MACFARLANE 1997; YI, CHANG et al. 1998) via modulation of  $I\kappa B-\alpha$  and  $I\kappa B-\beta$  and sustained activation of NF $\kappa$ B/c-Rel. Further studies on the identification and characterization of the sequence-dependent recognition mechanism of ODN within the cell are expected.

It is generally known that vaccines consisting of viable, attenuated bacterium or virus can induce strong and persistent immunity but produce relatively severe side effects. However, killed vaccines are relatively safe and induce antibody responses but induce cell-mediated immunity with difficulty. Efforts have been made to find effective and safe immunoadjuvants for devising vaccines consisting of components, recombinant subunits, or synthetic peptides. However, only aluminum salts and liposomes have been approved for use in humans. Recently, immunization with antigen-encoding plasmid DNA was found to induce immune responses to the encoded foreign proteins. At present, numerous numbers of plasmid-encoding immunogens of bacteria, viruses, parasites, and tumors are under investigation (GREGORIADIS 1998; TIGHE 1998). Many of them could induce immunity against infection, and clinical trials for treating or protecting against carcinoma and infection with herpes, influenza, hepatitis B, and human immunodeficiency viruses are in progress.

In 1996, Sato et al. reported that plasmid vectors expressing large amounts of gene product do not necessarily induce immune responses to the encoded antigens but require short ISS DNA (SATO 1996). Their effective plasmid contained two repeats of the palindromic CpG hexamer 5'-AACGTT-3'. Also, ISS DNA was reported to stimulate Th1 responses (ROMAN 1997). After these, many reports showed that the ISS within the plasmid backbone stimulates immune responses to the antigen encoded by the plasmid (CARSON 1997; CHU 1997; KLINMAN 1997; LIPFORD, BEAURE et al. 1997; DENNIS 1998; KRIEG 1998). ISS DNA stimulates immune responses to administered antigens, including  $\beta$ -galactosidase and HBs antigen (LECLERC 1997), the idiotype of surface IgM of a B lymphoma (WEINER 1997), intracelluar pathogens, such as *Listeria monocytogenes* (KRIEG, LOVE – HOMAN et al. 1998), and formalin-inactivated influenza virus (MOLDOVEANU 1998).

One of the greatest advantages of ISS DNA as an immunoadjuvant is that ISS DNA tends to stimulate the generation of Th1-biased immune responses to antigens that are encoded by the plasmid or are administered. ISS DNA suppresses IgM synthesis but promotes IFN- $\alpha$ , - $\beta$ , and - $\gamma$ , IL-12, and IL-18, all of which foster Th1 responses and enhance cell-mediated immunity.

Since ISS DNA is known to induce both IL-12 and TNF- $\alpha$ , attempts were made to find DNA sequences which differentially activate IL-12 versus TNF- $\alpha$ cytokine production in antigen-presenting cells (LIPFORD, SPARWASSER et al. 1997). They described a single-stranded 18-mer ODN with this characteristic and proposed it as the prototype for a useful ISS DNA that lacks harmful side effects; it contains a hexamer of GACGTT.

Krieg et al. pre-treated mice with ISS ODN and then challenged them ip with a lethal dose of *L. monocytogenes*. A 2–4 log decrease in the number of bacteria in the liver and spleen was brought about by  $10\mu g$  of the ODN. We pre-treated mice with ISS ODN by the nasal route, then challenged the mice with influenza virus via the same route, but failed to protect against the infection. Tighe et al. stressed the advantages of using DNA vaccines compared with conventional protein vaccines, particularly for neonates (TIGHE 1998).

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The dominance of the Th1 response generated by ISS DNA may be useful for protecting Th2-type allergic diseases (PISETSKY 1996; KLINE 1998; TIGHE 1998; ZIMMERMANN 1998). In two rodent experimental models, id gene vaccination with naked plasmid-encoding antigens reduced an ongoing IgE response to the antigen and lowered the levels of eosinophils observed in the lung after protein challenge (BROIDE 1998). Kline et al. observed that mice previously sensitized to an antigen have an established Th2 response; co-administration of ISS DNA and the antigen can redirect the immune responses to Th1 and can abolish allergic reactions on subsequent antigen challenge (KLINE 1998). Mice infected lethally with Leishmania major, which is known as a lethal Th2-driven disease model, were given ISS DNA having a GACGTC sequence (ZIMMERMANN 1998). When the ODN was given as late as 20 days after the lethal infection, it was curative, indicating that ISS ODN reverses an established Th2 response (ZIMMERMANN 1998). In a mouse model of asthma, airway eosinophilia, Th2 cytokine induction, IgE production, and bronchial hyper-reactivity were prevented by co-administration of ISS DNA and antigen.

Tumor immunotherapy may be one of the uses of ISS DNA. Many experimental trials have been reported (KRIEG 1998; TIGHE 1998). For instance, ISS ODNs enhance the efficacy of monoclonal-antibody therapy of murine lymphoma (WOOLDRIDGE 1997). We reported many experimental models in which intraregional injections with ISS ODN alone strongly inhibited tumor growth (TOKU-NAGA 1984; SHIMADA 1985, 1986; KATAOKA 1992; YAMAMOTO 1992). Clinical trials showed repeated intratumor injections with BCG-DNA to be remarkably effective (ISHIHARA 1988, 1990; JOHNO 1991).

It was stated that a 20-mer antisense ODN is currently in phase-III human clinical trials to treat cytomegalovirus retinitis (KRIEG 1996). The ODN used is an exceptionally potent inhibitor of virus replication and contains two GCG trinucleotides (KRIEG 1996).

Activation of the immune system with ISS DNA could have both beneficial and deleterious consequences (KRIEG 1996; PISETSKY 1996, 1997; ZIMMERMANN 1998). Schwartz et al. reported that instilling mice intra-tracheally with 20-mer ISS DNA or bacterial DNA (but not vertebrate DNA or methylated bacterial DNA) caused inflammation in the lower respiratory tract; a 50-fold increase in the concentration of TNF- $\alpha$  and a 4-fold increase of IL-6 were observed (SCHWARTZ 1997). Furthermore, DNA extracted from the sputum of patients with cystic fibrosis contained approximately 0.1–1% bacterial DNA, suggesting that ISSs of DNA may play an important pathogenic role in inflammatory lung disease. The possible role of immunostimulatory DNA in inflammation in other tissues exposed to bacteria must be considered seriously.

LPS of gram-negative bacteria has been known to cause sepsis syndrome. However, factors in addition to LPS must induce the syndrome, because bacteria that lack LPS still induce it. Cowdery et al. reported that intravenous injection with bacterial DNA or 20-mer ISS DNA (but not calf thymus DNA or 20-mer ODN without ISS) induced a rapid increase in IFN- $\gamma$  and, when followed by LPS, showed 75% mortality, while no death in mice treated with both calf thymus DNA and LPS was observed (COWDERY 1996). This mortality rate required the secretion of IFN- $\gamma$ , because mice lacking the IFN- $\gamma$  gene were protected. It was reported that mice given ODN having a -GACGTT- sequence suffered lethal toxic shock due to macrophage-derived TNF- $\alpha$ , resulting in fulminate apoptosis of liver cells (SPARWASSER 1997). LPS was synergized with ISS ODN, as measured by monitoring TNF- $\alpha$  release.

Systemic lupus erythematosus (SLE) is an autoimmune disease of unknown pathogenesis. ISS in bacterial DNA could promote the development of SLE by inducing polyclonal B-cell activation, over-expression of immune cytokines (such as IL-6) and resistance to apoptosis, thereby potentially allowing the survival of autoreactive cells. This suggests that ISSs in bacterial DNA are possibly involved in the pathogenesis of SLE (KRIEG 1995, 1996). SLE patients show deficient expression of antibodies specific for bacterial DNA (PISETSKY 1997).

Miyata et al. analyzed nine kinds of DNA clones from DNA – anti-DNAantibody-immune complexes of 11 SLE patients and found that the DNAs were rich in hexamer palindromic ISSs (MIYATA 1996). ODNs having –TTCGAA– in a cloned DNA augmented both the expression of intercellular adhesion molecule 1 on endothelial cells and the production of IL-2, IL-6, IL-8, and TNF- $\alpha$ , suggesting that DNA contained in the immune complex of SLE patients may exacerbate vasculitis.

Using a 10-mer palindromic DNA, TCAACGTTGA, Segal et al. reported that, when murine T cells that recognize the autoantigen myelin basic protein are exposed to either LPS or the palindromic ODN, the result is a conversion of the T cells, via an IL-12-dependent pathway, from quiescent cells into autoimmune-disease-effector Th1 cells (SEGAL 1997). Pre-autoimmune NZB/W mice have immunoregulatory defects that, due to immunization by bacterial DNA, allow activation of mammalian double-stranded-DNA-reactive B cells (WLOCH 1997). It was found that SLE patients have a marked deficiency in the production of antibodies specific for the antigen of *Micrococcus lysodeikticus* (PISETSKY 1997).

## **6** Conclusions

Both bacterial DNA and ISS ODNs having a 5'-ACGT-3' or 5'-TCGA-3' motif are more advantageous IFN- $\alpha$  inducers than are viruses. First, the ISS ODNs are stable, and it is easy to control their quality. Second, we can use the ISS ODNs without risk of infection. Third, since the action of the ISS ODN on IFN- $\alpha$ producing cells may be limited compared with the action of viral infection, it may be easier to determine the site of action of ISS ODN. Fourth, IFN- $\alpha$  induced by the ISS ODN can be determined without the cytopathic effects of the inducing virus.

The mechanism of IFN- $\alpha$  induction by the ISS ODN is still obscure at present. Since the minimum sequence requirement for ISS ODNs was reduced to 5'-ACGT-3' or 5'-TCGA-3', an antisense mechanism is unlikely to be involved,

because sequence-specific inhibition is exerted by antisense ODNs that are strictly complementary to the target mRNA. As is well known, many transcription factors, such as c-myc and AP-1, recognize short palindromic sequences. Hence, the most plausible hypothesis may be that the ODNs interact with transcription factors known to regulate IFN genes. However, the transcription factors known to regulate IFN-gene expression do not recognize the highly active sequences reported by us.

Our studies have shown the structure of ISS ODN essential to IFN- $\alpha$  induction (YAMAMOTO S, 1988; YAMAMOTO T, YAMAMOTO S, KATAOKA and KOHURO 1994). This study characterized the cells producing IFN- $\alpha$  via stimulation with the ISS ODN. From now on, our studies will deal with both the INF inducers and producers. Studies on the mechanism of IFN- $\alpha$  induction and production by ISS ODNs having 5'-ACGT-3' or 5'-TCGA-3' motifs will contribute to new developments in the field of IFN- $\alpha$  research. Since a large amount of DNA had never been administered to experimental animals or humans, it would be worthwhile for us to introduce clinical trials and pharmacological studies of BCG-DNA obtained in Japan in the late 1980s (TOKUNAGA 1999).

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# Macrophage Activation by Immunostimulatory DNA

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# **1** Introduction

Activation of macrophages/dendritic cells appears to be of crucial importance in vivo in immunostimulation by foreign DNA. Immune cell activation by bacterial

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DNA was first observed using mixed spleen cell cultures (SHIMADA et al. 1986; YAMAMOTO et al. 1992) and B cells (MESSINA et al. 1991), and the role of unmethylated CpG motifs in this activation was elucidated in B cells (KRIEG et al. 1995). This work showed that oligonucleotides containing an unmethylated CG dinucleotide in a certain sequence context (ACGT but not CCGG) could mimic the activity of bacterial DNA. A combination of methylation of the vertebrate genome and suppression of the frequency of activating sequences may prevent immune activation by self DNA. In mixed spleen cell culture, bacterial DNA or stimulatory oligonucleotides (here termed "CpG DNA") promote production of a range of cytokines, including interferons- $\alpha/\beta$  and - $\gamma$  (IFN- $\alpha/\beta$ , IFN- $\gamma$ ), interleukins-6 and -12 (IL-6, IL-12) and tumour necrosis factor- $\alpha$  (TNF- $\alpha$ ). A role for macrophages in this activation was suggested by the finding that the production of IFN- $\gamma$  by the non-adherent fraction of spleen cells was dependent on IL-12 and TNF-a production by the adherent fraction (HALPERN et al. 1996). Direct activation of macrophages by bacterial DNA and oligonucleotides was first shown by activation of nuclear factor (NF)-KB and induction of TNF-a mRNA and human immunodeficiency virus (HIV)-1 long terminal repeat (LTR) promoter activity (STACEY et al. 1996). Since then, it has become apparent that activation of macrophages and other antigen-presenting cells (APCs) is critical in the production of cytokines and the subsequent development of T-helper 1 (Th1)-biased T-cell lineages in response to CpG DNA. Responses of macrophages and dendritic cells will be addressed together in this review, as they can be considered to be different differentiative states of the same cell lineage (PALUCKA et al. 1998). This review will focus on the mechanism of macrophage activation, from DNA uptake to altered gene expression, and the CpG DNA-induced changes in macrophage function in vitro and in vivo.

## 2 Cellular DNA Uptake

## 2.1 Endocytic Uptake of DNA

DNA and oligonucleotides are internalised by mammalian cells, and most authors now seem to favour a requirement for DNA uptake in activation by CpG DNA, although an interacting role for signalling from a cell-surface receptor has not been excluded. In our experiments, even quite large DNA molecules, such as plasmids, can be taken up into a macrophage cell line, and some plasmids reach the nucleus in an intact state able to be expressed (STACEY et al. 1996). Uptake of naked DNA has been known for some time from studies of the fate of adenoviral DNA in culture (GRONEBERG et al. 1975). Recently, attention has been focussed on oligonucleotide uptake (reviewed in BENNETT 1993) and, at low oligonucleotide concentrations, the process seems to involve receptor-mediated endocytosis via clathrin-coated pits (BELTINGER et al. 1995). Internalised oligonucleotides were found in endosomes

and dense lysosome-like vesicles, with some escape to the cytoplasm and nucleus (BELTINGER et al. 1995). Phosphorothioate oligonucleotides taken up by the premyelocytic leukaemic line HL60 (TONKINSON and STEIN 1994) and the mouse macrophage cell line J774 (HÄCKER et al. 1998) were largely located within acidified vacuoles. Uptake of oligonucleotides seems to have little or no sequence specificity (KRIEG et al. 1995), and distinction between activating and non-activating oligonucleotides is proposed to take place at a stage later than cell-surface-receptor recognition. Receptor-mediated endocytosis, of course, requires a cell-surface receptor with affinity for DNA, and a recent paper gives evidence that the integrin Mac-1 (CD11b/CD18) may mediate oligonucleotide uptake (BENIMETSKAYA et al. 1997). Mac-1 is expressed largely on polymorphonuclear leukocytes and macrophages. In our own studies, it seems unlikely that Mac-1 is responsible for DNA uptake in macrophages, as its cell-surface concentration does not alter during DNA stimulation as would be expected for a receptor involved in endocytosis (Beasley and Hume, unpublished observations). Several studies have addressed DNA cellsurface binding (reviewed in Bennett 1993), with one crosslinking study showing DNA interacting with at least five unidentified proteins on K562 human leukaemia cells (Beltinger et al. 1995).

### 2.2 Clearance of DNA In Vivo

Macrophages are a major site of clearance of circulating DNA in vivo and, hence, may contribute more to the overall response than other cell types. Intravenously injected DNA was rapidly cleared from the bloodstream (90% cleared within 20min) and localised to the liver and spleen, suggesting a role for macrophages in clearance (CHUSED 1972). Single-stranded DNA taken up by the liver was primarily bound by Kupffer cells (liver macrophages) (EMLEN 1988). With the injection of phophorothioate-stabilised oligonucleotides, a wider tissue distribution was found, with a large component of clearance by the kidney; however, localisation to the liver and spleen was still apparent (SANDS et al. 1994). Interestingly, DNA need not be injected to gain access to the immune system, as uptake and even integration of foreign DNA into the genome of spleen cells has been demonstrated in an experiment where mice were fed M13 DNA (SCHUBBERT et al. 1997)! In vitro studies showed a difference in the ability of immune cells to take up oligonucleotide, with macrophages the most efficient, followed by B cells and then T cells, which had relatively low-efficiency uptake (ZHAO et al. 1996).

#### 2.3 DNA Uptake as a Requirement for Activation?

Evidence that uptake of foreign DNA is required for subsequent immune activation comes from several approaches. Two groups using immobilised oligonucleotides reported that internalisation of CpG DNA is necessary for the proliferative response of B cells, although data awaits publication (KRIEG et al. 1995; L. Manzel and

D. Macfarlane, personal communication). Such experiments have obvious difficulties in both the estimation of relative concentrations of free and immobilised oligonucleotides seen by cells and the problem of cleavage of oligonucleotides from the solid support. One published study on human B cells showed that immobilised oligonucleotides were active (LIANG et al. 1996), but the observed proliferative response was specific for some phosphorothioate oligonucleotides and not bacterial DNA and was not demonstrated to be CpG specific. This B-cell activation (LIANG et al. 1996) did not seem to be similar to the well-characterised mouse response and may, in fact, be a cell-surface-mediated phenomenon or the covalent coupling of oligonucleotide to sepharose may have allowed entry of oligonucleotides into cells, as found by other workers (L. Manzel and D. Macfarlane, personal communication). No direct studies on the requirement for internalisation have been performed on macrophages, but it is likely that the situation will be analogous to that for B cells. Findings that the potency of oligonucleotides is increased by lipofection into spleen cells (YAMAMOTO et al. 1994) and that lipofection makes possible a response of cells to six-base-long oligonucleotides (SONEHARA et al. 1996) has been cited as evidence that internalisation is required for activation; however, the data are ambiguous due to the possibility of synergy between oligonucleotides and lipofection reagents.

## 2.4 Endosomal Acidification Required for CpG DNA Responses

Less direct evidence of a requirement for internalisation has been obtained using compounds that prevent acidification of endosomes and block the action of DNA. Bafilomycin, chloroquine and monensin prevented J774 macrophage cell and spleen cell TNF-α, IL-12 and IL-6 production (HÄCKER et al. 1998; YI et al. 1998b). The effect was specific for the DNA response and did not prevent activation by phorbol 12-myristate 13-acetate + ionomycin or lipopolysaccharide (LPS). Chloroquine and related compounds also inhibited CpG DNA responses of B cells and unfractionated human peripheral blood monocytes (MACFARLANE and MANZEL 1998). Chloroquine and bafilomycin A did not block uptake of oligonucleotide or vesicular localisation (HÄCKER et al. 1998). These compounds act in different ways - bafilomycin A blocks the endosomal hydrogen-ion pump, and chloroquine is a strong base that localises to the vesicle. However, recent findings that chloroquine inhibits the action of CpG DNA at concentrations that do not prevent endosomal acidification (L. Manzel and D Macfarlane, personal communication) call into question the assumption that the effect of these inhibitors is necessarily related to change in endosomal pH. Further work is required to establish exactly how these inhibitors are acting. One possibility is that they inhibit release of oligonucleotides from the receptor mediating their uptake, and prevent them from reaching the cytoplasm. Although DNA "escape" from the vacuole into the cytoplasm has been suggested as a requirement for activation, a CpG-specific receptor could be located in the endosome. Escape from the vacuole, if required, is likely to be a regulated rather than an "accidental" event, as responses to exogenously added DNA are extremely rapid and reliable (see below).

## 3 Macrophage/Dendritic Cell Responses to CpG DNA

## 3.1 Signalling

The receptors for CpG DNA and initiation of signalling remain uncharacterised. Although it was proposed early on that CpG-containing DNA may induce transcription of cytokine genes by direct interaction with transcription factors (CowDERY et al. 1996), no evidence of this has emerged. Instead, a number of studies show activation of established intracellular signalling molecules. All signalling and gene inductions that have been studied are prevented by the inhibitors of endosomal acidification. This includes reactive oxygen species (ROS) production, NF- $\kappa$ B activation (Y1 et al. 1998b) and mitogen-activated protein kinase (MAPKinase) pathways (HÄCKER et al. 1998; Y1 and KRIEG 1998). Some of these responses occur within 15min of exposure to CpG DNA, showing that the inhibitors, such as chloroquine and bafilomycin A, interfere with a very early signalling step.

#### 3.1.1 Nuclear Factor KB

In macrophages, transcription factor NF- $\kappa$ B is activated by bacterial DNA and CpG-containing oligonucleotides but not by methylated or vertebrate DNA (STACEY et al. 1996; SPARWASSER et al. 1997b; Y1 et al. 1998b). NF- $\kappa$ B is bound to an inhibitory subunit, I $\kappa$ B, in the cytoplasm of cells until activation, when I $\kappa$ B is phophorylated and degraded, allowing NF- $\kappa$ B to be translocated to the nucleus (BALDWIN 1996). Activation of NF- $\kappa$ B is frequently initiated in situations of cellular stress, and it is implicated in induction of a wide range of inflammatory genes, including those involved in the CpG DNA response, such as TNF- $\alpha$  (DROUET et al. 1991), inducible nitric oxide synthase (iNOS) (XIE et al. 1994) and IL-12 (YOSHIMOTO et al. 1997). The use of inhibitors of I $\kappa$ B phophorylation or degradation suggests that active NF- $\kappa$ B is required for macrophage TNF- $\alpha$  production in response to CpG DNA (Y1 et al. 1998b). The upstream signalling involved in NF- $\kappa$ B activation in response to CpG DNA remains to be established but probably involves a redox-sensitive step (see below).

#### 3.1.2 Reactive Oxygen Species

Production of ROS has been implicated in signalling pathways leading to NF- $\kappa$ B activation (BAEUERLE and HENKEL 1994). CpG DNA induction of ROS was detected in the J774 macrophage cell line (YI et al. 1998b) and B cells (YI et al. 1996) by an increase in fluorescence of oxidation-sensitive intracellular dihydro-rhodamine123. The ROS production was prevented by inhibitors of endosomal acidification (YI et al. 1998b). The levels of ROS produced are much lower than those resulting from a respiratory burst response producing radicals for cytotoxic purposes, but they may have a role in signalling. The literature linking ROS with NF- $\kappa$ B activation relies heavily on the use of inhibitors, such as pyrrolidine

dithiocarbamate (PDTC) and other antioxidants (SCHRECK et al. 1992; BAEUERLE and HENKEL 1994). Antioxidant compounds are potentially capable of altering the redox states of many receptors and signalling molecules, and inhibition does not necessarily indicate a direct role for ROS as second messengers. PDTC has been shown to inhibit TNF- $\alpha$  mRNA induction in RAW264 macrophages by CpG DNA and submaximal but not maximal concentrations of LPS (K. Stacey, unpublished). PDTC also inhibited IL-6 production in response to CpG DNA in B cells, while protein kinase C and protein kinase A inhibitors had no effect (Y1 et al. 1996). Caution should be exercised in interpreting this as evidence for direct involvement of ROS in signalling. Some workers have presented evidence against ROS as second messengers involved in NF- $\kappa$ B activation (BRENNAN and O'NEIL 1995; SUZUKI et al. 1995), although the redox state of certain signalling molecules upstream of NF- $\kappa$ B must be important in allowing activation.

#### 3.1.3 Mitogen-Activated Protein Kinases

Many different stimuli (growth factors, cytokines, environmental stress) activate members of the MAPKinase superfamily, although the pattern of activation varies among stimuli. MAPKinases directly phosphorylate and activate certain transcription factors. The MAPKinases c-Jun N-terminal kinase (JNK) and p38 were activated by CpG DNA in macrophage cell lines, a B-cell line and in bone-marrow cells differentiated towards a dendritic cell phenotype with granulocyte/macrophage colony-stimulating factor (GM-CSF) (HÄCKER et al. 1998; YI and KRIEG 1998). This involves presumably sequential activation of the kinase JNK kinase 1 followed by the MAPKinase JNK, which is responsible for phosphorylation of the activator protein 1 (AP-1)-family transcription factor Jun within 15min of CpG DNA exposure (HÄCKER et al. 1998; YI and KRIEG 1998). MAPKinase p38 and its target transcription factor, activating transcription factor 2 (ATF-2), were also activated by CpG DNA (HÄCKER et al. 1998; YI and KRIEG 1998). This pathway is implicated in the CpG DNA-induced macrophage expression of IL-12 and TNF-a, as a specific p38 inhibitor reduced production of these cytokines (HÄCKER et al. 1998).

MAPKinases p44 and p42 (Erk-1 and -2) are downstream of ras in growthfactor-signalling pathways and are activated by the macrophage growth factor colony-stimulating factor-1 (CSF-1) and LPS (GEPPERT et al. 1994; LIU et al. 1994; BUSCHER et al. 1995; FOWLES et al. 1998). Although one report found little activation of Erk-1 and -2 by CpG DNA in J774 macrophages (Y1 and KRIEG 1998), we found that CpG DNA induced tyrosine phosphorylation of these MAPKinases in bone-marrow-derived macrophages (D. Sester, unpublished). Whilst CSF-1 and LPS gave rapid phosphorylation of Erk-1 and -2 within 5min, the response to DNA was slightly delayed, perhaps reflecting the difference between signalling from cellsurface receptors for LPS and CSF-1 and the requirement for internalisation of DNA. The common effects of LPS, CSF-1 and CpG DNA in activation of Erk-1 and -2 may relate to their shared anti-apoptotic activity (Sect. 3.6). An involvement of Erk-1 and -2 in anti-apoptotic responses (JARVIS et al. 1997; WANG et al. 1998) may explain why their activation by CpG DNA was detected in factor-dependent bone-marrow-derived macrophages starved of their growth factor, but not in factor-independent J774 macrophages.

#### 3.1.4 Transcription Factors

In addition to activation of NF- $\kappa$ B, Jun and ATF-2, mentioned above, CpG DNA and LPS both induce macrophage expression of mRNA for c-*myc* (INTRONA et al. 1986; MYERS et al. 1995; YI et al. 1998b) and Ets-2 (SwEET et al. 1998b). Ets-2 and c-*myc* are also up-regulated in response to tyrosine kinase and ras signalling induced by CSF-1 (XU et al. 1993; STACEY et al. 1995; FowLes et al. 1998). In addition to its induction, Ets-2 is activated upon phosphorylation by the MAP-Kinases Erk-1 and Erk-2 in response to LPS and CSF-1 (FowLes et al. 1998; SwEET et al. 1998b). Although it has not been demonstrated, a similar phosphorylation of Ets-2 is likely to follow the CpG DNA-mediated activation of Erk-1 and -2 (Sect. 3.1.3). The induction of c-*myc* and Ets-2 via MAPKinase-dependent routes in response to LPS, CSF-1 and CpG DNA may be related to anti-apoptotic effects of all three compounds (Sect. 3.6).

LPS and CpG DNA also both induce mRNA for CCAAT/enhancer-binding protein (C/EBP)- $\beta$  and - $\delta$  transcription factors. These factors bind to *cis*-acting elements of many myeloid-specific and inducible genes, and C/EBP- $\beta$  was first identified as NF-IL6, a factor involved in the expression of the IL-6 gene (AKIRA et al. 1990). Generation of a knockout mouse showed that NF-IL6 is essential for normal bacterial and tumour cell killing by macrophages (TANAKA et al. 1995).

### 3.2 Macrophage/Dendritic Cell Gene and Cytokine Induction by DNA

#### 3.2.1 Interleukin 12

One of the CpG-induced cytokines about which there is great interest is IL-12, given its role in early the induction of IFN- $\gamma$  in natural killer (NK) cells (CHACE et al. 1997) and the promotion of Th1-type immunity (SCOTT and TRINCHIERI 1997). Macrophages and dendritic cells are believed to be the primary source of IL-12 in response to CpG DNA, and IL-12 induction has now been detected in a variety of cells exposed to CpG DNA, including splenic macrophages (CHACE et al. 1997), J774 (LIPFORD et al. 1997) and ANA-1 murine macrophage cell lines, bone-marrow-derived dendritic cells (HÄCKER et al. 1998) and fetal-skin-derived dendritic cells (JAKOB et al. 1998).

#### 3.2.2 Tumor Necrosis Factor a

Together with IL-12, TNF- $\alpha$  may play a role in the induction of IFN- $\gamma$  in spleen cell cultures exposed to CpG DNA (HALPERN et al. 1996). Apart from its well-publicised role in pathological states, such as toxic shock, TNF- $\alpha$  is an important

part of the normal inflammatory response. TNF- $\alpha$  mRNA and protein are rapidly produced by CpG DNA in bone-marrow-derived macrophages (STACEY et al. 1996; Sweet et al. 1998a), bone-marrow-derived dendritic cells (HÄCKER et al. 1998), peritoneal macrophages (SPARWASSER et al. 1997b) and the murine macrophage cell lines RAW264 (STACEY et al. 1996; Sweet et al. 1998a), J774 (LIPFORD et al. 1997) and ANA-1 (SPARWASSER et al. 1997b). Strongly activating oligonucleotides induce both IL-12 and TNF- $\alpha$  efficiently. One group has described a phosphorothioate oligonucleotide that induced IL-12 and not TNF-a (LIPFORD et al. 1997), and this would implicate more than one mechanism in the detection of foreign DNA. However, we believe this finding may be a reflection of different genes having different threshold concentrations of DNA for activation. In our experiments, a phosphodiester version of this same oligonucleotide can induce TNF- $\alpha$  but is less potent than some other oligonucleotides (Sweet, unpublished observations). The dose-response curves for oligonucleotides are relatively steep, achieving maximal effect over a tenfold concentration range. Therefore, if different genes have different thresholds for activation, then at an appropriate concentration a weakly activating oligonucleotide could stimulate IL-12 and not TNF-a expression. The difference between our results and those of LIPFORD et al. (LIPFORD et al. 1997) warrant further investigation of dose responses to such oligonucleotides.

## 3.2.3 Other CpG DNA-Induced Cytokines/Genes

Macrophages (LIPFORD et al. 1997), bone-marrow-derived dendritic cells (SPAR-WASSER et al. 1998) and B cells (YI et al. 1996) produce IL-6 in response to bacterial DNA. IL-6 is a pleiotropic cytokine important in resistance to infection and involved in the acute phase response in liver (AKIRA and KISHIMOTO 1996). Induction of mRNA for the inflammatory cytokine IL-1 $\beta$  and a protease inhibitor, plasminogen activator inhibitor-2 (PAI-2), was detected in RAW264 macrophage cells (STACEY et al. 1996), but message levels were considerably lower than those induced by LPS. iNOS and its product, nitric oxide (NO, a free radical implicated in macrophage-mediated killing of micro-organisms and tumour cells), are produced by bone-marrow-derived macrophages in response to CpG DNA, but only in the presence of IFN- $\gamma$  (STACEY et al. 1996; SWEET et al. 1998a). This requirement for IFN- $\gamma$  priming is distinct from LPS induction of iNOS (Sect. 3.5).

Although the earliest characterised DNA response was production of type-I IFNs by spleen cells (TOKUNAGA et al. 1988), little work has been done to establish which cell types produce IFN- $\alpha/\beta$ . One recent study implicates APC production of type-I IFN in subsequent T-cell activation, as IFN-I-receptor<sup>-/-</sup> T cells failed to show normal stimulation with CpG DNA-activated APCs (SUN et al. 1998). Another study found production of IFN- $\alpha$  and - $\beta$  mRNA by human macrophages transfected with DNA (ROMAN et al. 1997). Confirmation of IFN- $\alpha/\beta$  production in the absence of the lipid transfection reagent and analysis of production by other cell types is yet to be presented. Similarly, IL-18 mRNA has been detected, after transfection, by the same group (ROMAN et al. 1997). IL-18 cooperates with IL-12

in the promotion of IFN- $\gamma$  production and the development of Th1-type immunity (TAKEDA et al. 1998).

#### 3.2.4 Responses of Human Monocytes

There is little published information on activation of human cells of any type in response to CpG DNA, a gap in the literature which needs to be filled before therapeutic applications can be designed or optimised. Some responses of human monocytes and macrophages have been observed. Using human peripheral blood mononuclear cells, bacterial DNA and CpG oligonucleotides were found to induce the cytokines IL-6 and TNF-a and intercellular adhesion molecule 1 (ICAM-1) in the monocyte fraction (HARTMANN and KRIEG 1999). However, the time course of expression was delayed compared with the rapid response to LPS, with TNF- $\alpha$  and IL-6 only being produced 18–24h after stimulation. In addition, relatively high levels of Escherichia coli DNA were required for this response. The delayed induction of normally rapidly induced genes suggests that this may be a differentiative response of monocytes to CpG DNA. Another group has found that CpG DNA specifically inhibits the adhesion of human monocytederived macrophages in vitro (D. Macfarlane, personal communication). One difficulty in the comparisons of human and mouse work is that mouse studies generally involve relatively mature macrophages, such as peritoneal or bonemarrow-derived macrophages, as opposed to the circulating monocytes most commonly available from human subjects. In addition, most human cell lines available are relatively immature. The use of monocyte-derived macrophages or other more mature human macrophages may well show similar DNA sensitivity to mouse macrophages. The induction of IL-18 and IFN- $\alpha$  and - $\beta$  mRNA in response to transfected DNA discussed above (Sect. 3.2.3) was performed in human macrophages purified from peripheral blood mononuclear cells by fibronectin adherence and occurred within 3h.

### 3.3 Negative Regulation of Macrophage Responses

Macrophage production of IL-12 in response to DNA leads to NK-cell IFN- $\gamma$  production, which in turn enhances the macrophage response to DNA, leading to increased production of TNF- $\alpha$ , NO (SwEET et al. 1998a) and, most likely, IL-12 itself. This gives the possibility of a self-amplifying loop, which must be kept under control in vivo if toxic shock is to be avoided. One cytokine which may play a role in this control is IL-10, which has been shown to inhibit macrophage IL-12 production in response to DNA (ANITESCU et al. 1997). Induction of IL-10 has been detected in splenocytes following bacterial DNA treatment and is probably B-cell derived (ANITESCU et al. 1997; REDFORD et al. 1998). Addition of anti-IL-10 antibodies increased spleen cell IL-12 production in response to CpG DNA (REDFORD et al. 1998) and, thus, IL-10 may be one of the cytokines acting to keep the response to foreign DNA under control.

### 3.4 Maturation of Cell Function

CpG DNA induces both acute responses, like induction of TNF- $\alpha$  mRNA expression in macrophages, which peaks at 1h (STACEY et al. 1996), and slower differentiative responses, in particular the maturation of APC function (JAKOB et al. 1998; SPARWASSER et al. 1998). Like LPS, CpG DNA treatment for 18h caused maturation of fetal-skin-derived dendritic cells to exhibit decreased adhesion, increased expression of major histocompatibility complex (MHC) class-II and co-stimulatory molecules CD86 and CD40 (JAKOB et al. 1998). Compared with the LPS response, CpG-treated cells made more IL-12 and less IL-6 and TNF- $\alpha$ . CpG-DNA-induced maturation of professional APC function has also been observed in bone-marrow-derived dendritic cells cultivated in GM-CSF (SPARWASSER et al. 1998).

### 3.5 Comparison of DNA and LPS Responses

The range of macrophage gene products induced by bacterial DNA is similar to an LPS response (STACEY et al. 1996), but the relative potency of the two products varies for different genes. Despite a similarity of action, an increasing number of studies are finding important differences in the actions of LPS and CpG DNA. Given that LPS is a powerful pyrogen and inducer of toxic shock and is unsuitable for use as an adjuvant in humans, establishment of a different mode of action for CpG DNA is a prerequisite to its use in human therapy.

One simple piece of evidence that LPS and DNA have substantially different actions in vivo is that the two compounds synergise in the induction of TNF- $\alpha$  (SPARWASSER et al. 1997b). This may require the cooperation of a mixture of cell types or may be post-transcriptional, as we are unable to show synergy at the level of TNF- $\alpha$  mRNA expression in macrophage culture (K. Stacey, unpublished observations). CpG DNA and LPS also synergised in the induction of IL-6 in human monocytes (HARTMANN and KRIEG 1999). This synergy in the induction of a gene within one cell precludes the activation of identical signalling pathways or transcription factors.

The CpG DNA response is separable from the LPS response by the use of LPS non-responder C3H/HeJ mice, which seem to retain all normal responses to DNA (SHIMADA et al. 1986; COWDERY et al. 1996; SPARWASSER et al. 1997b). The defect in the C3H/HeJ mouse has now been identified as a putative LPS receptor, Toll-like receptor 4 (POLTORAK et al. 1998), a molecule which is clearly unnecessary for the DNA response. A difference in early signalling is also shown by the inhibition of CpG DNA responses but not LPS responses by chloroquine and bafilomycin A, which may be interfering with normal endosomal pathways (Sect. 2.4; HÄCKER et al. 1998).

Signalling in response to LPS and DNA converges with the activation of MAPKinases (Häcker et al. 1998; YI and KRIEG 1998), activation by both agents of transcription factors NF-kB (STACEY et al. 1996; SPARWASSER et al. 1997b) and

AP-1 (HÄCKER et al. 1998; YI and KRIEG 1998) and induction of the transcription factors C/EBP- $\beta$ , C/EBP- $\delta$  and Ets-2 (Sweet et al. 1998b). However, there are clearly undiscovered differences in transcription-factor activation, as there are some separable functions of LPS and DNA in gene induction. In a comparison of the responses of two macrophage cell lines to LPS and DNA, one line produced IL-12 in response to both compounds, but RAW264 cells were responsive only to DNA (ANITESCU et al. 1997), although other LPS responses are perfectly intact in this cell line. As noted above, a difference we have detected in LPS and DNA signalling is that LPS, acting alone, can induce expression of iNOS and NO production by bone-marrow-derived macrophages and RAW264 cells, whilst CpG DNA requires IFN-γ priming for this response (STACEY et al. 1996; Sweet et al. 1998a; K. Stacey, unpublished). IFN- $\gamma$  induces Stat1 $\alpha$  transcription factor, which is required for iNOS-promoter activity. Induction of iNOS by LPS alone requires autocrine production of type-I IFNs, which subsequently induces Statla phosphorylation (GAO et al. 1998). We are currently investigating whether or not CpG DNA can mediate a similar activation of Stat $1\alpha$ .

The above examples highlight some differences between CpG DNA and LPS actions, but there are many similarities. The concept that prior exposure to substimulatory amounts of LPS induces tolerance to higher doses is well established. In experiments we have performed on activation of the HIV-1 LTR in RAW264 macrophages, overnight pretreatment with 0.1ng/ml LPS markedly reduced the response to subsequently added CpG DNA or LPS (S. Cronau and D. Hume, unpublished observations). The reverse is also true, with prior exposure to CpG DNA attenuating the response to both stimuli. Presumably, both agents are able to down-modulate a signalling molecule involved in the activation of the HIV LTR.

#### 3.6 Effects on Cell Survival and Growth

Bone-marrow-derived macrophages proliferate and differentiate in the presence of the growth factor CSF-1, and it is also required for survival in culture. In the absence of CSF-1 or another survival stimulus for 24h, the cells begin to commit to an apoptotic pathway. Addition of either LPS or CpG DNA prevents apoptosis (D. Sester, unpublished). CpG DNA also promotes survival of B cells (Y1 et al. 1998a) but, in that case, DNA and LPS are both mitogenic stimuli. In macrophages, DNA and LPS are not mitogenic stimuli, and they inhibit growth in response to CSF-1. The observed anti-proliferative and perhaps anti-apoptotic effects of CpG DNA are associated with downregulation of the CSF-1 receptor, which occurs between 10min and 60min after exposure to CpG DNA (S. Beasley and D. Hume, unpublished observations).

## 4 Responses to CpG DNA In Vivo

## 4.1 DNA and LPS in Toxic Shock

Macrophages are central to the production of cytokines in response to LPS, leading to toxic shock. Although it has been suggested that bacterial DNA may cause toxic shock through induction of TNF-α (SPARWASSER et al. 1997a,b), demonstration of a toxic effect required treatment of mice with D-galactosamine, which sensitises liver cells to TNF-α-mediated apoptosis (GUTIERREZ-RAMOS and BLUETHMANN 1997). In this model of sepsis, TNF- $\alpha$  is the major mediator of shock, whereas, in non-sensitised mice, ICAM-1 expression may be more important (GUTIERREZ-RAMOS and BLUETHMANN 1997), and a more complex array of cytokines may be involved. In fact, studies now show that, unlike LPS, a single dose of up to 500µg/mouse of phosphorothioate CpG oligonucleotide alone does not cause toxic shock, but two large doses (500µg/mouse) given within 1 week can cause a fatal sepsis-like condition (A. Krieg, personal communication). This may be mediated by a priming effect of cytokines induced by the first dose, since IFN- $\gamma$  and IL-12 knockout mice are resistant to this effect (A. Krieg, personal communication). Although elevated circulating IFN- $\gamma$  has only been detected for 24h after administration of bacterial DNA (COWDERY et al. 1996), IL-12 can remain high for a week (KRIEG et al. 1998). Administration of bacterial DNA 4h prior to LPS greatly sensitises the mice to LPSmediated shock, probably through production of IFN- $\gamma$  (Cowdery et al. 1996). As noted in Sect. 3.5, when added at the same time, LPS and DNA synergised induction of TNF- $\alpha$  in vivo (Sparwasser et al. 1997b), and DNA could play a synergistic role in promoting both Gram-negative and Gram-positive shock. Nevertheless, the difference in toxicity between LPS and DNA administered alone is apparent in vivo, which is encouraging for the therapeutic applications of CpG DNA.

## 4.2 Generation of Th1 Responses

Responses to foreign DNA sequences seem to drive development of Th1-type immune responses in studies using oligonucleotides as adjuvants (CHU et al. 1997; DAVIS et al. 1998) and in DNA vaccinations (SATO et al. 1996; KLINMAN et al. 1997; LECLERC et al. 1997; TIGHE et al. 1998; STACEY and BLACKWELL 1999). CpG DNA-induced early production of IL-12 by macrophages and subsequent induction of IFN- $\gamma$  in NK cells during first exposure to antigen appear to polarise subsequent development of T cells to the Th1 lineage. This response may well also involve CpG DNA induction of type-I IFNs and IL-18 from macrophages/APCs (ROMAN et al. 1997). Thus, macrophage/APC recognition of DNA may be seen as responsible for Th1 development, with DNA as an adjuvant, and may contribute to Th1 development in normal infections. The Th1 bias has applications not only in vaccination but also in therapy for allergy, which is a Th2-dominated process (GOODMAN et al. 1998; SPIEGELBERG et al. 1998).

#### 4.3 Activation of Macrophages/APCs in DNA Vaccination

Presentation of antigen by professional bone-marrow-derived APCs is required for cytotoxic T-lymphocyte generation in DNA vaccination (Corr et al. 1996). Activation of APCs by CpG sequences in the plasmid backbone is required for effective immunisation and development of Th1 responses in DNA vaccination (SATO et al. 1996; KLINMAN et al. 1997; LECLERC et al. 1997; TIGHE et al. 1998). However, the finding that addition of only one or two AACGTT sequences to a plasmid was critical for vaccination success (SATO et al. 1996) is very surprising given the number of potential activating sequences in any plasmid. Analysis of the capacity of mouse skin, grafted from sites that had been vaccinated by gene gun, to generate immune responses in recipient and donor mice showed that cells migrating from the site within 12h after vaccination are responsible for the generation of immulological memory (KLINMAN et al. 1998). Condon et al. (CONDON et al. 1996) showed that dendritic cells are directly transfected in gene-gun-mediated DNA vaccination and migrate from the skin to draining lymph nodes. These studies have not claimed that DNA alone induces APC migration, as at least some of the APCs involved were directly transfected by plasmid-coated gold particles (CONDON et al. 1996), but this is quite likely to be the case. Intradermal injection of CpG DNA caused activation of APCs as measured by increased MHC class-II, CD86 and IL-12 expression amongst epidermal cells 12h after injection (JAKOB et al. 1998). Study of the maturation of APCs within skin may be challenging, as they are likely to have migrated by the 12-h time point (KLINMAN et al. 1998), and they may have to be followed to the draining lymph node.

### 4.4 A Role for DNA Activation in Infection?

The involvement of unmethylated CpG sequences with macrophage/APC stimulation in DNA vaccination and with active oligonucleotides is well accepted, but the role of DNA as an immune stimulant during natural infection is difficult to assess. Some bacteria obviously contain other stimulatory molecules, such as LPS, which may have a quantitatively greater effect on the immune system than the foreign DNA to which the organism becomes exposed. However, detection of foreign DNA may be important in synergising with other activating molecules or in detection of bacterial strains that have avoided other means of surveillance. Another suggestion (PISETSKY 1996) is that the detection of foreign DNA may be of most importance in viral infections, since viruses do not contain the range of lipid and sugar molecules that generally identify bacteria and pathogens to the innate immune system.

In an infection, macrophages engulf and kill bacteria, with subsequent degradation of cellular components. Fragments of DNA would either have to escape to the cytoplasm or nucleus, or actually be detected within the phagosome in order to allow immune response. Alternatively, DNA escaping from the cell could stimulate other cells. One study looked at the fate of DNA from *E. coli* phagocytosed by macrophages and found that much of the genomic DNA was degraded

into short pieces and released into the medium, whereas plasmid DNA remained intact (ROZENBERG-ARSKA et al. 1984). Whether or not this DNA will stimulate the phagocytosing cell is difficult to assess, but it remains clear that exogenously added free CpG DNA is stimulatory. In order for the DNA of an infecting virus to be detected, it would be anticipated that the putative CpG DNA receptor would have to be located in the cytoplasm or nucleus rather than in the endosomes. Real assessment of the role of DNA as an immunostimulant in an infection awaits elucidation of the activating DNA receptor and generation of knockout mice or other approaches to interfering with its actions.

### **5** Summary

Macrophage/dendritic cells and B cells remain the only cell types where direct responses to CpG DNA are well established. The role of macrophages in vivo in DNA clearance and the potent cytokine induction in macrophages and dendritic cells places them in the central role in the in vivo response to foreign DNA. Although responses to DNA are unlikely to evolve and be retained if they are not significant in the immune response to infection, the relative contributions of DNA and other stimulators of the innate immune recognition of foreign organisms is difficult to assess. Although CpG DNA and LPS have similar actions, significant differences are emerging that make the use of DNA as a therapeutic immunostimulatory molecule feasible. The macrophage response to DNA generates cytokines favouring the development of Th1-type immunity, and active oligonucleotides now show promise as Th1-promoting adjuvants and as allergy treatments.

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# **Consequences of Bacterial CpG DNA-Driven Activation** of Antigen-Presenting Cells

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## **1** Introduction

Infectious pathogens that break through surface defences and reach underlying tissue or blood encounter an array of interior host defences. These fall basically into two classes: innate and adaptive defences. The innate system is the phylogenically older of the two branches and uses germline-encoded pattern-recognition receptors (PRR) to identify microbial invaders (PUGIN et al. 1994). An example is CD14, a receptor for bacterial cell-wall components (CWCs), such as lipopolysaccharides (LPS) and lipoteichoic acids. CD14 recognises carbohydrate constituents commonly expressed on pathogens (ESPEVIK et al. 1993; ZHANG et al. 1994; ULEVITCH and TOBIAS 1995; CLEVELAND et al. 1996) and signals "danger" to myeloid-lineage cells (MATZINGER 1994).

The reliance on carbohydrate recognition alone by innate defences would have the disadvantage of genetic inflexibility, since escape mutants lacking the target structures might arise. Because of this limitation, alternative mechanisms may exist

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that activate protective innate defences. The notion that bacterial DNA might be one candidate ligand is appealing for several reasons. Although considered immunologically inert for decades, bacterial DNA displays properties required for a perfect "danger signal", including structural diversity, species specificity and the ability to cross-link receptors (PISETSKY 1996). Importantly, bacterial and vertebrate DNAs differ in their relative abundances of CpG dinucleotides, termed CpG suppression, within vertebrate DNA. Furthermore, in contrast to bacterial DNA, vertebrate DNA displays a high degree of cytosine methylation (DOERFLER 1991; HERGERSBERG 1991; BIRD 1992). These unexplained structural differences (DOER-FLER 1991) may be viewed as the genetic code for recognition of bacterial DNA, independent of surface antigens, such as LPS (PISETSKY 1996). Specific conserved, static DNA sequence structures appear to be a key enabling the vertebrate immune system to distinguish danger from non-danger.

