## MYOCARDIAL ENERGY METABOLISM

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edited by

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For Lieneke

MYOCARDIAL ENERGY METABOLISM offers a scientific survey of the biochemistry of the heart, with an emphasis on the energy aspects. This approach may be looked upon as complementary to the classical haemodynamic view of the heart. Sections on *Basic mechanisms, Models and techniques, and on Clinical implications* make this book a worthwhile acquisition for cardiologists, cardiac surgeons, basic scientists in cardiovascular physiology, as well as for students with a more than average 'heart' for the organ.

Not all basic questions with regard to formation, breakdown and usage of energy (ATP) in the heart have been asked or answered, but invited experts from the fields of cardiac biochemistry, (electro)physiology, pharmacology, anesthesiology and clinical cardiological research have extensively covered the subject from their points of view. Furthermore, MYOCARDIAL ENERGY METABOLISM shows how basic scientific knowledge may be integrated into cardiological practice. Especially the potential that new techniques like NMR and PET-scanning offer to the cardiologist of the near future becomes clear when the heart is studied with regard to its energy metabolism.

I wish to thank the authors for their effort. I am especially grateful to Peter Achterberg for helpful discussion and contribution.

Rotterdam, April 1988

Jan Willem de Jong

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**Basic Mechanisms** 

### Chapter 1

# **ATP-Metabolism in Normoxic and Ischemic Heart**

#### J.W. de Jong and P.W. Achterberg, Cardiochemical Laboratory, Thoraxcenter, Erasmus University, Rotterdam, The Netherlands

ATP is the immediate fuel for the heart pump. The balance between ATP-synthesis and -utilization is delicate. Under normal circumstances the energy for ATPsynthesis derives from oxidative phosphorylation. Anaerobic glycolysis can only partly substitute for this process. During insufficient perfusion of cardiac tissue, ATP breaks down via ADP and AMP to adenosine, inosine, hypoxanthine, xanthine and urate. The nucleosides and purine bases leave the heart. They are useful for the diagnosis of ischemic heart disease. Various drugs, such as beta blockers and calcium entry blockers, can reduce ATP-catabolism. After restoration of blood supply to the ischemic heart, ATP-content remains low for a long time. During ischemia the decrease in ATP correlates with the decrease in heart function. This relationship is not so obvious in the reperfusion phase. Theoretically, a decreased myocardial ATP-content can be replenished by a) (ribo)phosphorylation of adenosine and adenine, b) salvage of inosine and hypoxanthine, and c) de novo synthesis from, a.o., amino acids. For therapeutic purposes inosine is a good candidate.

#### **Energy Sources**

The energy, needed for the continuous contraction of the heart, derives from the breakdown of ATP to ADP. Under aerobic conditions oxygen-dependent catabolism of nutrients produces most ATP in the numerous mitochondria in heart muscle (see Fig. 1). This oxidative phosphorylation is under triple control, i.e., by ADP, oxygen and substrate (4). Mitochondrial ATP moves via the adenine nucleotide translocator and creatine kinases to the cytosol, for use by the contractile proteins.

Heart supplies the other tissues with the prerequisites of normal cellular function. It pumps oxygen-rich and nutrient-containing blood through the body. The heart muscle itself receives oxygen and nutrients via the coronary circulation. Myocardial oxygen use (for contraction), oxygen supply and coronary flow are in delicate equilibrium. A shift occurs if, for instance, the amount of cardiac work varies. Catabolism of ATP to ADP during myocardial contraction happens at a very fast rate. About 4-8% of the total ATP (20-25 umol/g dry wt.) breaks down to ADP during a single contraction (8). The ADP formed can be reconverted back to ATP through the action of creatine kinase with breakdown of creatine phosphate to creatine. In this way creatine phosphate provides a temporary buffer against rapid ATP-depletion (Fig. 1). A more important function of creatine kinase is ATP transport from mitochondria to actomyosine (Chapter 7). Elevated ADP levels can also be reconverted to ATP by the enzyme adenylate kinase with simultaneous formation of AMP. This AMP can be deaminated to IMP [a major pathway in skeletal muscle (see Chapter 6)], or dephosphorylated to adenosine. The heart retains the major part of the nucleotides, in contrast to adenosine and other nonphosphorylated purines.

#### ATP catabolism

The heart releases adenosine and other purines as soon as ATP-breakdown exceeds ATP-production. Their appearance in the coronary circulation correlates with

increased coronary flow (Chapter 8) and thereby increased oxygen supply to the heart. This enhances ATP-formation, and decreases both adenosine release and coronary flow. Adenosine breaks down quickly to inosine and further to hypoxanthine, xanthine and urate (Fig. 2). These compounds also appear in the coronary effluent Xanthine and urate can only be formed if the enzyme xanthine oxidase/dehydrogenase is present in heart.