The first report detailing the immune-stimulating properties of bacterial DNA was by Tokunaga and co-workers, who succeeded in attributing the tumoricidal effects of *Mycobacterium bovis* bacille Calmette-Guerin (BCG) to natural killer (NK) cell activation by mycobacterial DNA (TOKUNAGA et al. 1984). Subsequently, the effects of bacterial DNA on cells of the adaptive immune system were observed. Pisetsky and co-workers demonstrated an LPS-like proliferation of murine B lymphocytes on in vitro stimulation with DNA from *Escherichia coli* (MESSINA et al. 1991, 1993). Consequently, unexpected immune-stimulatory effects of synthetic oligodeoxynucleotides (ODNs) used as antisense or control ODNs were reported; some ODNs also triggered polyclonal proliferation of murine and human B cells in a sequence-specific manner (HELENE and TOULME 1990; TANAKA et al. 1992; MCINTYRE et al. 1993; PISETSKY and REICH 1993; PISETSKY and REICH 1994; BRANDA et al. 1996a; LIANG et al. 1996). In 1995, KRIEG et al. defined bacterial DNA and synthetic, unmethylated CpG-DNA sequences as potent B-cell mitogens (KRIEG et al. 1995).

The discovery of bacterial immune-stimulating DNA sequences has two major implications: bacterial DNA is not inert but activates immune cells; thus, immune cells can sense pathogens via pathogen DNA. This review will focus both on the recognition of bacterial CpG DNA by cells of the innate immune system and on the pathophysiologic and possible therapeutic consequences associated with this pattern recognition.

## 2 Activation of Macrophages

Tokunaga and co-workers discovered that a nucleic-acid-rich fraction from BCG induces NK cell lytic activity and interferon- $\gamma$  (IFN- $\gamma$ ), - $\alpha$  and - $\beta$  secretion by murine spleen cells. The immune-stimulating capacity of bacterial DNA was DNase sensitive but RNase resistant and could not be mimicked by vertebrate DNA (S.E. Yamamoto, this issue). Comparable biologic activity was displayed by synthetic 30–45-mer single-stranded (ss) ODNs; these ODNs contained certain

palindromic hexamer motifs, such as AACGTT and GACGTC, and were derived from cDNA encoding for a BCG protein (TOKUNAGA et al. 1992; YAMAMOTO et al. 1992). In addition, similar sequences caused B-cell proliferation, and it was concluded that immunostimulatory bacterial "CpG-motifs" are characterised by a more liberal formula that does not require palindromic sequences but contains the motif 5'-purine-purine-CG-pyrimidine-pyrimidine-3' (KRIEG et al. 1995). If the central CpG motif was methylated or mutated, the immune-stimulating properties were lost. The efficacy of the CpG motif is influenced by flanking sequences (PISETSKY and REICH 1998). Interestingly, sequence rules that govern recognition of bacterial DNA by murine immune cells differ from the rules for human cells (LIANG et al. 1996; BAUER et al. 1999). While CpG DNA directly activated murine and human B cells, further studies revealed that the originally observed biological effects of bacterial DNA on NK cells were dependent on the presence of adherent spleen cells, pointing to a direct or indirect contribution of myeloid cells (HALPERN et al. 1996; CHACE et al. 1997).

In 1996, the first direct proof of the susceptibility of macrophages towards CpG DNA was published. Bacterial plasmid DNA was shown to activate the transcription factor nuclear factor  $\kappa B$  (NF $\kappa B$ ) and to induce tumour necrosis factor  $\alpha$  (TNF- $\alpha$ ) mRNA in murine bone-marrow-derived macrophages and in the macrophage cell line RAW 264. In contrast to LPS-inducible genes like inducible nitric oxide synthase (iNOS), induction of iNOS mRNA in RAW cells using CpG DNA as stimulus required IFN- $\gamma$  pretreatment, suggesting differences in the signal pathways (STACEY et al. 1996). In this study, initial experiments to elucidate the molecular mechanism of CpG DNA were performed. Plasmid DNA containing CpG sequences was shown to be taken up from the medium and to code for a reporter protein. While genomic DNA is known to be degraded within the lysosomes after receptor-mediated uptake, plasmid DNA apparently partly survives and reaches the nucleus (BENNETT et al. 1985). The question of whether recognition of bacterial DNA requires indirect interaction (after binding to cytoplasmic proteins) or direct interaction with transcription factors or if active DNA fragments bind e.g. endosomal receptors, thus initiating downstream signaling events, remained unanswered.

Subsequently, it was demonstrated that primary murine macrophages were activated in vitro and in vivo by genomic DNA from gram-positive (*Staphylococcus aureus, Streptococcus faecalis, Micrococcus lysodeikticus*) and gram-negative (*E. coli*) bacteria. In addition, synthetic oligonucleotides containing unmethylated CpG motifs were also effective in activating peritoneal macrophages and the macrophage cell line ANA-1. In vitro, bacterial DNA or CpG ODNs triggered nuclear translocation of NF $\kappa$ B, accumulation of pro-inflammatory cytokine mRNAs and subsequent release of substantial amounts of TNF- $\alpha$ , interleukin 1 $\beta$  (IL-1 $\beta$ ) and IL-6, which are known to be involved in the pathogenesis of septic shock (SPARWASSER et al. 1997a,b and unpublished data). Furthermore, cytokines [such as granulocyte–macrophage colony-stimulating factor (GM-CSF), IL-12 and IL-10] and chemokines [such as macrophage inflammatory protein 1 $\alpha$  (MIP-1 $\alpha$ ), MIP-1 $\beta$ , MIP-2 and monocyte chemoattractant protein 1 (MCP-1)] could be

induced in macrophage cell cultures (LIPFORD et al. 1997b and unpublished data). These observations could also be extended to human peripheral blood monocytes that produce IL-12, IFN- $\alpha$ , IL-6 and TNF- $\alpha$  upon stimulation with bacterial DNA or synthetic ODNs (ROMAN et al. 1997; BAUER et al. 1999). Activation of murine macrophages and human peripheral monocytes could also be demonstrated by upregulation of surface markers, such as CD86, CD40, and major histocompatibility complex (MHC) class-II molecules (BAUER et al. 1999 and our own unpublished data). In comparative studies, similarities and dissimilarities between LPS-and DNA-induced signal transduction pathways are evident (HÄCKER et al. 1998; YI and KRIEG 1998). However, the parallels between LPS and the structurally unrelated CpG DNA as strong inflammatory stimuli for macrophages are remarkable. Even though we have still a poor understanding of how CpG DNA mediates acute sequence-dependent cell activation, some questions about intracelular signaling are being unravelled (A. Krieg and H. Häcker, this issue).

## 3 Activation and Maturation of Dendritic Cells

Dendritic cells (DCs) originate from a common bone-marrow precursor cell giving rise to macrophages, granulocytes and DCs. The macrophage and myeloid DC lineages are not as strictly separated as was hypothesised in the past, e.g. because blood monocytes can differentiate under certain conditions into either DCs or macrophages (SZABOLCS et al. 1996). However, there are several essential differences between DCs and macrophages. DCs, as professional antigen-presenting cells (APCs), are most efficient in the activation of resting T cells and have the unique capacity to activate naive T cells. DCs exist in two functional states: immature antigen-capturing DCs and mature antigen-presenting DCs (STEINMAN 1991).

Immature DCs are abundant in peripheral non-lymphoid tissues, such as skin, and can be rapidly recruited to sites of local infection (Fig. 1). Immature DCs display a high endocytotic activity and high numbers of surface Fc receptors but low numbers of surface MHC class-II and co-stimulatory molecules. In order to initiate primary immune responses, immature DCs need to become activated and to differentiate into mature DCs able to present antigen in the MHC class-II and class-I contexts. This maturation/activation signal can be delivered by microbial products, such as LPS (BANCHEREAU and STEINMAN 1998). Interestingly, bacterial CpG DNA can substitute for this classic danger signal. Murine bone-marrow-derived DCs (BMDDCs) cultured in vitro in the presence of GM-CSF could be activated by CpG DNA, in a CD40 ligand-independent manner, to upregulate MHC class-II molecules and express co-stimulatory molecules, such as CD86 and CD40 (SPARWASSER et al. 1998). Low concentrations of CpG ODNs and bacterial genomic DNA efficiently induced secretion of pro-inflammatory cytokines, such as TNF-α, IL-12 and IL-6. The functional maturation of BMDDCs could be demonstrated in vitro both in mixed lymphocyte reactions and in staphylococcal



Site of antigen deposition: Pathogens, CWC, bCpG-DNA

Fig. 1. Antigen-presenting cells (APCs) as decisive interfaces between the innate and the adaptive immune systems. After recruitment of APC progenitor cells from the bone marrow (1) to the site of antigen challenge, pathogens and pathogen-derived stimuli, such as bacterial cell-wall components and bacterial CpG DNA, cause activation of macrophages and dendritic cells (DCs). Immature DCs, upon activation, maturate and migrate to secondary lymphoid organs, where they translocate to T-cell areas (2). Naive T cells enter the draining lymph nodes via high endothelial venules (3) and interact with mature antigenpresenting DCs. In contrast to antigen-capturing immature DCs, mature antigen-loaded DCs display high numbers of surface major histocompatibility complex class-II complexes and co-stimulatory molecules (CD80, CD86). Depending on the expression of these molecules and the local cytokine environment (interleukin 12), cellular T-helper (Th)1 or humoral Th2-oriented immune responses are triggered (4). Activated cytotoxic and helper T cells migrate to the site of infection, where they act, in concert with other effector cells of the adaptive and innate immune systems, to fight the microbial invaders (5)

enterotoxin B (SEB)-driven naive T-cell responses (SPARWASSER et al. 1998). A similar report showed activation of Langerhans' cell (LC)-like murine foetal-skinderived DCs upon stimulation with CpG ODNs and *E. coli* DNA in vitro. In vivo, a small subset of MHC-II-bearing LCs exhibited upregulation of MHC II and CD86 and intracellular accumulation of IL-12 (JAKOB et al. 1998).

Matured and activated DCs migrate to secondary lymphoid organs, where they translocate to the T-cell areas (DE SMEDT et al. 1996; REIS E SOUSA et al. 1997).

After subcutaneous injection of CpG ODNs or plasmid DNA, activation of DCs from the draining lymph nodes could be observed. Importantly, in addition to upregulation of CD80, CD86 and CD40, local IL-12 production within the T-cell areas could be detected (AKBARI et al. 1999 and our own unpublished data). Naive T cells, attracted and guided by chemokine/integrin-mediated mechanisms, enter the lymph nodes via high endothelial venules to interact with antigen-loaded DCs, leading to the generation of T-helper 1 (Th1)- or Th2-type effector cells (SALLUSTO et al. 1998). The interaction of activated DCs with T cells in the presence of IL-12 is thought to be responsible for a Th1-biased cellular immune response (TRINCHIERI 1995). A recent hypothesis proposes a 'dynamic' interaction between first DCs and CD4<sup>+</sup> T cells and subsequent CD8<sup>+</sup> T-cell activation triggered by the activated DC (RIDGE et al. 1998). In this model, CD40L-CD40-mediated interaction between DCs and T-helper cells is necessary to induce co-stimulatory molecules on DCs triggering cytotoxic T-cell (CTL) activation. The demonstration of CD40Lindependent DC activation suggests that it is possible to bypass T-helper celldependent mechanisms (JAKOB et al. 1998; SPARWASSER et al. 1998). In vaccination studies, for example, CD4-deficient mice can mount cellular immune responses if co-injected with CpG ODNs as adjuvants (our own unpublished data).

## 4 Pathophysiologic Consequences of Innate Immune Cell Activation

Bacterial CpG DNA operationally resembles bacterial LPS, a CWC known to be essential for morbidity and mortality caused by gram-negative bacteria. The question arises whether bacterial DNA, either liberated by lysosomal degradation within macrophages or released extracellularly from dead micro-organisms, can also trigger pathology.

## 4.1 Role of Bacterial DNA in Toxic Shock

Macrophages are regarded as key cells in the pathogenesis of septic shock (GLAUSER et al. 1991; RIETSCHEL and WAGNER 1996b). They are known to be activated by bacterial CWCs to release toxic amounts of pro-inflammatory cytokines that can lead to tissue damage, multi-organ failure and death (HACK et al. 1997). Cells of the innate system may be able to detect bacterial pathogens independent of CD14, the receptor for CWCs, such as LPS. This question is intriguing for several reasons. In clinical settings, about 50–60% of cases of septic shock are caused by gram-negative and about 35–40% by gram-positive bacteria (GLAUSER et al. 1991; BRANDTZAEG 1996). While the physiological significance of the CWCs of gram-negative bacteria (such as LPS, or endotoxin) is well established, the role of lipoteichoic acids and peptidoglycans in triggering gram-positive septic shock is less clear (WAKABAYASHI et al. 1991; DE KIMPE et al. 1995; KUSUNOKI et al. 1995; RIETSCHEL et al. 1996a). Some gram-positive bacteria produce exotoxins, like SEB. that can cause superantigen-mediated toxic shock syndrome and sensitise for sublethal doses of LPS (MARRACK and KAPPLER 1990; MIETHKE et al. 1992; BLANK et al. 1997). The majority of gram-positive shock cases are superantigen independent, implying that unknown components derived from gram-positive bacteria have a role in pathophysiology (GLAUSER et al. 1991). Interestingly, while the inflammatory response towards whole gram-negative or gram-positive bacteria in acute cytokine-release syndrome is comparable, purified LPS from gram-negative bacteria is more than 1000-fold more potent in activating macrophages than purified gram-positive CWCs (KUSUNOKI et al. 1995 and references within). Importantly, LPS-non-responder mice are susceptible to septic shock triggered by both heat-killed gram-positive and gram-negative bacteria (GALANOS and FRE-UDENBERG 1991). This observation suggests that additional signaling pathways are operative in septic shock, and one might claim that LPS-independent mechanisms trigger the activation of macrophages.

To investigate a possible role of bacterial DNA in septic shock, CpG ODNs and DNA from gram-positive and gram-negative bacteria were tested in a mouse shock model. These stimuli activated murine macrophages and macrophage cell lines in a manner similar to LPS. Bacterial CpG DNA triggered cytokine release by APCs, culminating in vivo in acute systemic release of pro-inflammatory cytokines and leading to TNF- $\alpha$ -mediated lethal toxic shock in D-galactosamine (D-GalN)sensitised mice (SPARWASSER et al. 1997a,b). Of note, LPS and active CpG ODNs synergized in vivo in their ability to trigger TNF- $\alpha$  release. This synergy may be mediated through the common activation of NF $\kappa$ B, and one might claim that there are unrecognised correlates in the gram-positive setting. In addition to the direct toxicity observed, bacterial DNA also sensitises mice (in a Shwartzman-like fashion via IFN- $\gamma$ ) to subsequent toxic shock from sub-lethal LPS doses (CowDERY et al. 1996 and our own unpublished results). This effect may be important in chronic infections.

#### 4.2 Bacterial CpG DNA and Inflammation

Unquestionably, systemic release of cytokines leading to septic shock associated with high lethality is the worst outcome of bacterial infections. However, it will be necessary to document whether bacterial DNA liberated in the course of infection is physiologically relevant in causing local tissue damage and inflammation. Schwartz et al. demonstrated that bacterial DNA, instilled intratracheally, causes significant inflammation in the lower respiratory tracts of mice. Increased numbers of mononuclear cells and increased levels of pro-inflammatory cytokines (TNF- $\alpha$ , IL-6, MIP-2) could be detected in whole-lung lavage after application of *E. coli* DNA or CpG ODNs, whereas LPS instillation caused a negligible inflammatory response (SCHWARTZ et al. 1997). Furthermore, up to 1% of DNA extracted from the sputum of cystic fibrosis (CF) patients suffering from increased and abnormal
mucus production and severe, recurrent bacterial infections was found to be of bacterial origin. Interestingly, these total DNA isolates induced acute inflammation in the mouse model similar to inflammation caused by E. *coli* DNA or CpG ODNs. One may speculate that, due to the local accumulation of bacterial DNA, tissue damage and inflammation persist in these patients even after antibiotic treatment (SCHWARTZ et al. 1997).

#### **5** Possible Therapeutic Applications

Bacterial immunostimulatory DNA sequences may trigger inflammatory disease or may even lead to septic shock. However, "beneficial" physiological host-defence functions, such as the activation and mobilisation of immune cells and the recognition of potentially dangerous pathogens, may be of major importance.

The tumoricidal effects of immune-stimulating DNA sequences were the first immunological effects of bacterial DNA to be reported and were considered to be "beneficial" (TOKUNAGA et al. 1984). The enhanced lytic anti-tumor activity of NK cells induced by prokaryote DNA could be demonstrated both in vitro and in vivo and was mediated by APC-derived cytokines, IL-12 and, to a minor extent, TNF- $\alpha$ and IFNs (HALPERN et al. 1996; CHACE et al. 1997). NK-stimulatory activity can be reproduced in the human system in vitro and enhances the efficacy of monoclonalantibody therapy in a mouse lymphoma model (BALLAS et al. 1996; WOOLDRIDGE et al. 1997). Here, further potentially therapeutic applications of CpG ODNs mediated by the activation of APCs will be discussed.

#### 5.1 Role of CpG DNA as an Adjuvant

The adaptive immune system partially depends on innate immunity to distinguish infectious non-self from non-infectious self (JANEWAY 1992; FEARON and LOCKSLEY 1996). Danger versus non-danger information is delivered by cells of the innate system, such as macrophages or DCs sensing pathogens via pattern recognition. If non-self antigens associated with pathogens are processed and recognised as dangerous, the APCs become activated. Once activated, APCs induce productive T-cell responses, while non-activated APCs can cause peripheral tolerance (EHL et al. 1998). The significance of this dynamic interaction between innate and adaptive immunity might have been underestimated in the past (FEARON and LOCKSLEY 1996; FEARON 1997). In fact, the need for bacterial products as APC stimuli to elicit humoral and cellular immune responses towards purified proteinaceous antigens, referred to as the "dirty little secret of immunologists", illustrates the importance of the innate system (JANEWAY 1989).

To enhance immunisation efficiency, various microbial products have served as adjuvants. It is believed that adjuvants primarily target APCs. These APCs have at

least two duties. First, they process and present antigen peptides to T cells via MHC molecules and, after activation, they express co-stimulatory molecules. Second, APCs secrete pro-inflammatory cytokines, thereby acquiring the ability to instigate productive T- and B-cell activation (JENKINS et al. 1991; GALVIN et al. 1992; JENKINS and JOHNSON 1993). Polarisation of activated T-helper cells into either Th1 or Th2 subsets is the result of secondary regulatory events and depends on APC secretion of effector cytokines, such as IL-12 and IL-10 (MOSMANN and COFFMAN 1989; ABBAS et al. 1996). Activation signals can be conveyed by pattern recognition of bacterial products, i.e. via CD14 or bacterial DNA.

The classical adjuvant CFA (complete Freund's adjuvant) contains dead mycobacteria in an oil-in-water emulsion (KE et al. 1995). This adjuvant is widely used in experimental animal protocols; however, massive local inflammatory response precludes its use in humans. Various adjuvants have been shown to differentially activate Th1 and Th2 subsets, with CFA tending to induce Th1 subsets, while IFA (incomplete Freund's adjuvant), which does not contain mycobacteria, produces Th2-like responses (FORSTHUBER et al. 1996). Strong macrophage activation caused by CpG DNA has been shown by different groups (STACEY et al. 1996; CHACE et al. 1997; SPARWASSER et al. 1997b). Two recent studies demonstrate the activation and maturation of DCs upon stimulation with CpG DNA (JAKOB et al. 1998; SPARWASSER et al. 1998). Because of the powerful APC-stimulatory qualities of bacterial DNA or CpG ODNs, one might ask if these substances could substitute for heat-killed mycobacteria as adjuvants.

The first studies using CpG ODNs as adjuvants were performed when only its capacity to upregulate B7 and MHC-II molecules in B cells was known. CpG ODNs were shown to enhance antibody production to tetanus toxoid (BRANDA et al. 1996b). Subsequently, CpG ODNs were evaluated in a protocol for vaccination against the soluble antigen ovalbumin (OVA) (LIPFORD et al. 1997a). Two effects were prominent. First, immunostimulatory ODNs increased specific immunoglobulin G2 (IgG2)-antibody titers by more than 100-fold, demonstrating a Th1 bias. Second, the use of CpG ODNs as adjuvants allowed the induction of primary CTL responses to either unprocessed protein antigen (OVA) or its immunodominant peptide (SIINFEKL) encapsulated in liposomes. The CTL induction was independent of B-cell activation, and B-cell bystander effects as demonstrated through the use of B-cell-deficient µMT mice. Whereas direct APCactivating effects are now generally accepted, the role of direct effects of CpG ODNs on T cells are still controversial. In vitro, CpG DNA co-stimulatory effects can be observed after T-cell-receptor engagement (BENDIGS et al. 1999). In contrast, in vivo data demonstrated T-cell activation only in the presence of APCs secreting type-I IFNs, suggesting that ODNs, as adjuvants, function primarily by potentiating APC function (Sun et al. 1998b).

The Th1-biasing effect of bacterial CpG sequences used as adjuvants has been described by various other groups (RAZ et al. 1996; CHU et al. 1997; ROMAN et al. 1997; WEINER et al. 1997; BRAZOLOT MILLAN et al. 1998; DAVIS et al. 1998; SUN et al. 1998a). Importantly, immunisation protocols leading to Th1-type cellular immunity can be essential for protective immune responses against infections. The

skewing of CpG-ODN-enhanced antibody responses towards the IgG2 isotypes indicates Th1-dependent cytokine-induced class switching (SNAPPER and PAUL 1987; FINKELMAN et al. 1990). In fact, the cytokine profile induced by CpG ODNs includes mainly IL-12, IL-18 (IFN- $\gamma$ - inducing factor: IGIF) and IFN- $\gamma$  and, thus, favours Th1-oriented T-cell responses (HALPERN et al. 1996; KLINMAN et al. 1996; LIPFORD et al. 1997b; ROMAN et al. 1997; MCCLUSKIE and DAVIS 1998; SPAR-WASSER et al. 1998). Interestingly, findings similar to those observed with synthetic ODNs can be demonstrated in DNA vaccination studies. Briefly, these vaccines consist of a bacterial plasmid backbone and a gene insert coding for the antigen. After intramuscular or intradermal injection of the plasmid, it is thought that host cells, including DCs, take up the plasmid and, thus, express the antigen. Because of its bacterial origin, these plasmids contain immunostimulatory CpG sequences. Operationally, "naked DNA" may, thus, harbour its adjuvant while acting as a translation unit. Indeed, recent evidence suggests that the immunogenicity of naked DNA is grossly influenced by the presence or absence of immunostimulatory CpG sequences, suggesting a key role in activating transfected APCs (AKBARI et al. 1999; KLINMAN et al. 1997; ROMAN et al. 1997; SATO et al. 1996). Apparently, the co-administration of synthetic CpG ODNs and plasmid DNA vaccines can reduce the efficacy of the vaccines, possibly due to competitive binding at cellular receptors (WEERATNA et al. 1998). Synthetic ODNs offer important advantages as adjuvants. They can be produced on a large scale with immense purity, combining the advantages of enormous stability and inexpensive synthesis as compared with protein or plasmid-based vaccines. Compared to a CpG-sequence-containing DNA vaccine, synthetic CpG ODNs, as adjuvants, may be more effective in inducing protective immune responses (BRAZOLOT MILLAN et al. 1998). In direct comparison, bacterial CpG DNA may even surpass CFA as an adjuvant (CHU et al. 1997; SUN et al. 1998a). Furthermore, since mucous membranes represent an important transmission site for many pathogens and since, to date, no adjuvant has been licensed for clinical application in mucosal vaccinations, its use as a mucosal adjuvant might be promising (McCluskie and Davis 1998; Moldoveanu et al. 1998).

Overall, recent data indicate that bacterial DNA and immunostimulatory ODNs convey an adjuvant effect, in that they activate professional APCs to express co-stimulatory molecules and secrete cytokines pivotal for the initiation of productive Th1 responses. The use of immunostimulatory DNA sequences in vaccination protocols could be an interesting therapeutic application, because vaccines are the most commonly used immunotherapeutics (KLINMAN 1998).

#### 5.2 CpG DNA as a Therapeutic Agent in Th2-based Pathology

APCs, as "sentinels at the gate", can strongly influence the differentiation of adaptive immune cells towards either cell-mediated (Th1) or antibody-mediated (Th2) forms of immunity. Intracellular pathogens, such as viruses, certain bacteria and protozoans, tend to induce cellular immunity, including CTL activity and the

production of pro-inflammatory cytokines, such as IFN- $\gamma$  and TNF- $\alpha$ . Extracellular pathogens, e.g. helminths, induce primarily humoral immune responses characterised by vigorous IgG1 and IgE responses and IL-4/IL-5/IL-10-dominated cytokine patterns (CONSTANT and BOTTOMLY 1997). In order to mount cellular immune responses, IL-12 needs to be present at the time of antigen recognition, driving NK cell- and Th1 cell-derived IFN- $\gamma$  production and thereby inhibiting Th2 development. In addition to the appropriate "cytokine environment", co-stimulatory signals delivered by innate immunity contribute to the decision of whether Th1- or Th2-biased adaptive responses result (SEDER and PAUL 1994; CONSTANT and BOTTOMLY 1997). Th2-based pathology can be caused by two different mechanisms: (1) inappropriate Th2 responses e.g. to environmental antigens or vaccines or (2) insufficient (humoral) immune responses against intracellular microorganisms.

Apart from vaccination, for which a Th1-directed situation mediating cellular immunity is often preferable, there are other clinical situations where a Th1 shift is of therapeutic use. Allergies and asthma are, for example, characterised by a Th2dominated status (FINKELMAN 1995). CpG ODNs may reduce the risk of anaphylactic shock, an extreme Th2 response and potentially fatal side effect of vaccination by its Th1 skewing effects (reduced IgE synthesis, inhibition of Th2 cells; CARSON and RAZ 1997). Co-administration of CpG ODNs can turn a Th2dominated immune response caused by IFA towards hen-egg lysozyme into a Th1type response, as defined by cytokine profile and IgG isotypes (CHU et al. 1997). In a murine asthma model, systemic administration of CpG ODNs appears to be beneficial even in animals already sensitised to the antigen (schistosome eggs), which normally causes airway eosinophilia, Th2 cytokine production, IgE production and bronchial hyper-reactivity (KLINE et al. 1998). In another mouse model of Th2-mediated pathology, CpG sequences could confer immediate and sustained protection against allergen-induced airway hyper-responsiveness (BROIDE et al. 1998). While the acute effects were caused by innate immune cell-derived Th1promoting cytokines, such as IL-12 and IFNs, the lasting effects over a time period of 1 week were explained by the inhibition of the bone-marrow production of eosinophils (BROIDE et al. 1998). These studies suggest that not only the initiation but also ongoing Th2 responses can be prevented and cured by the therapeutic use of CpG ODNs. Moreover, this treatment can confer lasting protection against inappropriate generation of Th2 responses.

Murine leishmaniasis is a well-characterised model of Th1/2-dependent pathology, rendering Th1-biased mouse strains resistant and Th2-biased strains sensitive to the intracellular pathogen (HEINZEL et al. 1989; HSIEH et al. 1995; REINER and LOCKSLEY 1995). Th2-biased BALB/c mice treated with immunostimulatory CpG sequences can be cured of normally lethal leishmania infection (ZIMMERMANN et al. 1998). Resistance is likely to be explained by APC-derived cytokines, such as IL-12 and IL-18, and NK cell-derived IFN- $\gamma$ , which promotes Th1-type cellular immunity (HEINZEL et al. 1993; SYPEK et al. 1993).

It might be more than a coincidence that, in developed countries, the decrease of Th1-promoting infectious diseases, such as tuberculosis and measles, correlates with a significant increase in Th2-driven atopic disorders (SHAHEEN et al. 1996; SHIRAKAWA et al. 1997). In addition to natural exposure to infectious diseases, other factors, such as dietary changes, "indoor life-style", increasing hygiene standards and Th2-promoting modern vaccinations, influence the fine-tuning of the T-cell repertoire and Th1/Th2 balance (ROMAGNANI 1994). Because "Westernisation" continually deprives the human immune system of its evolutionary input, it may be necessary to develop vaccination strategies that not only protect against specific pathogens but also maintain the correct cytokine balance and education of the adaptive immune system (ROOK and STANFORD 1998).

### 6 Conclusion

APCs, bridging the innate and adaptive immune systems, recognise prokaryotic DNA as a danger signal. This recognition of bacterial DNA mediates activation of macrophages and DCs, leading to secretion of pro-inflammatory cytokines, upregulation of co-stimulatory molecules and production of effector cytokines influencing the adaptive immune system's responses. Systemic toxicity of CpG DNA, triggering a cytokine-release syndrome culminating in toxic shock and local pathology caused by inflammatory responses, may represent examples of "harmful" consequences for APC activation. These same responses, however, contribute to CpG DNA's function as a therapeutic adjuvant, activating APCs and thereby enabling protective Th1-biased immune responses.

Bacterial CpG DNA is a good candidate for a conserved pattern-recognition molecule (KRIEG 1996; PISETSKY 1996; HEEG et al. 1998; WAGNER 1999). The mechanism of pattern recognition via non-self DNA also appears to extend to other invertebrate DNAs that have been reported to activate immune cells (SUN et al. 1996, 1998b; BROWN et al. 1998). For example, DNA from *Drosophila* causes upregulation of co-stimulatory molecules in murine B cells and activation of APCs leading to cytokine release (SUN et al. 1996, 1998b). The immunostimulatory potency of viral double-stranded RNA was recognised in the early 60s. At that time, the first suspicions about immunostimulatory prokaryotic DNA arose but was questioned later (VILCEK et al. 1969). It is also possible that the mechanism of pathogen-DNA pattern-recognition originally evolved as a way to recognise viruses lacking LPS-like surface structures, thus activating infected cells via "foreign" genetic material. The postulated CpG-PRR might have evolved to signal danger to APCs during infection with gram-positive bacteria, parasites or certain viruses.

To substantiate this proposal, there is a need to evaluate the physiological relevance of prokaryotic DNA (released either extra- or intracellularly during infection) in causing activation of APCs. Independent of whether sensing of foreign DNA represents an evolutionarily developed primordial recognition system signaling danger to cells of the innate immune system, its use as a natural adjuvant in promoting Th1 responses is promising.

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# Signal Transduction Pathways Activated by CpG-DNA

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# **1** Introduction

In the last few years, it has been shown that bacterial DNA stimulates and is recognized by cells of the immune system (WAGNER 1999). Cell types that have been shown to respond directly to bacterial DNA include cells of the innate immune system, such as macrophages and dendritic cells, as well as B cells. Notably, the pattern of responsive cell types strongly resembles the pattern of cells stimulated by other so-called pattern-recognition factors, such as lipopolysaccharides (LPS). In addition, the spectrum of effects induced by these biochemically different agents are remarkably similar to each other. It includes induction of a variety of soluble factors, upregulation of membrane proteins and regulation of cell proliferation and survival. Due to the profound alterations in the cells' behaviour upon stimulation

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by CpG-DNA, it has been compelling to postulate changes in the transcriptional and post-transcriptional activities. How can the information contained in unmethylated CpG-motifs in bacterial DNA be translated into gene expression? One possibility is that CpG-DNA, like other common ligands, engages a specific cellular receptor that transduces the signal by signal transduction pathways from the outside to the nucleus. Recently, it was shown that classic signal transduction pathways, such as the stress-kinase pathway and the nuclear factor  $\kappa B$  (NF- $\kappa B$ ) activation pathway, are switched on in response to CpG-DNA. The current knowledge about these pathways in the context of their activation by CpG-DNA and the upstream requirements for signal initiation are discussed in this chapter.

## 2 The NF-KB Activation Pathway

The transcription factor NF- $\kappa$ B is expressed in virtually all cell types of eukaryotes. In the immune system, it is involved in the inducible expression of proteins, including soluble factors like cytokines and chemokines, membrane proteins involved in co-stimulation and adhesion (GRILLI et al. 1993; KOPP and GHOSH 1995) and proteins regulating cell survival (KARIN 1998). In general, activation of NF-KB leads to an enhancement of immune responses. NF-kB is composed of dimers of the Rel family of DNA-binding proteins, RelA/p65, p50, p52, c-Rel, RelB and Bcl-3 (GHOSH et al. 1998). In most cell types, NF-KB activity is tightly controlled by inhibitory proteins, the I $\kappa$ Bs. Like the NF- $\kappa$ B family, I $\kappa$ B is a family of molecules; it includes IkBa, IkBb, IkBy and IkBe (Ghosh et al. 1998). Bcl-3 structurally resembles other IKB family members. However, because Bcl-3 primarily seems to stimulate NF- $\kappa$ B-dependent gene transcription (Bours et al. 1993), it is here included in the NF-kB family. By interaction with the pre-formed NF-kB complexes, I $\kappa$ Bs inhibit both the translocation of NF- $\kappa$ B to the nucleus and DNA binding. The best-characterized members of the IkB family are IkBa and IkBB. Cell stimulation, for example by pro-inflammatory cytokines, leads to rapid phosphorylation of two conserved serine residues within the regulatory N-terminal domain of IkB, followed by rapid ubiquitination and degradation of  $I\kappa B\alpha/\beta$  by the 26S proteasome. The released NF- $\kappa$ B dimer translocates to the cell nucleus, free to bind to NF- $\kappa$ B enhancers and activate gene transcription. Recently, two proteins –  $I\kappa B$  kinase  $\alpha$ (IKK $\alpha$ )/conserved helix-loop-helix ubiquitous kinase (Chuk) and I $\kappa$ B kinase  $\beta$  $(IKK\beta)$  – have been shown to contribute to a higher molecular weight complex of proteins with IkB-kinase activity (DIDONATO et al. 1997; REGNIER et al. 1997; ZANDI et al. 1997). This IKK complex specifically phophorylates IkBa and IkBβ at the serine residues critical for ubiquitination and degradation (KARIN 1998). For IKK $\alpha$ /Chuk, direct interaction with and activation by NIK, the NF- $\kappa$ B inducing kinase, has been described (REGNIER et al. 1997). Tumor necrosis factor (TNF)and interleukin 1 (IL-1)-induced activation of NF-kB seems to depend on this kinase activity (MALININ et al. 1997). With respect to IL-1-induced NF-KB activation, the proteins TNF-receptor-associated factor 6 (TRAF6), IL-1-receptorassociated kinase and myeloid differentiation marker 88 (MyD88) appear to connect the signalling pathway between IL-1 receptor and NF- $\kappa$ B activation with only minor gaps (Fig. 1) (WESCHE et al. 1997).

Recently, the IL-1-receptor-related human toll-like receptor 2 has been shown to be involved in LPS-induced signalling (KIRSCHNING et al. 1998; YANG et al. 1998). Using *trans*-dominant negative mutants of IKK $\alpha$ /Chuk, IKK $\beta$ , NIK, TRAF6 and MyD88, a critical role for these molecules in the LPS-induced activation of NF- $\kappa$ B has been demonstrated (Fig. 1) (KIRSCHNING et al. 1998).

#### 2.1 Plasmid DNA and CpG Oligodeoxynucleotides Activate NF-KB

Stacey and colleagues were the first to show that plasmid DNA (as a source of bacterial DNA) and synthetic oligodeoxynucleotides (ODNs) that contain a palindromic CpG-motif are inducers of NF-kB and NF-kB-dependent gene expression in macrophages (STACEY et al. 1996). They directly demonstrated enhanced NFκB-mediated DNA-binding activity of plasmid-treated RAW264 macrophages. Using a human immunodeficiency virus long terminal repeat (HIV-LTR)-driven luciferase reporter, which had been shown to be strongly dependent on NF-kB activity, they showed that plasmid DNA, but not CG-methylated plasmid DNA, induced transcriptional activity of this promoter. Furthermore, ODNs containing a palindromic AACGTT sequence, but not ODNs with the CpG palindrome ACCGGT, were able to induce HIV-LTR-dependent gene transcription. While these data show the importance of nucleotides surrounding the central CG, inversion of CG to GC also abolishes the NF-kB-inducing activity of ODN in RAW264 cells, directly proving the CpG dependency of CpG ODN (our own unpublished observation). Additionally, activation of other genes known to be induced by LPS and dependent on NF- $\kappa$ B activity, such as TNF, IL-1 $\beta$  and the HIV-1 LTR (DROUET et al. 1991; HISCOTT et al. 1993; POMBO et al. 1994), were activated by plasmid DNA (STACEY et al. 1996). Notably, the pattern of activated genes strongly resembles the one provoked by LPS, although, according to STACEY et al. (1996), LPS contamination of the various DNA stimuli could be ruled out.

Later on, it was directly shown that CpG ODN, dependent on a CpG motif, induces degradation of I $\kappa$ B $\alpha$  and I $\kappa$ B $\beta$  in WEHI-231 B cells (Y1 and KRIEG 1998b), thereby inducing translocation of p50/c-Rel (Y1 and KRIEG 1998b), p50/p65 and p50 homodimers (SPARWASSER et al. 1997). Degradation of I $\kappa$ B $\alpha$  is preceded by phosphorylation of Ser32 and Ser36 (our unpublished observation). Concurrent with degradation of I $\kappa$ B $\alpha$ , rapid induction of reactive oxygen species (ROS) has been demonstrated to occur in J774 macrophages and WEHI-231 B cells (Y1 et al. 1998b). Treatment of cells with the antioxidants *N*-acetyl-L-cysteine (NAC) or pyrrolidine dithiocarbamate (PDTC) suppressed TNF (Y1 et al. 1998b) and IL-6 secretion (Y1 et al. 1996), respectively, implying an important role for the redox state of the responding cell during stimulation. It is important to note that almost all inducers of NF- $\kappa$ B tested so far can be blocked by antioxidants (BAEUERLE and



Fig. 1. Model of nuclear factor  $\kappa B$  (NF- $\kappa B$ ) activation pathways induced by interleukin 1 (IL-1) and lipopolysaccharides (LPS). Solid arrows indicate where direct interaction of proteins has been shown, broken arrows are used where direct interaction has not been demonstrated. Arguments supporting the involvement of myeloid differentiation marker 88 (MvD88), tumor-necrosis-factor receptor-associated factor 6 (*TRAF6*), NF- $\kappa$ B-inducing kinase (*NIK*) and I $\kappa$  kinase  $\alpha/\beta$  in the LPS-induced NF- $\kappa$ B activation are exclusively based on experiments using dominant negative forms of the respective proteins. While the IL-1-receptor complex consists of two defined subunits, the situation of the LPS-receptor is more complicated and only partially elucidated. The glycosyl-phosphatidyl-inositol-linked CD14 molecule lacks a signal transduction moiety of its own and seems to be used to scavenge LPS - opsonized by LBP (LPSbinding protein) - for the actual signalling receptor. Toll-like receptor 2 (toll-2) has been shown to bind LPS and transduce the signal towards NF-KB (KIRSCHNING et al. 1998; YANG et al. 1998). However, LPSresistant mouse strains C3H/HeJ and C57BL/10ScCr both have mutations in another gene, toll-like receptor 4 (toll-4) (POLTORAK et al. 1998). Although biochemically not demonstrated yet, involvement of toll-4 in LPS signalling seems to be likely. Composition of the LPS receptor by different proteins - as indicated by "?" - is only one of many possible ways these findings might be brought in line (ULEVITCH 1999). AcP, accessory protein; IL-1R, IL-1 receptor; IRAK, IL-1R-associated kinase; IKK:, IK kinase complex:. IKK: is a complex of about 700 kDa and consists of several proteins, including IKKa/conserved helix-loop-helix ubiquitous kinase and IKKB. Other molecules of the complex have not been molecularly defined. All molecules in the figure are symbols only, and their shapes are not based on any biochemical data

HENKEL 1994; MUROI et al. 1994; MULLER et al. 1997). This could mean that induction of ROS is an important and general way of NF- $\kappa$ B activation. However, the critical step during signalling, which in fact depends on ROS generation, could

not be unequivocally demonstrated for any of the inducers described. While there is no doubt about the principal importance of the redox states of cells, the question of the significance of its altered dynamics during cell stimulation awaits further investigation.

The demonstration of CpG-DNA-induced specific phosphorylation of I $\kappa$ B $\alpha$  at Ser32 and Ser36 strongly suggests that the IKK complex, recently found to be activated by TNF, IL-1 and probably LPS (REGNIER et al. 1997; DIDONATO et al. 1997; ZANDI et al. 1997; KIRSCHNING et al. 1998), is also activated by CpG-DNA, although this has not been shown formally.

It will be interesting to see at which step CpG-DNA signalling enters the known pathways of NF- $\kappa$ B activation and whether this pathway can be traced back to an upstream receptor. Until now, every cell type shown to be directly responsive to CpG-DNA also activated NF- $\kappa$ B during stimulation (STACEY et al. 1996; YI et al. 1998b and our own unpublished observations). Therefore, this pathway seems to be a general mechanism through which CpG-DNA transduces its signal.

# 3 Mitogen-Activated Protein Kinase Pathways

Mitogen-activated protein kinases (MAPKs) are a group of serine/threonine-specific kinases that are activated in response to a variety of extracellular stimuli (LEWIS et al. 1998). MAPKs are involved in the regulation of specific transcription factors and other regulatory proteins and are themselves activated by upstream kinases called MAPK kinases or MAPK/extracellular-signal-regulated kinase (ERK) kinases (MAPKKs/MEKs) and MAPKK kinases/MEK kinases (MAP-KKKs/MEKKs) (Fig. 2). In vertebrates, at least three MAPK pathways can be differentiated. These pathways are able to sense and integrate incoming signals from different sources, for example membrane receptors, and transduce them into specific gene expression. Although some crosstalk between the different kinase cascades seems to exist, generally they are organized as sequentially working modules. Partially characterized scaffold proteins forming complexes with kinases belonging to a given cascade seem to contribute to the high degree of specificity (SCHAEFFER et al. 1998; WHITMARSH et al. 1998). According to the last MAPK in the corresponding cascade, they are referred to as the extracellular-signal-regulated ERK/MAPK pathway, the c-Jun N-terminal kinase (JNK)/stress-activated protein kinase (SAPK) pathway and the p38 MAPK pathway (Fig. 2). Members of both latter groups of kinases are sometimes also commonly referred to as SAPKs.

The ERK/MAPK pathway is primarily activated by growth factors and phorbol ester. Mitogenicity of growth factors largely depends on activation of the ERK pathway. Constitutive activity of members of this pathway contributes to the oncogenic transformation of cells (ROUSSEL 1998).

The JNK and p38 MAPKs are groups of kinases that are activated by different extracellular stimuli, including pro-inflammatory cytokines, such as TNF and IL-1,

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**Fig. 2.** Organisation of the mammalian mitogen-activated protein (MAP)-kinase signal-transduction pathways. The extracellular-signal-regulated kinase (ERK) pathway and the c-Jun *N*-terminal kinase (JNK)- and p38-stress-kinase pathways are illustrated schematically. Growth factors typically lead to the activation of the ERK subgroup of MAP kinases, while inflammatory cytokines and stress signals primarily activate the stress kinases JNK and p38. Lipopolysaccharides are able to activate both ERK- and stress-kinase pathways. At present, at least nine different MAP or ERK kinase kinases (MEKKs), including MEKK1-4, MAP kinase kinase 5, tumor progression locus 2, transforming-growth-factor-β-activated kinase 1, apoptosis-signal-regulating kinase 1 and nuclear-factor-κB-inducing kinase, have been identified in mammals with different specificities towards MAP kinase kinases (MKKs). The MLKs (mixed-lineage kinases) are a structurally different group of enzymes. For its family members MLK-2, MLK-3 and dual leucine zipper-bearing kinase (DLK), activation of MKKs has been observed (Lewis et al. 1998). JIP-1, the JNK-interacting protein, has recently been identified as a protein interacting with multiple components of the JNK signalling pathways (WHITMARSH et al. 1998) and has been postulated to be a Ste5p-like scaffold protein, facilitating and specifying JNK signalling (WHITMARSH and DAVIS 1998). *MAPK*, MAP kinase; *MAPKK*, MAP kinase kinase; *MAPKKR*, MAP kinase kinase inase inase

as well as LPS, ultraviolet (UV) light and stress signals (Fig. 2) (HIBI et al. 1993; BIRD et al. 1994; SLUSS et al. 1994; WESTWICK et al. 1994; HAMBLETON et al. 1996).

#### 3.1 CpG-DNA Induces the Transcription Factor Activating Protein 1

The first evidence that MAPKs might be involved in CpG-DNA-induced signalling came from transient transfection experiments using luciferase reporter plasmids with a triplet of activating protein 1 (AP-1)-binding sites. Stimulation of transfected ANA-1 macrophages with both plasmid DNA and CpG ODN induced transcriptional activity of AP-1 in a strictly CpG-dependent manner (Hacker et al. 1998).

The transcription factor AP-1 is comprised of members of the Fos, Jun and ATF (activating factor) families, which form homodimers and heterodimers. Its activity can be regulated at different levels (KARIN et al. 1997). Primarily, MAPK pathways control both the abundance of the AP-1 protein complex and its transcriptional activity. With respect to participation in AP-1 complexes, c-Fos, c-Jun and ATF-2 are especially well characterized. c-Fos, which is mainly regulated at the transcriptional level, contains two enhancer elements in its promoter region that directly

confer responsiveness to MAPK pathways. These two elements are: a CRE (cyclic adenosine monophosphate-responsive element) site and a serum-responsive element (SRE). The CRE site seems to be occupied by both CREB (CRE-binding protein) and ATF proteins. The SRE element binds dimers of the serum-response factors (SRF), recruiting ternary complex factors (TCF) like Elk-1 and stress-activated protein 1. The C-termini of these TCFs contain clusters of phosphorylation sites that are phoshorylated and thereby activated by ERK/MAP kinases and JNK/p38 stress kinases (HIPSKIND et al. 1994; PRICE et al. 1995). c-Jun is regulated by different mechanisms. Transcriptional regulation to most stimuli seems to be controlled through a c-Jun TRE (12-O-tetradecanoyl phorbol 13-acetate-responsive element) proposed to bind c-Jun/ATF-2 heterodimers (VAN DAM et al. 1993). In addition, c-Jun activity is regulated by specific phosphorylation at two sites, Ser63 and Ser73, which enhance the transactivating potency of the AP-1 complex (SMEAL et al. 1992). This phosphorylation is achieved by JNKs (SANCHEZ et al. 1994). In contrast to c-Fos and c-Jun, ATF-2 expression is constitutive. Its activity is regulated by phosphorylation by p38 and JNK (GUPTA et al. 1995; RAINGEAUD et al. 1996). Taken together, AP-1 integrates signals from different MAPK pathways.

Gel-shift analysis in CpG-DNA-stimulated ANA-1 macrophages revealed basal AP-1-binding activity but only a slight increase during the first 4h of simulation (HACKER et al. 1998). This AP-1 complex contained c-Jun. In contrast, c-Jun phosphorylation at Ser73 and Ser63 rapidly increased within 10min and stayed constant over at least 4h. The overall amount of c-Jun did not increase significantly during that time period. As transcriptional activity of AP-1 was correlated with specific phosphorylation of c-Jun, we concluded that, during the first hours of stimulation, phosphorylation might be the principal way of AP-1 regulation by CpG-DNA. Interestingly, using WEHI-231 B cells, Yi and Krieg (YI and KRIEG 1998a) also demonstrated phosphorylation of c-Jun and an additional significant increase in AP-1-binding activity. It is possible that the different cell types explain these differences.

#### 3.2 CpG-DNA Activates Stress Kinases

The kinases responsible for phosphorylation of c-Jun at Ser63 and Ser73 are the JNKs. Accordingly, JNKs and their upstream kinase, JNKK1, were found to be activated by CpG-DNA (HACKER et al. 1998). Additionally, p38, a stress kinase originally cloned as LPS-responsive kinase (HAN et al. 1994), has been found to be activated by CpG-DNA (HACKER et al. 1998; YI and KRIEG 1998a). Whether JNKK1 is also the upstream kinase of p38 in the cell types investigated so far is unknown. In general, p38 is efficiently activated by MKK3 and MKK6 (LEWIS et al. 1998). However, MKK4 has also been found to activate p38 in vitro and in vivo (SANCHEZ et al. 1994; DERIJARD et al. 1995; LIN et al. 1995) and could be involved in p38 activation (GANIATSAS et al. 1998).

Activation of JNK and p38 sufficiently explains phosphorylation of ATF-2 at Thr69/71, the critical site for transcriptional transactivation (GUPTA et al. 1995;

LIVINGSTONE et al. 1995). Interestingly, a specific inhibitor of p38, SB203580, blocked the DNA-binding activity of AP-1 in CpG-DNA-stimulated WEHI-231 cells (YI and KRIEG 1998a). Such p38 inhibition by SB203580 could affect AP-1 induction at different stages. As detailed above, c-Fos expression is regulated by a SRE site binding TCFs, which are phosphorylated and activated by different MAPKs. c-Jun is also transcriptionally regulated, involving ATF-2 and c-Jun themselves. DNA binding of ATF-2 also depends on its phosphorylation status (ABDEL-HAFIZ et al. 1992). It will be interesting to see which proteins are contained within this AP-1 complex and which components are affected at which stage by SB203580.

## 3.3 Downstream Effects of Stress Kinase Activation

While a panel of inducers of stress kinases in immune cells has been defined, effector functions clearly attributable to stress kinases are rare. One report demonstrated that mRNA stability of IL-2 is enhanced by JNK activity (CHEN et al. 1998). JNKK1 (MKK4)-deficient thymocytes exhibit a defect in IL-2 production in response to CD3/CD28 and phorbol 12-myristate 13-acetate (PMA)/Ca<sup>2+</sup> ionophore stimulation (NISHINA et al. 1997). However, these results are difficult to interpret, as – depending on the differentiation status of the cells – JNKK1 deficiency provokes additional phenomena, such as impairment of MKK3/6 phosphorylation (GANIATSAS et al. 1998). This implies that secondary effects must be considered in the interpretation of immune cell activation in knock-out animals. There is ample evidence that JNKs are involved in the regulation of apoptosis, although their specific role seems highly dependent on the cell type (IP and DAVIS 1998) and has not been evaluated for CpG-DNA-induced signalling.

A role for p38 with respect to the release of cytokines, such as TNF, IL-6 and IFN-γ (PRICHETT et al. 1995; BEYAERT et al. 1996; RINCON et al. 1998), during inflammatory responses in vitro and in vivo has been mainly deduced from experiments with a class of cytokine biosynthesis inhibitors called cytokine-suppressive anti-inflammatory drugs. Originally, p38 was defined by one of these specific inhibitors, SB203580 (CUENDA et al. 1995; LEE et al. 1994). As in the case of LPS-stimulated cells, CpG-DNA-induced TNF and IL-6 release is strongly suppressed by SB203580 (HACKER et al. 1998; YI and KRIEG 1998a). Remarkably, CpG-DNA-induced IL-12 release from macrophage cell lines and primary bone-marrow-derived dendritic cells was also significantly reduced (HACKER et al. 1998). However, as for many other effector functions affected by SB203580, it will be important to define the concrete target(s) downstream of p38.

# 3.4 Activation of the ERK/MAPK Pathway by CpG-DNA

Various observations strongly suggested that the ERK/MAPK pathway would also be activated by CpG-DNA. First, CpG-DNA induces strong proliferation in B cells

(MESSINA et al. 1991; KRIEG et al. 1995), obviously by stimulation of cell-cycle progression through G1 (YI et al. 1998a). This resembles the case for known mitogens, such as growth factors, which depend on ERK and MEK activity for their capacity to induce cell-cycle entry through G1 (ROUSSEL 1998). Second, LPS, which induces a very similar set of effector functions, including B-cell proliferation, induces ERK activity in different cell types (SWEET and HUME 1996).

Surprisingly, using phospho-specific antibodies against ERK1/2, Yi and Krieg reported that in WEHI-231 B cells and J774 macrophages JNK and p38, but not ERKs, are activated by CpG-DNA. Moreover, they found ERK1 to be activated by LPS in J774 to an extent comparable to that achieved by phorbol ester (PMA) (YI and KRIEG 1998a). If this holds true in more sensitive assays, it represents the first known qualitative difference between LPS and CpG-DNA-induced signalling.

Using another macrophage cell line, RAW264.7, and primary peritoneal macrophages for CpG-DNA stimulation, we observed substantial ERK2 kinase activity in in vitro kinase assays, comparable to that caused by LPS (unpublished observations). It will be interesting to see whether there exist differences in ERK activation among distinct cell types and, even more interesting, whether activation of different MAPK pathways qualitatively separates CpG-DNA from LPS.

# 4 CpG ODN-Induced Signalling Requires Non-Specific Uptake and Endosomal Acidification

#### 4.1 Endosomal Uptake of CpG ODN is CpG-Independent

CpG ODN and fluorescein-isothiocyanate-labelled CpG ODN are taken up rapidly from cells and move into an endosomal compartment (Tonkinson and STEIN 1994; HACKER et al. 1998). Based on competition analyses, it has been argued that ODN uptake is receptor mediated. Several cell-surface receptors lacking sequence specificity, including CD11b/CD18 integrins and scavenger receptors, have been shown to engage single-stranded DNA (KIMURA et al. 1994; BENIMETSKAYA et al. 1997). Our own studies showed that uptake of CpG ODN could be competed for by unrelated non-CpG ODNs. Although some ODNs were more potent than others, every non-CpG ODN was essentially able to compete for CpG ODN uptake. ODNs with poly-G stretches were especially potent competitors (G. Lipford, personal communication). These ODNs originally had been found to bind to the scavenger receptor, thus promoting efficient cellular uptake (KIMURA et al. 1994). Poly-G stretches, in the context of stimulatory CpG sequences, enhance the immunogenicity of CpG ODN; increased uptake has been proposed as probable reason for this enhancement (KIMURA et al. 1994).

Importantly, both uptake and CpG-induced signalling can be competed for by non-CpG ODN. Every effector function and signalling event investigated so far can

be blocked by non-CpG ODN, implying that binding of CpG DNA to a receptor represents a first, obligate step during signal induction (HACKER et al. 1998). This raises two possibilities: (1) this receptor mediates uptake *and* signalling or (2) this receptor mediates uptake, but signalling starts from a later point. The latter possibility is supported by experiments with endosomal acidification inhibitors (see below).

The observation that non-CpG ODNs compete for CpG-ODN-induced signalling raises interesting questions. Every CpG ODN without intrinsic activity will be able to act as a competitor for active CpG ODNs. Additionally, highermolecular-weight DNAs, like bacterial DNA and plasmid DNA, contain both activating sequences and non-activating sequences and, therefore, one attractive model would be to consider DNA as a composition of stimulatory and nonstimulatory (and hence inhibitory) sequences. According to this model, vertebrate DNA is able to inhibit cytokine release of plasmid-triggered spleen cells (WLOCH et al. 1998).

Recently, another possible explanation for the inhibitory effects of certain adenoviral DNAs has been proposed: the presence of specialized sequences surrounding central CGs (KRIEG et al. 1998). However, more work will be required to determine whether such sequences act as effective competitors or contain signalling qualities of their own in an inhibitory manner.

# 4.2 CpG-DNA-Induced Signalling can be Blocked by Inhibitors of Endosomal Maturation/Acidification

Endosomal maturation may be defined as pH-dependent evolution of early endosomes to lysosomal compartments (MELLMAN et al. 1986). Compounds like bafilomycin A and chloroquine block this evolution, the former by antagonising intravesicular hydrogen pumps (YOSHIMORI et al. 1991), the latter by moving into acidified vesicles and acting as a neutralising base buffer (OHKUMA and POOLE 1981). Neither compound significantly altered cellular uptake but modulated endosomal accumulation of ODN, possibly by altering the trafficking of lysosomal enzymes and receptors (CHAPMAN and MUNRO 1994) or by affecting efflux pathways. Endosomal vesicle compartments with varying ODN efflux rates have been described (TONKINSON and STEIN 1994).

Using a variety of 'read out' systems, such as cytokine production, cytokine promoter activity, AP-1 and NF- $\kappa$ B activation and induction of p38, it has been consistently observed that all these CpG-DNA-driven responses were sensitive to endosomal acidification inhibitors (HACKER et al. 1998; MACFARLANE and MANZEL 1998; YI and KRIEG 1998a; YI et al. 1998b). Notably, LPS-induced signalling was not affected by concentrations of chloroquine and bafilomycin A that efficiently blocked CpG-DNA-induced signalling (HACKER et al. 1998; YI and KRIEG 1998a).

As described above, CpG ODNs are taken up in endosomes in a CpG-independent but receptor-mediated way. The fact that these acidification inhibitors allow uptake of CpG ODN but are still able to block signalling strongly suggests that uptake of CpG ODN and signal induction are separable events. As such, these findings imply that both CpG-independent cellular uptake and endosomal maturation/acidification are required for signalling to be initiated.

The shift to low pH could be important for different reasons. Such pH changes could either trigger dissociation of CpG-DNA from a non-specific (cell-surface) receptor or enable binding of CpG-DNA to a specific (endosomal) receptor. Alternatively, a less well-defined molecular mechanism dependent on endosomal maturation may be required for signal transduction.

#### 5 The NFkB and Stress-Kinase Pathways are Activated in Parallel

As detailed above, separable signalling pathways are switched on in response to CpG-DNA. In this respect, it is important to know whether these pathways are turned on in parallel or sequentially. This is even more important in cells that after stimulation – secrete a variety of both defined and undefined cytokines able to act in an autocrine or paracrine fashion. We therefore compared the kinetics of the  $NF\kappa B$  and stress-kinase activation pathways, measuring the earliest defined events, i.e. IkBa degradation and JNKK1 phosphorylation, after stimulation by CpG ODN. Figure 3 shows that, using RAW264.7 cells, both pathways seem to be switched on in a parallel and not sequential way. This suggests that both pathways posses a common starting point upstream. These data argue against the possibility that activation of one pathway is the prerequisite for activation of the other. Another interesting point is the overall velocity of signal induction. CpG-ODNinduced signalling starts after about 5min and is easily detectable between 5min and 10min after stimulation (Fig. 3). LPS seems to be slightly faster (our personal observation), and signalling induced by classic transmembrane receptors, such as the IL-1R and TNF receptor p55, is significantly faster. The delay of CpG-ODNinduced signalling, in comparison to that induced by IL-1 and TNF, could be explained by the requirement for cellular uptake and endosomal maturation.



Fig. 3. The nuclear factor  $\kappa B$  activation pathway and the stress-kinase pathway are induced with comparable kinetics. Shown is a Western blot analysis of RAW264.7 macrophages stimulated with CpG oligodeoxynucleotides for different time periods and analysed by antibodies to I $\kappa$  kinase B $\alpha$ ( $\kappa B\alpha$ ) and the Thr223-phosphorylated form of c-Jun N-terminal kinase kinase I as indicated

# 6 Summary and Conclusion

While more and more attention has been paid to CpG-DNA with respect to its usefulness as an adjuvant, its molecular mechanism of action is less well defined. Over the last few years, at least two major signalling pathways have been shown to be utilized by CpG-DNA: the NF-KB activation pathway and the stress-kinase pathway. Direct downstream events of these pathways are induction of transcriptional activity of NF-kB and transcriptional activity of AP-1. As far as investigated, CpG-DNA uses signal transduction pathways originally described for other stimuli, such as LPS, IL-1 or TNF. Therefore, to us, the prime question is: where does CpG-DNA-induced signalling enter these known pathways? This raises questions about the existence of a CpG-DNA-sequence-specific receptor. Several points of evidence support the probability of the existence of a cellular receptor: There is a strong motif (unmethylated CpG) dependency for CpG-DNA-induced signalling. There is cell-type specificity. Dendritic cells, macrophages and B cells respond to CpG-DNA, but other cell types, such as fibroblasts and T cells, do not. In addition, classic signal-transduction pathways are rapidly switched on in a parallel manner, as is known for other receptors.

Using competing non-CpG ODNs and inhibitors of endosomal acidification, some evidence has been obtained that CpG ODNs are taken up into endosomes by a CpG-independent receptor, followed by a pH-dependent step before signalling starts. A model based on these findings is proposed in Fig. 4.

Nevertheless, other receptor-independent activities of CpG-DNA cannot yet be ruled out. Although unlikely, we should consider the possibility that CpG-DNA directly interacts with cellular nucleic acids either by direct hybridization with complementary nucleotides or by formation of DNA triplexes (VASQUEZ and WILSON 1998). While these possibilities have been explored by antisense technology, using a huge variety of ODNs, there is no experimental evidence that such interactions are important for the activity of CpG-DNA.

In this context, it is important to note that DNA, especially phosphothioatestabilized ODNs with poly-G stretches, have substantial CpG-independent activities, although these activities seem not to depend on specific, antisense-like DNA– DNA interactions (PISETSKY 1996). One good example comes from experiments using ODNs on primary T cells. Co-stimulation of CD3-primed T cells with CpG ODN leads to a significant increase of IL-2 secretion and proliferation; however, these effects are CpG independent (K. Heeg, personal communication). Remarkably, these poly-G stretches seem to be inactive when transferred to doublestranded DNAs, such as plasmid DNA (WLOCH et al. 1998). In contrast, to my knowledge, no immune-stimulatory effect of bacterial DNA has been described that can not be abolished by CpG-specific methylation.

Taken together, CpG-dependent and CpG-independent activities must be distinguished from one another. Among these effects, CpG-dependent signalling is better defined.



**Fig. 4.** Model of CpG-DNA-induced signal-transduction pathways. CpG oligodeoxynucleotides are taken up into endosomes by CpG-independent receptors. A pH-dependent step precedes signalling. The existence of a specific receptor in the endosomal compartment is speculative. All molecules in the figure are symbols only, and their shapes are not based on any biochemical data. *IKK*:, Ik kinase complex: IKK: is a complex of about 700 kDa consisting of several proteins, including IKK $\alpha$ /conserved helix-loophelix ubiquitous kinase (Chuk) and Ik kinase  $\beta$ . Other molecules of the complex have not been molecularly defined yet. As detailed in the text, transcription factors other than activating protein 1, as well as proteins not directly involved in transcriptional regulation, could also be activated by c-Jun *N*-terminal kinase/p38

Much effort is going into the investigation of the pharmacological applications of CpG-DNA. Once CpG-receptor-like structures are known, the question of the physiological role of CpG-DNA can be tackled.

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# CpG DNA Co-Stimulates Antigen-Reactive T Cells

K. Heeg

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# **1** Introduction

During the last several years, compelling evidence has accumulated that bacterial DNA is sensed as "infectious danger" by innate immune cells, such as macrophages and dendritic cells (DCs) (KRIEG 1996b; PISETSKY 1996a,b, 1997; LIPFORD et al. 1998). Those cells are thought not only to represent central cellular restriction points during induction and execution of innate immune responses but also to control activation of cells of the acquired immune system, such as T lymphocytes (MATZINGER 1994; FEARON and LOCKSLEY 1996; MEDZHITOV and JANEWAY 1997). Accordingly, processing and presentation of antigen (signal one) and the generation of permissive cytokine and co-stimulatory milieus (signal two) govern productive T-cell activation (BRETSCHER and COHN 1970; LAFFERTY and WOOLNOUGH 1977;

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MUELLER et al. 1989). Evidently, sensing "infectious danger" and, thus, activation of innate immune cells represents a major prerequisite for successful adaptive immune responses of the B- and T-cell lineages.

The observation that synthetic, single-stranded oligodeoxynucletides (ODNs) mimic the effects of bacterial DNA in a DNA-sequence-specific manner (YAMAMOTO et al. 1992; KRIEG et al. 1995) has led researchers to envisage its practical use (LIPFORD et al. 1997a; KLINMAN 1998; KRIEG et al. 1998), First, immunostimulatory properties of bacterial DNA and ODNs could be attributed to unmethylated CpG-dinucleotide motifs, which occur at expected frequencies in bacterial DNA but at low frequencies in mammalian DNA (BIRD 1980; KRIEG 1996a; PISETSKY 1996a; PESOLE et al. 1997). Second, ODNs displaying the classic canonical DNA motif 5'-purine-purine-CpG-pyrimidine-pyrimidine-3' were shown to be often immunostimulatory (CpG ODN) (KRIEG et al. 1995, 1996; KRIEG 1996a). Finally, modification of the ODNs' backbones by phosphothioate linkages made the ODNs resistant to the action of nucleases (AGRAWAL et al. 1991; GAO et al. 1992) and, thus, allowed their application in animal models in vivo. The large body of evidence obtained so far suggests that CpG ODNs might be suitable for adjuvant use in prophylactic and therapeutic vaccination formulas for both B-cell and T-cell vaccinations. The following review summarizes our current knowledge of indirect (extrinsic) and direct (intrinsic) effects of CpG and non-CpG ODNs on the immune reactivity of T lymphocytes.

## 2 CpG ODNs Exert Extrinsic Effects on T Cells

#### 2.1 CpG-Mediated Activation of APCs Augments T-Cell Responses in Vitro and in Vivo

Besides the mitogenic effects of CpG ODNs on B lymphocytes (KRIEG et al. 1995), it was soon recognized that bacterial plasmid DNA and certain palindromic, synthetic ODNs (YAMAMOTO et al. 1992) are taken up by macrophages, which in turn are activated, a process associated with the translocation of nuclear factor  $\kappa B$ (NF $\kappa B$ ) (STACEY et al. 1996; SPARWASSER et al. 1997). Further studies revealed that macrophages are directly activated by CpG ODNs to express co-stimulatory molecules (CD80, CD86, CD40) and produce pro-inflammatory [tumor necrosis factor (TNF), interleukin 6 (IL-6)] and effector cytokines (IL-12) (SPARWASSER et al. 1997; JAKOB et al. 1998; SPARWASSER et al. 1998). Modifications of the ODN sequence defined CpG ODNs that preferentially induced IL-12 (but not TNF) in vivo and in vitro (LIPFORD et al. 1997b). Of note, such ODNs would be of value in devising ODNs for therapeutic use, since they lack potentially harmful effects mediated by the release of toxic amounts of TNF.

DCs disseminated in non-lymphatic tissue are immature yet phagocytically active (Cella et al. 1997; VREMEC and SHORTMAN 1997; STEINMAN et al. 1998).

Upon maturation of DCs, phagocytic activity and antigen uptake is lost and co-stimulatory activity is gained. During this step, DCs translocate in lymphatic organs, express high levels of major histocompatibility complex (MHC) and co-stimulatory molecules (LUTZ et al. 1996; CELLA et al. 1997) and, thus, are equipped for the activation of T lymphocytes (SCHULER et al. 1997). In vitro, CpG ODNs were shown to induce both phenotypic maturation of bone-marrow-derived DCs (Sparwasser et al. 1998) and secretion of high levels of IL-12 (JAKOB et al. 1998; SPARWASSER et al. 1998). Thus, CpG ODNs are likely to induce a cellular and cytokine milieu that provides signals necessary for T-cell activation. These effects are primarily due to activation of antigen-presenting cells (APCs) by CpG ODNs (indirect or extrinsic effects). Examples of the extrinsic effects of CpG ODNs on Tcell activation in vitro are depicted in Fig. 1. When the number of APCs is limited in an allogeneic mixed-lymphocyte reaction, no T-cell proliferation can be induced. However, addition of CpG ODNs amplifies the stimulatory potency of the remaining APCs. As a consequence, T-cell responses to antigens are enhanced. In vivo, this effect might be responsible for the powerful activity of CpG ODNs as adjuvants in T-cell responses. Injection of "weak" proteinaceous antigens, together with CpG ODNs, allowed the generation of MHC-restricted peptide-specific CD8<sup>+</sup> T lymphocytes (LIPFORD et al. 1997a; EHL et al. 1998).

#### 2.2 CpG-Induced Cytokines Influence T-Cell Responses in Vivo

Unlike B cells, direct activation of T cells by bacterial CpG DNA or CpG ODNs has not been observed. Recently, however, it was found that lipopolysaccharides



**Fig. 1.** Extrinsic effect of CpG oligodeoxynucleotides on mixed lymphocyte reactions in vitro. T cells from C57Bl/6 mice (20,000/well) were stimulated with 1000 irradiated BALB/c spleen cells in microtiter plates as indicated. After 4 days, the proliferative response was recorded

(LPS), a known mitogen for murine B cells, also activated T cells in vivo and in vitro (MATTERN et al. 1994; TOUGH et al. 1997; KHORUTS et al. 1998) in the apparent absence of T-cell-receptor (TCR) ligation. Several lines of evidence indicated that responsible cytokines were type-I interferons (IFN), known to be induced by LPS or poly(I:C) (ZHANG et al. 1998). Moreover, these cytokines were found to induce the secretion of IL-15 in APC, which in turn caused bystander activation of memory (but not of naive T cells) in a TCR-independent fashion (ZHANG et al. 1998). Since CpG ODNs also induce type-I IFNs, these TCRindependent extrinsic effects of CpG ODNs should also be operative in vivo. Indeed, within 20 h after subcutaneous injection of CpG ODNs, upregulation of CD69 (but not of IL-2R $\alpha$ ) on T cells of the draining lymph nodes can be observed (SUN et al. 1998b, S. Bendigs, personal communication). It follows that CpG-ODN-induced type-I IFNs may influence the signal threshold of antigenreactive T cells in vivo. Again, this effect is due to the CpG-mediated activation of APCs.

Activation of T lymphocytes and subsequent differentiation into effector cells is guided by a distinctive milieu of cytokines. During T-helper cell development, IL-4 causes a bias for T-helper 2 (Th2) differentiation, while IL-12 instructs Th1 responses (MACATONIA et al. 1995; MOSMANN and SAD 1996). Immunizations performed with bacterial DNA, *Drosophila* DNA (SUN et al. 1998a) or CpG ODNs (CHU et al. 1997; LIPFORD et al. 1997a; ROMAN et al. 1997) revealed augmentation and a strong bias of the antibody response towards an immunoglobulin G2a (IgG2a) isotype characteristic of Th1 responses. These observations may be attributed to the strong induction of APC-derived IL-12 by CpG ODNs. The induction of antigen-specific CD8<sup>+</sup>-cytolytic T-cell responses also suggests the preferential induction of Th1-dominated responses by CpG ODNs (LIPFORD et al. 1997a).

The same conclusions are based on experiments using murine models for infectious diseases. For example, infection of BALB/c mice with *Leishmania major* is followed by disease, because a strong Th2 response is triggered (LOCKSLEY and LOUIS 1992). In contrast, C57B1/6 mice mount a Th1-biased immune response and control the infection via nitric oxide (NO) generation (STENGER et al. 1994). However, treatment of infected BALB/c mice with CpG ODNs resulted in a conversion of the disease, i.e., BALB/c mice mounted a strong Th1 response accompanied with a sustained expression of IL-12R<sup>β</sup>2 molecules and a T-helper cytokine pattern characteristic of a Th1 phenotype (ZIMMERMANN et al. 1998). CpG ODNs were not only effective in preventing lethal infections (when injected together with Leishmania) but were also able to cure an established Th2-dominated lethal disease when the CpG ODNs were administered up to 3 weeks after infection (ZIMMERMANN et al. 1998). Thus, CpG ODNs probably direct ongoing immune responses towards a Th1 phenotype due to the ODNs' ability to activate APCs and induce Th1-instructing cytokines. Similar conclusions can be drawn from models of murine experimental allergies (KLINE et al. 1998). The adjuvant activity and Th1promoting efficacy of CpG ODNs can be explained, at least in part, by their potent extrinsic activities on T cells.

# 3 Intrinsic Effects of CpG ODNs on T Cells

# 3.1 CpG-ODN-Mediated Co-Stimulation of T Cells in the Absence of APCs

To determine whether CpG ODNs can exert direct (intrinsic) effects on T cells, we prepared highly purified T lymphocytes (BENDIGS et al. 1999). To achieve this, T lymphocytes were enriched via several rounds of negative selection by magnetic cell separation and then stimulated with plate-bound anti-CD3 monoclonal antibodies (mAbs). A typical experiment is shown in Fig. 2. The fact that T cells could be stimulated neither by anti-CD3 alone nor in the presence of anti-CD3 and LPS or concanavalin A (used as a control) indicated that our cultures were indeed devoid of APCs. Since addition of IL-2 caused proliferation in anti-CD3 mAbstimulated cultures but not in the absence of anti-CD3, mAb signal 1 (TCR ligation) caused IL-2R expression and, thus, was operative. Of note, the requirement for IL-2 could be substituted by CpG ODNs (Fig. 2). This type of result allowed several conclusions. First, CpG ODNs did not directly activate T cells. In contrast to B cells and macrophages, purified T lymphocytes remained unresponsive to ODNs in the absence of TCR ligation. However, upon crosslinking of T-cell receptors (signal 1), addition of CpG ODNs induced proliferation of purified T cells in a dose-dependent manner (Fig. 2). In fact, upon TCR ligation, CpG ODNs induced expression of IL-2R, production of IL-2 and cytolytic activity in T cells (BENDIGS et al. 1999). T-cell proliferation was dependent on IL-2, since anti-IL-2 antibodies completely abrogated proliferative and cytolytic responses. CpG-medi-



**Fig. 2.** CpG oligodeoxynucleotides co-stimulate T cells. Purified T cells (20,000/well) from C57Bl/6 mice were stimulated on anti-CD3-antibody-coated microtiter plates in the presence of the reagents indicated. After 4 days, the proliferative response was recorded

ated co-stimulation required a functional TCR signal (signal 1), since cyclosporin A (CsA), known to block Ca<sup>++</sup>-dependent TCR signaling pathways (SIGAL and DUMONT 1992; McCAFFREY et al. 1993), completely abolished T-cell proliferation co-stimulated by CpG ODNs (BENDIGS et al. 1999).

Collectively, these data show that CpG ODNs co-stimulated T cells to express IL-2R, produce IL-2, proliferate and finally develop into cytotoxic effector cells. In this respect, co-stimulation mediated by CpG ODNs phenotypically resembles co-stimulation of T cells mediated by CD28 (HARDING et al. 1992; SAGERSTRÖM et al. 1993). This notion was further supported by the observation that CpG ODNs substituted, at least in part, for the co-stimulatory defects of T lymphocytes from CD28<sup>-/-</sup> mice in vitro (BENDIGS et al. 1999). CpG ODNs also provided signal 2 to T lymphocytes from TCR-transgenic mice stimulated with antigenic peptides. To exclude possible extrinsic CpG-ODN effects, in these experiments APCs were fixed with glutaraldehyde prior to loading with peptides. If peptide-loaded fixed APCs were added, CpG ODN induced proliferation of T lymphocytes (BENDIGS et al. 1999). These results demonstrated that co-stimulation by CpG ODNs not only applies for experimental TCR stimulation with mAbs but also holds true for physiological TCR ligation induced by MHC plus peptide.

The restricted action of ODNs on T cells may also be prototypic for other cell types. Although it was initially reported that natural killer (NK) cells were responsive towards CpG ODNs (BALLAS et al. 1996), it was later found that IFN-y release from NK cells was dependent on exogenous IL-12 (HALPERN et al. 1996; CHACE et al. 1997). Operationally, IL-12 could serve as signal 1 in NK cells to trigger sensitivity to CpG ODNs (signal 2). Similar principles might operate in B cells. B lymphocytes respond to ODNs in cultures of splenocytes with a short wave of cell divisions; nevertheless, proliferation, antibody production and cytokine secretion is several-fold enhanced after crosslinking of B-cell receptors (surfacebound Igs) (KRIEG et al. 1995). Thus, full sensitivity of B cells to CpG ODNs might develop only after receptor ligation (signal 1), as is the case for T cells. Overall, these results led us to conclude that APC-like macrophages or DCs are a priori sensitive to CpG ODNs, while cells of the lymphocytic lineage must develop sensitivity to CpG ODNs mediated by receptor ligation (signal 1). We anticipate that this functional divergence between APCs and lymphocytes might find its counterpart in different CpG-ODN receptors and/or distinct ODN-induced signal pathways.

### **3.2 ODN-Mediated Co-Stimulation Violates the Classical Rules of CpG-Mediated Stimulation of B Cells and APCs**

#### 3.2.1 The CpG Motif is Dispensable for Co-Stimulation of T Cells

When we analyzed the ODN-DNA-sequence requirements for T-cell co-stimulation, we were surprised by the finding that the rules of canonical CpG-DNA motifs, as defined for direct activation of B cells and APCs, did not apply for its co-stimulatory activity on T cells. First, in CpG motifs displaying ODNs like 1628, the central CpG could be methylated or replaced by a TpG without losing its co-stimulatory activity on T cells (Table 1). However, several ODNs (ODN 1720 or ODN ATT-Rep) were negative, indicating sequence dependency of the phosphothioate-modified ODNs towards T cells (BENDIGS et al. 1999). When we screened human DNA gene-sequence data for canonical CpG motifs, we also found CpG sequences in the human vertebrate genome (LIPFORD et al. 1997b, 1998). Some of these sequences were active to APCs [ODN IL-12p40, a CpG motif within the IL-12p40 gene (LIPFORD et al. 1997b)], yet some were inactive (ODN SP1, a CpG motif within the SP1-promoter site). However, ODN SP1 was active on T cells, i.e., it co-stimulated T cells (Table 1) (LIPFORD et al. 1999).

Designation	Sequence (5'–3')	APC			T cells	
		TNF	IL-12	Inhibition of CpG effects	Co-stimulation	Adjuvant for in vivo CTL induction
1668	tccaGACGTTcct	+ + +	+ + +		+	+ + +
IL-12p40	agetatGACGTT ccaagg	+	+ +		+	+ + +
1628	GGGGtcAACGTT	Ø	Ø	n.t.	+ + +	+
1628M	GGGGtcAAMGTT	Ø	Ø	n.t.	+ + +	n.t.
1628GC	GGGGtcAAGCTT	Ø	Ø	n.t.	+ + +	n.t.
SP1	tCGatCGGGGCG GGGCGaGC	Ø	Ø	n.t.	+ + +	+
SPIT	tTGatTGGGGTG GGGTGaGC	Ø	Ø	n.t.	+ + +	n.t.
SPIM	tMGatMGGGGM GGGGMGaGC	Ø	Ø	n.t.	+ + +	n.t.
1720	tccatgagetteetgatget	Ø	Ø		Ø	Ø
ATT-Rep	attattattattattattat	Ø	Ø		Ø	Ø
PZ31	ctcctattGGGGG tttcctat	Ø	Ø	+ + +	+ + +	+
PZ33	ctcctattGGGGT tttcctat	Ø	Ø	+ + +	+ + +	+
PZ34	ctcctattGGTGG tttcctat	Ø	Ø	Ø	Ø	Ø
SB	tccatGACGTT cctgaatggctGGGG	Ø	Ø	+ +	+ + +	n.t.
SBDG	tccatGACGTT cctgaatggctGAGT	+ + +	+ + +		+	n.t.
SBGC	tccatGAGCTT cctgaatggctGGGGG	Ø	Ø	+ +	+ + +	n.t.

Table 1. Effects of oligodeoxynucleotides (ODN) on antigen-presenting cells (APCs) and T cells

CTL, cytotoxic T lymphocyte; M, methylated; n.t., not tested.

#### 3.2.2 G-Rich ODNs Co-Stimulate T Cells Yet Fail to Activate APCs

We soon realized that ODNs active on T cells either possessed a classical CpG motif, like ODN 1668, or displayed stretches of guanines, like ODN 1628 or ODN SP1 (G-rich ODN) (Table 1). In order to define a putative motif, we synthesized ODNs where a central G stretch was permutated with thymidine residues (PZ3, PZ31, PZ34; Table 1). This approach led to the definition of new ODN motifs active on T lymphocytes. One motif (four Gs or 5'-G-X-G-G-G-3', where X is any base) characterizes ODNs that co-stimulate anti-CD3-triggered T cells. In retrospect, the active ODNs 1628 and SP1 also contain this G-rich motif (compare Table 1). Of note, ODNs embodying this G-rich motif failed entirely to activate APCs and did not trigger production of cytokines, such as IL-12 or TNF (LIPFORD et al. 1999). Further analyses revealed that G-rich ODNs actually antagonized the action of CpG ODNs on APCs (Häcker et al. 1998). Thus, G-rich ODNs effectively suppressed the activity of CpG ODNs on APCs. Detailed analyses with labeled ODNs revealed that G-rich ODNs are taken up into APCs with a much higher affinity and efficacy than were CpG ODNs. However, they competed for the uptake of CpG ODNs. Since G-rich ODNs fail to induce signaling cascades within APCs (Häcker et al. 1998) competition with CpG ODNs during uptake into APCs might explain the inhibitory effects of G-rich ODNs on CpG ODNs.

# 3.2.3 G-Rich Motifs in Hybrid ODNs are Dominant Negative on APCs Yet Co-Stimulate T Cells

When we synthesized hybrid ODNs consisting of a CpG motif (derived from ODN 1668) plus a G-rich motif at the 3' end of the ODN, we found that these ODNs (ODN SB in Table 1) failed to activate APCs. Disrupting the G stretch (ODN SB-DG) fully restored reactivity on APCs. Thus, G motifs have dominant negative effects as hybrid ODNs when tested on APCs. The very same ODNs, however, effectively co-stimulated T lymphocytes. Thus, ODNs containing the G motif are agonists on T cells yet antagonists on APCs. In retrospect, the peculiar activities of ODN 1628 and ODN SP1 may now be explained. Although ODN 1628 and ODN SP1 contain a CpG motif, they exhibit only marginal activity on APCs. In contrast, these ODNs strongly co-stimulate T cells (LIPFORD et al. 1999; Table 1). We suggest that their G motifs antagonize the effects on APCs yet trigger T-cell co-stimulation.

# 3.2.4 ODN-Induced Signal Pathways on T Cells are Distinct from Those on APCs

ODNs differ in their affinity kinetics for uptake into APCs (HÄCKER et al. 1998; G. B. Lipford, personal communication). When we tested uptake of ODNs into T cells, we noticed that only a minute number of ODNs, if any, are taken up by naive resting T cells (unpublished data). In contrast, if T cells were stimulated with anti-CD3 antibodies, uptake of ODNs was enhanced. However, the efficacy of uptake was lower than for the uptake of ODNs in APCs (unpublished data). Some investigators have suggested that APCs take up ODNs via surface receptors, such as Mac-1 or the scavenger receptor (BENIMETSKAYA et al. 1997). Since T cells express neither scavenger receptors nor Mac-1, different uptake mechanisms/ receptors might be operative in partially activated T cells.

Recently, aspects of the CpG-ODN-induced signaling cascade within APCs have been unraveled (HÄCKER et al. 1998). ODNs are taken up into early endosomes and, within minutes, trigger the stress-activated protein kinase/c-Jun *N*-terminal kinase stress-kinase pathway followed by transcriptional activity of activator protein 1 and NF $\kappa$ B (HÄCKER et al. 1998). A critical restriction point during this signaling process represents the maturation of endosomes. Agents that block maturation of endosomes (chloroquine or bafilomycin) prevent ODN-mediated signaling and, thus, block all downstream effects on APCs (HÄCKER et al. 1998; MACFARLANE and MANZEL 1998). When we tested the outcome of endosomal blockade on T-cell co-stimulation, we found that neither chloroquine nor bafilomycin inhibited co-stimulation of T cells mediated by either CpG ODNs or G-rich ODNs (LIPFORD et al. 1999).

In conclusion, the mode of action of ODNs on T cells differs in several respects from their action on APCs. First, ODNs exert no direct stimulation on resting T cells. In order to develop sensitivity to ODNs, T cells require partial activation via TCR occupancy (signal 1). Second, the ODN-sequence requirements for T cells differ from those for ODNs active on APCs. Methylated and un-methylated CpG ODNs and G-rich ODNs co-stimulate T cells effectively. Third, the signal pathways operative in T cells appear to be distinct. While cellular uptake of ODNs is only minute in T cells, as compared with APCs, endosomal maturation is dispensable for co-stimulation of T cells and a condition for APC activation. It will, therefore, be rewarding to define in detail the receptors/signal cascades in T lymphocytes.

### 4 Effect of ODN on T-Cell Responses in Vivo

Proteinaceous antigens usually fail to elicit strong cytolytic T-cell responses in vivo unless CpG ODNs are used as adjuvants (LIPFORD et al. 1997a). As discussed above, the potent adjuvanticity of CpG ODNs may be attributed to their capacity to activate APCs (an extrinsic effect on T cells). However, a direct (intrinsic) effect of CpG ODNs and G-rich ODNs on T cells might contribute to their adjuvant activities on T cells in vivo. Since G-rich ODNs fail to activate APCs, these ODNs should unravel the relative importance of intrinsic ODN effects on T-cell activation in vivo. We therefore tested various CpG, G-rich and hybrid ODNs for their adjuvanticity during induction of CD8<sup>+</sup> cytotoxic T lymphocytes in vivo. We found that all three classes served as adjuvants in vivo (LIPFORD et al. 1999). However, the relative efficacy was highest with CpG ODNs. These results suggest that G-rich ODNs can probably be used as T-cell adjuvants, since co-stimulation of T cells (in the absence of activated APCs) is sufficient to aid cellular immune responses in vivo. However, CpG ODNs activate (extrinsic effect) and co-stimulate (intrinsic effect) DCs. Therefore, CpG ODNs display the highest adjuvanticity.

An additonal discriminatory aspect of the action of CpG and G-rich ODNs emerged from these studies. In addition to their T-cell adjuvanticity, G-rich ODNs induced strong NK-like cytotoxic activity in vivo (unpublished data). Similar conclusions have previously been reported in the case of human NK cells (BALLAS et al. 1996). Active ODNs showed at least one stretch of G either at the 5' or 3' end of the ODN. Thus, G-rich ODNs also provoke effector functions of the innate immune system. It is obvious that this property of G-rich ODNs might be of decisive importance during infection with NK-sensitive organisms. Preliminary studies in a murine infection model with *Toxoplasma gondii* indicate that this might be indeed the case (S. Zimmermann, personal communication).

The ability of CpG and G-rich ODNs to co-stimulate T cells might be of practical relevance in situations where APCs are not available, e.g. in tumor tissues. According to the two-signal model of T-cell activation and anergy induction (MUELLER et al. 1989), tumor cells expressing antigen (signal 1) alone without co-stimulatory activity would be tolerogenic. Since CpG ODNs and G-rich ODNs co-stimulate T cells, these ODNs should bypass the tolerogenic effect and, thus, may activate anti-tumor T-cell responses. However, this beneficial outcome of T-cell co-stimulation by ODNs in vivo could include serious adverse consequences. ODNs might permit an APC-independent (re-)activation of autoreactive T cells.

# 5 Outlook

Analyses of the actions of ODNs on T cells enriched our knowledge in several respects. First, it was recognized that T cells respond to ODNs in an unperceived, restricted fashion. TCR ligation was necessary to render T cells permissive for the effects of ODNs. This mode of action could be prototypic for other cells of the lymphocytic lineage. Second, a new class of ODNs reactive on T cells and NK cells could be defined. The G-rich ODNs differ in many aspects from the classical CpG ODNs, yet they could be suitable for therapeutic use. In addition, the strong capacity of these ODNs to induce NK-like cytotoxicity could be advantageous in new therapeutic vaccines against NK-sensitive parasites. Third, beginning analyses of the signal cascade of ODNs on T-cell activation and immunization in vitro and in vivo could aid in the development of a new, safe and efficient adjuvant for therapeutic and prophylactic vaccines against infectious agents and cancer.

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# **Role of Type I Interferons in T Cell Activation Induced by CpG DNA**

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## **1** Introduction

As a result of early studies on mycobacteria (TOKUNAGA et al. 1992; YAMAMOTO et al. 1992), it is now well established that DNA from nonvertebrates, as well as certain synthetic oligodeoxynucleotides (ODNs), can cause strong activation of lymphoid cells, especially B cells, natural killer (NK) cells and antigen-presenting cells (APCs) (MESSINA et al. 1991; KRIEG et al. 1995; BALLAS et al. 1996; STACEY et al. 1996; SPARWASSER et al. 1998); such activation leads to B-cell proliferation and enhanced production of cytokines by APCs and NK cells. The immunostimulatory properties of DNA/ODNs are controlled by unmethylated CpG dinucleotide motifs with optimal flanking sequences (KRIEG et al. 1995). Reflecting the paucity of these motifs in vertebrate DNA – which is generally heavily methylated – purified DNA from humans, mice, frogs and fish (and also plants) are nonstimulatory for lymphoid cells, at least in terms of eliciting B-cell proliferation (SUN et al. 1997). By contrast, DNA from nonvertebrates, which is largely unmethylated, is strongly

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stimulatory for B cells (Sun et al. 1997); this applies to DNA from bacteria, insects, nematodes, yeasts and mollusks. For bacteria, insect and yeast DNA, selective methylation of CpG motifs abolished immunostimulatory activity (KRIEG et al. 1995; Sun et al. 1996, 1997).

The discovery that the immunostimulatory properties of DNA are restricted to unmethylated CpG motifs (plus appropriate flanking sequences) could explain the well-known phenomenon of "CpG suppression" found in vertebrate DNA (BIRD 1980). Thus, the low frequency of CpG motifs found in vertebrate DNA may reflect an evolutionary pressure to avoid DNA-induced stimulation of the immune system (SUN et al. 1997).

In the case of B cells and APCs, stimulation by CpG motifs in nonvertebrate DNA or ODNs (henceforth referred to as CpG DNA/ODNs) appears to involve endocytosis of the ligands, followed by initiation of various intracellular signalling pathways (HACKER et al. 1998; YI and KRIEG 1998; YI et al. 1998). Such signalling causes B cells to proliferate and differentiate into antibody-secreting cells, whereas APCs are induced to synthesize various cytokines, such as interleukin 1 (IL-1), IL-12, tumor necrosis factor  $\alpha$ , interferon  $\alpha$  (IFN $\alpha$ ) and IFN $\beta$ , and also to upregulate the expression of major histocompatibility complex (MHC) class II and co-stimulatory molecules (SUN et al. 1996; SPARWASSER et al. 1998). Information on the capacity of CpG DNA/ODNs to stimulate other types of lymphoid cells is still sparse.

For T cells, recent evidence suggests that CpG DNA/ODNs act as efficient adjuvants for stimulating antigen-specific proliferative responses of CD4<sup>+</sup> and CD8<sup>+</sup> cells (CHU et al. 1997; LIPFORD et al. 1997; ROMAN et al. 1997; WEINER et al. 1997; DAVIS et al. 1998; SUN et al. 1998a); for CD4<sup>+</sup> cells, the adjuvant effect promotes differentiation into T-helper 1 cells and preferentially stimulates production of immunoglobulin G2<sub>a</sub> (IgG2<sub>a</sub>) and IgG2<sub>b</sub> antibodies. Whether the adjuvant effect of CpG DNA/ODNs reflects a direct action on T cells, APCs or both is still unclear. Evidence favoring a direct effect on T cells has come from the finding that CpG ODNs can augment the proliferative response of purified T cells to anti-T-cell-receptor (TCR) monoclonal antibodies (mAbs) in vitro (LIPFORD et al. 1997), implying that CpG ODNs can provide a co-stimulatory signal to T cells. Alternatively, CpG DNA/ODNs may act largely by improving the function of APCs, e.g. by upregulating the expression of co-stimulatory molecules and/or inducing synthesis of stimulatory cytokines. On this second possibility, we have recently found that CpG DNA/ODNs can activate T cells via APC-production of type-I ( $\alpha$ , $\beta$ ) interferons (IFN-I) (Sun et al. 1998b). The data are summarized below.

## 2 T-Cell Activation in Vivo Following Injection of Insect DNA

In prior studies, we observed that proliferative responses of naïve CD4<sup>+</sup> and CD8<sup>+</sup> cells to specific antigens in vivo were augmented when the antigen was co-injected with CpG DNA (insect DNA) or CpG ODNs (SuN et al. 1998a); control ODNs

containing methylated rather than unmethylated CpG motifs (ZpG ODNs) were ineffective. This adjuvant effect on proliferation was not seen when CpG DNA/ODNs were injected without antigens. In fact, CpG DNA/ODNs proved incapable of inducing proliferation of naïve T cells. To our surprise, however, injection of CpG DNA/ODNs caused moderate to strong upregulation of a variety of cell-surface molecules, including CD69, B7–2, class I, Ly6C and intercellular adhesion molecule 1 (ICAM-1). This finding applied to CD4<sup>+</sup> and CD8<sup>+</sup> T cells, B cells and APCs (macrophages and dendritic cells) and was most prominent within the first 24h of injection; certain other molecules, e.g. CD25 (IL-2R $\alpha$ ), were not upregulated. Data illustrating upregulation of surface markers on CD4<sup>+</sup> and CD8<sup>+</sup> T cells and B220<sup>+</sup> B cells in the draining lymph nodes (LNs) of normal C57BL/6 (B6) mice injected with insect DNA 18h before are shown in Fig. 1.

For normal T cells, the expression of certain cell-surface markers, notably CD62L, CD45R and CD44, splits these cells into discrete subsets of naïve- and memory-phenotype cells (TOUGH and SPRENT 1994). Thus, for CD44 expression, typical naïve T cells express low to intermediate levels of CD44 (CD44<sup>lo/int</sup>), whereas memory cells express high levels of CD44 (CD44<sup>hi</sup>). As shown in Fig. 2, the upregulation of surface markers on T cells following injection of either insect DNA or CpG ODNs was not obviously skewed to particular subsets of T cells. Thus, both for CD4<sup>+</sup> and CD8<sup>+</sup> cells, upregulation of CD69 applies to naïve (CD44<sup>lo/int</sup>) T cells as well as to memory (CD44<sup>hi</sup>) cells.

### 3 The Role of APC

As in vivo, the capacity of CpG DNA/ODNs to induce upregulation of surface markers on T cells also occurs in vitro (SuN et al. 1998b). Significantly, however, this finding only applied when the cells cultured contained APCs, e.g. when CpG DNA/ODNs were added to unseparated spleen cells. Thus, when highly purified T cells were cultured with CpG DNA/ODNs, upregulation of surface markers on T cells was virtually undetectable. However, strong expression of these markers occurred when purified T cells were supplemented with APCs, e.g. spleen cells from



Fig. 1. DNA-induced upregulation of various cell-surface molecules on lymphoid subsets from B6 mice. B6 mice were injected with *Drosophila melanogaster* DNA ( $100\mu g/mouse$ ); 18h later, draining lymph-node cells were double stained for expression of the markers shown and also for B220, CD4 or CD8 expression. The data are expressed as the percentage increase in the mean fluorescence intensity ( $\Delta$ MFI) of staining for cells from DNA-injected mice versus phosphate-buffered-saline-injected mice. Adapted from SUN et al. (1998b)

DNA

+DNA



Fig. 2. CD69 upregulation of T cells in response to Drosophila melanogaster DNA or synthetic oligodeoxynucleotides (ODNs) in vivo. B6 mice were injected with D. melanogaster DNA (100µg/mouse) or ODNs (50µg/mouse); 18h later, draining lymph-node cells were triple stained for CD4 or CD8 and also for CD44 and CD69 expression. Adapted from SUN et al.

Fig. 3A,B. DNA-induced upregulation of CD69 on T cells requires antigen-presenting cells. Purified  $CD4^+$  or  $CD8^+$  cells (10<sup>6</sup> cells/well) were cultured overnight in the presence or absence of *Drosophila* melanogaster DNA (100µg/ml), either alone or in the presence of Rag-1<sup>-/-</sup> spleen cells (A) or B-celldepleted spleen cells from mahor histocompatibility complex  $\Gamma^{-1} \Pi^{-1}$  mice (B); spleen cells were added at 2×10<sup>6</sup> cells/well. Cells were then stained for CD8 and CD69. The data show the percentage of gated CD8<sup>+</sup> cells that were positive for CD69 expression relative to unstimulated CD8<sup>+</sup> cells. Adapted from SUN et al. (1998b)

-DNA

+DNA

RAG-1<sup>-/-</sup> mice (which lack T and B cells) (Fig. 3A) or B-depleted spleen cells from mice lacking MHC class-I and -II molecules (Fig. 3B).

These findings led to three conclusions. First, CpG DNA/ODNs appeared to have no effect on purified T cells. Second, CpG DNA/ODNs acted on T cells when accompanied by APCs. Third, to activate T cells, APCs did not have to express MHC molecules, implying that the function of APCs did not depend upon TCR/MHC interaction.

#### 4 The Role of IFN-I

Since CpG DNA/ODNs are known to induce APC to synthesize various cytokines, it seemed likely that APC-dependent activation of T cells by CpG DNA/ODNs was mediated by cytokines. In considering which particular cytokines might be involved, it is notable that the surface markers upregulated by CpG DNA/ODNs included Ly6C. This finding is of interest, because only one cytokine, namely IFN-I, is known to induce Ly6C upregulation (DUMONT and COKER 1986).

To seek direct evidence for the role of IFN-I, we examined the effects of injecting insect DNA into mice lacking receptors for IFN-I, i.e. IFN-IR<sup>-/-</sup> mice; since these mice were on a 129 background, normal 129 mice were used as a control. The results were clear-cut (Fig. 4). Thus, when insect DNA was injected in vivo (Fig. 4A) or added to a mixture of T cells and APCs in vitro (Fig. 4B), surface-marker upregulation on T cells was prominent with normal 129 cells but very low or undetectable with 129 IFN-IR<sup>-/-</sup> cells; this finding applied to B cells as well as to CD4<sup>+</sup> and CD8<sup>+</sup> T cells.

The above findings indicated that APC-dependent stimulation of T cells in response to CpG DNA was mediated largely or solely by IFN-I, presumably by a direct action of IFN-I on T cells. In favor of this idea, the failure of IFN-IR<sup>-/-</sup> T cells to respond to insect DNA in vitro could not be overcome by addition of normal IFN-IR<sup>+</sup> APCs, i.e. RAG-1<sup>-/-</sup> spleen cells (Fig. 4B).

A key prediction of the above findings was that adding IFN-I to purified normal T cells in vitro would induce upregulation of surface markers in the absence of APCs. This was indeed the case. Moreover, the range of surface markers upregulated when purified T cells were cultured with IFN-I (IFN- $\beta$ ) was virtually the same as when APC-containing T-cell populations were exposed to CpG DNA/ ODNs. The capacity of IFN-I to induce upregulation of CD69 and B7-2 on purified CD8<sup>+</sup> cells in vitro is shown in Fig. 5. In control studies, culturing purified T cells with IFN- $\gamma$  induced modest upregulation of ICAM-1 but no upregulation of CD69, B7-2, Ly6C or MHC class I.

### 5 The Adjuvant Effect of IFN-I

In prior studies, we observed that poly-I:C, a powerful inducer of IFN-I, acted as an adjuvant for the primary proliferative response of 2C TCR transgenic CD8<sup>+</sup>





**Fig. 4A,B.** DNA-induced upregulation of various cell surface molecules in interferon-I receptor (IFN-IR) knockout (KO) mice versus 129 wild-type control mice. **A** IFN-IR KO mice and 129 wild-type mice were injected with *Drosophila melanogaster* DNA (100µg/mouse); 18h later, draining lymph-node cells were double stained for expression of the markers shown and for B220, CD4 or CD8 expression. The data are expressed as the percentage increase in the mean fluorescence intensity ( $\Delta$ MFI) of staining for cells from DNA-injected mice versus phosphate-buffered-saline-injected mice. **B** Purified CD8<sup>+</sup> cells from 129 or IFN-IR KO mice (10<sup>6</sup> cells/well) were cultured with or without *D. melanogaster* DNA (100µg/ml), either alone or with RAG-1<sup>-/-</sup> spleen antigen-presenting cells (2 × 10<sup>6</sup> cells/well), then stained for CD8 and CD69 or B7-2 18h later. The data are expressed as the MFI of staining for the markers shown, gating on CD8<sup>+</sup> cells. Adapted from SUN et al. (1998b)

cells responding to specific antigens (TOUGH et al. 1996). Similar results were found with CpG DNA/ODNs (SUN et al. 1998a). In both situations, co-injection of poly-I:C or CpG DNA/ODNs considerably augmented the T proliferative responses to specific antigens. In light of these findings, we have been assessing the adjuvant effects of purified IFN-I (IFN- $\beta$ ), using IFN- $\gamma$  as a control. A summary of our unpublished data on this topic is given below.

In initial experiments, we examined whether IFN-I could amplify the in vitro proliferative response of 2C TCR transgenic CD8<sup>+</sup> cells to specific peptides. This was indeed the case. Thus, when 2C CD8<sup>+</sup> cells were cultured with peptides presented by normal spleen APCs, addition of IFN-I considerably augmented the T proliferative response; IFN- $\gamma$ , by contrast, had no effect. The adjuvant effect of IFN-I also applied in vivo. Thus, as found previously with poly-I:C (TOUGH et al.



**Fig. 5.** Upregulation of CD69 and B7-2 on purified CD8<sup>+</sup> cells in response to interferon  $\beta$  (IFN $\beta$ ) in vitro. Purified CD8<sup>+</sup> cells (2 × 10<sup>6</sup> cells/well) were cultured with or without IFN $\beta$  (10,000 units/ml) overnight; cells were then triple stained for CD8, CD44 and CD69 or B7-2. The data shown are for gated CD8<sup>+</sup> cells

1996) and CpG DNA/ODNs (SUN et al. 1998a), injection of IFN-I enhanced the proliferative response of 2C CD8<sup>+</sup> cells to specific peptides in vivo. To our surprise, however, under certain experimental conditions, IFN-I did not act as an adjuvant but instead inhibited the T proliferative response.

### 6 Anti-Proliferative Effects of IFN-I

It should be noted that the adjuvant effect of IFN-I was only apparent during the later stages of the T proliferative response, i.e. on day 3 and thereafter; this applied both in vivo and in vitro. At earlier stages, i.e. on day 2, T proliferative responses were markedly inhibited by IFN-I (but not by IFN- $\gamma$ ). Under certain conditions, IFN-I also inhibited late (day 3, 4) proliferative responses. Here, the presence or absence of viable APCs seemed to be the key. Thus, with T proliferative responses driven by peptides presented by normal viable spleen APCs, IFN-I inhibited early responses but augmented late responses. By contrast, when responses occurred in the absence of viable APCs, IFN-I inhibited both early and late proliferative responses. This latter finding applied to naïve T-cell proliferative responses elicited by cross-linked anti-TCR plus anti-CD28 mAbs; it also applies to proliferation of memory-phenotype (CD44<sup>hi</sup>) CD8<sup>+</sup> cells in response to a cytokine, IL-15. Similar findings applied to proliferation of 2C CD8<sup>+</sup> cells to peptides presented by transfected *Drosophila* cells as APCs; these cells die rapidly at 37°C (CAI et al. 1996) and are presumably unresponsive to IFN-I.

Our interpretation of these findings is that IFN-I has two opposing effects on T-cell function. In the absence of viable APCs, IFN-I acts directly on T cells,

presumably via IFN-IR, and downregulates the capacity of the cells to mount a proliferative response. When viable APCs are present, however, stimulation of APCs by IFN-I somehow counters the inhibitory effects of IFN-I on T cells and leads to enhanced proliferative responses.

#### 7 Concluding Comments

As with other nonspecific stimuli, such as lipopolysaccharides (LPS) and poly-I:C, the effects of exposing lymphoid cells to CpG DNA/ODNs are highly complex. For some cell types, especially B cells and dendritic cells, CpG DNA/ODNs appear to act directly and cause upregulation of certain cell-surface markers (including co-stimulatory molecules, such as B7) and synthesis of various cytokines.

Direct activation of APCs would seem the most likely explanation for the strong adjuvant function of CpG DNA/ODNs. This notion rests on the assumption that resting APCs function poorly and need to be activated in order to present antigens effectively to naïve T cells. However, the precise difference between "efficient" and "inefficient" APCs is still unclear. The simplest idea is that the efficiency of APCs is a direct reflection of the range and density of the co-stimulatory/ adhesion molecules on the cell surface. The data on the APC function of transfected Drosophila cells are in accordance with this idea. Thus, in terms of presenting specific peptides to purified naïve 2C transgenic CD8<sup>+</sup> cells, class-I-(L<sup>d</sup>)-transfected Drosophila cells are highly efficient APCs when these cells co-express B7 (B7-1 or B7-2) and ICAM-1 but are totally nonfunctional without these co-stimulatory molecules (CAI et al. 1996). Interestingly, these data only apply when purified T cells are used. With unseparated 2C spleen cells as responders, the results are quite different (SUN et al. 1996). Here, significant T proliferative responses to peptides occur, with Drosophila cells expressing only class-I molecules alone. The explanation for this paradox is that CpG DNA released from the Drosophila cells acts on the B cells and APCs in spleen and induces upregulation of B7 and ICAM-1. Provided that high concentrations of peptides are used, these activated cells then provide bystander co-stimulation for the responding CD8<sup>+</sup> cells. It is worth noting that, even with high concentrations of peptides, Drosophila APCs expressing class-I molecules alone are totally non-stimulatory for purified CD8<sup>+</sup> cells. Hence, in this system, we have not found evidence that CpG DNA can provide a direct co-stimulatory signal for T cells (SUN et al. 1996; LIPFORD et al. 1997).

In considering the above data, the point to emphasize is that co-expression of just two co-stimulatory molecules, B7 and ICAM-1, enables class-I-transfected *Drosophila* cells to display strong APC function for purified T cells in the absence of added cytokines. Since the endogenous "cytokines" made by *Drosophila* cells are unlikely to affect mammalian cells (T cells), it would seem to follow that, for normal APCs, efficient presentation of peptides to T cells simply requires that the APCs express high levels of appropriate co-stimulatory molecules. Hence, if

adjuvants are able to induce direct upregulation of costimulatory molecules on APCs – which seems to be the case – concomitant production of cytokines by APC may be redundant. But is this the case in a normal physiological setting?

With normal APCs in intact animals, assessing whether the adjuvant effect of CpG DNA/ODNs is due in part to cytokine production is clearly difficult, because the range of cytokines released by APCs after activation by adjuvants is considerable. One approach to this problem is to investigate whether individual APC-derived cytokines have adjuvant activity. Based on the effects of IFN-I considered earlier, this is indeed the case. Thus, both in vitro and in vivo, purified IFN-I acts as a strong adjuvant for APC-induced T proliferative responses (unpublished data of the authors). In light of this finding, it would seem quite likely that the adjuvant function of co-stimulatory molecules on APCs but also involves the production of stimulatory cytokines, such as IFN-I.

How IFN-I exerts its adjuvant activity is still unclear. It seems that the most likely possibility is that, like CpG DNA/ODNs, IFN-I functions by augmenting upregulation of co-stimulatory molecules on APCs. This idea may be an over-simplification, however, because in our experiments, IFN- $\gamma$  induced strong upregulation of co-stimulatory molecules on APCs but, at least in vitro, failed to display adjuvant activity. Hence, one has to consider other possibilities. For example, IFN-I may induce APCs to produce cytokines with direct co-stimulatory functions for T cells, e.g. IL-15 and IL-6. IFN-I could also act at the T-cell level, e.g. by causing T cells to express receptors for stimulatory cytokines released by APCs.

Although direct evidence for these and other possibilities is still sparse, it is of interest that IFN-I does act directly on T cells, including naïve cells, and causes these cells to enter a state of partial activation. Such activation is associated with upregulation of various cell-surface markers but does not lead to cell division. Indeed, in the absence of viable APCs, it is striking that the partial activation of T cells induced by IFN-I is associated with a reduced capacity to mount T proliferative responses. The cells are not simply "anergic", however, because we have found no evidence that IFN-I inhibits T-cell production of IL-2. Based on studies with cell lines (GRANDER et al. 1997), the anti-proliferative function of IFN-I on naïve T cells may reflect intracellular production of cell-cycle inhibitors. If so, the paradox remains that the presence of viable APCs reverses the anti-proliferative effect of IFN-I and causes enhanced T proliferative responses. How APCs negate the anti-proliferative function of IFN-I has yet to be resolved.

As a whole, the data on the adjuvant effects of CpG DNA/ODNs suggest that these compounds act primarily at the APC level and lead both to direct upregulation of co-stimulatory molecules and release of various cytokines. For naïve T cells responding to specific antigens, the combination of APC activation and cytokine production elicited by CpG DNA/ODNs and other adjuvants enhances the T proliferative response and promotes differentiation into long-lived memory T cells. It is important to emphasize that T-cell activation and proliferation in this situation is heavily antigen dependent; without antigens, T cells undergo partial activation, e.g. in response to IFN-I, but do not enter the cell cycle. However, this scenario applies only to naïve T cells. Thus, when memory-phenotype T cells are exposed to adjuvants, the cytokines released from activated APCs can act directly on T cells and induce proliferation in the apparent absence of antigen (TOUGH et al. 1996). Such "bystander" proliferation of memory-phenotype T cells is especially prominent for  $CD8^+$  cells ( $CD44^{hi}$   $CD8^+$  cells) and can be evoked not only by CpG DNA/ODNs but also by other compounds with adjuvant activity, notably poly-I:C and LPS (TOUGH et al. 1996, 1997). How these agents – and also certain cytokines, such as IFN-I – induce proliferation of  $CD44^{hi}$   $CD8^+$  cells in vivo is still unclear, but secondary production of stimulatory cytokines, such as IL-15, is a likely possibility.

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# Hematopoietic Remodeling Triggered by CpG DNA

G.B. LIPFORD and T. SPARWASSER

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## **1** Introduction

During acute bacterial infection, significant hematological alterations occur, most of which are simply monitored as blood leukocytosis (SELIG and NOTHDURFT 1995). Hematopoietic growth factors acting as regulators maintain the steady state between production and consumption of mature blood cells. Particularly under stress conditions, such as infection, these factors play a major role in cellular adaptation processes (CANNISTRA and GRIFFIN 1988). Many of these growth factors not only influence the proliferation and commitment at the stem cell and progenitor cell level, but also signal for a rapid mobilization of hematopoietic progenitor cells from the bone marrow (BM) into the blood and distant hematopoietic organs, thus remodeling cellular compartmentalization (MORRISON et al. 1995). Additionally, growth factors and other cytokines play a major role in functional adaptation at the level of the mature cell compartment, stimulating metabolic, cytotoxic, or phagocytic activities. Because, at any given time, progenitors in different hematopoietic organs are exposed to different cytokines and growth factors, the adaptive demand induced by infection on cellular composition and status can be met.

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The mobilization of stem cells is evolutionarily conserved in all species tested thus far, including mouse, dog, monkey, and human. However, the selective forces for the conservation of this phenomenon are not immediately apparent (MORRISON et al. 1995). It has long been recognized, in experimental animals, that splenomegaly, mobilization of hematopoietic precursors into the blood, and increased numbers of colony-forming cells (CFCs) are induced not only by gram-positive and gram-negative bacteria but also by bacterial subcellular fractions, such as the mycobacterial component of complete Freund's adjuvant (CFA), lipopolysaccharides (LPSs), and muramyl dipeptide. Additionally, synthetic double-stranded polyribonucleotides, such as Poly I–Poly C, show enhancement in CFCs. We add to this list bacterial CpG DNA, recently shown to trigger hematopoietic events after injection in mice. Apparently, recognition of bacterial CpG DNA as a foreign bacterial product signals infectious danger and induces the release of a variety of colony-stimulating factors (CSFs), cytokines, and chemokines that facilitate hematopoietic events.

#### **2** Infection Danger Induces Hematopoietic Mobilization

Macrophages and dendritic cells (DCs) are equipped with pattern-recognition receptors (PRRs), which recognize conserved molecular structures shared by large groups of pathogens. The main difference between PRRs and the clonally distributed antigen receptors of T and B cells is that the specificities of PRRs are germline encoded. Thus, one parameter imposed on PRRs is the recognition of non-self through pathogen-associated structural patterns. The innate immune system controls the initiation of the adaptive immune response by regulating the expression of co-stimulatory activity on antigen-presenting cells (APCs) and the release of effector cytokines, which in turn instruct the adaptive immune system to develop a particular T-helper 1 (Th1)- or Th2-effector response (T. Sparwasser and G.B. Lipford, this issue). Here, we focus on an additional responsibility of the innate immune system: elaboration of hematopoietically active cytokines and growth factors. Upon stimulation with bacterial products, macrophages and DCs are capable of releasing granulocyte CSF, monocyte CSF, granulocyte-monocyte CSF (GM-CSF), interferon  $\alpha$  (IFN- $\alpha$ ), tumor necrosis factor  $\alpha$  (TNF- $\alpha$ ), transforming growth factor β, interleukin 1 (IL-1), IL-6, IL-8, IL-10, IL-12, macrophage inflammatory protein 1 (MIP-1), MIP-2, and others. Many of these substances have dramatic hematopoietic consequences (BROXMEYER 1995).

It has long been known that certain adjuvants induce hematopoietic events. McNeill demonstrated that intraperitoneal (i.p.) injection of CFA stimulated the multiplication and affected the distribution of cells in vivo that form granulocytic and monocytic colonies in vitro (McNEILL 1970). The results showed that a 100-fold rise in spleen CFCs occurs at the same time as a CFC rise in the blood, but the level of CFCs in the BM is only two- to threefold greater than control levels. The rise in CFCs in the spleen was associated with leukocytosis in the peripheral

blood. It was hypothesized that, following injection of foreign materials granulocyte-monocyte progenitor cells in the marrow are stimulated to divide, and many of these cells leave the marrow and are trapped in the spleen. These cells then undergo further differentiation into more mature cells, which appear in the peripheral blood and cause leukocytosis. A likely explanation was given by the observation that CFA induced increased serum CSF levels within 6–8h after injection. Similar observations and explanations were subsequently provided when either LPS or muramyl dipeptide was used as a stimulant (MOATAMED et al. 1975; APTE and PLUZNIK 1976a,b; STABER and METCALF 1980; WUEST and WACHSMUTH 1982; GALELLI and CHEDID 1983). In an early study utilizing LPS, it was shown that LPSnon-responder mice (C3H/HeJ) had strongly diminished CFC levels due to their reduced ability to produce CSFs in response to LPS (APTE and PLUZNIK 1976b). These observations imply that several bacterial cell products stimulate the release of hematopoietically active factors that expand and mobilize progenitor cells.

Several bacterial infections have been shown to lead to very similar phenomena. These micro-organisms include both gram-positive and gram-negative bacteria. Salmonella infection induces a marked increase in splenic and BM CFCs 2-3 days after infection (WILSON et al. 1982). Others have claimed, however, that BM CFCs were depressed after Salmonella infection (MIYANOMAE et al. 1983; KIRIKAE et al. 1986). Of additional interest, the hematopoietic response to Salmonella infection could be monitored in LPS-unresponsive C3H/HeJ mice (MIYANOMAE et al. 1983). Similar observations have been made after Listeria monocytogenes, Brucella abortus, and Escherichia coli infections (WING et al. 1985; CHEERS and YOUNG, 1987; ROTHSTEIN et al. 1987). Bacillus Calmette-Guerin (BCG), injected intravenously (i.v.), was noted to quickly push the quiescent stem cells of BM into the S phase of the cell cycle (POUILLART et al. 1975). In later studies, it was noted that i.v. injection of BCG into C57Bl/6 mice resulted in a rapid development of transient anemia associated with an increased number of granulocytes and monocytes (MARCHAL and MILON 1986). In IFN-y-deficient mice infected with mycobacteria, a dramatic remodeling of the hematopoietic system was noted (MURRAY et al. 1998). Myeloid cell proliferation proceeded unchecked throughout the course of infection, resulting in a transition to extramedullary hematopoiesis. The splenic architecture was altered by expansion of macrophages, granulocytes, and extramedullary hematopoietic tissue. These features coincided with splenomegaly, an increase in splenic myeloid CFC, and granulocytosis in the peripheral blood. Systemic levels of cytokines were elevated, particularly IL-6 and GM-CSF.

## 3 CpG DNA Signals Infectious Danger

Yamamoto, Tokunaga and colleagues discovered immunostimulatory DNA by a series of studies originally aimed at analyzing BCG-mediated tumor resistance in mice. A fraction extracted from BCG (designated MY-1) was shown to exhibit anti-

tumor activity in vivo, augment natural killer (NK) cell activity, and trigger type-1 and type-II IFN release from murine spleen cells or human peripheral blood lymphocytes (PBL) in vitro (TOKUNAGA et al. 1984, 1988; MASHIBA et al. 1988; YAMAMOTO et al. 1988). These activities can be destroyed by DNase pre-treatment of MY-1, but not by RNase treatment. Pisetsky and co-workers independently observed that normal mice, as well as humans, respond to bacterial DNA (but not vertebrate DNA) by producing anti-DNA antibodies (MESSINA et al. 1991). They realized that bacterial DNA was mitogenic for murine B cells and postulated that this activity resulted from "non-conserved structural determinants". The differential stimulative capacity of bacterial DNA versus vertebrate DNA was also demonstrated for induction of NK cell activity by Yamamoto et al. (YAMAMOTO et al. 1992b).

High-performance liquid chromatography analysis of BCG extracts showed that the MY-1 fraction was composed of a broad size range of DNA fragments, with a peak at 45 bases. Synthetic 45-mer oligodeoxynucleotides (ODNs) derived from BCG cDNA sequences were positive for IFN-inducing capacity and augmentation of NK cytotoxicity (KATAOKA et al. 1992; YAMAMOTO et al. 1992a, 1992b). Subsequently, Krieg et al. formulated a hypothetical framework for understanding the pattern recognition of bacterial or synthetic DNA (KRIEG et al. 1995). Using sequence-specific CpG-containing ODNs mediated mitogenicity to B cells as an assay, they discovered that CpG dinucleotides with selective flanking bases were important and, specifically, that DNA motifs displaying a 5'-Pu-Pu-CpG-Pyr-Pyr-3' base sequence were biologically active. Thus, it was speculated that immune cells sense unique base sequences of pathogen-associated DNA. The realization that these sequences are under-represented in vertebrate DNA offers an explanation for several biological observations in the context of non-self pattern recognition by the immune system (WAGNER 1999).

CpG ODNs directly activate immature DCs and macrophages (STACEY et al. 1996; SPARWASSER et al. 1997b, 1998; JAKOB et al. 1998). In utilizing PRRs, macrophages recognize non-self DNA through CpG motifs and initiate inflammatory responses. In early work by Yamamoto et al. it was concluded that IFN- $\alpha/\beta$  produced by immunostimulatory-ODN-stimulated spleen cells might have originated in an adherent cell population (TOKUNAGA et al. 1988; YAMAMOTO et al. 1988). We and others have discovered that DNA from gram-negative and gram-positive bacteria, plasmid DNA, or synthetic CpG ODNs triggered macrophages to activate the transcription factor nuclear factor (NF $\kappa$ B), to transcribe cytokine mRNAs, and to secrete pro-inflammatory cytokines, such as GM-CSF, TNF- $\alpha$ , IL-1, IL-6, II-12, and IL-18 (HALPERN et al. 1996; SATO et al. 1996; STACEY et al. 1996; CHACE et al. 1997; LIPFORD et al. 1997b; ROMAN et al. 1997; SPARWASSER et al. 1997a,b).

### 4 Induction of Splenomegaly by CpG ODNs

Splenomegaly is a well-recognized phenomenon accompanying some oligonucleotide injections. Branda et al. observed that mice developed massive splenomegaly and polyclonal hypergammaglobulinemia within 2 days after intravenous injection of a phosphorothioate oligomer that was antisense to a portion of the rev region of the human immunodeficiency virus 1 genome (BRANDA et al. 1993). Histological examination of spleens from injected animals showed marked expansion of a uniform-appearing population of small lymphocytes. Flow-cytometry analysis indicated that the responding cells were predominantly B lymphocytes. Mojcik et al. observed that injection of mice with antisense to the initiation region of the env gene resulted in (i) increased spleen cell numbers, primarily due to an increase in splenic B cells, (ii) increased class-II major histocompatibility complex expression on B cells, (iii) increased RNA and DNA synthesis, and (iv) increased numbers of immunoglobulin (Ig)-producing cells (MOJCIK et al. 1993). They concluded that products of certain endogenous retroviral sequences regulate lymphocyte activation in vivo. McIntyre et al. in efforts to test the efficacy of NFkB p65 oligonucleotides in vivo, unexpectedly observed that the control p65-sense (but not the p65-antisense) oligonucleotides caused massive splenomegaly in mice (MCINTYRE et al. 1993). In this study, they demonstrated a sequence-specific stimulation of splenic cell proliferation, both in vivo and in vitro, by treatment with p65-sense oligonucleotides. Cells expanded by this treatment were primarily B-220+, sIg+B cells. The secretion of Ig by the p65-sense oligonucleotide-treated splenocytes was also enhanced. In addition, the p65-sense-treated splenocytes (but not several other cell lines) showed an upregulation of NFkB-like activity in the nuclear extracts, an effect not dependent on new protein or RNA synthesis. Zhao et al. concluded that phosphorothioated-ODNs induce splenomegaly due to B-cell proliferation (ZHAO et al. 1996b). In a follow-up, they observed that administration of the 27-merphosphorothioate oligonucleotide in mice resulted in splenomegaly and an increase in IgM production 48h after administration (Zhao et al. 1996a).

Bacterial DNA and synthetic ODNs containing unmethylated CpG dinucleotides induce murine B cells to proliferate and secrete Ig in vitro and in vivo (KRIEG et al. 1995). This activation is very much enhanced by the antigen-receptor ligation. Optimal B-cell activation requires a DNA motif in which an unmethylated CpG dinucleotide is flanked by two 5' purines and two 3' pyrimidines. ODNs containing this CpG motif induce more than 95% of all spleen B cells to enter the cell cycle. In a study by Monteith et al., treatment of rodents with phosphorothioate ODNs induced a form of immune stimulation characterized by splenomegaly, lymphoid hyperplasia, hypergammaglobulinemia, and mixed mononuclear cellular infiltrates in numerous tissues (MONTEITH et al. 1997). Immune stimulation was evaluated in mice with in vivo and in vitro studies, including a review of historical data. All phosphorothioate ODNs evaluated induced splenomegaly and B-lymphocyte proliferation. Splenomegaly and B-lymphocyte proliferation increased with the dose or concentration of ODN. The overriding evidence provided by the literature thus concludes that the phenomena of splenomegaly induced by ODNs is probably sequence dependent and explained by B-cell mitogenicity.

Zhao et al. administered to mice a 27-mer phosphorothioate oligonucleotide (sequence 5'-TCG TCG CTG TCT CCG CTT CTT CTT GCC-3'), which had

previously been shown to cause splenomegaly and hypergammaglobulinemia on in vivo administration in mice, and studied the pattern and kinetics of cytokine production at both the splenic mRNA and serum protein levels (ZHAO et al. 1997). Following i.p. administration of high doses (50mg/kg) of oligonucleotide, significant increases in the splenic mRNA levels of IL-6, IL-12p40, IL-1β, and IL-1Ra and serum levels of IL-6, IL-12, MIP-1β, and MCP-1 were observed. In contrast, no significant differences in splenic mRNA levels of IL-2, IL-4, IL-5, IL-9, IL-13, IL-15, IFN-γ, or migration inhibition factor or serum levels of IL-2, IL-4, IL-5, IL-10, IFN- $\gamma$ , or GM-CSF were detected. These studies show a distinct pattern and kinetics of cytokine production following oligonucleotide administration and further demonstrate that cytokine induction is not a general property of phosphorothioate oligonucleotides, but is dependent on a given sequence and dose of the oligonucleotides. Serum release of IL-1, IL-6, IL-12, and TNF-a has been confirmed (LIPFORD et al. 1997b and unpublished data). Contrary to Zhao et al. Klinman et al. have demonstrated GM-CSF production by macrophages in vitro, and the production of IFN- $\gamma$  has been demonstrated by several authors (KLINMAN et al. 1996; Y1 et al. 1996).

## 5 Splenomegaly is Associated with Extramedullary Hematopoiesis

The cytokine repertoire induced by low doses of CpG ODNs is Th1 in nature, and ample evidence suggests that this strongly biases subsequent immune-response development to Th1 (T. Sparwasser and G.B. Lipford, this issue). This type of cellular-oriented immune response becomes particularly important in infections caused by intracellular pathogens. CpG ODNs have been demonstrated to protect mice efficiently in different infection models, such as listeriosis, leishmaniasis, and pathology caused by *Francisella tularensis* (KRIEG et al. 1998; ZIMMERMANN et al. 1998; ELKINS et al. 1999). Interestingly, in addition to acute protective effects primarily mediated by activation of APCs, lasting effects up to 2 weeks after injection of ODNs have been observed (KRIEG et al. 1998; our own unpublished results). These findings suggest a propensity for CpG DNA to induce longer-term changes in the immune system.

Pro-inflammatory cytokines secreted by innate immune cells upon stimulation with bacterial CpG DNA, such as TNF- $\alpha$  and IL-1, can trigger CSF production by accessory cells locally as well as in the bone-marrow microenvironment. Compared to hematopoietic growth factors induced in the BM, CSFs generated at peripheral sites of infection have different functions. Here, CSFs are most likely to influence activation and maturation of myeloid cells necessary for the acute defense against the invading pathogen, whereas the proliferation and differentiation of myeloid progenitor cells and hematopoietic remodeling provide for an adequate second line of defense (CANNISTRA and GRIFFIN 1988).

When we examined the protective effects of CpG ODNs, we observed a dramatic but transient splenomegaly, with a maximum increase of spleen weight at day 6 (SPARWASSER et al. 1999). This increase of spleen weight by more than three times the normal weight was dose- and sequence-dependent and normalized after 12-14 days. In contrast to previous studies by others analyzing the splenic cellular composition within the first 4 days after ODN challenge, we could not explain our observations by B-cell proliferation. Histologically, splenic architecture was altered, showing large immature blast within the red pulp. Facs staining of day-6 spleen cells revealed a more than tenfold expansion of a non-B-/non-T-cell fraction. This population of spleen cells was highly enriched for hematopoietic progenitors. Colony assays measuring granulocyte-macrophage colony-forming units (GM-CFUs) demonstrated an increase in splenic myeloid progenitors paralleling the increase in splenic cell count. Kinetically, a discrete increase of GM-CFUs in BM preceding the splenic changes could be detected, as if a mobilization of BM-derived progenitor cells to the spleen may have taken place. A single application of CpG ODN (60µg/injection) far exceeded the documented hematopoietic stimulus LPS. Furthermore, the number of early erythrocyte progenitors (burst-forming units, BFU-Es) in the spleen was also increased after i.p. injection of CpG ODNs. Transfer of spleen cells from CpG-ODN pre-treated animals into lethally irradiated syngenic mice yielded increased spleen colony forming units (CFU-Ss). CFU-Ss are an indicator for the existence of primitive stem cells, and it is controversial whether these cells respond to cytokines (MORRISON et al. 1995). These data suggested the possibility of reconstituting lethally irradiated mice by an adoptive transfer of CFU-Ss contained in the spleens of CpG-DNA-treated animals.

## 6 CpG-DNA-Mediated Protection from Immunosuppression

Proliferation, mobilization and differentiation of hematopoietic progenitor cells are of major therapeutic importance in clinical situations of immunosuppression. Recently, we described that the hematopoietic effects of CpG ODNs could be used to mitigate irradiation-induced damage to the hematopoietic system (SPARWASSER et al. 1999). CpG-DNA challenge of sublethally irradiated mice caused radioprotective effects, in that recovery of GM-CFU and cytotoxic T-cell (CTL) function was enhanced. Interestingly, a single i.p. injection after sublethal irradiation caused sufficient hematopoietic remodeling to compensate for radiation-induced damage to the lympho-hematopoietic system. CpG-ODN challenge within 30min after sublethal irradiation led to a fourfold increase of splenic GM-CFUs after 2 weeks. Irradiated mice immunized with soluble ovalbumin (OVA) exhibited significantly enhanced OVA-specific CTL activity if CpG DNA was given therapeutically. The increase in GM-CFUs and CTL function correlated with enhanced resistance to *Listeria* infection. Mice were infected at day 14 after irradiation, and survival was recorded for 30 days. If CpG ODNs were applied, these mice were protected against a normally lethal challenge with the pathogen, implying an increased resistance due to CpG-DNA-triggered cellular-adaption processes. Preliminary data suggest that these results can also be extended to immunosuppression caused by chemotherapeutic agents, such as 5-fluorouracil (our own unpublished data).

#### 7 Conclusion

CpG DNA signals infectious danger not only to immune cells but also to hematopoietic cells and thereby communicates the need for increased hematopoiesis necessary to combat infections. APC-derived cytokines and CSFs appear to be responsible for a transient phase of hematopoietic remodeling comprising mobilization and differentiation of BM progenitor cells. To date, it is not clear which of these factors, which can also act synergistically, are relevant for the hematopoietic effects observed in mice. It is also feasible that CpG ODNs directly target BM stroma cells, thereby inducing the production and release of hematopoietically active growth factors, cytokines, and chemokines.

The strong hematopoietic potency of bacterial CpG-DNA motifs, which far exceeds that of high doses of LPS in promoting hematopoietic changes in the murine system, could explain why e.g. Salmonella infections have similar effects in LPS-susceptible and LPS-resistant mice (MIYANOMAE et al. 1983). Furthermore, the well-known hematopoietic effect of CFA containing dead mycobacteria may depend strongly on the presence of bacterial DNA (McNeill 1970). Most importantly, CpG ODNs appear to display a dual function. When used as a "new class of adjuvants" with proteinaceous antigen (LIPFORD et al. 1997a), CpG-ODNs cause an acute and strong Th1-biasing effect on emanating immune responses via DC activation and local production of cytokines, such as IL-12. However, longterm effects include crucial changes in the cellular compartmentalization of innate and adaptive immune cells; these changes are antigen independent. As a consequence, the system is "primed" for protective Th1-polarized immune responses, as shown in CTL responses to OVA and resistance of BALB/c mice against Leishmania major (our own unpublished data). Protective immunity against various pathogens has been demonstrated to depend on acute APC activation caused by the pattern recognition of pathogen DNA (KRIEG et al. 1998; ZIMMERMANN et al. 1998; ELKINS et al. 1999). Conversely, its ability to cause the release of hematopoietic factors that control the steady state between production and consumption of immune cells may be a critical factor contributing to the observed lasting priming effects (Th1) of CpG DNA. The cellular and molecular basis of the priming effect is poorly understood. However, because of its possible importance in immunosuppressive situations, the therapeutic use of CpG ODNs to mitigate damage to the hematopoietic system by e.g. chemotherapy, radiotherapy, or accidental radiation exposure might be an interesting alternative to combinations of recombinant growth factors.

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# CpG DNA Augments the Immunogenicity of Plasmid DNA Vaccines

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## 1 Introduction: Immunostimulatory Properties of CpG Oligodeoxynucleotides

Vaccine development has been revolutionized by the use of antigen-encoding DNA plasmids to induce cellular and humoral immune responses against pathogenic viruses, parasites, bacteria and tumors (Cox et al. 1993; ULMER et al. 1993; WANG et al. 1993; SEDEGAH et al. 1994). DNA vaccines are composed of an antigenencoding gene whose expression is regulated by a strong mammalian promoter incorporated into the plasmid backbone of bacterial DNA (WOLFF et al. 1990; MANTHORPE et al. 1993; ULMER et al. 1993). When injected intramuscularly or intradermally, DNA vaccines are transcribed and translated, and the protein they encode is presented to the immune system in the context of self major histocompatibility complex (MHC) (WOLFF et al. 1990; Cox et al. 1993; ULMER et al. 1993).

Although the nature, magnitude and duration of the immune response elicited by DNA vaccines is influenced by multiple factors, it has been repeat-

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edly shown that intramuscular delivery stimulates a T helper 1 (Th1)-driven response characterized by cytotoxic T lymphocyte (CTL) induction and the release of interferon  $\gamma$  (IFN $\gamma$ ) and antigen-specific immunoglobulin G2a (IgG2a) antibodies (ULMER et al. 1993; MOR et al. 1995; KLINMAN et al. 1998). Numerous animal studies (particularly in mice) demonstrate that DNA-vaccineinduced immunity can confer protection against pathogen challenge (ULMER et al. 1993; SEDEGAH et al. 1994). Nevertheless, it has proven difficult to elicit equally strong and protective immune responses in primates (CALAROTA et al. 1998; WANG et al. 1998). This has prompted widespread efforts to improve the immunogenicity of DNA vaccines, including (i) altering membrane expression of the encoded gene, (ii) substituting stronger promoters, (iii) targeting the plasmid to transfect specific cell types, and/or (iv) co-administering DNA vaccines with adjuvants (including plasmids encoding cytokines and/or co-stimulatory molecules).

Investigators in the field of autoimmunity were the first to demonstrate that bacterial DNA has immunostimulatory properties (GILKESON et al. 1989; MESSINA et al. 1991). Yamamoto et al. were the first to report that synthetic oligodeoxynucleotides (ODNs) with sequences patterned after those found in bacterial DNA could activate natural killer (NK) cells to secrete IFN $\gamma$  (YAMAMOTO et al. 1992). They hypothesized that palindromic sequences present in the synthetic ODNs were responsible for this stimulation. In collaboration with Dr. Krieg, my lab demonstrated that specific sequence motifs present in bacterial DNA consisting of an unmethylated CpG dinucleotide flanked by two 5' purines (optimally, GpA) and two 3' pyrimidines elicited an "innate" immune response (KRIEG et al. 1995; HALPERN et al. 1996; KLINMAN et al. 1996; SATO et al. 1996). These motifs are 20 times less common in mammalian than in microbial DNA, due to differences in frequency of utilization and methylation pattern of CpG dinucleotides in eukaryotes versus prokaryotes (RAZIN and FRIEDMAN 1981; CARDON et al. 1994). We showed that these CpG motifs activate a variety of immunologically relevant cells to proliferate and/or secrete, including macrophages and B, T and NK cells (KRIEG et al. 1995; KLINMAN et al. 1996; SATO et al. 1996). ODNs containing immunostimulatory CpG motifs induced a significant rise in the number of cells secreting interleukin 6 (IL-6), IL-12 and IFN $\gamma$  within 10h and a rise in the number of IgM-secreting cells within 36h. In contrast, mammalian DNA and ODNs in which the critical CpG dinucleotide was eliminated by inversion or methylation did not stimulate cytokine or Ig secretion (Table 1; KLINMAN et al. 1996). Other investigators subsequently showed that IL-18 and tumor-necrosis-factor production were also induced by this motif (HALPERN et al. 1996). The immunostimulatory capacity of CpG motifs may be of use in a variety of therapeutic applications. This review will examine the contribution of CpG motifs as immune adjuvants when used with DNA vaccines, emphasizing the extensive studies conducted in mice.

**Table 1.** Immunostimulatory effect of CpG DNA. BALB/c spleen cells incubated with  $50\mu$ g/ml of heatdenatured *Escherichia coli* DNA or calf thymus DNA, or with  $1\mu$ M of stimulatory or control phosphorothioate oligodeoxynucleotides (ODNs)

	Fold increase in cytokine-secreting cell number			
	IL-6	IL-12	IFNγ	IgM
E. coli DNA	$3.2 \pm 0.2$	$3.8 \pm 0.4$	4.7 ± 2.3	3.9 ± 1.1
Calf-thymus DNA	$0.8 \pm 0.2$	$1.1 \pm 0.2$	$0.8 \pm 0.3$	$0.7 \pm 0.2$
CpG ODN	$5.5 \pm 1.1$	$8.3 \pm 1.7$	$4.7 \pm 1.1$	$4.2 \pm 1.6$
CpG ODN (methylated)	$0.9 \pm 0.2$	$1.2 \pm 0.3$	$0.8 \pm 0.2$	$1.1 \pm 0.2$
CpG ODN (DNase)	$1.3 \pm 0.2$	$0.8 \pm 0.2$	$1.1 \pm 0.2$	$0.9 \pm 0.2$
GpC ODN	$1.2 \pm 0.3$	$1.3 \pm 0.3$	$1.2 \pm 0.3$	$1.3 \pm 0.3$

The effect on cytokine production was determined after 10h by ELIspot assay (Mor et al. 1995; KLINMAN et al. 1996). Data represent the fold increase in the number of cytokine-secreting cells over background. Results represent the mean  $\pm$  SD of at least three independent experiments.

IFN, interferon; Ig, immunoglobulin; IL, interleukin.

#### 2 Immunostimulatory Properties of DNA Plasmids

#### 2.1 Cytokine Production

A considerable body of evidence indicates that plasmid DNA vaccines primarily elicit IFN $\gamma$ - and IgG2a-dominated immune responses when injected intramuscularly (Mor et al. 1995; SATO et al. 1996). We examined whether this reflected the activity of CpG motifs in the bacterial plasmid backbones of these vaccines.

Initially, the ability of DNA plasmids (with and without protein-encoding inserts) to induce cytokine production was studied in vitro. Two different plasmid constructs were examined: a vaccine encoding the circumsporozoite protein (CSP) of *P. yoelli* malaria known as 1012/PyCSP and manufactured by VICAL, Inc. (Mor et al. 1995) and a vaccine encoding the gp160 envelope protein of the human immunodeficiency virus 1 virus, known as pCMV160. When incubated in vitro with spleen cells from normal mice, both DNA vaccines induced a 5–8-fold increase in the number of cells secreting IgM, IL-6, IL-12 and IFN $\gamma$  (Table 2 and data not shown). Similar levels of immune stimulation were observed when the experiment was repeated using plasmid vectors from these vaccines (i.e., devoid of antigenencoding insert DNA). Of interest, each vector contained over a dozen immuno-stimulatory CpG motifs.

Further experiments were undertaken to determine whether the CpG motifs in the backbones of these DNA vaccines contributed to the observed cytokine production. To reduce the possibility that some form of bacterial contamination was responsible for the observed immunostimulation, we tested three different plasmids prepared in three different laboratories. Equivalent amounts of each plasmid elicited similar levels of cytokine production in vitro (data not shown). Treating these plasmids with DNAse or Sss I methylase (the latter selectively methylates the cytosine of CpG dinucleotides) uniformly eliminated cytokine production, indi-

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	Fold increase in cytokine secreting cell number			
	IL-6	IL-12	IFNγ	IgM
1012 vector	5.3 ± 1.1	7.6 ± 1.4	$5.0 \pm 1.2$	$4.6 \pm 0.6$
1012/PyCSP DNA vaccine	$5.9 \pm 1.3$	$8.1 \pm 1.9$	$4.6 \pm 1.5$	$5.2 \pm 1.6$
DNAse treated 1012/PyCSP	$0.9 \pm 0.2$	$1.0 \pm 0.1$	$1.1 \pm 0.2$	$1.2 \pm 0.2$
20% methylated 1012/PyCSP	$4.7 \pm 1.3$	$7.7 \pm 1.6$	$4.0~\pm~0.9$	ND
80% methylated 1012/PyCSP	$1.9 \pm 0.3$	$2.2 \pm 0.4$	$2.6 \pm 0.4$	ND
97% methylated 1012/PyCSP	$1.2 \pm 0.2$	$1.3 \pm 0.1$	$0.9~\pm~0.1$	$0.8~\pm~0.2$

Table 2. Immunostimulatory effect of CpG motifs in DNA plasmids

BALB/c spleen cells were incubated with  $50\mu g/ml$  of plasmid DNA. The effect on cytokine production was determined as described in the legend to Table 1.

CSP, cirsumsporozoite protein; IFN, interferon; Ig, immunoglobulin; IL, interleukin; ND, not determined.

Table 3. Addition of CpG motifs increases DNA-vaccine immunogenicity

	Fold increase		
	Ab titer	G2a:G1 ratio	IFNγ
50µg of 1012/PyCSP	680	3.6	4.6
4µg of 1012/PyCSP	64	1.4	1.2
4μg of 1012/PyCSP + 50μg of CpG ODN	410	2.7	3.9
4µg of 1012/PyCSP + 50µg of GpC ODN	58	1.3	1.7
4µg of 1012/PyCSP + 100µg of 1012 vector	440	2.9	5.2
2μg of 2534/PyCSP	260	3.4	3.5

BALB/c mice were primed and boosted with each plasmid. Serum immunoglobulin G (IgG) anti-CS.1 levels and frequency of P16-responsive interferon- $\gamma$  (IFN $\gamma$ )-secreting T cells was monitored 3 weeks post boost.

Ab, antibody; CSP, cirsumsporozoite protein; ODN, oligodeoxynucleotides.

cating that DNA, rather than some unknown contaminant, was responsible for the cytokine release (Table 2).

These findings led us to examine the nature of the cytokines produced by antigen-specific T cells derived from DNA-plasmid-vaccinated mice. We found that mice primed and boosted with DNA vaccines mount an IFN $\gamma$ -dominated Th1 immune response (MoR et al. 1995), an observation confirmed in other systems (SATO et al. 1996). We then took spleen cells from animals immunized with 1012/ PyCSP and re-stimulated them in vitro with P16, an immunodominant T-cell epitope present on the CSP. As seen in Table 3, spleen cells from vaccinated mice responded to antigen re-stimulation by a significant increase in IFN $\gamma$  (but not IL-4) production, consistent with a CpG-mediated effect.

#### 2.2 Antibody Production

The isotype of antigen-specific antibodies induced by plasmid DNA vaccination was also examined. Animals immunized with CS1 protein emulsified in complete

Freund's adjuvant primarily produce IgG1 anti-CSP antibodies (IgG1:IgG2a ratio of 5.6). In contrast, intramuscular injection of CSP-encoding DNA vaccines preferentially stimulated IgG2a-antibody production [IgG1:IgG2a ratios less than 0.7 for three different CSP-encoding plasmids (data not shown)]. Since IFN $\gamma$  promotes IgM-to-IgG2a isotype switching (SNAPPER and PAUL 1987; FINKLEMAN et al. 1988), these findings are consistent with CpG-motif-induced IFN $\gamma$  production contributing to the preferential production of IgG2a antibodies in immunized mice.

# **3** Co-Administration of CpG ODNs Improves the Immunogenicity of DNA Vaccines

If CpG motifs present in DNA plasmids contribute to vaccine immunogenicity, then co-administering additional motifs (in the form of CpG ODNs) would be expected to boost the immune response elicited by DNA vaccines. To examine this hypothesis, we primed and boosted mice with only 4µg of 1012/PyCSP (a suboptimal vaccine dose was utilized to maximize the likelihood of detecting a CpG effect) (KLINMAN et al. 1997). As seen in Table 3, this sub-optimal dose stimulated a detectable but reduced anti-CSP response. Co-administering 50µg of CpG ODNs significantly improved both IgG anti-CSP serum levels in vivo and P16-dependent IFNy production in vitro (Table 3). This observation has since been confirmed using CpG ODNs with other DNA vaccines (DAVIS et al. 1998). Control ODNs lacking immunostimulatory CpG motifs had no effect. Of interest, co-administering 100µg of vector alone (without CSP-encoding inserts) also improved the immune response elicited by 1012/PyCSP (KLINMAN et al. 1997). Presumably, CpG motifs present in the vector backbone acted as adjuvants in a fashion similar to the CpG ODNs. This observation raises the interesting possibility that higher doses of plasmid vaccine or co-administration of multiple DNA vaccines encoding different antigens may boost the immune response to each element due to the synergistic immunostimulatory effect of the additional CpG motifs.

In addition to their direct effects on cytokine and Ig-secreting lymphocytes, CpG ODNs also contribute to the development of an immune response by upregulating cell-surface expression of MHC class-II molecules (KRIEG et al. 1995; JAKOB et al. 1998; SPARWASSER et al. 1998). Originally demonstrated in B cells, this effect has also been observed in professional antigen-presenting cells. Indeed, CpG ODNs up-regulate the expression of a variety of co-stimulatory molecules in dendritic cells, including CD40 and CD86 (JAKOB et al. 1998; SPARWASSER et al. 1998), with the fraction of stimulated antigen-presenting cells APCs rising as a function of CpG-ODN concentration. Of particular importance, CpG-ODNmediated activation of these APCs increased their functional capacity, as reflected by an improved ability to stimulate alloreactive T cells (JAKOB et al. 1998).

## 4 Engineering CpG Motifs into the Plasmid Vector Improves DNA-Vaccine Immunogenicity

A more direct approach to assessing whether CpG motifs contribute to the immunogenicity of DNA vaccines involved engineering more CpG motifs into the DNA-vaccine backbone. This approach was pioneered by Sato et al., who substituted a CpG-containing *ampR* gene for a *kanR* selectable marker in a  $\beta$ -galactosi-dase-encoding plasmid. They found that the re-engineered plasmid elicited a higher IgG-antibody response, more CTLs and greater IFN $\gamma$  production than the original vector (SATO et al. 1996). Our lab utilized a series of vectors engineered by VICAL, Inc., who inserted multiple AACGTT motifs into the 1012 plasmid vector. Mice were primed and boosted with 50, 10 or 2µg of the 1012/PyCSP- or CpG-enriched 2534/PyCSP plasmid. As seen in Table 3, both vaccines stimulated strong antibody responses at optimal doses of 50µg/mouse. However, low doses of 2534/PyCSP elicited an IgG anti-CSP response significantly higher than that caused by a similar dose of 1012/PyCSP. These findings indicate that additional CpG motifs decreased the amount of vaccine required to induce antigen-specific antibody production.

In the same way, spleen cells from mice immunized with each vaccine were tested for reactivity to P16 stimulation in vitro. Optimal doses of both 1012/PyCSP and 2534/PyCSP generated P16-responsive IFN $\gamma$ -secreting cells. At a low dose, only cells from 2534/PyCSP-immunized mice were responsive to P16. As expected, no cytokine production resulted when cells from non-immunized mice (or mice treated with vector alone) were exposed to P16, nor did cells from immunized mice respond to stimulation by an unrelated peptide (data not shown).

It is noteworthy that CpG motifs were limited in their ability to augment antibody and cytokine production in vivo. For example, the immune response induced by the 2534/PyCSP plasmid was no greater than that of the 1020/PyCSP plasmid when both were administered at an optimal dose of 50µg. Similarly, increasing the concentration of CpG ODN to above 1µg/ml had little additional effect on the number of spleen cells activated to secrete cytokine in vitro (data not shown). Work by Krieg et al. indicates that adding too many CpG motifs to the plasmid backbone may actually reduce immunogenicity. In that work, introducing 16 additional CpG motifs into a plasmid improved the humoral immune response of the DNA vaccine, while introducing 50 such motifs reduced the response (KRIEG et al. 1998). These findings suggest that the maximal stimulatory effect of CpG motifs may require relatively low doses of DNA.

## 5 Deleting CpG Motifs from the Plasmid Vector Reduces DNA-Vaccine Immunogenicity

When the CpG motifs present in bacterial DNA or synthetic ODNs are methylated, their ability to induce cytokine production is significantly reduced (Tables 1, 2;

KRIEG et al. 1995; KLINMAN et al. 1996). To further examine the contribution of CpG motifs to the immunogenicity of DNA vaccines, we treated the 1012/PyCSP plasmid with Sss I CpG methylase. This reagent selectively methylates the cytosines of CpG dinucleotides (NUR et al. 1985). The duration of Sss-I treatment could be adjusted to achieve different levels of CpG methylation. Increasing the percent methylation of CpG dinucleotides reduced the ability of the 1012/PyCSP vaccine to activate cytokine-secreting cells in vitro. For example, 80% methylated vaccine induced 59–83% less IL-6, IL-12 and IFN $\gamma$  than did native 1012/PyCSP, whereas 97% methylated material induced virtually no cytokine production (Table 4 and data not shown). Similarly, highly methylated DNA vaccines are less immunogenic in vivo. As seen in Table 4, the IgG anti-PyCSP antibody response induced by 97% methylated material in vivo was only 3% of the control values.

The effect of CpG methylation on vaccine immunogenicity could have been due to either (i) the elimination of immunostimulatory CpG motifs or (ii) reduced activity of the plasmid's promoter region, resulting in decreased antigen expression (HuG et al. 1996). To evaluate the latter possibility, HeLa cells were transfected in vitro with native or methylated 1012/PyCSP. As seen in Table 5, cells transfected with 20, 80 and 97% methylated vaccine produced 21, 42 and 75% less CSP protein, respectively, than cells treated with the same amount of native plasmid. Thus, methylation reduced but did not eliminate promoter-region activity. Based on these in vitro results, we calculated that 40 $\mu$ g of 80% methylated 1012/PyCSP would induce the same level of in vivo antigen expression as 23 $\mu$ g of native vaccine. However, dose-titration experiments showed that BALB/c mice immunized and boosted with 23 $\mu$ g of 1012/PyCSP mounted an IgG anti-CSP antibody response nearly fivefold greater than that induced by 40 $\mu$ g of 80% methylated plasmid (Table 5). These findings support the conclusion that immunostimulatory CpG motifs play a role in vaccine immunogenicity.

To further establish that decreased promoter region function did not account for the full effect of methylation on plasmid immunogenicity, we co-administered the methylated 1012/PyCSP vaccine with 1012 vector alone or with CpG-containing ODNs. Although neither the vector nor the ODN encoded CSP protein, both were able to boost the CPS-specific immune response induced by native and

	Fold increase			
	Ab titer	G2a:G1 ratio	IFNγ	
Plasmid DNA vaccine	39	3.1	5.1	
DNAse treated DNA vaccine	2	ND	1.1	
80% methylated DNA vaccine	6	ND	1.8	
97% methylated DNA vaccine	1	ND	0.9	

Table 4. Elimination of CpG motifs reduces DNA-vaccine immunogenicity

BALB/c mice were immunized with  $50\mu g$  of plasmid DNA. The effect on serum immunoglobulin G anti-CS.1 antibody titer and on the number of T cells activated by in vitro stimulation by P16 is shown 3 weeks post immunization.

Ab, antibody; IFN, interferon; ND, not determined.

% Methylation of 1012/PyCSP	In vitro protein production (units)	Equivalent dose of DNA vaccine	Predicted IgG anti-CSP titer	Observed IgG anti-CSP titer	Difference
0% (Native)	100	40	3200	$3200 \pm 280$	0
20% Methylated	79	32	3000	$3000~\pm~340$	0
80% Methylated	58	23	2800	$610 \pm 110$	4.6
97% Methylated	25	10	2100	$160~\pm~60$	13.1

Table 5. Effect of methylation on promoter-region function and vaccine immunogenicity

An 80% confluent monolayer of HeLa cells was transfected overnight with  $l\mu g$  of plasmid DNA. Production of the encoded circumsporozoite protein (CSP) was monitored by incorporation of sulfur-35methionine followed by immunoprecipitation with anti-CSP antibodies (Abs). Relative protein content was determined by Phospho-Imager analysis of sodium dodecyl sulfate-polyacrylamide gels. The dose of native vaccine that was equivalent to  $40\mu g$  of methylated plasmid (in terms of protein production) was calculated. The titer of immunoglobulin G (IgG) anti-CSP Ab produced by BALB/c mice immunized and boosted with that dose of native vaccine was compared to the titer obtained when mice were immunized and boosted with 40 $\mu g$  of methylated vaccine (3 mice/group). The difference between the calculated and observed anti-CSP Ab response presumably reflects the contribution of CpG motifs to vaccine immunogenicity.

methylated 1012/PyCSP (KLINMAN et al. 1997). This finding demonstrates that partial methylation of the plasmid's promoter region does not prevent expression or recognition of the encoded CSP antigen and confirms that CpG motifs contribute to vaccine immunogenicity.

## 6 CpG ODNs Improve the Immunogenicity of Protein Antigens

We examined whether CpG ODNs could boost the immune response elicited by protein antigens. CpG- and non-CpG-containing ODNs were mixed with ovalbumin (OVA) and injected intraperitoneally into BALB/c mice. As seen in Table 6, soluble OVA alone stimulated a modest IgG anti-OVA response. Antibody production was increased approximately threefold when the OVA was co-administered with CpG (but not control) ODNs.

In these experiments, both the OVA and ODNs could freely diffuse from the site of injection. We reasoned that the adjuvant-like effect of CpG ODNs might be improved if their proximity to antigen were maintained. Towards that end, ODNs and OVA were emulsified in incomplete Freund's adjuvant (IFA). As expected, inclusion of CpG ODNs with OVA in IFA boosted IgG anti-OVA production by tenfold (Table 6). This effect was not observed with OVA in IFA alone (KLINMAN et al. 1998). Other laboratories have made similar observations, showing that co-delivery of CpG ODNs with antigens in a lipid/liposome formulation induced a substantial increase in the immune response (CHU et al. 1997; LIPFORD et al. 1997; WEINER et al. 1997).

To eliminate the requirement for IFA in this system, we sought to conjugate synthetic ODNs to these antigens. Unfortunately, efficiently linking DNA to pro-

	Fold increase			
	Ab titer	G2a:G1 ratio	IFNγ	
OVA alone	18	0.1	2.3	
OVA + CpG ODN	46	0.4	4.7	
OVA + GpC ODN	16	0.1	2.1	
OVA conjugated to GpC ODN	180	0.9	9.4	
(OVA + CpG ODN) emulsified in IFA	140	0.8	8.8	

Table 6. CpG-oligodeoxynucleotides (ODN) increase the immune response to a protein antigen

BALB/c mice were immunized and boosted with  $20\mu g$  of ovalbumin (OVA) plus  $50\mu g$  of ODN. The ODN were either mixed in the same syringe, conjugated to the OVA via biotin–avidin bridges, or emulsified in incomplete Freund's adjuvant (IFA). Three weeks after treatment, antigen-specific serum antibody (Ab) titers were determined by enzyme-linked immunosorbent assay. At the same time, the number of cells stimulated to secrete interferon  $\gamma$ (IFN $\gamma$ ) was monitored by ELIspot assay.

tein proved difficult. As an alternative, we biotinylated both CpG ODNs and OVA and used multivalent avidin cross-linkers to create stable complexes between these molecules. As seen in Table 6, OVA conjugated to CpG ODN was extremely immunogenic, with IgG anti-OVA antibody production being increased tenfold over OVA-avidin or OVA alone (KLINMAN et al. 1998). This effect was eliminated when the complexes were treated with DNAse, demonstrating that the effect was due to the CpG-containing DNA (data not shown).

Consistent with results involving DNA vaccines, CpG ODNs altered the isotypes of antibodies elicited by OVA immunization. OVA alone elicited a primarily IgG1 antibody response (IgG2a:IgG1 ratio of 0.1, Table 6). Addition of CpG ODNs significantly increased the production of IgG2a antibodies and increased the IgG2a:IgG1 ratio from 0.1 to 0.9. A similar shift in isotype profile has been observed using a variety of other antigens (including polysaccharide antigens) administered in the context of CpG ODNs by our laboratory and others (CHU et al. 1997; LIPFORD et al. 1997; WEINER et al. 1997).

Finally, we examined the effects of CpG ODNs on the activation of antigenspecific IFN $\gamma$ -producing cells. Co-administration of CpG ODNs with OVA (emulsified in IFA or complexed through biotin–avidin bridges) increased the number of spleen cells actively secreting IFN $\gamma$  in vivo by twofold when compared to mice immunized with OVA alone (KLINMAN et al. 1998). To establish that this effect was antigen specific, cells from immunized mice were stimulated in vitro with 20µg/ml of OVA. There was a significant dose-related increase in IFN $\gamma$  production accompanying the co-administration of CpG ODNs with OVA (Table 6). Coadministration of CpG ODNs with OVA emulsified in IFA had a similar effect on IFN $\gamma$  production (data not shown). In contrast, non-CpG ODNs did not elicit the development of cytokine-producing spleen cells.

Several other laboratories have also observed that CpG motifs have adjuvantlike properties. Consistent with our results, those labs found that adjuvant activity was improved by linking ODNs directly to antigens (either by covalently linking ODNs to proteins or by adding motifs to the backbones of DNA vaccines) (ROMAN et al. 1997). These results confirm the intuitive expectation that optimal stimulation occurs when antigen and adjuvant are presented to the immune system in close spatial and temporal proximity.

#### 7 Immunosuppressive Motifs in DNA Vaccines

Whereas CpG-containing bacterial DNA causes immune stimulation in vivo and in vitro, when mammalian DNA is co-administered with CpG DNA, it can block such activation in a dose-dependent manner. This suppression may account for the inability of mammalian DNA, which contains some CpG motifs (albeit at much lower frequency than bacterial DNA), to induce immune stimulation. Several labs have shown that a subset of non-stimulatory ODNs actually suppress the activation induced by CpG ODNs. Recent work by Krieg et al. indicates that eliminating suppressive motifs from the backbone of a DNA vaccine can improve its immunogenicity by threefold (KRIEG et al. 1998). These observations suggest that DNA can both stimulate and suppress the immune system.

## 8 Conclusion

In the studies reviewed above, several different immunostimulatory CpG ODNs were analyzed (AACGTTGAACGTTCGC, GCTAGACGTTAGCGT and TCAACGTT). All of these exhibited adjuvant-like properties, whether co-administered with DNA vaccines or proteins. It is unlikely that the effects observed were due to non-DNA contaminants, since (i) CpG-containing and control ODNs were prepared, processed and administered under identical conditions, yet only the former were active, (ii) all of the reagents used in these studies were carefully purified (endotoxin contamination was under 0.4IU/µg) and (iii) the activity of these CpG-containing ODNs was eliminated by methylation and by DNAse treatment.

Our findings, and those of several other investigators, indicate that the CpG motifs present in DNA vaccines serve an immunostimulatory function, triggering an innate immune response that promotes humoral and/or cell-mediated responses against environmental and plasmid-encoded antigens. This model is consistent with that of Fearon and Locksley, who postulated that an innate immune response could create an immune milieu conducive to the development of antigen-specific immunity (FEARON and LOCKSLEY 1996).

DNA vaccines have shown enormous promise in animals studies, yet their activity in humans has been less impressive. Thus, vaccine researchers continue to search for additional methods of improving DNA-vaccine efficacy. Evidence sug-

gests that the motifs that induce optimal immune stimulation in mice differ from those that are most active in humans. As our understanding of the processes underlying CpG-induced immune stimulation mature, we will be better able to harness their activity for the improvement of DNA vaccines.

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# The Role of Bacterial DNA in Autoantibody Induction

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## **1** Introduction

Antibodies to DNA (anti-DNA) are prototypic autoantibodies that occur prominently in systemic lupus erythematosus (SLE). These antibodies target sites on single- as well as double-stranded DNA and serve as markers of diagnostic and prognostic significance. In addition, anti-DNA play a direct role in disease pathogenesis, provoking renal injury through immune-complex deposition (PISETSKY 1992). Because of their close association with SLE, anti-DNA have been considered virtually synonymous with autoimmunity and key to understanding disease mechanisms (EILAT and ANDERSON 1994; RADIC and WEIGERT 1994).

While anti-DNA are a cardinal feature of SLE, immune reactivity to DNA extends beyond this disease. As shown by provocative studies from many laboratories, DNA, depending on base sequence, can induce responses in normal as well as aberrant immunity, with DNA from bacteria displaying potent immunostimulatory activities. These activities include induction of specific antibodies as well as the stimulation of cytokine production and B-cell activation (PISETSKY 1996a,

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1996b). In its effects on innate immunity, bacterial DNA resembles endotoxin, suggesting a role as a danger signal in eliciting host defense (PISETSKY 1997).

Since the immunostimulatory properties of bacterial DNA are discussed in detail elsewhere in this volume, this chapter will focus on their relevance to the induction of anti-DNA autoantibodies. As this account will indicate, bacterial DNA displays unique immunological properties that could allow it to drive both anti-self as well as anti-foreign responses to DNA antigens. These properties reflect a variety of structural elements and suggest novel mechanisms accounting for the serological features of SLE.

#### 2 DNA as an Autoantigen

Until recently, analysis of the immunological properties of DNA was guided by two main ideas. The first idea is that the relevant immunological determinants of DNA are conserved determinants present on all DNA. Indeed, most studies on lupus serology have dichotomized DNA antigens into either single stranded (ss) or double stranded (ds) forms, with antibodies to dsDNA the most diagnostic of disease. In the case of dsDNA, the epitopes recognized by autoantibodies have been considered charge arrays along the phosphodiester backbone. While this binding is dependent on charge–charge interactions, it requires the spatial orientation of the classic double helical structure, or B-DNA (STOLLAR 1992).

In contrast to dsDNA, ssDNA is structurally flexible, because the rigidity conferred by base pairing of the two strands is lost. Nevertheless, autoantibody binding to ssDNA also appears to be a consequence of charge, albeit in a less ordered format. Many antibodies to DNA crossreact with ss- and dsDNA, suggesting a common determinant exposed on both DNA forms. While interactions with phosphate groups in the backbone provide the preponderance of bond energy for antibody binding, interactions with bases themselves may also contribute.

From the molecular analysis of SLE antibody binding, DNA emerges as a relatively uniform antigen, with a charged backbone providing the major site of immune recognition. As such, anti-DNA autoantibodies can bind foreign as well as self DNA as long as the backbone structure is present. This notion is implicit in the technology of anti-DNA assays. Assays for anti-DNA have used many different DNA preparations, essentially interchangeably, with assays based on mammalian and bacterial DNA (including supercoiled plasmids). These assays all perform well clinically, suggesting that the major autoantibody specificities are directed to conserved determinants rather than non-conserved sites exclusive to self DNA (PISETSKY 1998).

The second idea underlying much research in SLE concerns the immunogenic potential of DNA. Following the recognition of DNA's role as a major autoantigen, investigators attempted to replicate lupus in normal animals by immunizing them with DNA. These studies showed that naturally occurring DNA is a poor immunogen. When administered alone, DNA fails to stimulate antibody production although, when complexed with a protein carrier and presented in adjuvant, it can elicit a limited response to ssDNA. Immunization experiments, with some exceptions (see below), failed to induce antibodies to dsDNA, the serological hallmark of lupus (MADAIO et al. 1984).

In contrast to natural DNA, certain synthetic DNA and chemically modified DNA are immunogenic. For example, normal mice can generate a robust antibody response to Z-DNA. Z-DNA is a helical structure formed by tracts of alternating purine and pyrimidine bases; the presence of certain divalent cations promotes transition to this structure. In Z-DNA, the helix displays a zig-zag structure with a left-handed orientation, in contrast to the right-handed orientation of conventional B-DNA. When used as an immunogen, Z-DNA induces antibodies that are specific for this helical form and do not crossreact with B-DNA (LAFER et al. 1981; STOLLAR 1994). Other unusual DNA structures (DNA cruciforms) can also elicit responses in normal animals, although the induced antibodies are specific for the immunizing DNA and do not crossreact with natural DNA (FRAPPIER et al. 1989).

The immunization experiments have suggested that natural dsDNA is immunologically inert and unable to induce a significant antibody response. This paltry activity could reflect two phenomena. The first concerns tolerance and suggests that, because of anergy or clonal deletion, B cells that recognize dsDNA are absent from the normal B-cell repertoire. As such, the normal animal would be unable to produce anti-DNA even when stimulated with DNA in the context of a protein carrier providing T-cell help. The failure to produce autoantibodies to DNA by immunization contrasts with the generation, in normal animals, of autoantibody responses to protein antigens, such as collagen or myelin basic protein. With protein autoantigens, immunization can induce both autoimmune responses and clinical manifestations of disease. This result suggests that tolerance for nucleic-acid antigens may be more complete than for proteins, perhaps reflecting their structure or in vivo metabolism.

As noted above, while natural DNA is a weak immunogen, unusual DNA structures can elicit appreciable responses under conditions in which B-DNA is inactive. The difference in response patterns among nucleic acids may reflect the concentration of unusual DNA structures and their ability to induce tolerance. Z-DNA exists at most transiently in cells and is present in only low concentrations because of sequence requirements. As a result, the contact of Z-DNA with immune cells may be ephemeral or below a threshold necessary to establish tolerance. When presented in a stabilized form in high concentration, Z-DNA may, therefore, induce responses.

While tolerance mechanisms could explain the poor immunogenicity of natural DNA, other properties of DNA could influence the induction of responses. DNA is readily digestible in vivo because of nucleases in serum and cells. Furthermore, DNA in the circulation is rapidly cleared by cells of the reticulo-endothelial system, suggesting that DNA used for immunization may be short lived (EMLEN and MANNIK 1984; YOSHIDA et al. 1996). Indeed, there is evidence that the immunogenicity of DNA is related to its stability, with immunogenicity enhanced by

backbone modifications that protect against nuclease digestion. Among these modifications, compounds with phosphorothioate backbones are more effective immunogens than phosphodiesters; phosphorothioates are nuclease resistant (BRAUN and LEE 1988).

In evaluating the immunogenicity of DNA, the protein carrier has received little attention as a factor influencing responsiveness. Compounds such as methylated bovine serum albumin (mBSA) and methylated gammaglobulin have been commonly used in these experiments, because they can complex DNA by charge. Studies of a synthetic protein called Fus-1 indicate that the carrier may affect the immunogenicity of natural DNA. Fus-1 is 27 amino acids long and bears a sequence corresponding to a protein from *Trypanosoma cruzi*. Complexes of Fus-1 with sheared mammalian DNA can induce antibodies to dsDNA in normal mice. Since these induced antibodies are similar to lupus anti-DNA in their binding properties, these results suggest that normal mice may produce anti-DNA auto-antibodies under certain circumstances (DESAI et al. 1993; KRISHNAN and MARION 1993).

The impact of Fus-1 on responsiveness to DNA is not well understood, though it will be important to later considerations on the response to bacterial DNA (see below). Assuming that normal mice have at least some capacity for anti-DNA production, the ability of Fus-1 complexes to induce anti-dsDNA could reflect at least three factors: formation of complexes that protect dsDNA from degradation; induction of more vigorous T-cell help; and more effective presentation of a dsDNA epitope for stimulation of B cells. Even with a carrier like Fus-1, however, the immunogenicity of mammalian DNA appears limited and dependent on physical-chemical form.

## **3** Antibody Response to Bacterial DNA

The categorization of DNA as ss or ds is an obvious simplification that ignores the structural richness imparted by the base sequence. Theoretically, such sequences could serve as epitopes for antibody binding just as amino acid sequences serve as epitopes for antibody binding to protein antigens. In our laboratory, we have been interested in the possibility that anti-DNA antibodies can bind non-conserved sequences and conserved backbone conformations. Our efforts to find antibodies to DNA sequences provided some of the initial evidence for the immunological heterogeneity of DNA and the unique properties of bacterial DNA antigens.

To identify antibodies directed to base sequences as opposed to backbone, we tested as antigens a series of mammalian and bacterial DNA. These DNA differed in base composition and presumably presented a range of sequences to allow detection of antibodies to sequence motifs. While many different DNA preparations have been used as antigens in clinical assays, there have been few studies directly comparing the antigenicities of natural DNA varying in species origin (STOLLAR et al. 1962). In these experiments, with ssDNA as an antigen, sera from SLE patients showed similar levels of binding to the various DNA preparations in an enzyme-limited immunosorbent assay (ELISA) (KAROUNOS et al. 1988). These findings indicate predominant reactivity to a conserved determinant widely expressed among natural DNA. This determinant could involve both base sequences and backbone; the particular sequences, however, would have to be ubiquitous among the DNA.

Although anti-DNA responses have been widely viewed as exclusive to SLE, we nevertheless tested sera of normal human subjects (NHS) using the same antigen preparations. Results of these studies were entirely unexpected and refute the notion that anti-DNA are exclusive to SLE (KAROUNOS et al., 1988). As these data indicated, the reactivity of NHS sera with DNA is not uniform (Fig. 1). Thus, NHS sera, while non-reactive to mammalian DNA, bind significantly to DNA from certain bacterial DNA, including *Micrococcus lysodeikticus* (MC) and *Staphylococcus epidermidis* (SE). The antigenicity of these preparations is fully sensitive to digestion with DNAse, indicating that DNA is the relevant antigen. In NHS sera, as in sera from SLE patients, the anti-DNA antibodies display the immunoglobulin G (IgG) isotype and differ from natural autoantibodies, which are IgM. These results demonstrate clearly that anti-DNA responses are a feature of both normal and aberrant immunity.

As shown by subsequent experiments, anti-DNA antibodies in NHS differ from SLE antibodies in both specificity and immunochemical properties. Whereas SLE anti-DNA bind DNA independent of species origin, NHS anti-DNA bind



Fig. 1. Anti-DNA antibodies in sera of normal human subjects. Sera of 12 normal human subjects were screened by enzyme-linked immunosorbent assay for binding to single-stranded DNA from calf thymus (CT), human placenta (HP), Micrococcus lysodeikticus (MC), Pseudomonas aeruginosa (PA), Staphylococcus epidermidis (SE), and Escherichia coli (EC). Results are presented as mean  $\pm$  SD. Reproduced with permission from Journal of Immunology

selectively to bacterial DNA and recognize epitopes with limited distribution among bacterial species. This selectivity suggests predominant reactivity with bases as opposed to the phosphodiester backbone. Indeed, binding to NHS antibodies to bacterial DNA is much less sensitive to salt concentration than binding to SLE anti-DNA, consistent with less dependence on ionic interactions (PISETSKY and ROBERTSON 1992; WU et al. 1997).

In addition to epitope specificity, antibodies to DNA in NHS and SLE sera differ in isotype. Whereas SLE anti-DNA are primarily IgG1 and IgG3, NHS anti-DNA display a preponderance of the IgG2 isotype (ROBERTSON et al. 1992; WU et al. 1997). In normal immunity, the IgG2 isotype characterizes the response to bacterial carbohydrate antigens. These findings suggest that bacterial DNA may resemble polysaccharides in immune activity, displaying a T-independent antibody induction in contrast to the T-dependent induction of anti-DNA auto-antibodies.

While not predicted by prevailing ideas on SLE, immune responses to bacterial DNA can be readily explained using concepts derived from the studies of protein antigens and unusual DNA structures. Thus, bacterial DNA, like foreign proteins, displays sequences that are species-specific and absent from the host. These sequences cannot establish tolerance and can, therefore, induce responses when presented in an immunogenic form during either colonization or infection. Since NHS sera contain antibodies directed to DNA from only certain bacteria, other factors must determine the response; at present, the rules for antigenicity are not clear. Interestingly, among bacterial DNA, *Escherichia coli* (EC) DNA does not show appreciable antibody levels in NHS sera, although this organism is ubiquitous in the gut ( $W_U$  et al. 1997).

In addition to DNA from bacteria, DNA from certain viruses show reactivity with normal sera (REKVIG et al. 1992). For both viral and bacterial DNA, the antibodies in NHS sera appear to be highly specific for each DNA. Together, these findings indicate that antibody responses to foreign DNA are a common feature of normal immunity, with previous failure to recognize these responses reflecting the range of DNA antigens tested. While certain bacterial or viral DNA show similar levels of antibodies in normal individuals and lupus patients, responses to B-DNA are nevertheless specific for lupus and retain their diagnostic utility.

## 4 Induction of Anti-DNA Antibodies by Immunization with Bacterial DNA

Although studies on normal humans strongly suggest that bacterial DNA is immunogenic, more direct proof for this idea was sought from animal experiments. Normal mice were, therefore, immunized with preparations of EC dsDNA complexed with mBSA in Freund's adjuvant, and their sera was assayed for antibodies both to the immunizing bacterial DNA and mammalian calf thymus (CT) DNA. This format was similar in design to previous studies indicating that mammalian DNA is immunologically inactive.

As these studies indicated, EC dsDNA can induce responses in normal mice under conditions in which mammalian DNA is without effect (GILKESON et al. 1989a). The induced antibodies, however, are highly specific for bacterial DNA and do not crossreact with mammalian DNA. In this respect, the induced antibodies resemble antibodies in NHS sera that target non-conserved sites on bacterial DNA; they differ from SLE anti-DNA that target conserved sites present on both bacterial DNA and mammalian DNA. The epitopes recognized by the induced antibodies are most likely sequences, although charges from the backbone may contribute to binding energy.

Immunization with bacterial ssDNA produces a somewhat different response pattern than dsDNA (GILKESON et al. 1989b). Whereas, in normal mice, bacterial dsDNA elicits antibodies that bind exclusively to the immunizing antigen, bacterial ssDNA stimulates antibodies that crossreact with mammalian ssDNA. The induction of these crossreactive antibodies may suggest that-B cell tolerance for ss- and dsDNA differ, with normal individuals retaining a greater capacity for producing autoantibodies to the ssDNA form. It is important to note, however, that the levels of antibodies induced by bacterial ssDNA are far greater than those induced by mammalian ssDNA, indicating that bacterial DNA enhances responsiveness even to conserved determinants.

In view of the immunological activities of bacterial DNA, the immunogenicity of this nucleic acid appears related to its intrinsic immunostimulatory activity and its content of antigenic sequences that are exclusive to bacterial DNA. As discussed elsewhere in this volume, bacterial DNA is mitogenic for B cells and can stimulate a variety of cytokines, including interleukin 12 (IL-12), tumor necrosis factor  $\alpha$ , IL-6, interferon  $\alpha/\beta$  (IFN- $\alpha/\beta$ ) and IFN- $\gamma$ . These activities result from short six base motifs called CpG motifs or ISS (immunostimulatory sequences) that have the general structure of two 5' purines, an unmethylated CpG motif and two 3' pyrimidines. These sequences occur much more commonly in bacterial than in mammalian DNA because of CpG suppression in mammalian DNA and the frequent methylation of cytosine in this position in mammalian DNA (YAMAMOTO et al. 1992a; KRIEG et al. 1995). The content of ISS differs among bacterial DNA, suggesting that bacterial DNA vary in immunogenicity (YAMAMOTO et al. 1992b).

Taken together with the serological studies, these considerations suggest that DNA displays three immunologically relevant determinants: conformational determinants on ds- or ssDNA that function as B-cell epitopes in SLE; sequential determinants on bacterial ss- or dsDNA that function as B-cell epitopes in normal immunity; and ISS on bacterial DNA that serve as internal adjuvants. While an ISS could theoretically be a target of antibody reactivity, the reactivity of NHS suggests that these sequences are not commonly recognized. Instead NHS antibodies appear to target sites that are species-specific rather than targeting an ISS, which would be expected to have a broader distribution.

## 5 The Role of Bacterial DNA in Driving Autoantibody Production

The immune properties of bacterial DNA, while unanticipated by previous research, can nevertheless explain a conundrum that has long influenced SLE research: the mechanism by which a putatively non-immunogenic molecule can drive autoantibody production. As shown clearly in studies on both mouse and man, DNA is neither uniform nor inert. Rather, bacterial DNA, unlike mammalian DNA, is immunologically potent and can drive production of antibodies to sequential determinants. The intriguing question is whether it can also drive the production of autoantibodies to conformational determinants. If so, this finding would suggest that bacterial DNA could either initiate an autoantibody response to DNA or boost its levels during active SLE.

To explore the capacity of bacterial DNA to drive autoantibody production, we used as a model immunization of NZB/NZW mice. These animals develop a lupus-like illness characterized by high levels of anti-DNA production in association with glomerulonephritis. Using protocols developed from normal mice, we immunized female NZB/NZW mice with bacterial or mammalian dsDNA prior to the onset of spontaneous disease. Sera were assayed by ELISA for antibodies to both bacterial and mammalian dsDNA (GILKESON et al. 1995).

The results of these studies provide important insight into the potential of bacterial DNA to promote SLE antibody production. In the preautoimmune mice, EC dsDNA elicited significant production of anti-dsDNA antibodies. Unlike antibodies in normal mice, however, the induced antibodies had autoantibody activity and bound both mammalian and bacterial dsDNA. This crossreactivity is reminiscent of the SLE response and signifies binding to a conserved determinant. In contrast, immunization of NZB/NZW mice with mammalian dsDNA was without effect, suggesting that unique structural features of bacterial DNA are required for autoantibody production (GILKESON et al. 1995). The failure of mammalian DNA to induce anti-DNA in NZB/NZW mice is surprising, because these mice are destined to produce such antibodies and would be expected to have appropriate B-cell precursors.

Two conclusions from these studies are relevant for models for the mechanisms of anti-DNA production in SLE: (1) the induction of crossreactive antibodies in autoimmune mice occurs under conditions in which normal mice generate a response exclusive for foreign DNA; and (2) bacterial DNA can stimulate anti-DNA autoantibody responses under conditions in which mammalian DNA is inactive. As suggested above, the production of autoantibodies to DNA may reflect abnormalities in the B-cell repertoire in lupus mice. Because of tolerance defects in autoimmunity, this repertoire may retain precursors that can bind dsDNA and be stimulated to autoantibody production under appropriate immunization conditions. In this view, the difference between normal and autoimmune mice would reside in the B-cell-precursor population with the autoimmune repertoire skewed to reactivity to conserved conformational determinants (as opposed to non-conserved sequential determinants) (GILKESON et al. 1993). The failure of mammalian DNA to induce these precursors in NZB/NZW mice is more difficult to explain. Since the immunizations involve a protein carrier and adjuvant, induction of adequate T cells would be expected. In this situation, either mammalian or bacterial DNA should be able to stimulate the B-cell response, because both DNA sources would bear the crossreactive backbone determinant. If, however, the generation of T-cell help in NZB/NZW mice was limiting, the adjuvant properties of the bacterial DNA could be a critical ingredient allowing induction of adequate levels of help. It is also possible that bacterial DNA show less degradation in vivo than mammalian DNA, providing a higher antigen content.

The studies using Fus-1 as a carrier may be relevant to these considerations. As described previously, Fus-1 as a carrier may allow induction of responses under conditions in which mBSA fails. Fus-1, because of its sequence, may induce a more robust T-cell response and allow emergence of an autoantibody response from rare precursors. Similarly, bacterial DNA may facilitate these responses, because its adjuvant activity can enhance T-cell responses to mBSA and thereby allow adequate help for anti-DNA with that carrier. Whether the induction of T-cell help for anti-DNA results from properties of the carrier or adjuvant properties of the DNA, autoantibody production appears to occur much more readily in mice from an autoimmune background than in those of a normal strain.

#### 6 Antibody Responses to Bacterial DNA in SLE

Theoretically, an individual with lupus can produce two types of anti-DNA: anti-DNA autoantibodies that bind both bacterial and mammalian DNA and anti-DNA antibodies that bind exclusively to bacterial DNA. The balance between these responses would be a function of the composition of both the B-cell precursor population and the role of foreign vs self DNA in driving these responses. Determination of the nature of this balance should, therefore, help distinguish the role of different DNA antigens in disease induction.

To assess the expression of different types of anti-DNA in SLE, we performed absorption experiments using DNA affinity columns to deplete anti-DNA populations (PISETSKY and DRAYTON 1997). SLE sera with high titers of anti-DNA autoantibodies were absorbed with CT DNA and then tested for binding to both CT and MC DNA. Control NHS sera were similarly absorbed. The results of these experiments were clear-cut. Absorptions that removed antibodies to CT DNA from SLE sera essentially completely removed antibodies to MC DNA; this absorption had no effect on the binding of NHS sera to MC DNA. Furthermore, direct comparison of absorbed SLE and NHS sera showed that SLE sera have markedly lower levels of antibodies specific for bacterial DNA than the levels found in NHS sera (Fig. 2).

These observations document a reduction in the specific response to bacterial DNA in NHS and raise two mechanistic possibilities. The first is that a shift in



**Fig. 2.** Specificity analysis of antibodies in sera of normal human subjects (NHS) and patients with systemic lupus erythematosus (SLE). A panel of 20 NHS and 16 SLE sera were absorbed with either control cellulose (*open bar*) or calf thymus (CT) single-stranded-DNA cellulose (*closed bar*) and then tested for binding to either *Micrococcus lysodeikticus* or CT DNA. Asterisks indicates statistically significant differences (P < 0.001). Reproduced with permission from Proceedings of the Association of American Physicians

antibody specificity in SLE leads to the expression of antibodies to conserved rather than non-conserved determinants whether the inducing antigen is foreign or self. This shift could result from distortion in the B-cell repertoire because of impaired tolerance. The second interpretation is that SLE patients have an antibody-deficient state that prevents them from mounting a specific antibody response to bacterial DNA. In the absence of such specific antibodies, bacterial DNA, because of its content of ISS, could stimulate the immune system sufficiently to cause autoantibody production.

At present, there are insufficient data to distinguish these possibilities, although reference to other autoantibody responses suggests that an aberrant pattern of epitope recognition may mark the immune systems of SLE patients. In the response to protein autoantigens, SLE sera tend to react to conformational epitopes rather than sequential determinants of the kind recognized by antibodies induced by immunization in animals (TAN 1989; YAMANAKA et al. 1989). The lack of binding to peptide antigens is striking and is contrary to predictions for antigenicity based on experimental immunization models. A failure to bind peptide determinants may reflect the process of antigen presentation, although skewing in the B-cell repertoire could also affect the epitopes recognized.

While bacterial DNA has the potential to drive anti-DNA production, its role in the setting of human disease is unknown at present. In other studies on anti-DNA production, chromatin has been implicated as the driving antigen, because antibodies to DNA frequently coexist with antibodies to histones and other chromatin components. Furthermore, in autoimmune mice, antibodies to chromatin may precede the expression of antibodies to isolated DNA, suggesting that antibodies to DNA may emerge from a response directed to a larger complex. The ability of nucleosomes to stimulate T cells is further evidence that chromatin functions as a key autoantigen to drive SLE autoantibody production (BURLIN-GAME et al. 1993; MOHAN et al. 1993; BURLINGAME et al. 1994).

Even if chromatin is a major autoantigen, bacterial DNA may nevertheless have an important role in stimulating anti-DNA in SLE. As shown in longitudinal studies, levels of anti-DNA fluctuate dramatically over time and show patterns of temporal expression that differ from antibodies to other nuclear antigens, such as Sm (McCARTY et al. 1982). If self-antigens were the exclusive driving antigens, more coordinate appearance of autoantibodies would be expected. However, a foreign antigen that periodically impinges on the immune system could lead to the flares of serological activity that characterize the course of SLE. We are currently performing experiments to address these possibilities.

#### 7 Summary

Bacterial DNA has potent immunological properties that can stimulate the immune system in SLE in both specific and non-specific ways. As such, this molecule may play an important role in disease pathogenesis, because it can exert immunomodulatory activity and function as a molecular mimic. Future studies will hopefully both determine the role of foreign nucleic acids in the induction of autoantibodies and lead to strategies for their elimination.

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# CpG DNA in Cancer Immunotherapy

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## **1** Introduction

Anecdotal reports of tumor regression following systemic bacterial infection have been observed for centuries. The first reported attempt to explore this effect in a systematic fashion took place in the 1890s, when Dr. William Coley, a New York surgeon, performed a series of studies evaluating anti-tumor therapy with bacteria and bacterial products. Dr. Coley's original attempt to use bacteria as an antitumor agent involved the use of live cultures of streptococci injected directly into tumor masses (COLEY 1893). This resulted in tumor regression in some cases but proved to be toxic, with the first patient almost dying of erysipelas. Subsequent studies by Coley involved a mixture of heat-killed streptococci and serratia (then known as *Bacillus prodigiosus*) (COLEY 1894). This preparation was still quite toxic, but did result in some anti-tumor responses. It is this preparation that is now referred to as Coley's toxin. Much of the antitumor activity of Coley's toxin is attributed to endotoxin (WIEMANN and STARNES 1994). However, it is curious to note that Coley's original success was with an organism that does not produce

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endotoxin. Thus, additional bacterial components, such as bacterial DNA, may well have played a role in the observed responses. Indeed, recent reports have confirmed that cancer can regress following bacterial infection (MORI et al. 1997).

Despite the fact that Coley performed his intriguing experiments over a century ago, immunotherapy still has little impact on the management of patients with cancer. The recent reawakening of interest in cancer immunotherapy results from a more complete understanding of the immune response in general and the interaction between the immune system and tumors in particular. We are currently making rapid progress in these areas. Nevertheless, we are still in the infancy of scientific cancer immunotherapy and are only now beginning to understand the true promise of specific agents, such as immunostimulatory DNA, as potentially useful agents for the treatment of cancer.

The complex immunologic effects of immunostimulatory DNA in general and CpG DNA in particular are outlined elsewhere in this volume and will not be discussed here. Instead, we will focus on a number of promising approaches to cancer immunotherapy and how CpG DNA might play a role in these approaches. Some convincing data is reviewed below; however, evaluation of CpG DNA as a cancer therapy is only beginning, and much of what follows is based on circumstantial evidence with little or no direct experimental data. Nevertheless, our basic understanding of tumor immunology, combined with an increasing knowledge of the immunologic effects of CpG DNA, is enough to warrant discussion of CpG DNA as a cancer therapy using these approaches. Approaches discussed below include:

- 1. CpG DNA as a single agent for the treatment of cancer
- 2. CpG DNA for the enhancement of the efficacy of antibody therapy
- 3. CpG DNA for the enhancement of the efficacy of immunization with tumorassociated antigens
- 4. CpG DNA for the enhancement of the efficacy of cellular immunotherapy
- 5. CpG DNA for the enhancement of the anti-tumor response following DNA immunization

## 2 CpG DNA as a Single Agent for the Treatment of Cancer

As outlined above, much of the effect of Coley's toxin has been attributed to endotoxin, which is known to induce production of a number of cytokines that have anti-tumor effects in vitro and in animal models. This has led, over the past several years, to the clinical evaluation of a number of recombinant cytokines as cancer treatments (HEATON and GRIMM 1993). Some recombinant cytokines have been shown to have clear, although not dramatic, anti-tumor activity in clinical trials (ROSENBERG et al. 1985; BUKOWSKI 1997; KANTARJIAN et al. 1998). While such studies demonstrate that cancer immunotherapy can be effective, overall responses to such agents have been disappointing given the expectations in the 1960s and 1970s, when cytokines, such as the interferons (IFNs), were first being discovered and proposed as cancer cures.

In retrospect, the expectation that single cytokines might result in significant anti-tumor responses was quite naive. The immune response normally involves the integrated production of a variety of cytokines, which work in concert both locally and systemically. Agents that are able to orchestrate the production of cytokines by the host both temporally and spatially may be more effective at inducing an antitumor response compared with passive administration of recombinant cytokines. CpG is capable of inducing production of a number of cytokines that have been shown to have anti-tumor activity, including tumor necrosis factor  $\alpha$  (TNF $\alpha$ ), interleukin 12 (IL-12) and IFN- $\gamma$  (BALLAS et al. 1996; COWDERY et al. 1996; LIPFORD et al. 1997). In addition, CpG DNA has direct effects on immune cell subpopulations that play an important role in anti-tumor immunity such as B cells (KRIEG et al. 1995), natural killer (NK) cells (BALLAS et al. 1996), monocytes and macrophages (CHACE et al. 1997; SPARWASSER et al. 1997), dendritic cells (JAKOB et al. 1998; LIU et al. 1998; SPARWASSER et al. 1998), and possibly T cells (LIPFORD et al. 1997). The complex cascade of effects on the immune system induced by CpG DNA would seem to point towards the ability of CpG DNA to enhance nonspecific anti-tumor immunity.

Limited data suggest that CpG DNA by itself can have anti-tumor effects. Smith et al. used a murine model of lymphoma to evaluate the anti-tumor effects of an antisense phosphorothioate DNA designed to block the c-myc oncogene. While their antisense DNA inhibited tumor growth, so did a control DNA that did not have antisense activity but was immunostimulatory (SMITH and WICKSTROM 1998).

Ballas has evaluated the effect of CpG DNA on the growth of B16 melanoma cells in syngeneic mice (manuscript in preparation). All mice treated with non-CpG DNA died within 24 days, while those treated with CpG DNA had 40% long-term survival. Surviving mice re-challenged with melanoma died similar to control, suggesting that there is no memory. Studies done in severe combined immunode-ficiency mice gave identical results to immunocompetent mice. Taken together, these results suggest that innate immunity (probably NK cells) and not T cells are responsible for the therapeutic effect.

While toxicity from CpG DNA in animal models has been limited, toxicity from anti-cancer agents that work in a non-specific manner often limits their clinical efficacy. Clinical trials will be required before we can know whether CpG DNA has promise as a single agent for the treatment of cancer.

# **3** CpG DNA for the Enhancement of the Efficacy of Antibody Therapy

Kohler and Milstein's seminal technology provided monoclonal murine antibodies for both preclinical and clinical evaluation over 20 years ago (KOHLER and MILSTEIN 1975). Such murine antibodies were effective in migrating to their designed target. However, unmodified murine antibodies were limited in their ability to manipulate the human immune system, had relatively short half-lives and proved to be immunogenic in themselves (LINK and WEINER 1998). Since that time, protein chemists and molecular biologists have used a variety of approaches to "humanize" antibodies and so overcome some of the limitations of the original murine antibodies. Studies of antibodies with largely human frameworks demonstrate they are more effective at activating human effector mechanisms, have longer half-lives and are less immunogenic than their murine counterparts. Recent clinical trials with such agents have demonstrated monoclonal antibodies (mAbs) can have significant anti-tumor effects in patients (MALONEY et al. 1997; PEGRAM et al. 1998), although there continues to be significant room for improvement. The mechanisms responsible for the clinical response to antibody therapy are not yet clear, although antibody-dependent cellular cytotoxicity (ADCC) mediated by NK cells and monocytes/macrophages likely plays a large role.

A number of investigators have explored approaches to enhancing the activation of NK cells with cytokines to increase ADCC in vitro and the efficacy of mAb therapy in vivo. For example, IL2 has been shown to increase the efficacy of mAbs in a mouse lymphoma model (BERINSTEIN and LEVY 1987). This approach is now being evaluated in clinical trials. CpG DNA activates both NK cells and macrophages and, therefore, could enhance the efficacy of antibody therapy by increasing effector cell killing of antibody-coated tumor cells. We found that, in vitro, CpG-DNA-activated splenocytes induced tumor cell lysis more effectively than did unactivated splenocytes. Such effector cells were also superior to unactivated splenocytes or cells activated with a control methylated DNA at inducing ADCC. In vivo, CpG DNA alone had no effect on survival of mice inoculated with lymphoma. However, a single injection of CpG DNA enhanced the anti-tumor response to anti-tumor mAb therapy. These anti-tumor effects were less pronounced when treatment consisted of an identical DNA containing methylated cytosines. A single dose of CpG DNA was more effective than multiple doses of IL-2 at inhibiting tumor growth when used along with anti-tumor mAbs (WOOLD-RIDGE et al. 1997). We have found similar results in vitro using human effector cells and human tumor cells (manuscript in preparation). Thus, use of CpG DNA to enhance the efficacy of antibody therapy remains a real possibility. A clinical trial designed to assess this possibility is currently being planned.

## 4 CpG DNA for the Enhancement of the Efficacy of Immunization with Tumor-Associated Antigen

Vaccinations for a variety of deadly infectious diseases, such as small pox and diphtheria, have had a major impact on world-wide public health over the past century. However, cancer has proven to be less responsive to vaccination. Recent

advances in the field of immunology are allowing us to understand why development of a successful vaccination strategy has been so difficult for cancer. In broad and simplified terms, the major problems include: finding an antigen that is adequately tumor specific yet expressed by all the malignant cells; breaking immune tolerance against that antigen; and inducing an immune response strong enough to induce tumor cell destruction (SPOONER et al. 1995; GILBOA 1996; HELLSTROM et al. 1997).

A number of tumor-associated antigens have been identified that appear to be acceptable, although not ideal, targets for cancer-immunization strategies. Examples include the idiotype (Id) expressed by B-cell malignancies (Hsu et al. 1997) and a number of proteins associated with melanoma (MAEURER et al. 1996). Induction of an immune response against antigens that have been used for years as tumor markers, such as carcinoembryonic antigen (PERVIN et al. 1997) and prostate-specific antigen (CORREALE et al. 1997; HODGE et al. 1995), is also showing early promise.

The question of whether a humoral or a cellular anti-tumor response is most desirable is still debated and likely depends on the tumor type and the antigen being targeted. Many cancer-vaccine studies use antibody production as an indication of a successful immune response. This is based in large part on the relative ease of quantitating the antibody response compared to the cellular response. There is also evidence that passive antibody therapy can have anti-tumor activity and that the ability to induce a humoral response following immunization in some cancervaccine trials correlates with clinical outcome (LIVINGSTON 1998). Obviously, an antibody response can only be effective if the target antigen is a surface molecule. In addition, numerous solid-tumor animal models demonstrate that a cellular immune response is required if a significant anti-tumor response is desired. Overall, there is now general agreement that induction of a potent anti-tumor cellular response is required if cancer immunotherapy is to be effective against most tumor types.

Unfortunately, the immune adjuvants used in most cancer-vaccination trials to date enhance the T helper 2 (Th2) response, and so have a sub-optimal effect on cellular immunity. Aluminum hydroxide, which is used in many commercial vaccine preparations, has been shown to selectively block activation of CD8<sup>+</sup> cytotoxic T lymphocytes (CTLs) in mice immunized against hepatitis-B surface antigen (SCHIRMBECK et al. 1994). Other adjuvants that induce a more extensive Th1 response are currently being evaluated in both pre-clinical and clinical studies (BALDRIDGE and WARD 1997), and are reviewed elsewhere in this volume. These include threonyl-muryl dipeptide (Hsu et al. 1993, 1997), a variety of attenuated or killed bacteria (CHEN et al. 1993; BALDRIDGE and WARD 1997) and bacterial derivatives (JOHNSTON and BYSTRYN 1991), BCG (MASTRANGELO et al. 1996) and Quillaja saponaria 21 (LIVINGSTON et al. 1994). While each of these has shown some efficacy, none are ideal due to toxicity (including both systemic toxicity and local inflammation after repeated immunization), minimal efficacy at stimulating a cellular response or difficulties associated with production. Thus, new adjuvants are needed for vaccine approaches designed to enhance the Th1 response.

Synthetic DNA containing immunostimulatory sequences are particularly attractive as adjuvants for tumor-antigen immunization because of their ease of production and potent ability to induce a Th1 response. In a number of murine systems, CpG DNA have been evaluated as immune adjuvants and shown to be effective. The studies outlined below were not all performed in tumor models, but the results point to the promise of CpG DNA as an adjuvant in tumor-antigen immunization. Chu et al. used hen-egg lysozyme (HEL) as a model antigen and found that immunization with HEL in incomplete Freund's adjuvant (IFA) resulted in Th2-dominated immune responses characterized by HEL-specific secretion of Th2 cytokines (IL-5 but not IFN- $\gamma$ ). In contrast, the addition of CpG DNA to immunization with IFA-HEL switched the immune response to a Th1dominated cytokine pattern (high levels of IFN- $\gamma$  and decreased IL-5) (C<sub>HU</sub> et al. 1997). CpG DNA also enhanced production of anti-HEL immunoglobulin G2a (IgG2a, a Th1-associated isotype) when compared with IFA-HEL alone. This Th1 response was more marked than that seen with complete Freund's adjuvant (CFA). Using hepatitis-virus-B surface antigen as an immunogen, Davis et al. demonstrated a marked increase in antigen-specific IgG and IgG2a (DAVIS et al. 1998). Importantly, they also found development of an antigen-specific cytotoxic-T-cell response. A similar humoral and cellular response was found by Lipford et al. using ovalbumin as the target antigen (LIPFORD et al. 1997). In addition, these investigators detected an enhanced CTL response to both unprocessed protein antigen and to major histocompatibility complex (MHC) class-I-restricted peptide.

We utilized the Id from the 38C13 murine lymphoma model as the target antigen. The Id served as the tumor antigen. CpG DNA was as effective as CFA at inducing an antigen-specific antibody response, but was associated with less toxicity. CpG DNA induced a higher titer of antigen-specific and tumor-specific IgG2a than did CFA. Mice immunized with CpG DNA as an adjuvant were protected from tumor challenge to a degree similar to that seen in mice immunized with CFA but with less toxicity (WEINER et al. 1997). We also explored synergy with granulocyte-monocyte colony-stimulating factor (GM-CSF). Immunization using antigen (Id), CpG DNA and soluble GM-CSF enhanced production of antigen-specific antibody and shifted production towards the IgG2a isotype. This effect was most pronounced after repeat immunizations with CpG DNA and antigen/GM-CSF fusion protein. A single immunization with CpG DNA and antigen/GM-CSF fusion protein 3 days prior to tumor inoculation prevented tumor growth, whereas other approaches to vaccination in this model are ineffective under those conditions (LIU et al. 1998).

Studies in these varied systems indicate that CpG DNA enhances an antigenspecific Th1 response after immunization with soluble antigen and can lead to development of antigen-specific cellular immunity. CpG DNA have been well tolerated and have been equal or superior to standard adjuvants, such as CFA, at inducing an antibody response. Perhaps most importantly, CpG DNA has also been superior at inducing a therapeutic effect. Additional studies that compare CpG DNA to the other adjuvants outlined above are needed in order to determine whether synergy exists between these adjuvants and to determine the efficacy of CpG DNA as an adjuvant in humans.

Classic immunologic teaching states that intracellular proteins are processed and presented in class-I molecules, and this leads to a cellular immune response. In contrast, extracellular antigens are presented in class-II molecules, which leads to a humoral response. There is increasing evidence that there is also cross-talk between the class-I and class-II pathways. Some extracellular antigens are taken up by antigen-presenting cells (APCs) and processed in a manner that leads to presentation in class-I molecules and development of a CTL response termed "crosspriming" (Rock 1996). The results outlined above support the concept that an effective cellular response can be induced by immunization with an intact tumor antigen plus an adjuvant, such as CpG DNA, that activates cells that can process exogenous antigen and present peptides in class-I MHC (DAVIS 1998).

It is important to note the CpG DNA have yet to be evaluated as immune adjuvants in humans. Although individual CpG DNA differ somewhat in their ability to activate various immune cell populations and induce cytokine production in human and murine systems, it is now clear that both human and murine leukocytes respond to this novel pathway of immune activation (KRIEG et al. 1996). It is, therefore, likely that sequences that induce activation of human antigen presenting cells in vitro will be effective adjuvants in vivo, as is observed in the mouse models.

## 5 CpG DNA for the Enhancement of the Efficacy of Cellular Immunotherapy

The 1980s saw the development of recombinant cytokines, such as IL-2, that allowed for the ex vivo expansion of various populations of lymphocytes. Cells expanded in this manner were called "lymphokine-activated killer" or LAK cells and had significant in vitro tumoricidal activity. Initial clinical trials of "adoptive immunotherapy" using LAK cells were promising (LOTZE and ROSENBERG 1986). However, it soon became apparent that LAK cells were not migrating well to tumor (BASSE 1995). In addition, large doses of cytokines were needed to keep the LAK cells functional once they were re-infused into the patient. This resulted in significant toxicity. Finally, LAK cells expanded from the peripheral blood were not tumor specific. "Tumor-infiltrating lymphocytes", or TILs, were obtained from the tumor mass itself and expanded in a similar manner in an attempt to enhance the number of tumor-specific cells. Despite enhanced specificity in vitro, clinical trials of TILs demonstrated little anti-tumor activity (SCHWARTZENTRUBER et al. 1994). Although the lymphocytes were harvested from the tumor mass, many may have been bystander cells and not truly specific for the tumor. There is also evidence that the signaling pathways in TILs are abnormal and that this may limit the cytotoxic capabilities of TILs (FINKE et al. 1993). Adoptive immunotherapy studies using cytotoxic lymphocytes are continuing in both the laboratory and the clinic, but problems related to target specificity and migration of lymphocytes to the target remain.

If these problems can be overcome, CpG DNA might have a role in adoptive immunotherapy approaches. Animal models suggest much of the anti-tumor activity observed with LAK and TIL cells may actually be related to NK activity. CpG DNA is known to enhance NK function in vitro and in vivo (BALLAS et al. 1996). CpG DNA might, therefore, be useful in expanding and activating NK cells intended for adoptive immunotherapy. The effect of CpG DNA on T-cells remains controversal. In studies using highly purified T cells, Lipford et al. found that CpG DNA enhances proliferation of T-cells when the T-cell receptor is engaged, suggesting that CpG DNA is capable of supplying a co-stimulatory signal directly to the T cell (LIPFORD et al. 1997). In contrast, we have been unable to demonstrate a direct co-stimulatory effect of CpG DNA on T-cell activation or proliferation. Further studies are needed to determine whether CpG DNA does indeed have a co-stimulatory effect on T-cell activation and whether such an effect can be used to enhance T-cell-mediated lysis of tumors.

While it is unclear whether CpG DNA can impact directly on T-cell activation, there is no doubt that CpG DNA can enhance T-cell activity indirectly by improving antigen presentation. This effect fits in well with the recent focus of adoptive-transfer studies on the use of APCs to induce an active immune response in the host. Among the most promising recent cancer-immunotherapy studies are those involving dendritic cells (DCs), which are extremely potent APCs (GILBOA et al. 1998; Lotze et al. 1997; Morse and Lyerly 1998). Of particular interest is the ability of DCs to induce a cellular immune response (HAMBLIN 1996; STEINMAN 1996; GIROLOMONI and RICCIARDICASTAGNOLI 1997; MCCANN 1997). Levy and colleagues have demonstrated induction of an antigen-specific response following treatment with antigen-pulsed DCs in both an animal model (TIMMERMAN and LEVY 1997) and humans (HSU et al. 1996). In a small clinical trial, four of four treated patients demonstrated a cellular immune response to Id following immunization with Id-pulsed DCs, and clinical regression of tumor was seen, to some degree, in all patients. Interestingly, no humoral response was seen. This was in stark contrast to the studies, reported by the same group, which demonstrated that immunization of patients with Id-keyhole limpet hemocyanin leads to an intense humoral response (Hsu et al. 1993; NELSON et al. 1996). Of particular interest is the suggestion of prolonged survival in the small number of patients treated with Id-pulsed DCs (Hsu et al. 1997).

DCs can be generated using a variety of cytokines and from a variety of organs, including the bone marrow, lymph nodes and skin. The resulting DCs have varied phenotypes and functions depending upon their site of origin and the cytokines that are used to expand and activate them in vitro. The ideal source of DCs or approach to in vitro expansion, activation, exposure to antigen and reinfusion has yet to be determined.

CpG DNA has effects on DCs; these effects enhance the ability of the DCs to present antigen and induce an antigen-specific cellular response. Sparwasser et al.

have shown that CpG DNA induces maturation of immature DC obtained from murine bone marrow and activates mature DC to produce cytokines, including IL-12, IL-6 and TNF- $\alpha$  (SPARWASSER et al. 1998). Jakob et al. found that treatment of DCs derived from murine fetal skin decreased E-cadherin-mediated adhesion, upregulated MHC class-II and co-stimulatory molecules and enhanced accessory cell activity. Injection of CpG DNA into murine dermis led to enhanced expression of MHC class II and CD86 by the Langerhans' cells (JAKOB et al. 1998).

We also found that CpG DNA markedly enhances the production of cytokines, including IL-12, from bone-marrow-derived DCs (LIU et al. 1998). In addition, the effect of CpG DNA on DCs was synergistic with that induced by GM-CSF. Recent studies in our laboratory suggest that the effect of CpG DNA on cytokine production by DCs is largely responsible for their enhanced antigen-presenting capabilities. In particular, the effect of CpG DNA is due largely, but not fully, to enhanced production of IL-12 by the DCs. We have also found that CpG DNA can enhance the survival, maturation and differentiation of human DCs in vitro (HARMANN et al. 1999).

Given the increasing interest in DC-based cancer immunotherapy and the potent effect of CpG DNA on DCs, it is clear that the combination of CpG DNA and DCs requires further evaluation. However, before a rational clinical trial can begin, we need to have a better understanding of the cellular response of DCs to CpG DNA so that the optimal sequence of DC generation, antigen pulsing and CpG-DNA administration can be used.

## 6 CpG DNA for the Enhancement of the Anti-Tumor Response Following DNA Immunization

Another area of intense interest in the field of cancer immunotherapy is the use of DNA immunization using naked DNA constructs containing sequences that encode for the antigen of interest (SPOONER et al. 1995; ULMER et al. 1996; COHEN et al. 1998). The intent of such therapy is to have host cells take up the DNA, produce protein based on the infused DNA, and express peptides derived from that protein in host class I, thereby inducing a cellular immune response directed towards that antigen. It is unclear what cell type is responsible for uptake of the DNA, production of the protein and presentation of peptide in class I following DNA immunization. Although initial thoughts were that any cells (such as myocytes) could perform these functions, there is now evidence that professional APCs are involved (DoE et al. 1996; Fu et al. 1997). Irrespective of which cells are responsible, this approach has been evaluated with some success in tumor models (SYRENGELAS et al. 1996).

The importance of professional APCs in DNA immunization is consistent with recent studies demonstrating that the presence of CpG motifs in DNA-vaccine constructs impacts on their effectiveness as vaccines. Sato et al. found that human

monocytes transfected with plasmid DNA or double-stranded oligonucleotides containing CpG sequences transcribed larger amounts of IFN- $\alpha$ , IFN- $\beta$ , and IL-12 when compared with cells transfected with DNA that did not contain such sequences (SATO et al. 1996). Krieg and colleagues have also found that modifying the CpG content of vectors intended for DNA immunization can have a significant impact on their ability to induce development of a cellular response (BRAZOLOT MILLAN et al. 1998). The ability to construct vectors that encode for a specific protein and enhance a Th1 response to peptides derived from that protein would have clear implications in the area of tumor vaccination and surely will be evaluated extensively in the years ahead.

## 7 Conclusion

Advances in our understanding of the relationship between the immune system and malignancy have resulted in a reawakening of interest in the field of cancer immunotherapy (Table 1). Recent recognition of the potent immunostimulatory effects of CpG DNA suggest that such agents may well be important if cancer immunotherapy is to play a major role in our treatment and prevention of malignancy. Preliminary studies suggest that CpG DNA can be effective, alone or in combination with mAbs, at inducing tumor regression. Perhaps even more prom-

	CpG as a single agent	CpG to enhance effect of antibody therapy	CpG as an adjuvant with protein immunization	CpG to enhance efficacy of adoptive immunotherapy with dendritic cells	CpG as an important component of DNA vaccines
Most likely mechanism	Enhanced NK activation	Enhanced ADCC by monocytes and NK cells	Enhanced production of antibody and improved activation of cellular immunity	Increased production of Th1 cytokines, including IL-12	Enhanced presentation of antigen by antigen- presenting cells
Evidence	Enhanced in vitro killing and in vivo efficacy in a mouse melanoma model	Enhanced ADCC in vitro and improved therapeutic effects in a mouse lymphoma model	Effective at inducing a cellular responses and improving efficacy in mouse models	Enhancement of dendritic cell function	CpG content of vectors impacts on efficacy of DNA immunization

Table 1. Summary of potential uses of CpG DNA in cancer immunotherapy

ADCC, antibody-dependent cellular cytotoxicity; IL, interleukin.

ising is the potential use of CpG DNA to enhance development of an active cellular anti-tumor response in the host through its use as a vaccine adjuvant, as an agent capable of stimulating APCs or as part of DNA immunization strategy. It remains to be determined whether CpG DNA has a future as a single agent in the treatment of NK-sensitive malignancies, such as melanoma or renal cell cancer, as an agent in combination with antibodies currently being used to treat lymphoma and breast cancer, such as Rituximab and Trastuzumab, or as a part of vaccine therapy for a variety of malignancies. Further work in both the laboratory and the clinic is needed before we can know the true promise of this exciting new class of agents.

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# Use of CpG DNA for Enhancing Specific Immune Responses

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## **1** Introduction

## 1.1 CpG DNA and the Vertebrate Immune System

Bacterial DNA, in contrast to vertebrate DNA, has the capacity to act as a potent immune stimulator (TOKUNAGA et al. 1984; MESSINA et al. 1991; YAMAMOTO et al. 1992b). This is now known to be due to unmethylated CpG dinucleotides that are immunostimulatory when present within the context of certain flanking bases (CpG-S motifs) (KRIEG et al. 1995). CpG dinucleotides are present at the expected frequency of 1/16 bases in bacterial DNA but are under-represented (1/50–1/60 bases) in vertebrate DNA, where they are almost always methylated (BIRD et al. 1987). CpG-S motifs are virtually absent from vertebrate DNA, and the rapid immune activation in response to CpG DNA appears to be an evolutionary adaptation whereby CpG-S motifs act as a "danger" signal that the vertebrate innate immune defense mechanisms can recognize and respond to (KRIEG et al. 1996).

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CpG DNA can induce proliferation of almost all (>95%) B cells and further protects B cells from apoptosis. It also triggers polyclonal immunoglobulin (Ig), interleukin 6 (IL-6) and IL-12 secretion from B cells and can drive isotype shifting in vitro (KRIEG et al. 1995; KLINMAN et al. 1996; YI et al. 1996a,b). This B-cell activation by CpG DNA is T-cell independent and antigen non-specific although, as discussed below, these effects have strong synergy with antigen-specific responses. In addition to its direct effects on B cells, CpG DNA also directly activates monocytes, macrophages and dendritic cells to secrete interferon  $\alpha/\beta$  (IFN- $\alpha/\beta$ ), IL-6, IL-12, granulocyte-monocyte colony-stimulating factor (GM-CSF), chemokines and tumor necrosis factor  $\alpha$  (TNF- $\alpha$ ) (Cowdery et al. 1996; HALPERN et al. 1996; KLINMAN et al. 1996; SCHWARTZ et al. 1997). These cytokines stimulate natural killer (NK) cells to secrete IFN- $\gamma$  and have increased lytic activity (YAMAMOTO et al. 1992a; BALLAS et al. 1996; COWDERY et al. 1996; KLINMAN et al. 1996; YI et al. 1996; CHACE et al. 1997). Overall, CpG induces a T-helper type 1 (Th1)-like pattern of cytokine production dominated by IL-12 and IFN- $\gamma$  with little secretion of Th2 cytokines (KLINMAN et al. 1996). CpG ODN also induces co-stimulatory molecule expression in vitro and in vivo (DAVIS et al. 1998) and upregulates expression of major histocompatibility complex (MHC) molecules. Similar effects to those induced by bacterial DNA may be obtained with synthetic oligodeoxynucleotides (ODN) containing CpG-S motifs (CpG ODN).

Not all DNA sequences containing unmethylated CpG dinucleotides are stimulatory, and there is some species specificity to such effects (see below). CpG-S motifs are typically preceded on the 5' side by an ApA, GpA, or GpT dinucleotide and followed on the 3' side by two pyrimidines, especially TpT. Many sequences containing CpG dinucleotides that are non-stimulatory for a given species appear to have no effect on the immune system; however, some DNA sequences containing CpG dinucleotides may actually counteract the stimulatory effects of CpG-S motifs. Such "neutralizing" CpG motifs (CpG-N), which are over-represented in certain adenoviral genomes (types 2 and 5), typically include CCG, CGG, or CGrepeats in their sequences (KRIEG et al. 1998). Such sequences are typical of the socalled "CpG islands" in vertebrate DNA, described by BIRD et al. (1987), which are usually unmethylated but nevertheless non-stimulatory.

#### 1.2 Adjuvant Properties of CpG DNA

In addition to the activation of innate immune responses, CpG DNA can serve as a potent adjuvant for the enhancement of antigen-specific responses. Enhancement of humoral responses appears to be due to the strong synergy between the direct activation of B cells by CpG DNA and the signals, delivered through the B-cell antigen receptor, for both B-cell proliferation and Ig secretion (KRIEG et al. 1995). In addition, antigen-specific humoral responses are likely enhanced by the induction of cytokines that could have indirect effects on B cells via T-helper pathways. Overall, CpG DNA induces a Th1-like pattern of cytokine production dominated by IL-12 and IFN- $\gamma$ , with little secretion of Th2 cytokines (KLINMAN et al. 1996).

Th1-type immune responses are desirable because (i) they are associated with the production of predominantly IgG2a antibodies, which are thought to have better neutralizing capabilities than the Th2-associated IgG1 antibodies, (ii) antigen-specific cell-mediated immunity (CMI) is enhanced by better antigen presentation with the enhanced expression of MHC and co-stimulatory molecules (DAVIS et al. 1998), and (iii) the Th1-type cytokines can aid the maturation of cytotoxic T lymphocytes (CTL).

CpG DNA appears to enhance immune responses against any type of antigen and, as such, it could be used to improve the efficacy of any prophylactic vaccines against infectious disease. The strong Th1 bias of CpG DNA and the ability to induce strong CMI also opens the possibility for use of CpG DNA in therapeutic vaccines against, for example, chronic infections or cancer.

For the purpose of in vivo enhancement of antigen-specific responses in humans, it is preferable to use synthetic short ODN, which can be well characterized, rather than bacterial DNA. As well, for in vivo use, the ODN should be made with a synthetic phosphorothioate backbone to render them more nuclease-resistant without losing the immune-stimulatory effects (KRIEG et al. 1996; ZHAO et al. 1993, 1996; BOGGS et al. 1997). In vivo adjuvant effects of CpG ODN with a native phosphodiester backbone (O-ODN) or a chimeric backbone composed of a phosphodiester center and phosphorothioate ends (SOS-ODN) are diminished or lost (Davis, unpublished observation), presumably due to more rapid destruction by lymphocyte nucleases (ZHAO et al. 1993). In practice, we have found 18–24-mer ODN to be optimally effective but of a manageable size for easy and inexpensive manufacturing.

CpG-S motifs may also be used to enhance immune responses with DNA vaccines. In this case, for reasons discussed below, it is best to clone the sequences into the backbones of the plasmid vectors.

#### 1.3 Species-Specificity of CpG DNA

CpG-S motifs are species-specific, to a certain extent. For example, our optimal mouse motif with a single CpG dinucleotide in a phosphorothioate ODN (TGACGTT) does not activate human cells. It is possible to induce more potent immune activation by having more than one CpG motif in the ODN, in which case the spacing between motifs is critical (Davis and Krieg, unpublished observations). Furthermore, the same CpG sequence in phosphodiester or phosphorothioate backbones may not show the same stimulatory effects or work at the same optimal concentration. Thus, factors that seem to influence the stimulatory effect of a particular CpG ODN in a given species include the backbone of the DNA and the sequence, number, and spacing of the CpG motifs.

Mouse cells will respond to a wide variety of CpG motifs, including human motifs, whereas human cells are stimulated by a much more restricted subset of CpG-S motifs. Therefore, it is not possible to predict the in vivo effects of CpG-S motifs in humans from in vivo results in mice. Furthermore, it may not even be

possible to make such predictions from primate studies. For example, we have found that immune cells from such closely related species as the human and the chimpanzee don't necessarily respond to the same CpG-S ODN, even when the sequence is very similar to that of another ODN that does activate cells from both species (Krieg and Davis, unpublished results).

While the mechanism of the surprisingly complex role of the bases flanking CpG dinucleotides has yet to be elucidated, it is important to take them into consideration when designing in vivo studies to evaluate CpG ODN as a vaccine adjuvant. In particular, it is absolutely necessary to use immune cells from the desired target species and test them with a panel of in vitro tests (i.e. B-cell proliferation, cytokine secretion, NK activity) to determine which are immunostimulatory for that species before attempting to use them as vaccine adjuvants in vivo.

## 2 Experience with CpG DNA and Antigen-Based Immunization

CpG ODN have been shown to be potent adjuvants for augmenting both humoral and cell-mediated immune responses against a wide variety of antigens. These include protein antigens of a diverse nature and polysaccharide (PS) antigens. Furthermore, CpG ODN has been delivered with antigens both parenterally as well as mucosally.

The CpG ODN is best given mixed together with the antigen. While this may not be absolutely essential, we have found that enhancement of the humoral response is greatly abrogated if the CpG ODN is given at a different time (i.e. one day before or after) or in the contralateral leg (Davis and Krieg, unpublished observations). This effect may be partly antigen dependent, since others have reported adjuvant effects when the CpG ODN was given up to 30 days prior to a model antigen (Lipford and Wagner, personal communication) or when it was given in the contralateral leg to the site of injection of whole killed influenza vaccine (B. Wahren, personal communication).

#### 2.1 CpG DNA with Protein Antigens

The greatest amount of experience using CpG ODN to augment antigen-specific immune responses has been with protein antigens. Many of these are infectiousdisease-related antigens, including live attenuated measles virus (Kovarik et al. 1999), whole killed influenza virus (MOLDOVEANU et al. 1997; ROMAN et al. 1997), hepatitis B surface antigen (HBsAg, a virus-like particle) (BRAZOLOT MILLAN et al. 1998; DAVIS et al. 1998), and a peptide derived from tetanus toxoid (Kovarik et al. 1999). Enhanced immune responses have also been demonstrated using CpG ODN with model antigens, such as ovalbumin (KLINMAN et al. 1997; LIPFORD et al. 1997a,b), hen-egg lysozyme (CHU et al. 1997), β-galactosidase (ROMAN et al. 1997; HORNER et al. 1998) and fowl  $\gamma$ -globulin (SUN et al. 1998). Finally, CpG ODN has been shown to be highly effective for both the enhancement of Th1-like immune responses against a tumor antigen, the idiotype of the surface IgM of 38C13 mouse lymphoma cells, and for protection against subsequent tumor challenge (WEINER et al. 1997; LIU et al. 1998). The ability to use CpG ODN with virtually any type of antigen is particularly attractive, because alum, the only adjuvant currently licensed for human use in most countries, cannot be used with live attenuated pathogens and has not been found to be particularly useful with other vaccines, such as those containing whole killed influenza virus.

The findings of the above studies have shown that, in virtually all cases, CpG ODN causes significant enhancement of antigen-specific antibody titers. These can be as much as 2–3 log higher than titers obtained with antigen alone or with a non-CpG control ODN. In several studies, strong Th1 bias of the CpG-induced response is also shown by the induction of potent CTL responses (LIPFORD et al. 1997a,b; BRAZOLOT MILLAN et al. 1998; DAVIS et al. 1998), preferential secretion of Th1-type cytokines, such as IFN- $\gamma$  or IL-12 (Chu et al. 1997; HORNER et al. 1998; KOVARIK et al. 1999), and increased relative proportion of IgG2a antibodies (CHU et al. 1997; ROMAN et al. 1997; BRAZOLOT MILLAN et al. 1998; DAVIS et al. 1998; SUN et al. 1998).

CpG ODN, which is water soluble, has a strong synergy with other adjuvants, such as alum (DAVIS et al. 1998), GM-CSF cholera toxin (CT) (McCLUSKIE and DAVIS 1998), and incomplete Freund's adjuvant (IFA). Indeed, these combinations give responses equal or superior to those with complete Freund's adjuvant (CFA), with much less toxicity. The synergy may be related to different mechanisms of the immune modulators (GM-CSF, CT) or in the case of alum and IFA, the CpG ODN may benefit from the depot effect. We have not detected any signs of toxicity in mice, even with doses 50 times higher than those required for strong adjuvant effects.

The strong adjuvant effects of CpG ODN may also allow the use of reduced doses of antigen. For example, reducing the dose of HBsAg (alone or with alum) by tenfold likewise reduces the antibody titers in mice by tenfold but, when given with CpG ODN, there is no loss of the high titers of anti-HBs antibodies (Davis, unpublished results).

#### 2.2 CpG DNA for Enhancing Immune Responses in Early Life

Vaccination of infants is often delayed, because the neonatal immune system is considered too immature to respond appropriately (HUNT et al. 1994). In particular, B-cell responses are weak and preferentially generate IgM/IgG1 antibody isotypes, and cytotoxic responses are poor (KOVARIK et al. 1998). Nevertheless, newborns are at risk for exposure to many infectious diseases, and it would be highly desirable to have vaccines that are effective even with an immature immune system. In particular, the neonatal immune system is strongly Th2 biased, and this results in very poor CMI, which is crucial for the prevention or control of many diseases that neonates are at risk of developing.

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Young mice are useful models in which to test immunization strategies for newborn humans, because their responses to protein antigens have similar limitations (BARRIOS et al. 1996). CpG ODN has been found to be effective in very young mice for inducing or enhancing antibodies and CTLs against HBsAg (BRAZOLOT MILLAN et al. 1998), live attenuated measles virus, and a tetanus-toxoid-derived peptide (KOVARIK et al. 1999). Not only does CpG ODN allow the induction of immune responses at younger ages than does antigen alone or antigen with alum, but these responses are more Th1-like. This is particularly important, because the immature immune system is strongly Th2 biased.

#### 2.3 CpG DNA as a Mucosal Adjuvant

Parenteral immunization can induce systemic (i.e. circulating antibodies and CTLs), but only rarely mucosal, immunity (i.e. secretory IgA antibodies). In contrast, antigens delivered at a mucosal surface can trigger both mucosal immunity at local and distant sites and systemic responses (HANEBERG et al.1994; GALLICHAN and ROSENTHAL, 1995). While live attenuated organisms have proven effective for mucosal immunization, results have been disappointing with subunit vaccines, in large part due to the lack of a safe and effective adjuvant.

CT and the closely related *Escherichia coli* heat-labile enterotoxin (LT) are effective mucosal adjuvants in animal models (LYCKE et al. 1992; SPANGLER et al. 1992); however, they are highly toxic, especially in humans. In efforts to produce effective but safe mucosal adjuvants for use in humans, two strategies have been adopted to render CT and LT less toxic: (i) the use of the non-toxic B subunits that lack enzymatic activity (HOLMGREN et al. 1993; VERWEIJ et al. 1998) or (ii) genetically detoxified mutants that have little or no enzymatic activity. Such compounds, which have greatly reduced toxicities, retain some adjuvanticity in animal models (FONTANA et al. 1995; RAPPUOLI et al. 1995; DOUCE et al. 1998; GUILIANI et al. 1998; KOMASE et al. 1998).

CpG ODN has been found to be a highly effective mucosal adjuvant when administered intranasally to mice. For example, both systemic (IgG) and mucosal (secretory IgA) humoral responses against whole killed influenza virus were significantly enhanced by the addition of CpG ODN (MOLDOVEANU et al. 1997). Furthermore, CpG ODN was able to induce antibodies and CTL against HBsAg, which by itself did not induce any detectable responses even with high doses and boosting (McCLUSKIE and DAVIS 1998). In fact, this study showed CpG ODN to be as effective as CT, and to have a strong synergy with CT when the two adjuvants were co-administered. In addition, CpG ODN gave a more Th1-like response than CT, which gave a Th2-type response.

Effective doses of CpG ODN for intranasal delivery  $(1-10\mu g)$  are lower than those required for parenteral immunization  $(10-50\mu g)$ . In addition, CpG ODN appears to be safe and well tolerated with intranasal delivery to mice, even at very

high doses (up to 500µg). Mice inhaling CpG showed no short-term signs of distress over those receiving HBsAg alone, and all recovered quickly, with no apparent long-lasting effects. In contrast, mice receiving high doses of CT (>10µg) show signs of toxicity, such as ruffling of fur and diarrhea. CT is even more toxic in humans, where a dose as low as 1–5µg can cause diarrhea (JERTBORN et al. 1992).

Thus, CpG ODN appears to be a promising new mucosal adjuvant that could be administered with a variety of antigens. It might also be used in combination with mucosal toxins to obtain better immune responses with a lower dose and less adjuvant-related toxicity. The ability to induce mucosal immunity that can prevent entry of pathogens into the body is highly desirable, but equally important is the ability to deliver vaccines in an easy and needle-free fashion. This could radically change the prospect of delivering vaccines to less developed areas of the world and, even in more developed nations, avoidance of the risk of needle-stick injury is highly desirable.

#### 2.4 Th1 Influence of CpG DNA

The strong Th1-bias of CpG ODN has been demonstrated with in vitro experiments and in vivo delivery of antigens by parenteral and mucosal routes to adult and young mice. The best marker of this is the type of cytokine induced (high IFN- $\gamma$  and IL-12 and low IL-4 and IL-5); however, it is also indicated by presence of CTL and a preponderance of IgG2a relative to IgG1 antibodies.

For vaccination against infectious diseases, a Th1-like response is generally desirable. Furthermore, CTLs appear to be essential for protection against certain diseases, especially those where the pathogen resides intracellularly. It is generally accepted that induction of CTL is important for immunotherapy of chronic infections, such as hepatitis B virus (HBV), hepatitis C virus, *Mycobacterium* (tuberculosis), and human immunodeficiency virus. In addition, the Th1 cytokines, such as IFN- $\gamma$  and TNF- $\alpha$ , have been shown to be important for non-lytic control of chronic HBV infection (CHISARI 1997).

While strong Th1 CMI may not be necessary to protect against all infectious diseases, it is clearly a requirement for cancer vaccines where the intent is to induce CTL that can attack and kill cancer cells, for example those remaining after surgical removal of a solid tumor or after chemotherapy. In this regard, CpG ODN has been found to be a more potent Th1 adjuvant than many other adjuvants, including CFA (Davis, unpublished data).

A Th1-like response may be preferential in the lungs, because allergic asthma is generally associated with Th2-like responses (KAY 1996; HOGG 1997). Asthma has become an increasingly prevalent disease in the developed world, and this has been has been linked to the reduced incidence of bacterial infections owing to better hygiene and widespread use of antibiotics (COOKSON and MOFFATT 1997). It is also possible that the strong Th2 bias of alum, which is currently used in many pediatric vaccines, may contribute to the rising incidence of asthma. The ability to induce

Th1 responses in babies and young children with CpG ODN may help reduce the incidence of asthma. Indeed, CpG ODN has recently been shown, in mice, to prevent allergen-induced asthmatic responses, which include airway eosinophilia, Th2 cytokine induction, IgE production, and bronchial hyperactivity. Furthermore, CpG ODN can even "cure" pre-established asthmatic responses in mice through redirection of the immune response towards a more Th1-like profile (BROIDE et al. 1998; KLINE et al. 1998).

## 2.5 Polysaccharide Antigens

PS antigens are particularly difficult, because immune responses against them are largely T-cell independent. As such, there is no CMI, antibody responses are short lived, and there is no memory response. One published report found that CpG ODN actually reduced the level of antibodies against a high-molecularweight PS in mice (THREADGILL et al. 1997). However, others have found modest to good increases in antibodies, including those recognized as being T dependent (IgG2a), such as CpG ODN and pneumococcal PS (Siegrist, personal communication), meningitis type-C PS (Westerink, personal communication), and trinitrophenyl-Ficoll (Waldschmidt, personal communication), although some of these studies failed to demonstrate induction of memory responses. The poor results in the first study may be related to the dose of  $500\mu g$  CpG ODN, which is greatly in excess of an optimal dose for mice (about 10µg). Nevertheless, results obtained with CpG ODN and PS antigens have been less dramatic relative to those obtained with protein antigens. This is likely related the T-independent nature of anti-PS responses, since many of the effects of CpG-S motifs are involved with antigen presentation to T-cells or induction of cytokines that serve T-helper functions.

## 2.6 CpG DNA with Antigen–Antibody Complexes

Antigen–antibody complexes have been shown to induce significantly better immune responses than antigen alone in a number of different animal models (POKRIC et al. 1993; HANKE et al. 1994; HADDAD et al. 1997). It has also recently been found that CpG ODN can significantly enhance these responses in mice when administered parenterally with complexes of hepatitis B surface antigen (HBsAg) mixed with antibodies against HBsAg (anti-HBs; HBsAg/Ab complexes) (Wen, personal communication).

We have also shown that HBsAg/Ab complexes can induce strong immune responses when given by a mucosal route (McCLUSKIE et al. 1999). Although CpG ODN is a highly effective mucosal adjuvant with some antigens (discussed below), in the case of antigen–antibody complexes, CpG ODN does not enhance antibody titers, although it can shift the humoral response to be more Th1-like.
#### **3** Role of CpG Motifs in DNA Vaccines

DNA vaccination has emerged as an exciting new approach to vaccination that promises significant advantages over current vaccination strategies. Numerous groups have demonstrated the induction of potent and long-lasting immune responses, including induction of CTL, against viral, bacterial, or parasitic antigens encoded in DNA plasmids (DAVIS 1998; DONNELLY et al. 1998). The strength of the immune response with DNA vaccines is impressive considering the very small amount of antigen produced. This appears to be due to the in vivo synthesis of antigens (and, thus, efficient MHC presentation of antigens) over an extended period of time, as well as to the CpG-S motifs inherent in the backbone of the plasmid DNA vectors, such as those commonly used for DNA-based immunization. For example, removal or methylation of potent CpG-S motifs has been found to reduce or abolish immune responses against encoded antigens (SATO et al. 1996; KLINMAN et al. 1997)

Other investigators working in the field of DNA vaccines have also observed that the DNA backbone itself provides a Th1-type adjuvant effect when administered with a protein antigen (LECLERC et al. 1997). This adjuvant effect of the DNA backbone appears to require unmethylated CpG motifs, because it is lost if the DNA vector is treated with CpG methylase (KLINMAN et al. 1997). Moreover, a non-expressing plasmid enriched in CpG motifs is a more effective adjuvant for protein vaccines than a non-optimized plasmid (SATO et al. 1996; ROMAN et al. 1997).

In order to use CpG-S motifs to enhance immune responses with DNA vaccines, it is really essential to clone them into the vector backbone. We have shown that it is not possible to directly mix S-ODN with plasmid DNA, because this will result in an ODN-dose-dependent reduction in gene expression from the plasmid (WEERATNA et al. 1998), possibly due to competitive interference at binding sites on the surfaces of target cells (ZHAO et al. 1993). While ODN with a phosphorothioatephosphodiester chimeric backbone (SOS-ODN) do not adversely affect the level of gene expression (except those with poly-G sequences, which have greatly increased binding to cell membranes) (HUGHES et al. 1994), this is not useful, because SOS-ODN are apparently not sufficiently nuclease resistant to exert a strong CpGadjuvant effect. It is also not possible to augment responses to DNA vaccines by administering the CpG S-ODN at a different time or site than the plasmid DNA (Weeratna and Davis, unpublished results). Thus, at least for the present, it appears necessary to clone CpG motifs into DNA vaccine vectors in order to take advantage of their adjuvant effects.

Optimization of the CpG content of DNA vaccines is further complicated by the presence of numerous CpG-N motifs, which can neutralize the effects of the CpG-S motifs. We have shown that it is possible to improve the efficacy of a DNA vaccine in mice by using site-directed mutagenesis to remove many of the CpG-N motifs. Further improvement was then obtained by cloning additional CpG-S motifs into the non-coding backbone (KRIEG et al. 1998). Such CpG-optimized 180 H.L. Davis

DNA-vaccine vectors may be particularly advantageous for use in humans, where the more restricted subset of effective CpG-S motifs will negatively skew the balance of neutralizing and stimulatory motifs.

# 4 Summary

CpG ODN, owing to its wide range of immunostimulatory effects has been found to be a potent Th1-type adjuvant that is effective with virtually any type of antigen, although responses are less impressive with PS than protein antigens. The use of CpG ODN as an adjuvant may allow the development of vaccines against a wider range of diseases, which could include therapeutic vaccines for chronic infections or cancer, effective pediatric vaccines for newborns, and easily delivered mucosal vaccines.

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# Immunostimulatory-Sequence DNA is an Effective Mucosal Adjuvant

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# **1** Introduction

The respiratory, gastrointestinal, vaginal, and rectal mucosa are major portals of entry for infectious agents and foreign proteins (CZERKINSKY and HOLMGREN 1995; STAATS and McGHEE 1996). Locally produced and secreted immunoglobulin A (IgA) and cytotoxic T lymphocytes (CTL) within the mucosal tissue and draining lymph nodes protect against microbial infection (GALLICHAN et al. 1993; CZERKINSKY and HOLMGREN 1995; STAATS and McGHEE 1996; ADA and McELRATH 1997). Because the mucosal immune system serves a front-line role in protecting us from our environment, there is a great deal of interest in developing strategies for stimulating protective mucosal immune responses. Unfortunately, traditional vaccination methods are not effective and, at present, there is no simple, safe, and generally applicable approach for the induction of mucosal immune responses.

Mucosal delivery of replicating immunogens, such as live attenuated vaccines, elicits robust and long-lasting immune responses, including the production of mucosal IgA, serum antibody, and A CTL response (CZERKINSKY and HOLMGREN

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1995; STAATS and McGHEE 1996). However, simple protein antigens are often used for research purposes and clinical applications, and they are much less immunogenic. Systemic [i.e. intradermal (i.d.)] vaccination with monomeric proteins elicits serum antibody production, but many arms of the immune response, such as mucosal IgA synthesis and CTL activity are not elicited. However, mucosal vaccination with protein alone generally will not elicit any immune response and may even induce tolerance (CZERKINSKY and HOLMGREN 1995; GUPTA and SIBER 1995; STAATS and McGHEE 1996). In general, to elicit a mucosal IgA response, monomeric protein antigens need to be delivered to mucosal surfaces with an adjuvant (CZERKINSKY and HOLMGREN 1995; STAATS and McGHEE 1996). Cholera toxin (CT) is the best-studied and most potent mucosal adjuvant identified to date. Unfortunately, like many adjuvants, it has unacceptable toxicity for use in humans (GUPTA and SIBER 1995). In addition, CT biases systemic immune responses toward a T-helper 2 (Th<sub>2</sub>) phenotype, potentially leading to IgE production and allergic hypersensitivity toward the antigen (SNIDER et al. 1994; MARINARO et al. 1995).

Immunostimulatory-sequence DNA contained within plasmids and oligodeoxynucleotides (ISS-ODN) has previously been shown to have potent adjuvant activity for systemic immune responses to protein antigens. The immune responses seen after systemic vaccination with simple proteins and immunostimulatory-sequence DNA shares many features with the immune responses seen after viral infections. This immune response characteristically has a Th<sub>1</sub> bias, is robust and long lasting, and generally includes CTL activity (SATO et al. 1996; ROMAN et al. 1997; DAVIS et al. 1998; ZIMMERMANN et al. 1998). Here we review our recent observations, which lead us to conclude that ISS-ODN is a potent mucosal adjuvant that induces both secretory IgA, and a strong Th<sub>1</sub>-biased systemic immune response following intranasal (i.n.) delivery with antigen (HORNER et al. 1998).

# 2 ISS-ODN is an Adjuvant for Th<sub>1</sub>-Biased Systemic Immune Responses

ISS-ODN containing the hexamer 5'-pyrimidine-pyrimadine-CpG-purine-purine-3' have been shown to provide effective adjuvant activity for the induction of systemic Th<sub>1</sub>-biased immune responses toward protein antigens co-administered via i.d. and intramuscular routes. The immune response includes the induction of a Th<sub>1</sub>-cyto-kine profile [interferon  $\gamma$  (IFN $\gamma$ ) but not interleukin 4 (IL-4)], the production of high IgG2a and low IgG1 titers (IFN $\gamma$  and IL-4-dependent isotypes respectively), and a CTL response. In contrast, immunization with protein alone leads to a relatively Th<sub>2</sub>-biased immune response without CTL activity (ROMAN et al. 1997; DAVIS et al. 1998; HORNER et al. 1998; ZIMMERMANN et al. 1998).

#### 2.1 Insights into How ISS-ODN Functions as an Adjuvant

The mechanisms underlying the Th<sub>1</sub>-biased adjuvant activity of ISS-ODN are not well understood at the molecular level. However, at the cellular level, we do have

some mechanistic insights. ISS-ODN can induce the production of IFN  $\alpha + \beta$ , IL-12, and IL-18, from macrophages and IFN $\gamma$  from natural killer (NK) cells in an antigen-independent manner (YAMAMOTO et al. 1992; KLINMAN et al. 1996; PISETSKY 1996; ROMAN et al. 1997). These cytokines are known to promote Th<sub>1</sub>-biased immune responses (COFFMAN and MOSMANN 1988; MOSMANN and COFFMAN 1989; ROMAN et al. 1997). In addition, ISS-ODN stimulates IL-6 production from B cells and IL-10 production from macrophages (KLINMAN et al. 1996; ANITESCU et al. 1997; personal observations). IL-6 stimulates polytypic antibody production, while IL-10 promotes IgA synthesis specifically (MURAGUCHI et al. 1988; DEFRANCE et al. 1992). However, neither IL-6 nor IL-10 are considered to be characteristic Th<sub>1</sub> cytokines (COFFMAN and MOSMANN 1988; MOSMANN and COFFMAN 1989).

Along with the induction of an innate cytokine response, ISS-ODN induces or increases, in an antigen-independent manner, the expression of a variety of cellsurface proteins important for productive immune responses. Incubation of naive splenocytes with ISS-ODN leads to expression of B7.1 and B7.2 on B cells and macrophages. These molecules provide important co-activation signals to T cells via CD 28 engagement (JAKOB et al. 1998; SPARWASSER et al. 1998; SHAHINIAN et al. 1993; Martin-Orozco et al. manuscript submitted). In addition, ISS-ODN increases CD40 expression on B cells and macrophages. CD40 ligation by CD154 (CD40 ligand) stimulates isotype switching by B cells and activates antigen-presenting cells (APCs) (Sparwasser et al. 1998; FULEIHAN et al. 1993; Martin-Orozco et al. manuscript submitted). ISS-ODN also increase B-cell and macrophage expression of class-I and class-II molecules, further increasing their capacity to function as APCs (JAKOB et al. 1998; SPARWASSER et al. 1998; KARLSSON et al. 1996; Martin-Orozco et al. manuscript submitted). Lastly, cytokine-receptor expression (IL2R and IFN $\gamma$ R) is increased on B cells and macrophages, increasing their potential to respond to cytokines in their local environments (Martin-Orozco et al. manuscript submitted).

A central dogma in immunology is that T cells direct the Th bias of antigenspecific immune responses. However, while cytokine production and expression of cell-surface molecules on B cells and macrophages are promoted by incubation with ISS-ODN, purified T cells do not demonstrate these responses when incubated with ISS-ODN (Martin-Orozco et al. manuscript submitted). We believe that the antigen-independent innate responses of B cells, NK cells, macrophages, and other APCs toward ISS-ODN provide important upstream signals, which both promote and bias subsequent antigen-dependent and -specific T-cell and B-cell immune responses. If correct, this model implies that the Th<sub>1</sub>-biased immune response which develops toward antigens delivered with ISS-ODN is initiated and shaped by B cells, macrophages, and possibly other APCs, but not T cells.

# 3 The Mucosal Immune System

Mucosal immunity clearly plays an important front-line role in protecting man from his/her environment. However, adjuvants and mucosal delivery are needed for the induction of mucosal immune responses to many antigens, including most monomeric proteins (CZERKINSKY and HOLMGREN 1995; STAATS and McGHEE 1996). Research suggests that lymphocytes involved in mucosal immune responses express a unique set of surface proteins, such as integrin  $\alpha_4\beta_7$ , which direct these cells to mucosal organs. These homing receptors allow lymphocytes from the site of primary contact with antigen to traffic to multiple mucosal sites, leading to both local and distal mucosal immunity (BERLIN et al. 1993; QUIDING-JARBRINK 1997). Therefore, the mucosal immune system can be considered to be made up of a unique population of lymphocytes that can function semi-independently of lymphocytes involved in systemic immune responses.

# 3.1 Mucosal Immunization with ISS-ODN Induces a Secretory IgA Response

To further characterize the adjuvant potential and clinical utility of ISS-ODN, we assessed its ability to function as a mucosal adjuvant. CT is the best-studied and most potent known mucosal adjuvant for the induction of secretory IgA. Therefore, the IgA response of mice mucosally immunized with protein and ISS-ODN was compared to the IgA response when mice were mucosally immunized with protein and CT. Both ISS-ODN and CT were found to be effective mucosal adjuvants for a variety of antigens and mouse strains. In Fig. 1, the mucosal IgA response of BALB/c mice i.n. immunized with  $\beta$ -galactosidase ( $\beta$ -gal) is presented (HORNER et al. 1998). While direct comparison of the IgA levels from different

Fig. 1A–D. Immunoglobulin A (IgA) responses. Mice received a single immunization with  $\beta$ -galactosidase (β-gal; 50µg) alone, with oligodeoxynucleotide immunostimulatory sequence (ISS-ODN; 50µg), mutated oligodeoxynucleotide (50µg), or cholera toxin (CT; 10µg) via intranasal (i.n.) or intradermal routes. Immunostimulatory sequences and mutated oligodeoxynucleotides used in these experiments have the sequences 5'-TGACTGTGAACGTTCGAGATGA-3', and 5'-TGACTGTGAACCTTAGA-GATGA-3', respectively. Results were obtained by enzyme-linked immunosorbent assay and represent mean values for four mice per group; error bars reflect the standard errors of the means. Results are representative of three similar and independent experiments. A Bronchoalveolar lavage fluid (BALF) IgA. Bronchial lavage was carried out at sacrifice during week 7 with 800µl of phosphate-buffered saline (PBS). There was no significant difference in BALF anti-β-gal IgA levels between i.n. β-gal/ISS-ODN- and  $\beta$ -gal/CT-immunized groups. **B** Vaginal IgA. Vaginal lavage with 50µl of PBS was carried out at sacrifice. There was no significant difference in vaginal anti- $\beta$ -gal IgA levels between i.n.  $\beta$ -gal/ISS-ODN- and  $\beta$ gal/CT-immunized groups. C Fecal IgA. Feces were collected at 2, 4, and 7 weeks, and IgA was extracted as previously described (HANEBERG et al. 1994). There was no significant difference in fecal anti-β-gal IgA levels between the i.n.  $\beta$ -gal/ISS-ODN and  $\beta$ -gal/CT groups except at 2 weeks (P = 0.03). **D** BALF, fecal, vaginal, and serum IgA. Serum was obtained at sacrifice during week 7, and IgA content was compared to week-7 BALF, fecal, and vaginal IgA. In i.n.  $\beta$ -gal/ISS-ODN- and  $\beta$ -gal/CT-immunized mice, the mucosal compartments demonstrated consistently higher anti-β-gal IgA levels then the serum



mucosal sites can not be made due to differences in sample-collection techniques, the data clearly demonstrate that  $\beta$ -gal/ISS-ODN- and  $\beta$ -gal/CT-immunized mice had equivalent antigen-specific IgA levels in bronchial, vaginal, and intestinal secretions. To establish that a mucosal adjuvant was needed for the induction of mucosal IgA, mice were i.n. immunized with  $\beta$ -gal alone or with a mutated oligodeoxynucleotide (M-ODN). However, i.n. vaccination without mucosal adjuvant resulted in no detectable IgA. To evaluate whether contact with the respiratory mucosa was required for ISS-ODN to have mucosal adjuvant activity, mice were immunized with  $\beta$ -gal and ISS-ODN via intragastric (i.g.) and i.d. routes. These routes of immunization did not lead to measurable IgA in mucosal secretions (data for i.g. immunization not shown). To confirm that the IgA detected in bronchoalveolar lavage fluid (BALF), fecal material, and vaginal washes of immunized mice was actively secreted by mucosal tissue and did not passively diffuse from serum, anti-β-gal IgA levels in serum, feces, vaginal washings, and BALF were compared. It should be noted that initial acquisition of BALF, fecal samples, and vaginal washes requires an unmeasurable dilution of the IgA contained in the material, which does not occur when obtaining serum. Despite this fact, i.n. β-gal/ISS-ODN- and i.n. β-gal/CT-immunized mice produced higher levels of anti-β-gal IgA in feces, vaginal washes, and BALF than in serum, demonstrating the active secretion of antigen-specific IgA from the mucosal surfaces of these mice.

These results show that ISS-ODN and CT have equivalent mucosal adjuvant activity with a test antigen that has no capacity to induce mucosal IgA production when delivered alone. In addition, mucosal delivery of antigen with ISS-ODN is shown to lead to a secretory IgA response, while i.d. delivery does not. Taken together, these findings demonstrate that ISS-ODN is an excellent adjuvant for the induction of both local and distal mucosal immunity when co-delivered with antigen via the nose.

# 3.2 Mucosal Immunization with ISS-ODN Induces a Th<sub>1</sub>-Biased Systemic Immune Response

Although systemic immune responses can occur in isolation, immune responses involving mucosal lymphocytes generally occur with concomitant stimulation of the systemic immune system (CZERKINSKY and HOLMGREN 1995; STAATS and MCGHEE 1996). Systemic-immune responses can be subdivided into Th<sub>1</sub>- and Th<sub>2</sub>-biased responses based on the specific cytokine and antibody profile elicited. Th<sub>1</sub>-biased immune responses are seen following viral infection and are generally associated with a CTL response (MOSMANN and COFFMAN 1988, 1989). In contrast, Th<sub>2</sub>-biased immune responses are associated with allergic hypersensitivities and the synthesis of IgE (SNIDER et al. 1994). The adjuvants ISS-ODN and CT are considered to promote Th<sub>1</sub>- and Th<sub>2</sub>-biased immune responses, respectively (SNIDER et al. 1994; MARINARO et al. 1995; ROMAN et al. 1997; DAVIS et al. 1998; MCCLUSKIE and DAVIS 1998; HORNER et al. 1998). Therefore, the magnitude and phenotype of the systemic immune response induced by i.n.  $\beta$ -gal/ISS-ODN and

i.n.  $\beta$ -gal/CT immunization was evaluated. Splenocytes from immunized mice were incubated with  $\beta$ -gal, and culture supernatants were assayed for the production of IFN $\gamma$  and IL-4, cytokines classically associated with Th<sub>1</sub> and Th<sub>2</sub> immunity, respectively (Fig. 2) (MOSMANN and COFFMANN 1989). Splenocytes



**Fig. 2A,B.** Splenocyte antigen-induced cytokine profiles. Mice received a single immunization with βgalactosidase (β-gal; 50µg) alone, with oligodeoxynucleotide immunostimulatory sequence (ISS-ODN; 50µg), mutated oligodeoxynucleotide (50µg), or cholera toxin (CT; 10µg) via intranasal (i.n.) or intradermal (i.d.) routes. The sequences of the oligodeoxynucleotides are provided in the Fig. 1 legend. Splenocytes were harvested from sacrificed mice during week 7, cultured in media with or without β-gal (10µg/ml), and 72-h supernatants were assayed by enzyme-linked immunosorbent assay. Splenocytes cultured without β-gal produced no detectable interferon  $\gamma$  (IFN $\gamma$ ) or interleukin 4 (IL-4; data not shown). Results represent the mean for four mice in each group, and similar results were obtained in two other independent experiments. *Error bars* reflect standard errors of the means. A IFN $\gamma$  levels. IFN $\gamma$ levels were equivalent in i.n. and i.d. β-gal/ISS-ODN versus β-gal/CT mice). B IL-4 levels. IL-4 levels above background were detected only in mice immunized with i.n. β-gal/CT (P = 0.04 versus i.n. β-gal/ ISS-ODN) from mice immunized with  $\beta$ -gal and ISS-ODN via i.n. and i.d. routes produced similarly high levels of IFN $\gamma$  but no detectable IL-4. In contrast, i.n. vaccination with  $\beta$ -gal and CT led to low-level splenocyte production of IFN $\gamma$  but significant levels of IL4. Intranasal immunization with  $\beta$ -gal alone or with M-ODN led to



**Fig. 3A,B.** Serum immunoglobulin G (IgG) subclass profiles. Mice received a single immunization with β-galactosidase (β-gal; 50µg) alone, with oligodeoxynucleotide immunostimulatory sequence (ISS-ODN; 50µg), mutated oligodeoxynucleotide (50µg), or cholera toxin (CT; 10µg) via intranasal (i.n.) or intradermal (i.d.) routes. The sequences of the oligodeoxynucleotides are provided in the Fig. 1 legend. Serum was collected at 2, 4, and 7 weeks from immunized mice and assayed by enzyme-linked immunosorbent assay. Results represent mean values for four mice per group, and *error bars* reflect standard errors of the means. Results are representative of three similar and independent experiments. A Serum IgG2a. Serum IgG2a levels were equivalent in i.n. and i.d. β-gal/ISS-ODN but statistically higher than in other immunization groups at 7 weeks (P = 0.005 for i.n. β-gal/ISS-ODN - but statistically lower than in i.n. β-gal/CT-immunized mice at all time points (P = 0.003, P = 0.02, and P = 0.02 for i.n. β-gal/ISS-ODN versus β-gal/CT i.n. β-gal/ISS-ODN versus β-gal/CT mice at 2, 4, and 7 weeks, respectively)



**Fig. 4.** Serum immunoglobulin E (IgE). Mice received a single immunization with  $\beta$ -galactosidase ( $\beta$ -gal; 50µg) alone, with oligodeoxynucleotide immunostimulatory sequence (ISS-ODN; 50µg), mutated oligodeoxynucleotide (50µg), or cholera toxin (CT; 10µg) via intranasal (i.n.) or intradermal routes. The sequences of the oligodeoxynucleotides are provided in the Fig. 1 legend. Serum was collected from mice during week 7 and anti- $\beta$ -gal IgE levels were assayed by enzyme-linked immunosorbent assay. Results represent mean values for four mice per group, and *error bars* reflect standard errors of the means. Results are representative of three similar and independent experiments. Serum IgE was only detectable in i.n.  $\beta$ -gal/CT immunized mice (P = 0.004 for i.n.  $\beta$ -gal/CT- versus i.n.  $\beta$ -gal/ISS-ODN-immunized mice)

much lower or undetectable cytokine production in splenocytes (HORNER et al. 1998).

IFN $\gamma$  is a switch factor for IgG2a production, while IL-4 is a switch factor for IgG1 and IgE (COFFMAN and MOSMANN 1988). Given the previously noted splenic cytokine profiles, it would, therefore, be predicted that i.n.  $\beta$ -gal/ISS-ODN co-administration would lead to higher IgG2a and lower IgG1 and IgE levels than i.n.  $\beta$ -gal/CT immunization. Indeed, i.n. and i.d.  $\beta$ -gal/ISS-ODN-immunized mice produced equivalent Th<sub>1</sub>-biased serum antibody responses, while i.n.  $\beta$ -gal/CT vaccination led to a Th<sub>2</sub>-biased antibody profile. As can be seen in Fig. 3, both i.n. and i.d. vaccination with ISS-ODN and  $\beta$ -gal led to high levels of IgG2a and low levels of IgG1 when compared to i.n.  $\beta$ -gal/CT vaccination (HORNER et al. 1998). In addition, IgE production in i.n.  $\beta$ -gal/ISS-ODN- and i.n.  $\beta$ -gal/CT-immunized mice was evaluated. As expected and seen in Fig. 4, only  $\beta$ -gal/CT-immunized mice produced IgE (previously unpublished data).

Cumulatively, these observations demonstrate that, under the conditions employed in this series of experiments, i.n. and i.d. delivery of antigens with ISS-ODN leads to equivalent systemic  $Th_1$ -biased cytokine and antibody profiles, while i.n.  $\beta$ -gal/CT co-administration leads to a  $Th_2$ -biased immune response that includes the production of IgE. Considered in conjunction with the IgA data previously presented, the data further demonstrate that induction of mucosal IgA can occur in the context of both  $Th_1$ - and  $Th_2$ -biased systemic immune responses.



**Fig. 5.** Splenic cytotoxic T-lymphocyte (CTL) responses. Mice received a single immunization with β-galactosidase (β-gal; 50µg) alone, with oligodeoxynucleotide immunostimulatory sequence (ISS-ODN; 50µg), mutated oligodeoxynucleotide (50µg), or with cholera toxin (CT; 10µg) via intranasal (i.n.) or intradermal (i.d.) routes. The sequences of the oligodeoxynucleotides are provided in the Fig. 1 legend. Splenocytes were harvested from mice at week 7. They were cultured for 5 days with interleukin 2, β-gal peptide, and mitomycin-C-treated splenocytes from naïve mice, and CTL responses were subsequently determined. Results represent mean values for four mice per group, and *error bars* reflect standard errors of the means. Results are representative of three similar and independent experiments. CTL responses were equivalent in i.n. and i.d. β-gal/ISS-ODN immunized mice at all enzyme:target (E:T) ratios but were statistically higher then in i.n. β-gal/CT-immunized mice at E:T ratios of 25:1 and 5:1, respectively)

#### 3.3 Mucosal Immunization with ISS-ODN Induces a Splenic CTL Response

Although development of antigenspecific CTL activity is associated with Th<sub>1</sub>biased immunity, not all Th<sub>1</sub>-biased immune responses include the development of cytotoxic T cells (STAATS and MCGHEE 1996; ADA and MCELRATH 1997). Therefore, the ability of i.n. co-delivery of  $\beta$ -gal and ISS-ODN to induce a CTL response was evaluated. As demonstrated in Fig. 5, mice immunized with  $\beta$ -gal and ISS-ODN by either the i.n. or i.d. route displayed vigorous splenic CTL activity. However, i.n.  $\beta$ -gal/CT immunization resulted in a poor CTL response. Likewise, i.n. immunization with  $\beta$ -gal alone or with M-ODN led to poor or undetectable CTL responses (HORNER et al. 1998). Considered in conjunction with the splenic cytokine and serum antibody responses previously presented, this data confirms that i.d. and i.n. vaccination with  $\beta$ -gal and ISS-ODN leads to Th<sub>1</sub>-biased systemic immune responses of equivalent magnitude. In addition, the CTL assay results further demonstrate the dichotomy between the Th<sub>1</sub>- and Th<sub>2</sub>-biased systemic immune responses seen when  $\beta$ -gal is co-delivered i.n. with ISS-ODN and CT, respectively.

# **4** Discussion

In summary, these results demonstrate that ISS-ODN is a potent mucosal adjuvant. Intranasal delivery of antigen with either ISS-ODN or CT leads to an equivalent and vigorous mucosal IgA response, while i.d. co-delivery of antigen with ISS-ODN does not lead to mucosal IgA production. However, under the experimental conditions employed, i.d. and i.n. vaccination with target antigen and ISS-ODN induces equivalent systemic Th<sub>1</sub>-biased immune responses characterized by high levels of antigen specific IFN $\gamma$  but no IL-4 production from cultured splenocytes, high IgG2a and low IgG1 and IgE serum concentrations, and vigorous CTL responses. In contrast, i.n. co-delivery of antigen with CT leads to a Th<sub>2</sub>-biased systemic immune response characterized by low IFN $\gamma$  but substantial IL-4 production from in vitro antigen-stimulated splenocytes, high IgG1 and IgE and low IgG2a serum concentrations, and a poor CTL response. While initial work utilized BALB/c mice and  $\beta$ -gal as antigen, subsequent studies have demonstrated the generalizability of this phenomenon to other mouse strains and protein antigens (unpublished data).

Consistent with our findings, Dr. Moldoveanu and colleagues have shown that ISS-ODN is an effective mucosal adjuvant for IgA production against a formalin-inactivated influenza vaccine (MOLDOVEANU et al. 1998). Interestingly, under the experimental conditions utilized in these studies, the inactivated influenza vaccine stimulated secretory IgA production even without an adjuvant. The multimeric nature of the antigen preparation and the epithelial-binding properties of the influenza virus may help explain this observation (SMITH 1998). These investigators were unable to elicit a CTL response against influenza virus. Formalin inactivation leads to cross-linking of protein. This, in turn, could limit peptide processing and presentation by major histocompatibility complex molecules, leading to relatively weak T-cell responses, as suggested by the authors (MOLDOVEANU et al. 1998). McCluskie and colleagues have also recently shown that ISS-ODN serves as an effective mucosal adjuvant for the induction of an IgA response, using the hepatitis-B surface antigen (McCluskie and Davis 1998). Unlike the influenza vaccine, but consistent with antigens used in our studies, hepatitis B surface antigen did not induce an IgA response when mucosally delivered without adjuvant. However, in contrast with our findings, these investigators were also unable to elicit a significant CTL response if ISS-ODN was the only adjuvant used to immunize mice. Our experience suggests that the low doses of antigen and ISS-ODN used in these experiments may have played a role in the poor CTL responses reported.

The fact that equivalent mucosal IgA levels can develop in the context of  $Th_1$ and  $Th_2$ -biased systemic immune responses with i.n.  $\beta$ -gal/ISS-ODN and  $\beta$ -gal/CT immunizations, respectively, is consistent with other published results. Marinaro and colleagues recently demonstrated that oral delivery of tetanus toxoid with CT led to mucosal IgA production in conjunction with a  $Th_2$ -biased immune profile. However, when tetanus toxoid and CT were co-administered with oral IL-12, the systemic immune response was skewed toward a  $Th_1$  phenotype, while mucosal IgA production was unaffected (MARINARO et al. 1997). Taken together, these findings document that synthesis of mucosal IgA can occur in the context of both  $Th_1$ - and  $Th_2$ -biased systemic immunity.

Mucosal IgA and CTL responses are known to provide protection against a number of infectious agents. Human immunodeficiency virus is but one important example (Gallichan et al. 1993; CZERKINSKY and HOLMGREN 1995; STAATS and MCGHEE 1996; ADA and MCELRATH 1997; LETVIN 1998; VANCOTT et al. 1998). There are a number of strategies available for the development of vaccines that induce these immune parameters. However, none appear to be globally applicable. Live attenuated vaccines produce robust immune responses, including mucosal IgA synthesis and CTL activity. Unfortunately, difficulty in attenuating many pathogens and the risk of iatrogenic disease limits the use and development of live attenuated vaccines (CZERKINSKY and HOLMGREN 1995; GUPTA and SIBER 1995; STAATS and McGHEE 1996; LETVIN 1998). However, recombinant proteins from infectious agents are generally safe but induce relatively poor immune responses and generally induce none when delivered to mucosal surfaces. However, mucosal adjuvants can improve immune responses towards co-administered protein antigens substantially (CZERKINSKY and HOLMGREN 1995; GUPTA and SIBER 1995; STAATS and MCGHEE 1996).

CT is an extremely potent mucosal adjuvant but is inherently toxic and induces a Th<sub>2</sub>-biased immune response that can include the development of IgE and consequent allergic sensitization toward the target antigen (SNIDER et al. 1994; MARINARO et al. 1995; personal observations). At present, such toxicity and other technical problems have kept many adjuvants from becoming available for use in humans (GUPTA and SIBER 1995). Alum is essentially the only adjuvant in clinical use today. It is relatively weak, does not work with a number of antigens, does not induce CTL activity, and because it must be delivered systemically, does not induce mucosal IgA (GUPTA and SIBER 1995). A safe and effective mucosal adjuvant would be of great value in the development of better vaccines. ISS-ODN is a potent adjuvant which works with a wide range of protein antigens and generally induces a Th<sub>1</sub>-biased immune response with CTL activity (SATO et al. 1996; ROMAN et al. 1997; DAVIS et al. 1998; ZIMMERMANN et al. 1998). Intranasal and i.d. administration of protein with ISS-ODN leads to vigorous and equivalent Th<sub>1</sub>-biased systemic immune responses, while only i.n. delivery induces a mucosal immune response (HORNER et al. 1998). Therefore, i.n. delivery of relevant antigens with ISS-ODN may well prove superior to i.d. delivery for the induction of protective immunity against mucosal pathogens.

# **5** Future Prospects

ISS-ODN are easy to manufacture, stable, and lack identified toxicity at immunogenic doses in mice and primates (unpublished observations). Additionally, use of antisense ODN in monkeys and human clinical trials have demonstrated no significant toxicity with daily doses of up to fivefold more per kilogram then those used in the present study (WEBB et al. 1997). Moreover, we and others have shown that human and mouse immunocytes display similar immunologic responses to ISS-ODN (BALLAS et al. 1996; ROMAN et al. 1997). Because ISS-ODN can be utilized as an adjuvant for both mucosal and systemic Th<sub>1</sub>-biased immune responses toward simple monomeric protein antigens, it may well prove to be a valuable reagent for clinical applications. Thus, future investigations in primates and humans are needed to establish whether ISS-ODN has utility in the development of vaccines that induce protective immune responses against mucosal pathogens.

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# CpG DNA Switches on Th1 Immunity and Modulates Antigen-Presenting Cell Function

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# **1** Introduction

DNA containing CpG motifs, which are present in bacterial DNA but suppressed in mammalian DNA, can stimulate immune cells and modulate immune responses. A CpG motif consists of an unmethylated CpG dinucleotide flanked by two 5' purines and two 3' pyrimidines (KRIEG et al. 1995). CpG motifs are similar to palindromic sequences identified earlier in bacille Calmette-Guerin DNA, which increase the cytolytic function of natural killer (NK) cells (YAMAMOTO et al. 1992), and immunostimulatory sequences, which increase the immunogenicity of DNA vaccines (SATO et al. 1996). Bacterial DNA or oligodeoxynucleotides (ODN) containing CpG motifs can stimulate B cells (KRIEG et al. 1995; SUN et al. 1997), macrophages (STACEY et al. 1996; SPARWASSER et al. 1997b), dendritic cells (JAKOB et al. 1998; SPARWASSER et al. 1998), and NK cells (YAMAMOTO et al. 1992; BALLAS et al. 1996).

ODN containing CpG motifs (CpG ODN) have important potential utility as vaccine adjuvants due to their ability to stimulate cytokines (including Th1-associated cytokines), which may enhance vaccine efficacy. In vitro, CpG ODN stimulate splenocytes to produce IL-6, IL-12, and IFN- $\gamma$  (KLINMAN et al. 1996). In particular, macrophages produce IL-12 in response to CpG ODN, and production of IFN- $\gamma$  by NK cells is increased by CpG ODN both directly and indirectly via macrophage-derived IL-12 produced in response to CpG ODN (COWDERY et al. 1996; HALPERN et al. 1996; CHACE et al. 1997). These observations suggest that CpG ODN could affect the differentiation of T-cell responses by controlling the

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cytokine milieu present during the generation of T-cell responses. Experiments that address this hypothesis are discussed below, in Sect. 3.

In addition to their effects via modulation of the cytokine milieu, CpG DNA may influence T-cell responses by altering antigen processing and presentation functions of APCs, such as dendritic cells, macrophages, and B cells. Stimulation of antigen presenting cells (APCs) with CpG DNA can alter the expression of MHC molecules and co-stimulator molecules (DAVIS et al. 1998; JAKOB et al. 1998; SPARWASSER et al. 1998; ASKEW et al. 1999; CHU et al. 1999). Our recent studies of the effects of CpG ODN on antigen-processing functions of macrophages and dendritic cells are discussed below, in Sect. 4.

# 2 Th Subsets and Their Importance in Disease States

The CD4 Th cell plays a crucial role in the adaptive immune response, providing activation and differentiation signals in the form of cytokines and direct cell–cell contact to B cells,  $CD8^+$  T cells, and dendritic cells. The ability of Th cells to influence multiple aspects of immune responses is facilitated by the existence of different Th subsets with particular effector functions. One current delineation of different Th subsets is the Th1/Th2 model, which divides mature Th cells by the types of cytokines that they secrete (MOSMANN and COFFMAN 1989; ABBAS et al. 1996). Specifically, Th1 cells secrete cytokines, such as IFN- $\gamma$ , IL-2, and lymphotoxin, while Th2 cells secrete IL-4, IL-5, IL-6, IL-10, and IL-13. Certain immune responses and disease states may involve predominance of Th1 or Th2 effects. Much evidence has been generated to support this Th-cell dichotomy, although it is clear that this is a simplified model, and in many situations a mixture of T-cell subsets may occur, with secretion of cytokines associated with both Th1 and Th2 cells (KELSO 1995).

Th1 cells play an important role in responses to microbes. The secretion of IFN- $\gamma$  by Th1 cells allows this subset to activate and increase the microbicidal activity of macrophages and to recruit other inflammatory cells. IFN- $\gamma$  also promotes the development of cytotoxic-T-lymphocyte responses and Ig class switching to the IgG2a and IgG3 isotypes (in mice). These isotypes are effective in complement fixation and opsonization, thus facilitating the antimicrobial response. In contrast, IL-4-secreting Th2 cells are important mediators of B-cell differentiation and cause Ig class switching to the IgG1 isotype (in mice). IgG1, while less effective in complement fixation, is the predominant isotype in antibody responses against protein antigens. The production of IL-4 by Th2 cells also promotes secretion of IgE, while IL-5 from Th2 cells causes eosinophil activation. Thus, Th2 responses promote hypersensitivity responses. Also, some of the cytokines produced by Th2 cells, such as IL-10, are important in negatively regulating inflammatory responses by suppressing macrophage activation.

The development of Th cells towards either subset is mainly dependent upon the cytokine environment at the time of activation of uncommitted Th cells (also called Th0 cells). Other factors, such as antigen dose and particular co-stimulatory molecules involved in Th activation, may also play a role, but the overall importance of these factors is uncertain. The presence of IL-12 at the time of antigen presentation, however, has been clearly shown to stimulate the differentiation of Th1 cells, while IL-4 is the predominant cytokine which promotes the differentiation of Th2 cells (O'GARRA 1998). The cytokines produced by one subset are often inhibitory toward the development or activity of the other. For example, IL-4 and IL-10 are capable of inhibiting IL-12 production by APCs and thus inhibit the development of Th1 cells (O'GARRA 1998).

Because the effector functions of Th1 and Th2 cells are varied and often antagonistic toward those of the opposite subset, it would be desirable, in many disease states, to be able to direct immune responses to differentiate toward one subset versus the other (immune deviation). For example, in many infections involving intracellular microbes, a Th1-dominated response is crucial for resistance. In contrast, a Th2-dominated response may be more protective during infections with some parasites and helminths. In addition, various autoimmune diseases, such as multiple sclerosis and insulin-dependent diabetes mellitus, have been proposed to be a result of Th1-driven autoimmune responses, and patients with such diseases may benefit from therapy that promotes development of opposing Th2 cells (LIBLAU et al. 1995). Th2 cells are also implicated in disease states, specifically atopic disease and asthma. Clinically, immune deviation towards Th1 responses might be useful in alleviating allergic disease (FINKELMAN 1995).

# 3 CpG ODN Act as Th1-Directing Adjuvants

Adjuvants are used to enhance immune responses to immunogens. Different adjuvants have been found to direct Th-subset differentiation in different ways. Alum, an adjuvant that is used for clinical purposes in humans, has been noted to induce weak Th2 responses in mice (RAz et al. 1996; ADA and RAMSAY 1997). Immunization with antigen in incomplete Freund's adjuvant (IFA) also generates Th2-type responses, with antigen-specific memory T cells secreting IL-5, while immunization with complete Freund's adjuvant (CFA) generates Th1 responses with T cells that secrete IFN- $\gamma$  (FORSTHUBER et al. 1996). IL-12 has also been used as an adjuvant for Th1 responses in experimental systems (BLISS et al. 1996). Some of these adjuvants (such as IFA and CFA), however, are difficult to use in human vaccines because of their toxic potential.

In the interest of anti-microbial vaccine design, there is a need for Thldirecting adjuvants that can elicit protective cellular immunity while being safe for human use. CpG ODN are potential vaccine adjuvants. CpG ODN are potent activators of macrophages and dendritic cells, causing to them to secrete IL-12 and other cytokines (see above and other chapters in this volume). The ability of CpG ODN to induce IL-12 suggested the hypothesis that CpG ODN would act as Thldirecting adjuvants in vivo, and a murine experimental system was developed to test this hypothesis.

To test the ability of CpG ODN to direct specific Th1 responses in vivo, BALB/ c mice were injected with hen-egg lysozyme (HEL) in IFA with or without the addition of CpG ODN or non-CpG ODN (see Table 1 for ODN sequences). Figure 1 shows that immunization with HEL in IFA alone produced a Th2 response against HEL, characterized by a high level of IL-5 secretion with little or no IFN- $\gamma$ production by splenocytes upon in vitro antigen challenge (CHU et al. 1997), consistent with prior observations (FORSTHUBER et al. 1996). However, the addition of CpG ODN 1826 to the HEL/IFA immunization caused a marked increase in antigen-specific IFN- $\gamma$  production by splenocytes and a decrease in IL-5 secretion compared to the response induced by HEL/IFA alone (CHU et al. 1997) (Fig. 1). The in vitro splenocyte response to HEL was shown to be CD4<sup>+</sup> T-cell dependent, as addition of anti-CD4 blocking antibodies abrogated cytokine production. In addition to CpG ODN 1826, two other CpG ODN 1760 and 1585, had similar effects. In

ODN	Sequence <sup>a</sup>	Motif	
1826	TCCATGACGTTCCTGACGTT	CpG	
1745	TCCAATGAGCTTCCTGAGTCT	non-CpG	
1760	ATAATCGACGTTCAAGCAAG	CpG	
1908	ATAATAGAGCTTCAAGCAAG	non-CpG	
1585	GGGGTCAACGTTGAGGGGGG	CpG	
1972	GGGGTCT <u>GTGCTT</u> TTGGGGGGG	non-CpG	

Table 1. Sequences of synthetic oligodeoxynucleotides (ODNs)

<sup>a</sup> The CpG motifs or corresponding non-CpG motifs are underlined.



**Fig. 1A,B.** Immunization with CpG oligodeoxynucleotide (ODN) as an adjuvant induces a T helper cell 1 (Th1)-dominated cytokine profile, with increased interferon  $\gamma$  (IFN- $\gamma$ ) and decreased interleukin 5 (IL-5). BALB/c mice were injected i.p. with complete Freund's adjuvant/hen-egg lysozyme (HEL) (a control for a Th1-dominated response), incomplete Freund's adjuvant/HEL (a control for a Th2-dominated response), or IFA/HEL with 100µg (**A**) or 30µg (**B**) of ODN (non-CpG ODN 1745 or CpG ODN 1826). After 3 weeks, splenocytes were isolated and incubated with HEL (*closed circles*) or medium alone (*open circles*). Cytokine production by individual cells was assessed by an enzyme-linked immunosorbent assay spot assay for IFN- $\gamma$  (**A**) or IL-5 (**B**). Each *point* represents one mouse; *horizontal bars* indicate the mean of points for each group of mice. Figure adapted from Chu et al. (CHU et al. 1997) and reproduced with permission from the Journal of Experimental Medicine

contrast, addition of control non-CpG ODN (1982 1908, or 1972; Table 1) to the HEL/IFA immunization did not alter the cytokine profile of HEL-specific memory cells, indicating the importance of the CpG sequence in generating a Th1 response.

These immunization studies showed that CpG ODN are very potent Th1 adjuvants. Optimal induction of Th1 responses was seen with  $30\mu g$  of CpG ODN per mouse, and doses of as little as  $3-10\mu g$  of ODN per mouse were also effective at inducing Th1 differentiation. The levels of antigen-specific IFN- $\gamma$  secretion generated by CpG ODN were relatively high and even exceeded those induced by immunization with HEL in CFA.

Immunization with CpG ODN also induced changes in humoral responses consistent with enhanced levels of antigen-specific IFN- $\gamma$  production. Immunization with HEL in IFA, IFA plus non-CpG ODN, and IFA plus CpG ODN all produced similar total levels of HEL-specific antibodies (Fig. 2A). In contrast, immunization with HEL in IFA plus CpG ODN induced production of HELspecific IgG2a, an isotype associated with Th1 responses, whereas immunization with HEL in IFA plus non-CpG ODN did not induce HEL-specific IgG2a (Fig. 2B). The levels of IgG2a, similar to the in vitro splenocyte IFN- $\gamma$ responses, were consistently higher with CpG ODN than with CFA as an adjuvant (CHU et al. 1997).

Thus, CpG ODN act as immunomodulators and produce strong Th1 responses against co-administered antigens, manifested by both cytokine and antibody isotype profiles. The addition of non-viable *Mycobacterium tuberculosis* to IFA, producing CFA, is an established method used to induce Th1 responses. In our



**Fig. 2A,B.** T helper cell 1 (Th1)-associated antigen-specific immunoglobulin G2a (IgG2a) responses are induced by immunization of BALB/c mice with incomplete Freund's adjuvant (IFA)/hen-egg lysozyme (HEL)/CpG oligodeoxynucleotide (ODN) but not IFA/HEL/non-CpG ODN. Mice were immunized as in Fig. 1. Sera were collected from mice 15–18 days after injection and assayed by enzyme-linked immunosorbent assay for: anti-HEL total antibody response (A) and anti-HEL IgG2a, an isotype associated with Th1-dominated responses (B). Figure adapted from Chu et al. (CHU et al. 1997) and reproduced with permission from the Journal of Experimental Medicine

studies, preparations of IFA plus CpG ODN caused more effective induction of Th1 responses than did CFA. Strong Th1 responses were seen after a single immunization with CpG ODN. Furthermore, the Th1 adjuvant activity of CpG ODN was observed in both BALB/c mice, which have a Th2 bias in some experimental systems, and B10.D2 mice, which show stronger Th1 responses.

In addition to the use of immunomodulatory substances, many adjuvants, such as alum and IFA, create an antigen depot that is critical for retention of antigens and immunomodulators, allowing for persistent stimulation of the immune system. The IFA in these studies provided an important depot function that enhanced the adjuvant effect. In comparison, an injection of antigen plus CpG ODN in saline generated a weaker Th1 response, as measured by antigen-specific IFN- $\gamma$  secretion in vitro, than immunization with antigen plus CpG ODN in IFA, although both adjuvant preparations similarly induced antigen-specific IgG2a (R.S. Chu, unpublished data). Future vaccines using CpG ODN as Th1-directing adjuvants may include less toxic components, such as biodegradable oils or alum, which can generate an antigen depot in vivo. It has already been shown that the use of alum in conjunction with CpG ODN is effective in generating strong humoral responses against hepatitis-B surface antigen, including induction of specific IgG2a (DAVIS et al. 1998). Alternatively, multiple injections of CpG ODN may obviate the need for an additional depot-forming adjuvant. Such an approach was used by Roman et al. in the generation of Th1-biased humoral and cellular responses against proteins encoded by DNA vaccines (Roman et al. 1997).

One concern for vaccine adjuvants is toxic potential. Repeated administration of large doses of CpG ODN (150mg/kg) can cause mortality in mice (SARMIENTO et al. 1994), and the induction of tumor necrosis factor  $\alpha$  (TNF- $\alpha$ ) secretion by CpG ODN has been shown to cause fatal shock in mice that have been previously sensitized with D-galactosamine (SPARWASSER et al. 1997a). However, in our studies using low doses of ODN (30–100µg/mouse), we did not observe any significant toxicity in mice treated with CpG ODN. There were no observable changes in mouse appearance, behavior, or body weight. Administration of CpG ODN alone, at the doses used in our studies, produced mild splenomegaly and hyperplasia of draining lymph nodes that were reversible within 10–14 days, while a single injection of up to 1mg of CpG ODN does not alter mouse feeding, grooming, physical activity, or behavior (A.M. Krieg, personal communication). Thus, CpG ODN provide potent adjuvant activities at doses that do not produce significant toxicity.

Another concern regarding CpG ODN as Th1-directing adjuvants is the potential induction of Th1-dominated autoimmunity, which has been observed with other manipulations that enhance Th1 responses. For example, lipopolysac-charide (LPS)- or CpG-DNA-induced IL-12 production causes in vitro differentiation of antigen-specific lymph-node cells that produce IFN- $\gamma$ . When injected into normally experimental allergic encephalitis (EAE)-resistant mice, these cells are subsequently able to cause EAE, a disease associated with autoreactive Th1 effector cells (SEGAL et al. 1997). When CpG ODN are used in humans, their potent Th1 adjuvant activity will warrant careful monitoring of patients prone to Th1-dominated autoimmune diseases.

There are many exciting potential clinical applications for CpG ODN. The use of CpG ODN to produce protective Th1 responses has been explored in disease models involving Th1 or Th2 subsets, such as asthma (KLINE et al. 1998) and murine leishmaniasis (ZIMMERMAN et al. 1998), as discussed elsewhere in this volume. In addition to promoting Th1 differentiation, CpG ODN have general adjuvant effects in other systems, increasing total specific antibody levels (DAVIS et al. 1998; SUN et al. 1998) and augmenting cytotoxic-T-lymphocyte responses after immunization with protein alone or with protein encapsulated in liposomes (LIPFORD et al. 1997; DAVIS et al. 1998).

# 4 Modulation of MHC-II Antigen Processing and Presentation by CpG ODN

The adjuvant activities of CpG DNA include activation of cells via both direct and indirect mechanisms. The direct effects of CpG DNA result from signals transduced in cells, such as macrophages, dendritic cells, B cells, and NK cells, which directly recognize CpG DNA (YAMAMOTO et al. 1992; STACEY et al. 1996; SPARWASSER et al. 1997b, 1998; JAKOB et al. 1998; YI and KRIEG 1998). The direct effects of CpG DNA include the induction of cytokines, including IL-6, IL-12, and IFN- $\gamma$ , which are expressed by macrophages, dendritic cells, B cells, and NK cells. These cytokines then act on many cell types to promote the indirect effects of CpG DNA, such as enhancement of T-cell responses and Th1 differentiation, as discussed above. Cytokine-induced effects may occur via signaling in T cells to alter T-cell differentiation or via activation of APCs to enhance antigen presentation and T-cell responses.

Many questions about the impact of CpG DNA on antigen presentation remain. In vivo, CpG DNA may indirectly influence antigen presentation by inducing cytokines (IFN- $\gamma$ ) that are known to activate APCs. In addition, CpG DNA may directly signal APCs to alter antigen processing or presentation. Induction of an increase in processing and presentation of exogenous antigen may contribute to the adjuvant activity of CpG DNA. Such an upregulation of antigen presentation would benefit the host in the context of bacterial infection, allowing more bacterial epitopes to be presented to T cells. However, the direct effects of CpG DNA on antigen processing and presentation remain poorly characterized, and this topic has been the focus of our recent studies.

In order to examine the direct effects of CpG DNA on intracellular antigen processing, we studied the in vitro effects of CpG ODN on processing and presentation of exogenous antigens by isolated macrophages. Macrophages were isolated from peritoneal exudate cells by adherence and were treated with CpG ODN overnight (18–24h). The macrophages were then pulsed briefly (1–2h) with protein antigen and fixed with paraformaldehyde. Antigen processing and presentation were measured using T-cell hybridomas specific for HEL or bovine ri-

bonuclease (RNase), including T hybridomas that recognize the following complexes: RNase(42–56):I-A<sup>k</sup>, RNase(90–105):I-E<sup>k</sup>, HEL(34–45):I-A<sup>k</sup>, and HEL(48–61): I-A<sup>k</sup>.

After overnight treatment with CpG ODN ( $1\mu$ g/ml), macrophage processing of all four epitopes was actually inhibited to varying degrees rather than upregulated, as measured by IL-2 production by the antigen-specific T-cell hybridomas (CHU et al. 1999). Interestingly, similar results were seen when the macrophages were treated with another bacterial product, LPS (100ng/ml), but not with non-CpG-ODN controls.

Additional studies examined the effects of CpG ODN on binding and presentation of exogenous peptides, which do not require intracellular protein antigen processing. The presentation of exogenous peptides by CpG ODN- or LPS-treated macrophages was also decreased compared to untreated cells, but to a lesser extent than presentation of peptides derived from processed whole antigen. These results suggested that CpG ODN and LPS downregulate expression of major histocompatibility complex class II (MHC-II) molecules. Surface I-A<sup>k</sup> levels were analyzed by flow cytometry and were found to be decreased after treatment of macrophages with CpG ODN or LPS, but not after treatment with non-CpG ODN. Northern analysis revealed that the level of I-A<sup>k</sup> mRNA was lower in macrophages that were treated with CpG ODN than in untreated macrophages. Thus, the downregulation of MHC-II expression mediated by CpG ODN occurred via transcriptional regulation (or possibly by regulation of mRNA stability). In contrast, the endocytic function of macrophages, as measured by uptake of fluoresceinated dextran, was not specifically affected by CpG ODN (CHU et al. 1999). In summary, CpG ODN and LPS caused a downregulation in macrophage antigen processing and presentation, which was primarily mediated by a decrease in the synthesis of MHC-II molecules. These results are reminiscent of previous studies demonstrating an inhibitory effect of LPS on MHC-II expression by IFN-y-stimulated macrophages (STEEG et al. 1982; KOERNER et al. 1987; SICHER et al. 1994). The effects of LPS on MHC-II synthesis were proposed to be a result of increased levels of prostaglandin (STEEG et al. 1982) or nitric oxide (SICHER et al. 1994). These substances may also be involved in the effects on MHC-II synthesis mediated by CpG ODN.

It is unclear how the downregulation of MHC-II expression and antigen processing produced by CpG ODN in vitro relates to either the actual in vivo effects of bacterial DNA in the physiological context of an infection or the in vivo effects of CpG DNA as a vaccine adjuvant. There is a significant delay between the exposure to CpG DNA and the onset of inhibition of antigen processing, and bacterial antigens could be effectively processed during this interval. In addition, our in vitro system examines the direct effects of CpG ODN on processing and presentation functions of isolated macrophages. In vivo, the interactions of multiple cell types must be considered, and the in vivo effects of CpG DNA may include indirect effects on macrophages via stimulation of other cell types, e.g. NK cells, to produce IFN- $\gamma$ . For example, the ability of CpG DNA to upregulate transcription of inducible nitric oxide synthase in macrophages is detected only when the macrophages are first treated with IFN- $\gamma$  (STACEY et al. 1996). Similarly, other indirect effects may occur in vivo to counteract the direct inhibitory effects of CpG DNA on antigen processing. Alternatively, an inhibitory effect may be the predominant effect of CpG ODN on macrophage antigen processing in vivo. As such, the macrophage response to bacterial DNA may have evolved to produce a macrophage activation state characterized by other beneficial anti-microbial functions, such as cytokine secretion and microbicidal activity, rather than an upregulation in antigen presentation.

Even if macrophage antigen processing is not enhanced by CpG ODN, it is necessary to consider the multiplicity of APC types available in vivo. Some of the adjuvant properties of CpG ODN may still be explained by enhanced antigen processing and presentation by other types of APCs. In fact, previous studies have shown that CpG ODN can increase expression of co-stimulator molecules on B cells (DAVIS et al. 1998) and increase expression of both MHC-II and co-stimulator molecules on dendritic cells (JAKOB et al. 1998; SPARWASSER et al. 1998). We have recently investigated the effects of CpG ODN on the antigen processing and presentation functions of dendritic cells, and these studies have shown effects quite different from those we have observed in macrophages.

Dendritic cells exhibit extensive changes in their ability to process and present antigen as a function of their maturation. Immature dendritic cells reside in the periphery, express a high level of endocytic activity; however, they express lower levels of MHC-II than do mature dendritic cells, and they lack co-stimulatory molecules required for antigen presentation to primary T cells (CELLA et al. 1997a,b; PIERRE et al. 1997; BANCHEREAU and STEINMAN 1998). Upon introduction of a maturation signal, such as that initiated by bacteria or bacterial products (LPS or CpG DNA) (CELLA et al. 1997a), dendritic cells migrate to lymphoid tissues and mature. Maturation of dendritic cells involves decreased endocytic activity, decreased expression of the mannose receptor, FcR (CD32 and CD23), CD14, and intracellular MHC-II molecules, increased expression of cell surface MHC-I and MHC-II molecules, and increased expression of co-stimulatory molecules, such as CD40, CD80, and CD86 (Sallusto et al. 1995; Henderson et al. 1997; Winzler et al. 1997; JAKOB et al. 1998; RESCIGNO et al. 1998; SPARWASSER et al. 1998). Mature dendritic cells also have increased production of cytokines, such as IL-1, IL-6, IL-10, IL-12, and TNF (HENDERSON et al. 1997; SPARWASSER et al. 1997a; JAKOB et al. 1998; RESCIGNO et al. 1998).

Recent studies by Askew et al. (manuscript in preparation) have examined the effects of CpG ODN on the maturation of bone-marrow-derived dendritic cells and their ability to process and present protein antigens. Murine bone-marrow-derived dendritic cells were cultured in a medium containing GM-CSF with or without the addition of CpG ODN. CpG ODN induced the differentiation of cells with dendritic morphology and increased expression of MHC-II, CD80, CD86, and CD40. In addition, CpG ODN enhanced the processing of HEL or RNase when the antigen was added at the same time as the CpG ODN. After exposure to CpG ODN for 24–48h, however, dendritic cells displayed decreased antigen-processing function. More detailed antigen-processing studies suggest that the long-term

inhibitory effects on dendritic cell antigen processing particularly involve an inhibition of processing mechanisms that use newly synthesized MHC-II molecules, whereas processing mechanisms that use recycling MHC-II appear to be maintained in mature dendritic cells. In summary, CpG ODN induced dendritic cell maturation that was accompanied by a transient increase in antigen-processing function, followed by a decrease in certain antigen-processing activities. However, the mature dendritic cells maintained high levels of expression of MHC-II and co-stimulator molecules and had the ability to present antigens that were processed earlier. Thus, the ability of CpG ODN to transiently increase antigen-processing activity by dendritic cells may contribute to the adjuvant activity of CpG ODN.

# 5 Conclusion

Cpg ODN activate immune cells and induce production of IL-6, IL-12, and IFN-g by splenocytes in vitro. As development of Th1 cells is dependent upon the presence of IL-12 at the time of antigenic stimulation, we investigated whether CpG ODN could act as Th1-directing adjuvants in vivo. Administration of CpG ODN with protein antigen caused a Th1-dominated cytokine response by splenocytes, with increased production of IFN-g and decreased production of IL-5 upon antigenic recall stimulation in vitro. In addition, CpG ODN induced production of antigen-specific lgG2a, a Th1-associated isotype. These data indicate that CpG ODN direct Th1 responses against simultaneously delivered antigen.

One mechanism for the adjuvant activity of CpG ODN may be modulation of antigen processing and presentation by antigen presenting cells such as macrophages, dendritic cells, or B cells. We investigated the effects of CpG ODN on macrophages and dendritic cells. Overnight treatment of macrophages with CpG ODN resulted in an inhibition of antigen processing and presentation that was due in part to a decrease in synthesis of MHC-II molecules, and no enhancement of antigen processing by macrophages was detexted at any time point. In contrast, treatment of dendritic cells with CpG ODN induced dendritic cell maturation and a transient increase in antigen processing, followed by a decline in antigen processing with maintained presentation of previously processed antigen.

CpG ODN have been shown to act as powerful adjuvants for the development of Th1 responses. They are, therefore, potentially useful in vaccine design for the induction of protective cellular immunity or in clinical situations where IFN- $\gamma$ dominated Th responses are desirable (atopic disease). Future studies in humans will determine the usefulness of these agents in altering human disease states and will, at the same time, provoke few toxic or harmful effects. Additional studies are also needed to elucidate the mechanisms of the Th1 and general adjuvant effects of CpG ODN. In addition to the modulation of cytokine secretion, one of these mechanisms may be the upregulation of antigen processing and presentation by dendritic cells.

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# Effects of CpG DNA on Th1/Th2 Balance in Asthma

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# **1** Introduction

Asthma is a disease whose prevalence, morbidity, mortality, and cost have increased in the past two decades worldwide. These changes have led to increased attention to the causes of asthma and asthma exacerbations; it is now appreciated that airway inflammation is central to the problem of asthma. Recent international guidelines for the management of asthma emphasize that anti-inflammatory therapies are necessary for most asthma patients, and this focus is slowly improving compliance from asthmatics and their health-care providers. Unfortunately, current anti-inflammatory treatment in asthma is non-specific. In this chapter, the Th1/Th2 paradigm of inflammation, which may govern the eosinophilic inflammation found in asthma, is described. Data are presented that demonstrate that

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CpG oligodeoxynucleotides (ODN) are capable of modulating the Th2-mediated inflammatory response in a murine model of asthma. These studies suggest that CpG DNA may be useful in the prevention and treatment of asthma.

# 2 Background

# 2.1 Asthma: Statement of the Problem

Asthma has attracted a great deal of attention from both the public and from the medical community in the past few years. It has been termed an "epidemic" and has been the subject of cover stories in major newspapers and magazines. This is primarily due to the observation that the disease is worsening, particularly in Western, industrialized nations. In the past three decades, the prevalence, severity, and mortality of asthma have increased significantly (Evans et al. 1987). A recent study estimated the total annual cost in the United States at almost \$6 billion (SMITH et al. 1997). Asthma is a disease characterized by eosinophilic airway inflammation, bronchospasm, and bronchial hyper-reactivity (National Asthma Education and Prevention Program 1997). Once thought to be due to airway muscle spasm, asthma is now known to be an inflammatory disorder; during an asthma exacerbation, inflammation precedes bronchospasm. In acute asthma, eosinophils may form up to half of the cellular infiltrate (ROBBINS and COTRAN 1979), and bronchoalveolar eosinophilia invariably follows allergen inhalation in asthma attacks (METZGER et al. 1985, 1986, 1987). Eosinophils cause inflammation and bronchial hyper-reactivity through release of mediators, such as leukotrienes, major basic protein, eosinophilic cationic protein, and eosinophilic peroxidase (BRUIJN-ZEEL 1994).

# 2.2 Th1 and Th2 Balance in Asthma

The number and activity of eosinophils are controlled by cytokines released from activated T cells, especially interleukin 4 (IL-4), IL-5, and IL-13. T-lymphocytes can be divided, on the basis of cytokine production, into T helper 1 (Th1) and Th2 (MOSMANN et al. 1986). Th1 cells produce IL-2 and interferon  $\gamma$  (IFN- $\gamma$ ) but not IL-4 or IL-5, and Th2 cells produce IL-4, IL-5, IL-6, IL-10, and IL-13 but not IL-2 or IFN- $\gamma$ . Th1 and Th2 cells interact in a counter-regulatory fashion: IL-4 and IL-10 promote Th2 development (SWAIN et al. 1990; PARRONCHI et al. 1992) and inhibit Th1 cell and cytokine production (MOORE et al. 1990), and IFN- $\gamma$  inhibits the proliferation of Th2 cells (GAJEWSKI and FITCH 1988) and promotes the development of Th1 cells (PARRONCHI et al. 1992). IL-12, mainly a product of activated macrophages, is also a strong promoter of Th1 responses (BLISS et al. 1996) and is often considered a Th1 cytokine; many of the activities ascribed to IL-12 are

due to induction of IFN- $\gamma$  (MICALLEF et al. 1996). Th1 and Th2 cells have been identified in humans in vivo as well as in vitro (WIERENGA et al. 1990; DEL PRETE et al. 1991; ROMAGNANI 1991; SALGAME et al. 1991; FIELD et al. 1993).

The Th2 cytokines, IL-4, IL-5, and IL-13 (ROBINSON et al. 1992; GRUNIG et al. 1998: WILLS-KARP et al. 1998), have been increasingly implicated in the inflammation of asthma. IL-4 amplifies allergic responses by inducing immunoglobulin E (IgE) production by uncommitted B cells (DEL PRETE et al. 1988) and is a growth factor for mast cells (SAITO et al. 1988). IL-5 also stimulates Ig secretion (TAKATSU et al. 1980; Swain et al. 1990) as well as stimulating the proliferation and activation of eosinophils (CLUTTERBUCK et al. 1988; LOPEZ et al. 1988; WALSH et al. 1990) and basophils (HIRAI et al. 1990). IL-13 has recently been shown to cause airway hyperresponsiveness and inflammation independently of eosinophils or IL-4 (GRUNIG et al. 1998; WILLS-KARP et al. 1998). In vitro, allergen-specific T-cell clones from atopic donors release Th2 cytokines after stimulation by specific allergens (PAR-RONCHI et al. 1991). During asthma exacerbations, peripheral T-cell activation and increased serum IL-5 correlate with eosinophilia and asthma symptoms (CORRIGAN et al. 1993), and bronchoalveolar lavage (BAL) T-cells release cytokines in a Th2like pattern (ROBINSON et al. 1992, 1993a,b). Non-atopic asthmatics also have increased levels of Th2-like cytokines; increased IL-5 release from BAL T lymphocytes is characteristic of both atopic and non-atopic asthmatics (WALKER et al. 1992), and expression of peripheral T-cell IL-5 mRNA from non-atopic asthmatic subjects correlates with increased BAL (MARINI et al. 1992) and peripheral blood (WALKER et al. 1992) eosinophilia.

#### 2.3 Inflammation and the Treatment of Asthma

Because of these observations, the focus of treatment in asthma has shifted from primarily addressing bronchospasm to modulating inflammation. Recent guidelines for the management of asthma recommend that anti-inflammatory therapy be used for all but the most intermittent and benign cases of the disease (National Asthma Education and Prevention Program 1997). Current anti-inflammatory therapy, however, remains disappointingly broad; corticosteroids are the "gold standard" for asthma treatment, and inhaled corticosteroids are only incrementally better than they were 25 years ago. The much-touted leukotriene pathway antagonists, which have been released in the last 5 years, have been helpful only in a subset of asthmatics. Thus, the need remains for novel, effective treatments of inflammation in asthma.

#### 2.4 DNA and Th1 Responses

The observation that ODN containing CpG motifs (the dinucleotide CpG in specific base-sequence contexts) have remarkable immunostimulatory properties, including induction of Th1 cytokines, has given rise to a series of studies examining how these agents may be useful in the modulation or abrogation of Th2-mediated

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inflammation in asthma. Bacterial, but not vertebrate, DNA causes immune activation (KRIEG et al. 1995); the dinucleotide CpG is markedly under-represented in vertebrate DNA and, when present, is almost invariably methylated (KRIEG et al. 1995). ODN containing unmethylated CpG motifs reproduce the effect of bacterial DNA strongly and rapidly; CpG DNA induces cytokine mRNA within 30min of administration (Y1 and KRIEG 1998), and IL-6 and IL-12 protein secretion is increased within 4h after in vivo treatment with CpG DNA (KLINMAN et al. 1996). CpG-DNA-induced IL-12 production precedes and is responsible for the subsequent natural killer cell IFN- $\gamma$  secretion (BALLAS et al. 1996; HALPERN et al. 1996).

# 2.5 Early-Life Infections and Th1 Responses

The possibility that CpG DNA of prokaryotic origin may be protective against asthmatic inflammation is counterintuitive; indeed, it has been suggested that the pathogenesis of asthma may be related to poor control of airway infections. However, a number of studies have examined the association between childhood infection (resulting in early exposure to viral or prokaryotic DNA) and asthma, bolstering the hypothesis that this exposure may result in the conversion of a propensity for a Th2 response to allergen into a Th1 or non-atopic response. Von Mutius and colleagues hypothesized that worsening air pollution may be responsible for the increased morbidity of asthma in Western nations and compared the prevalence of asthma and atopy between children of Leipzig, in polluted former East Germany, and Munich, in less polluted former West Germany (VON MUTIUS et al. 1992). Surprisingly, a lower rate of asthma and allergic disorders was found among children from Leipzig than in Munich children. The investigators postulated that this finding might result from differences in living conditions rather than environmental exposures. In a follow-up study, they examined the relationship between atopy and number of siblings and found that children with increased numbers of siblings were less likely to express atopy (von MUTIUS et al. 1994). As children with multiple older siblings were more likely to develop childhood illnesses earlier in life than those without siblings (the older siblings bring home infections from school or daycare), it is possible that the early-life infections prevent the later development of atopy. These results suggested an interpretation of their earlier study: the Leipzig children (who were more likely to be in out-of-home daycare than the Munich children) may also have been protected against atopy by an increase in early-life childhood infections.

A more direct examination of the relationship between infection and atopy was carried out by Shaheen and colleagues (SHAHEEN et al. 1996). A measles epidemic occurred in Guinea-Bissau in 1979. Subsequently, children were evaluated for the development of atopy, which was defined by reactivity to skin testing for common environmental allergens. After controlling for potentially confounding variables, such as numbers of older siblings, history of breast feeding, and in-home exposure to livestock, a history of measles infection was found to be associated with a

significant reduction in the risk of atopy (relative risk 0.20, P < 0.01) (SHAHEEN et al. 1996). Finally, in a much-quoted recent study, Shirakawa and colleagues found that a positive response to tuberculin predicted a lower incidence of asthma and atopy in a cohort study of Japanese schoolchildren (SHIRAKAWA et al. 1997). Moreover, the tuberculin responsiveness correlated with induction of Th1 (IFN- $\gamma$ ) and suppression of Th2 (IL-4, IL-13) serum cytokine levels. This exciting study, while not definitive evidence, strongly supports a link between childhood infection and protection against the development of atopy.

# **3** Results

## 3.1 CpG ODN Prevent Eosinophilic Airway Inflammation in Asthma

Based on these epidemiological studies and observations that CpG ODN were potent inducers of the Th1 cytokines IL-12 and IFN- $\gamma$ , we elected to examine the effects of CpG DNA in an animal model of asthma. For the initial studies, we adapted a murine model in which C57BL/6 mice were sensitized to Schistosoma mansoni proteins by intraperitoneal injection of schistosome eggs (5,000 eggs). The mice subsequently received two airway challenges with soluble egg antigen from the schistosome eggs (SEA; 10µg) (LUKACS et al. 1994; KLINE et al. 1998). Following exposure to the second airway challenge, mice were studied. Some mice received ODN ( $30\mu g$ , either CpG or control, on a phosphothioate backbone) by intraperitoneal injection at the time of sensitization. For these studies, the CpG ODN used was TCCATGACGTTCCTGACGTT, and the control ODN was TCCAT-GAGCTTCCTGAGTCT. In comparison with saline control mice, we found that significant eosinophilic pulmonary inflammation developed following airway challenge in the sensitized mice (Fig. 1). Although all cell types increased in lung lavage samples, a majority of the cells in the influx were eosinophils. This pleocytosis and eosinophilia were almost completely abrogated in the mice who received CpG ODN; interestingly, mice who received control ODN at the time of sensitization demonstrated a slight but consistent diminution of airway inflam-



Fig. 1. CpG oligodeoxynucleotides (ODN) prevent the development of airway eosinophilia in a murine model of asthma. C57BL/6 mice who received CpG ODN at the time of sensitization, but not those who received control ODN, develop significantly less airway eosinophilia than mice sensitized to schistosome eggs in the absence of ODN. \*P < 0.001 vs the schistosome-egg-sensitized group
mation, suggesting the existence of a non-specific ODN effect. Histopathologic examination of the lungs of the mice confirmed the pronounced eosinophilic inflammation induced by schistosome sensitization and airway challenge; a marked peribronchial and perivascular inflammatory exudate included eosinophils, mast cells, macrophages, and lymphocytes, and the airway epithelium was thickened, with prominent mucus production (Fig. 2). These inflammatory changes were markedly reduced in those mice who received pretreatment with CpG ODN.

### 3.2 CpG ODN Prevent Bronchial Hyper-Reactivity

As asthma is defined by the presence of bronchial hyper-reactivity to non-specific stimuli in addition to eosinophilic airway inflammation, we next wanted to evaluate the effect of CpG ODN on the airway pathophysiologic responses found in this model. For these studies, we evaluated the response of the mice to inhaled methacholine and used a computer-interfaced whole-body plethysmograph (Buxco, Troy, NY, USA), which non-invasively analyzes the response to increasing doses of inhaled methacholine. "Enhanced pause" (Penh) is calculated as a function of expiratory time, relaxation time, and peak inspiratory and expiratory flow (Penh =  $[(T_e/0.3T_r)-1] \times [2P_{ef}/3P_{if}]$ ); Penh correlates with directly measured



**Fig. 2A–H.** CpG oligodeoxynucleotides (ODN) prevent the development of peribronchial eosinophilia in a murine model of asthma. Compared with saline control mice (A, E), those mice who are sensitized to schistosome eggs and then are challenged with soluble egg antigen in the airway (B, F) develop marked peribronchial-eosinophilic inflammation and epithelial activation, which is markedly reduced in CpG-ODN-treated (C, G) but not control-ODN-treated (D, H) mice

pronchospasm (HAMELMANN et al. 1997; KLINE et al. 1998). Although C57BL/6 nice are relatively insensitive to inhaled methacholine, those mice who are sensitized and later challenged with schistosome proteins have a marked increase in response to methacholine (Fig. 3); pretreatment with CpG ODN, but not control DDN, returned the bronchial reactivity nearly to baseline. These studies demonstrated that CpG ODN could prevent the physiologic sequellae of inflammation and the inflammatory response itself.

### 3.3 CpG ODN Suppress Pulmonary and Systemic Th2 Responses

Because induction by CpG ODN of Th1 cytokines, such as IFN- $\gamma$  and IL-12, had previously been demonstrated, we next examined the effects that CpG ODN have on the regulation of cytokines in the lung. For these studies, we measured the concentration of IL-4, IFN- $\gamma$ , and IL-12 in lung lavage fluid. We found (Table 1) that IL-4 was strongly induced in the sensitized/challenged mice. This induction was greatly diminished in mice that were treated with CpG ODN. Conversely, both IFN- $\gamma$  and IL-12 were greatly upregulated in the lavage fluid of mice treated with CpG



Fig. 3. CpG oligodeoxynucleotides (ODN) inhibit the development of bronchial hyper-reactivity to innaled methacholine. Mice that are sensitized to schistosome eggs and challenged with soluble egg antigen n the airway develop bronchial hyper-reactivity (demonstrated as increased Penh index or fold increase in Penh, a measure of airway resistance) to inhaled methacholine, compared with saline-control mice. This response is markedly blunted in mice treated with CpG ODN but not in those mice treated with control DDN. \*P < 0.05 vs the schistosome-egg-sensitized group

Fable 1. Th1 and Th2 cytokine levels in bronchoalveolar lavage fluid

Condition of sensitization	IL-4 (pg/ml)	IFN-γ (pg/ml)	IL-12 (pg/ml)
Saline	5.3 ± 2.1*	8.4 ± 3.1	$6.9 \pm 3.4$
CpG ODN + egg	$23.6 \pm 4.7*$	$403.2 \pm 107.5^{**}$	$57.6 \pm 19.7^{**}$
Control ODN + egg	$82.3 \pm 17.6$	$107.6 \pm 82.4$	$20.4 \pm 7.1$
Schistosome egg	$115.8 \pm 24.9$	$15.2 \pm 9.9$	$18.3 \pm 4.6$

[FN, interferon; IL, interleukin; ODN, oligodeoxynucleotide.

\*P < 0.05 vs schistosome-egg-sensitized mice.

\*\*P < 0.01 vs schistosome-egg-sensitized mice.

ODN (Table 1). These data suggested that induction of the Th1 cytokines, IFN- $\gamma$  and IL-12, were responsible for the anti-inflammatory effects of CpG ODN in the murine model of asthma and that their effect was downregulation of Th2 responses.

Humoral Th1 and Th2 responses also differ from one another. Humoral Th2 responses are characterized by induction of IgE antibodies, and Th1 responses are characterized by induction of IgG2a and IgG2b antibodies. In the schistosome murine model of asthma, total IgE is induced, and there is no change in the total IgG2a or IgG2b. Treatment with CpG ODN significantly reduced the induction of IgE (from 4.10  $\pm$  0.54µg/ml to 1.04  $\pm$  0.32µg/ml, P < 0.05) but had no effect on IgG subtypes.

# 3.4 The Effect of CpG ODN in Murine Asthma is Neither Strain nor Model Dependent

Because we wished to ascertain whether the effect of CpG ODN on atopic airway inflammation was strain-specific, we next examined the effect of the ODN on the development of airway inflammation and bronchial hyper-reactivity using the schistosome egg/SEA protocol on BALB/c mice. These mice developed less airway eosinophilia than C57BL/6 mice do, but their eosinophilia was completely abrogated by the CpG ODN, demonstrating that the protection offered by CpG ODN is not strain specific ("asthmatic" BALB/c mice developed  $1.2 \pm 0.4 \times 10^6$  eosinophils/lavage; CpG-treated mice developed  $0.06 \pm 0.04 \times 10^6$  eosinophils/lavage, P < 0.001). Moreover, using an ovalbumin murine model of asthma, we also examined whether this protection was model specific. For these studies, C57BL/6 mice were sensitized to ovalbumin with a single intraperitoneal injection of ovalbumin (10µg) heat-precipitated with alum; this was followed by a period of ovalbumin inhalation (6% solution aerosolized in a chamber for 30 minutes daily on days 7-11 and 14-18). Using this model, we found that the eosinophilia induced by ovalbumin in sensitized mice was significantly reduced by CpG ODN, but not by control ODN (ovalbumin-sensitized mice: 2.6  $\pm$  0.7  $\times$  10<sup>6</sup> eosinophils; CpG ODNtreated mice:  $0.03 \pm 0.02 \times 10^6$  eosinophils, P < 0.001; control ODN-treated mice:  $1.9 \pm 0.8 \times 10^6$  eosinophils, P = no significance).

### 3.5 The Effects of CpG ODN are Persistent

To examine whether the effects of CpG ODN are long-lasting, we next sensitized a cohort of mice to schistosome eggs in the presence or absence of CpG or control ODN and challenged them as per our protocol at days 7 and 14; 4 weeks later, we repeated the process, but this time all the mice received schistosome eggs without ODN. Lavage of the mice after the second series of SEA inhalation revealed that those mice who received CpG ODN along with the sensitizing antigen received long-lasting protection against sensitization (Fig. 4). These mice were also protected against the development of bronchial hyper-reactivity (data not shown).



Fig. 4. Mice that receive CpG oligodeoxynucleotides (ODN) at the time of sensitization are protected against subsequent sensitization. C57BL/6 mice were sensitized to schistosome eggs in the presence or absence of CpG ODN or control ODN or received saline alone as a control, and were then challenged with soluble egg antigen (SEA) in the airway. Four weeks later, all mice received a second intraperitoneal injection of schistosome eggs (no ODN) followed by two weekly airway challenges with SEA. Those mice that received the CpG ODN (but not those that received the control ODN) were initially protected against the development of eosinophilic inflammation even when challenged after the second exposure to schistosome eggs. \*P < 0.01 vs the schistosome-egg-sensitized group



**Fig. 5.** CpG oligodeoxynucleotides (ODN) are effective in prevention of eosinophilic airway inflammation and bronchial hyper-reactivity in the presence of antibodies to interferon  $\gamma$  (IFN- $\gamma$ ) or interleukin 12 (IL-12). Administration of anti-IFN- $\gamma$  or anti-IL-12 blocking antibodies does not significantly impair the effectiveness of CpG ODN in prevention of eosinophilic airway inflammation. \*P < 0.005 vs the egg group. There is no significant difference between the three CpG groups

### **3.6** Neither IFN-γ nor IL-12 is Required for the Protective Effects of CpG ODN

To evaluate whether the Th1 cytokines, IFN- $\gamma$ , and IL-12 are required for the protective effects of CpG ODN, we first examined whether they would prevent the development of airway eosinophilia and bronchial hyper-reactivity in the absence of IFN- $\gamma$ . For these studies, we administered anti-IFN- $\gamma$  antibodies (R4–6A2) at the time of the administration of the ODN and the schistosome eggs. These anti-

bodies are known to specifically block the effects of IFN- $\gamma$  in vivo (SPITALNY and HAVELL 1984). To our surprise, we found that CpG ODN, in the absence of IFN- $\gamma$  activity, were still effective at preventing airway eosinophilia (Fig. 5) and bronchial hyper-reactivity (not shown). To definitively evaluate whether CpG ODN could prevent the development of asthma-like inflammation in the absence of IFN- $\gamma$ , we next used IFN- $\gamma$  knockout (KO) mice (DALTON et al. 1993) in our asthma model. We found that these mice were also protected from the development of eosinophilic inflammation (Fig. 6A) and airway hyper-reactivity (Fig. 6B).

We performed similar studies to evaluate the role of IL-12 in the mediation of the CpG effect. These studies used the blocking antibody TOSH [a rat anti-murine IL-12 IgG1 antibody (WENNER et al. 1996)] or an isotype-specific control antibody; systemic blockade of IL-12 only modestly reduced the protective effects of CpG ODN on the prevention of airway eosinophilia (Fig. 5). Using IL-12 KO mice (MAGRAM et al. 1996), we found that these mice were also protected from the development of schistosome/SEA-induced airway eosinophilia by administration of



**Fig. 6A,B.** CpG oligodeoxynucleotides (ODN) are effective in prevention of eosinophilic airway inflammation and bronchial hyper-reactivity in interferon  $\gamma$  (IFN- $\gamma$ ) knockout (KO) mice. IFN- $\gamma$  KO mice were sensitized to schistosome eggs in the presence or absence of CpG or control ODN. KO mice that received CpG, but not those who received control ODN at the time of sensitization, were protected against the development of (A) airway eosinophilia and (B) hyper-reactivity to inhaled methacholine. \*P < 0.01 vs the schistosome-egg-sensitized group

CpG ODN but not control ODN (Fig. 7A). The CpG-treated IL-12 KO mice exposed to schistosome egg and then SEA also had diminished bronchial hyperreactivity to inhaled methacholine compared with sensitized and challenged IL-12 KO mice who did not receive CpG ODN (Fig. 7B).

Since the protection conferred by CpG ODN on both IFN- $\gamma$  KO and IL-12 KO mice was qualitatively similar to the protection offered to wild-type C57BL/6 mice, we next examined whether it was quantitatively identical. In the preceding studies, we used CpG ODN in an amount previously found to be well within the protective range, 30µg (KLINE et al. 1998). Therefore, we proceeded to examine whether the dose-response relationship between the amount of CpG ODN used and protection against eosinophilia was altered in either or both of the cytokine KO mice. For these studies, we used the same protocol described above, but we administered varying amounts of CpG ODN (between 0.3µg and 30µg) at the time of sensitization to schistosome eggs (Fig. 8). We found that, at lower doses of CpG ODN ( $<10\mu$ g), both of the KO mice were offered significantly less protection against the development of airway eosinophilia than wild-type C57BL/6 mice.



**Fig. 7A,B.** CpG oligodeoxynucleotides (ODN) are effective in prevention of eosinophilic airway inflammation and airway hyper-reactivity in interleukin 12 (IL-12) knockout (KO) mice. IL-12 KO mice were sensitized to schistosome eggs in the presence or absence of CpG or control ODN. CpG ODN, but not control ODN, significantly reduces the development of (A) airway eosinophilia and (B) methacholine-induced bronchospasm in IL-12 KO mice. \*P < 0.01 vs the schistosome-egg-sensitized group



**Fig. 8.** In the absence of interferon  $\gamma$  (IFN- $\gamma$ ) or interleukin 12 (IL-12), mice are relatively deficient in response to CpG oligodeoxynucleotide (ODN). Varying amounts of CpG ODN (0, 0.3, 3, 10, or 30µg) were administered to wild-type C57BL/6, IFN- $\gamma$  knockout (KO), and IL-12 KO mice (each on a C57BL/6 background) at the time of sensitization to schistosome eggs. All mice who received 30µg and the wild-type mice who received 10µg of CpG ODN were fully protected against airway eosinophilia. There was no difference in airway eosinophilia between any of the groups that received no CpG ODN. A significant difference in the development of airway eosinophilia was seen between the wild-type and KO mice groups that received between 0.3µg and 10µg of CpG ODN. \*\*P < 0.005 vs the egg group; n = 4 mice in each group

### 4 Summary

Thus, in our studies, we demonstrated that CpG ODN are effective in preventing the development of eosinophilic airway inflammation and bronchial hyper-reactivity in a murine model of asthma. Antigen-associated elevation of serum IgE levels is also suppressed. CpG ODN, administered in conjunction with antigen, is also effective in down-regulation of established Th2 responses. This protection is neither murine strain-dependent nor model-dependent. Although these effects of CpG ODN are associated with the induction of the Th1 cytokines IFN- $\gamma$  and IL-12, neither cytokine is absolutely required for the protection. These results suggest that CpG ODN may be an effective immunomodulatory agent in the treatment, and possibly prevention, of asthma.

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# **Responses of Human B Cells to DNA and Phosphorothioate Oligodeoxynucleotides**

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### **1** Introduction

Until recently, DNA was thought to be neither immunogenic nor immunostimulatory, but emerging information has documented that specific forms of DNA can be both. It has been found that certain DNA structures (Z-DNA) can induce significant antibody responses when administered to normal mice (STOLLAR 1975, 1986; RICH et al. 1984; ZARLING et al. 1984; FRAPPIER et al. 1989). In addition, studies in mice demonstrated that DNA from various micro-organisms (bacterial *Escherichia coli* DNA) induced B-cell proliferation, immunoglobulin (Ig) production and cytokine secretion both in vitro and in vivo. In contrast, mammalian DNA did not stimulate murine B-cell responses (MESSINA et al. 1991, 1993; KRIEG 1995; KRIEG et al. 1995; PISETSKY 1995, 1996; KLINMAN et al. 1996b; SUN et al. 1996, 1997). Comparison of the sequences of mammalian and microbial genomes indicated that CpG dinucleotides are ten times more frequent in bacterial than in vertebrate DNA, and 80% of the cytosines within CpG dinucleotides in vertebrate genomes are methylated, whereas those in microbes and synthetic ODNs are non-

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methylated (BIRD 1986, 1987; CARDON et al. 1994). After testing a battery of ODNs for their immunostimulatory activities, KRIEG and his colleagues found that the stimulatory activity of microbial DNA and sODNs could be attributed to a CpG motif, in which an unmethylated CpG dinucleotide is flanked by two 5' purines and two 3' pyrimidines (KRIEG et al. 1995; KRIEG 1996a,b).

The immunostimulatory activity of DNA has also been demonstrated with certain polynucleotides, especially those synthesized from phosphorothioate derivatives of nucleotides. Phosphorothioate oligodeoxynucleotides (sODNs) have a sulfur substitution for a non-bridging oxygen in the backbone (Fig. 1) and differ from natural phosphodiester-linked polynucleotides in a number of properties, such as nuclease resistance and melting temperature (STEIN and COHEN 1988; STEIN and CHENG 1993; TONKINSON and STEIN 1996). Among 18–27-bp sODN antisense constructs designed to inhibit the expression of specific target genes, some induced activation of murine B lymphocytes both in vivo and in vitro (KRIEG et al. 1989; BRANDA et al. 1993b, 1996b; MCINTYRE et al. 1993, MOJCIK et al. 1993; PISETSKY and REICH 1993; MONTEITH et al. 1997). B-cell stimulation by the antisense sODNs did not appear to result from inhibition of specific genes by the compounds but rather from direct stimulation of B cells. Although some stimulatory sODNs contained the CpG motifs, murine B-cell activation could also be induced by sODNs containing no CpG at all (BRANDA et al. 1996b, MONTEITH et al. 1997).

In addition to the activation of murine B cells, bacterial DNA and ODNs containing the CpG motif also directly activate murine macrophages and dendritic cells and induce the secretion of a variety of cytokines, which subsequently stimulate natural killer (NK) cells to secrete interferon  $\gamma$  (IFN- $\gamma$ ) and enhance NK lytic activity (BALLAS et al. 1996; COWDERY et al. 1996; HALPERN et al. 1996; STACEY et al. 1996; CHACE et al. 1997; LIPFORD et al 1997b; SPARWASSER et al. 1997, 1998; JACOB et al. 1998). Moreover, because of their ability to induce production of interleukin 12 (IL-12) and IFN- $\gamma$ , bacterial DNA and ODNs containing the CpG motif can stimulate a selective T helper 1 (Th1) response both in vivo and in vitro (RAZ et al. 1996; CARSON and RAZ 1997; CHU et al. 1997; ROMAN et al. 1997).



**Fig. 1.** Structure of oligodeoxynucleotides (*ODNs*) and phosphorothioate ODNs (sOD-Ns)

Because of the variety of activities on the immune system leading to antibody production and Th1 responses, the immunostimulatory DNA or ODNs containing the CpG motif could be important adjuvants for conventional protein vaccination and necessary components for DNA vaccination against a variety of targets (KLINMAN et al. 1996a; PISETSKY 1996; SATO et al. 1996; LIPFORD et al. 1997a; ROMAN et al. 1997; WEINER et al. 1997; DAVIS et al. 1998; KRIEG 1998; MOLD-OVEANU et al. 1998; SUN et al. 1998). The immunostimulatory DNA or ODNs containing the CpG motif can also have important therapeutic effects, including an enhancement of the immune response to tumor cells and protection from allergic disease (WOOLDRIDGE et al. 1997; KLINE et al. 1998).

These effects have been noted in mice and suggest that immunostimulatory DNA or sODNs may have important roles in vaccination, tumor resistance and modification of Th2-mediated responses in humans. However, there is little information available on the potential immunostimulatory effects of DNA or sODNs in humans. Because there are significant differences in the responsiveness of human and murine lymphocytes, it is not possible to extrapolate results obtained from mouse directly to man. Therefore, investigation of the capacity of immunostimulatory DNA and ODNs to induce activation of human lymphocytes is necessary to determine both the potential impact of these materials on human immune responsiveness and their potential role as immune modulators.

### 2 Polyclonal Activation of Human B Cells by DNA and sODNs

The first suggestion of an immunostimulatory activity of DNA in man came from studies on the binding of human sera to a variety of mammalian and nonmammalian DNA. Sera from normal subjects were noted to contain antibodies that bound DNA from two bacterial species, Micrococcus lysodeikticus and Staphylococcus epidermidis. These antibodies had high affinity and specificity for bacterial and non-mammalian DNA, which suggested that they bound to structural epitopes unique to these DNAs (ROBERTSON et al. 1992; BUNYARD and PISETSKY 1994; PISETSKY 1997; WU et al. 1997; KORANS et al. 1998; PISETSKY 1998). It was also noted that sera from normal subjects contained antibodies that bound to DNA from BK polyomavirus (FREDRIKSEN et al. 1993). Moreover, it was noted that antisense sODNs complementary to parts of the hepatitis-B-virus genome stimulated proliferation of B cells from chronic hepatitis patients (CHEN et al. 1996). Furthermore, it has been demonstrated that 27bp and 21bp antisense sODNs to the rev region of the human immunodeficiency virus (HIV) genome increased proliferation and Ig production of peripheral blood mononuclear cells (PBMCs) from normal subjects as well as those with common variable immunodeficiency and chronic lymphocytic leukemia (BRANDA et al. 1993a; BRANDA et al. 1996a). The stimulation indices of PBMCs from patients were similar to those from normal human subjects, suggesting that responsiveness did not require antecedent exposure to HIV (BRANDA et al. 1996a). Taken together, these reports suggest that certain DNA or sODNs can induce lymphocyte proliferation and antibody production from patients and normal human subjects. However, the mechanism of the human B-cell activation and the question of whether DNA and sODNs can directly activate normal human B cells remain unclear.

Many studies have been carried out recently to investigate the immunostimulatory activity of CpG-containing DNA and ODNs in mouse and other species, but minimal information on the responsiveness of human B cells is available. To address this issue, a series of experiments was carried out to determine whether bacterial DNA and various ODNs (Table 1) could activate human B cells directly (LIANG et al. 1996). These studies employed highly purified human peripheral blood B cells so that a direct stimulatory activity on human B cells could be assessed. As shown in Fig. 2, three of the various ODNs tested consistently induced significant human B-cell proliferation. Each of these stimulatory polynucleotides was a phosphorothioate. Of note, neither *E. coli* DNA nor calf thymus DNA was stimulatory. Other sODNs (poly G, poly C, poly T, randomers) and ODNs induced

**Table 1.** Sequence, length and description of representative oligodeoxynucleotides (ODNs). The categories were defined according to their ability to activate human B cells. There was no significant difference in the magnitude of activation by any pair of synthetic ODNs (sODNs) within each category, although there were significant differences between activation induced by sODNs from different categories (P < 0.05) and between activation induced by any sODN and the response of B cells alone. *HIVas*, antisense human immunodeficiency virus; *HSVas*, antisense herpes simplex virus; *NK*, natural killer

Name	Sequence	Length	Description
Most active			
HSVas	5'-GCCGAGGTCCATGTCGT ACGA-3'	21bp	Antisense to a translation initiation region of HSV
20-mer	5'-TTGCTTCCATCTTCCTC GTC-3'	20bp	C
HIVas	5'-TCGTCGCTGTCTCCGCTTC TTCTTGCC-3'	27bp	Antisense to rev of HIV
TCG₄	5'-TCGTCGTCGTCG-3'	12bp	
MCMT <sup>+</sup>	5'-TGACGTTTGACGTTT- GACGTT-3'	21bp	Murine CpG motif
$MCMT^+C\Delta T$	5'-TGATGTCTGATGTCT- GATGTC-3'	21bp	Murine CpG motif with $C\Delta T$
$MCMT^+C\Delta G$	5'-TGAGGTCTGAGGTCT- GAGGTC-3'	21bp	Murine CpG motif with $C\Delta G$
Moderately active			
NKNSO	5'-GGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGG	30bp	NK nonstimulatory sODN
NKSO	5'-GGGGGGGGGGGGGGAACGT TGGGGGGGGGGGGGGG-3'	30bp	NK stimulatory sODN
Minimally active			
G <sub>20</sub>	5'-GGGGGGGGGGGGGGGG GGGG-3'	20bp	Poly G
C <sub>20</sub>	5'-CCCCCCCCCCCCCCC CCC-3'	20bp	Poly C
CG <sub>7.5</sub>	5'-CGCGCGCGCGCGCGCGC-3'	15bp	Poly CG



**Fig. 2.** B-cell proliferation induced by phosphorothioate oligodeoxynucleotides (sODNs). Highly purified B cells  $(50 \times 10^3/\text{well})$  were cultured with various concentrations of ODNs  $(1\mu g/ml - 50\mu g/ml)$  for 4 days, and proliferation was assessed by <sup>3</sup>H-thymidine incorporation. The data indicate results with the optimal stimulatory concentrations, usually  $5\mu g/ml$  (except  $1\mu g/ml$  for antisense human immunodeficiency virus and  $25\mu g/ml$  for Rmer10). Data with herpes simplex virus antisense oligodeoxynucleotide (HSVaso) were obtained by incubation of B cells with  $5\mu g/ml$  antisense herpes simplex virus and then adding  $5\mu g/ml$  of HSVaso to culture daily. A single addition of HSVaso induced no response. The data are the means of triplicate cultures with a standard error of the mean of less than 10%, and are representative of three separate experiments, each carried out with B cells from a different donor

minimal, but nonetheless detectable, responses that were statistically different from the <sup>3</sup>H-thymidine incorporation manifested by B cells cultured alone. Poly A and three 8bp sODNs did not stimulate B cells at all. Moreover, the diester form of the herpes simplex virus (HSV) antisense (HSVas) was much less stimulatory than its phosphorothioate counterpart within this concentration range, even when added to culture repetitively (the mean maximum response to the diester form of HSVas was  $615 \pm 22$ cpm vs.  $3113 \pm 93$ cpm for the phosphorothioate form of HSVas). Although modest, the response induced by the diester form of HSVas was significantly greater than the tritiated thymidine incorporation of control B cells (99  $\pm$  7cpm, P < 0.01) (LIANG et al. 1996). These results indicated that certain sODNs induced marked human B-cell proliferation far more efficiently than their phosphodiester counterparts did.

In addition, the sODNs that induced maximal proliferation also induced production of IgM, IgG and IgA from purified B cells in the absence of exogenous cytokines or T cells. The other ODNs and DNA induced little or no Ig production (LIANG et al. 1996).

IgM anti-DNA antibody was also induced by the sODNs that stimulated maximal proliferation, and the anti-DNA antibodies secreted bound all sODNs tested. Therefore, maximal stimulatory sODNs activated B cells committed to the production of anti-DNA antibodies. However, specificity of the anti-DNA antibody for the activating sODNs was not apparent (LIANG et al. 1996).

Of note, certain cytokines (IL-2) enhanced responses of B cells activated by the most stimulatory sODNs (LIANG et al. 1996). Other cytokines tested, including IL-4 and IL-10, had no augmenting effects. Purified human T cells also enhanced the responses of human B cells to sODNs (LIANG et al. 1996), although T cells did not bind sODNs (Fig. 3B) and did not proliferate when cultured with sODNs (LIANG et al. 1996). These results indicated that IL-2 and intact T cells enhanced B-cell responses to the most stimulatory sODNs, but they were not required for the stimulatory activity of sODNs.



Fig. 3A–D. Antisense human immunodeficiency virus (HIVas) binds to B cells, natural killer cells and CD14-positive myeloid cells but not T cells. Peripheral blood mononuclear cells were incubated with HIVas–fluorescein isothiocyanate (FITC;  $4\mu g/sample$ ) at 37°C for 30min. After washing to remove unbound HIVas–FITC, cells were stained with monoclonal antibodies (mAbs) against CD19 (A), CD3 (B), CD14 (C) and CD16 (D) or their isotype-matched controls. The FACscan plots show mAb staining on the y-axis and HIVas-FITC staining on the x-axis and are representatives of one of three experiments, each using cells from a different donor. HIVas–FITC is bound by 67% of CD19-positive cells, 3% of CD3-positive cells, 96% of CD14-positive cells and 76% of CD16-positive cells

sODNs also induced the expression of activation markers by highly purified B cells. Of interest, a much larger number of B cells expressed activation markers than might have been anticipated from the moderate degree of B-cell proliferation noted. Thus, the sODNs that induced maximal proliferation activated more than 80% of B cells to express activation markers, including CD69, CD86 and CD25. In contrast, sODNs that stimulated minimal proliferation also induced either minimal or no increased expression of activation markers. The finding that sODNs induced activation-marker expression by more than 80% of B cells implies that the response is polyclonal. This was confirmed by documenting that B cells expressing all six major  $V_{\rm H}$  families were activated by stimulatory sODNs (LIANG et al. 1996).

The large number of B cells induced to express activation markers, compared to the modest degree of proliferation induced by sODNs, implied that many B cells were initially activated, but few underwent clonal expansion in response to sODNs. Additional experiments directly analyzing cell-cycle entry and progression confirmed that active sODNs stimulated a few cells to enter the cell cycle, and these cells underwent multiple rounds of proliferation.

Taken together, certain sODNs induced T-cell-independent polyclonal activation of human B cells, resulting in polyclonal expansion of a small number of activated B cells and differentiation of some stimulated B cells into Ig secreting cells. Thus, certain sODNs provide sufficient signals to induce human B-cell responses in vitro in the absence of T cells and other cells capable of facilitating B-cell activation.

### 3 Mechanism of Activation of Human B Cells by sODNs

#### 3.1 sODNs Activate Human B Cells by Engaging Surface Receptors

The mechanism of human B-cell activation by sODNs has not been completely delineated. To examine whether sODNs activated human B cells by engaging surface receptors or whether cellular entry of sODNs was necessary for the stimulatory activities, active sODNs were covalently coupled to Sepharose beads and tested for their ability to activate B cells. The results showed that Sepharose-bound sODNs stimulated B cells as effectively as soluble sODNs (LIANG et al. 1996). Additional experiments documented that the stimulatory activity did not result from soluble sODNs released from the beads. These results indicate that human B-cell stimulation results from engagement of sODN-binding surface receptors.

Examination of binding of fluorescein-isothiocyanate-labeled sODNs indicated that binding was rapid, saturable, specific and initially temperature independent, suggesting that the binding is likely to be receptor mediated. Most sODNs tested appeared to bind to the same or to a limited set of receptor(s). However, the nature of the sODN-binding stimulatory receptor remains unknown. These data indicate that sODNs directly bind to surface receptors on human B cells and suggest that binding to specific receptors initiates polyclonal B-cell activation. Although a number of DNA or sODN-binding receptors have been found on a variety of cells, and binding of sODNs to some of these receptors may have functional consequences (BENNETT et al. 1985, 1987, 1991; LOKE et al. 1989; YAKUBOV et al. 1989; HEFENEIDER et al. 1990, 1992; GESELOWITZ and NECKERS 1992; PEARSON et al. 1993; KIMURA et al. 1994; BELTINGER et al. 1995; BENIMETSKAYA et al. 1997), the nature of the sODN-binding receptor on human B cells that eventuates in polyclonal activation remains unclear.

### 3.2 The Nature of the Stimulatory sODNs

A number of possible characteristics could account for the stimulatory properties of sODNs. First, there might be unique sequence motifs accounting for the stimulatory activity of sODNs. Alternatively, sODNs might assume unique secondary or tertiary structures that are necessary to activate human B cells.

To identify potential sODN sequences necessary for B-cell responses, one of the most stimulatory sODN, antisense HIV (HIVas), was examined in detail. Three 15bp sODNs that contained the sequences corresponding to the 5', central, and 3'regions of HIVas were synthesized. The 5' 15bp of HIVas stimulated B-cell activation to a degree comparabe to that induced by the full-length 27-mer, whereas the central and 3' 15-mers were less stimulatory (LIANG et al. 1996). Comparison of the sequences indicated that TCGTCG was present only in the 5' 15bp of HIVas, but not in the others. To determine whether this motif was stimulatory, the activity of additional 15bp sODNs was tested. Of these compounds, (TCG)<sub>5</sub> induced B-cell activation equal to or greater than that induced by HIVas, whereas a 5' TCG doublet followed by a 3' G nonamer failed to stimulate B cells. Additional sODNs, in which the TCG repeat was altered to ACG, TCC or TGG, had no stimulatory activity (LIANG et al. 1996). In order to examine the minimal length of active sODNs, additional sODNs that contained two to four repeats of TCG were synthesized. sODNs containing three TCG repeats were stimulatory, but a tandem TCG had minimal stimulatory activity. Taken together, these data indicate that  $(TCG)_n$ , where n > 3, is a minimal stimulatory element. In addition to TCG repeats, repeats of the entire CpG motif that stimulated murine B cells also activated human B cells (LIANG et al. 1996). However, these motifs are not necessary for human B-cell activation, since sODNs containing no CpG also induce maximal human Bcell activation. Thus, after testing 62 ODNs and sODNs, a unique sequence accounting for the stimulatory activity of sODNs could not be identified.

To examine the second hypothesis that secondary or tertiary structure rather than a specific sequence might be involved in stimulating human B cells, <sup>32</sup>P-labeled sODNs were resolved by polyacrylamide gel electrophoresis (PAGE) and detected by autoradiography. The results showed that HIVas, HSVas and MCMT<sup>+</sup> form tertiary structures, as indicated by their aberrant mobilities in non-denaturing gel. This finding suggested that structure rather than primary sequence might be an

important determinant of the stimulatory capacity of sODNs, but the structures that could be necessary for stimulating B cells remain unknown.

### 4 Human Monocytes Enhance Response of Human B Cells to sODNs

In addition to binding human B cells, most stimulatory sODNs also bound to almost all the monocytes and most NK cells in the peripheral blood, but they did not bind to T cells (Fig. 3). Moreover, highly purified human monocytes augmented B-cell Ig production induced by most stimulatory sODNs. These results suggested that, in addition to a direct stimulatory effect on human B cells, sODNs can also co-stimulate human B cells by an indirect effect mediated by monocytes.

### 5 Differences Between Human and Murine B Cells in the Responsiveness to DNA and sODNs

The current data suggest that human B-cell activation induced by sODNs is governed by different principles than those observed in the mouse (Table 2). First, the response of human and murine B cells to bacterial DNA is different. It has been demonstrated in mice that *E. coli* genomic DNA activated B cells both in vivo and in vitro. In contrast, human B cells failed to respond to *E. coli* DNA in vitro (LIANG et al. 1996). Whereas sera from normal subjects contained antibodies that bound DNA from two bacterial species, *M. lysodeikticus* and *S. epidermidis*, it is not known whether DNA from these micro-organisms directly stimulate human B cells.

	Murine B cells	Human B cells
Stimulation by bacterial DNA	E. coli	None
Activation motifs	CpG motif (necessary)	CpG motif (not necessary); TCG repeats
Cellular entry of stimulus	Necessary	Not necessary; activation is receptor mediated
Cellular activation	95% of B cells in cell cycle	<20% of B cells in cell cycle
Mechanism	IgM production requires IL-6	IgM production is independent of stimulation of IL-6

**Table 2.** Differences between human and murine B cells in responsiveness to DNA and synthetic oligodeoxynucleotides. IgM, immunoglobulin M; IL, interleukin

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Secondly, the basic activation motif appears to be different in man and mouse. The CpG motif has been shown to be sufficient and necessary for inducing murine B-cell activation (KRIEG et al. 1995). In contrast, for stimulation of human B cells, the CpG motif is sufficient but not necessary. In addition, (CpG)<sub>n</sub>, which is a potent stimulator of murine B cells, does not appear to be capable of inducing maximal human B-cell activation, although it did stimulate human B cells minimally (LIANG et al. 1996). Of note, TCG repeats activate human B cells comparably to CpG motifs (LIANG et al. 1996), but this sequence has not been tested in murine systems. Of importance, sODNs containing no CpG dinucleotides are also capable of activating human B cells. It should be noted that sODNs containing no CpG motifs also have been found to activate murine B cells (BRANDA et al. 1996); MONTEITH et al. 1997), suggesting that the difference between mouse and human in this regard may be more relative than absolute.

Thirdly, the mechanism of B-cell activation also appears to be different between mice and humans. It has been suggested that binding to surface receptors is not involved in murine B-cell activation, whereas cellular uptake of the sODNs seemed to be required (KRIEG et al. 1995). In contrast, sODNs immobilized so as to prevent uptake induced maximal activation of human B cells (LIANG et al. 1996). Binding of sODNs to human B cells appears to be receptor mediated, indicating that sODNs stimulate human B-cell activation by binding to surface receptors. The nature of these receptors remains to be delineated.

Fourthly, the character of B-cell activation seems to be different in man and mouse. It has been reported that DNA or sODNs containing the CpG motif induced more than 95% of murine B cells to enter the cell cycle and also rescued murine B cells from apoptosis (KRIEG et al. 1995; YI et al. 1996b, 1998; MAC-FARLANE et al. 1997; WANG et al. 1997; YI and KRIEG 1998). In contrast, less than 20% of human B cells were in the cell cycle after stimulation with active sODNs. Whether this represents a species difference or is related to the responsiveness of human peripheral blood B cells relative to murine splenic B cells remains to be determined.

Finally, the cellular mechanism of B-cell activation induced by sODNs seems to be different between mice and humans. It has been shown that DNA or sODNs containing the CpG motif induced marked IL-6 secretion from murine B cells and that IL-6 was necessary for IgM production induced by sODNs (KLINMAN et al. 1996b, YI et al. 1996a,c). In contrast, limited IL-6 secretion was detected in the supernatant of human B cells cultured with stimulatory sODNs, and neutralizing antibodies to IL-6 only modestly inhibited the IgM production of human B cells. These results suggest that sODNs directly stimulate human B-cell proliferation and differentiation and that IL-6 production plays only a modest role. Taken together, human and murine B cells appear to differ significantly in their responses to DNA and sODNs, implying that results derived from studies of murine B cells can not be directly extrapolated to human B-cell responses.

### 6 Summary and Conclusion

Emerging information has documented that certain DNA and sODNs can be both immunogenic and immunostimulatory. sODNs, but not DNA, induce T-cellindependent polyclonal activation of human B cells by engaging cell-surface receptors. Manifestations of sODN-induced human B-cell activation include expression of activation markers, proliferation, Ig production and anti-DNA antibody production. IL-2 and intact T cells enhanced B-cell responses to sODNs but were not required. Monocytes also provided a modest enhancement of human Bcell responses induced by sODNs. The chemical nature of sODNs capable of stimulating human B cells and the specific cell-surface receptors involved have not been completely delineated. Further studies will be necessary to elucidate the potential role of stimulatory sODNs in disease pathogenesis and to develop a means to employ ODNs as therapeutic agents in humans.

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