As mentioned before, AMP breakdown to inosine occurs either via adenosine or IMP. 5'-nucleotidases, present in the cell membrane and the cytosol, catalyze the former pathway. A soluble AMP deaminase catalyzes the latter pathway. Regulation of these enzymes takes place by ATP and the adenylate energy charge (11,18). Adenosine forms also from S-adenosylhomocysteine (SAH). The enzyme involved, SAH-hydrolase, is rather inactive, but it is important for adenosine production during normoxia (1). Inosine derives from either adenosine (catalyzed by an active adenosine deaminase), or from IMP [catalyzed by 5'-nucleotidase (2,18)].

In turn inosine breaks down to hypoxanthine, a reaction catalyzed by the reversible nucleoside phosphorylase reaction (2,10,12). The next step is hypoxanthine catabolism to xanthine and further to urate. Xanthine reductase catalyzes both reactions (16). There is considerable species variation in activity of the cardiac enzyme; it is virtually absent from rabbit heart (15). During ischemia conversion of the reductase to the oxidase form can take place; the oxidase generates superoxide radicals during urate formation (14). Clinical trials with cardiac xanthine oxidase inhibitors should wait until one has properly demonstrated the enzyme in human heart. Relatively little information is available on cardiac XMP and GMP catabolism (19).



Fig. 1. Major pathways of ATP-formation for myocardial contraction. Mitochondria produce ATP by oxidative phosphorylation; the so-called adenine nucleotide translocator transports it to the cytosol. Immediate transfer of the "high-energy" phosphate group of ATP to creatine phosphate (CrP) occurs via the mitochondrial creatine kinase (CK), bound to the mitochondria (a). Cytosolic creatine kinase (b) reconverts this creatine phosphate to ATP. Myocardial contraction consumes ATP. Fatty acids, ketone bodies, glucose and lactate break down to carbon dioxide and water in the presence of sufficient amounts of oxygen (see Chapters 3 and 4). During hypoxia or ischemia, lactate leaves the heart. ATP-production via anaerobic glycolysis is insufficient. Ox, red = oxidized and reduced components of the respiratory chain, respectively.

The release of adenosine and catabolites (i.e., inosine, hypoxanthine, xanthine and urate) is useful to get an indication of the energy status of the heart. Similarly, with cardiac ATP and creatine phosphate determination, for example in perioperative biopsies, one can assess the "energetic quality" of heart tissue (Chapters 11 and 22). Both in experimental and clinical studies, the increased release of purines is a good indicator of myocardial hypoxia or ischemia (Chapter 21).

Drugs, found to protect the ischemic heart, decrease ischemia-induced purine release. Calcium entry blockers reduce net ATP-breakdown and purine release (5,6) by either increasing ATP-formation through coronary vasodilation, by decreasing ATP-breakdown through overload reduction, or through negative inotropy (Chapter 24). Beta-blockers decrease purine release (20) induced by ischemia. Cardiac purine release is a valuable tool in pharmacological studies.

#### **ATP** Anabolism

Several of the purines mentioned above can be incorporated into the myocardial ATP-pool, which diminishes severely during prolonged ischemia. In the heart only a small amount of adenosine forms constantly from AMP, and rephosphorylates immediately to AMP (3). An active adenosine kinase in the cytosol catalyzes the phosphorylation of adenosine (7). The enzyme has a low  $K_m$  (0.5 uM). This pathway is useful to raise a decreased myocardial adenine nucleotide pool (13), but hypotension and renal vasoconstriction are potential problems. In addition, adenosine elicits electrophysiological effects (Chapter 9).

Salvage of hypoxanthine, or its precursor inosine, is feasible through its reaction with 5-phosphoribosyl-1-pyrophosphate (PRPP), catalyzed by a reversible hypoxanthine phosphoribosyltransferase. The product, IMP, converts to AMP via adenylosuccinate (enzymes involved: adenylosuccinate synthase and lyase). The PRPP concentration is low (Chapter 20), and adenylosuccinate metabolism is slow. Therefore, the overall conversion of inosine and hypoxanthine to AMP is not a rapid process, but it is promising for cardioprotection (see Chapter 26). Almost nothing is known about xanthine and guanine salvage in heart tissue (19). Hypoxanthine accomplishes active regeneration of GTP in post-ischemic hearts (10).



Fig. 2. Scheme depicting AMP catabolism and anabolism. AICAR = 5-amino-4-imidazolecarboxamide ribotide, AMPS = adenylosuccinate, SAH = S-adenosylhomocysteine.

In principle, ribophosphorylation of adenine [catalyzed by adenine phosphoribosyltransferase (17)], could also increase the cardiac AMP level. However, adenine feeding is toxic (21).

Another pathway leading to AMP - de novo synthesis of the purine skeleton from small molecules, such as glycine (22) - is very slow. It is a multistep synthesis, involving many ATP's. PRPP is rate-limiting. A trial to short-cut this pathway with 5-amino-4-imidazolecarboxamide riboside (AICAriboside), which can be phosphory-lated to the intermediate AICAR (see Fig. 2), has met with little success (19). Ribose speeds up salvage and de novo synthesis (Chapter 10).

#### **ATP and Function**

The ATP-level after hypoxia and reperfusion of the heart correlate well with myocardial function (see Chapter 5). Stimulation of myocardial ATP-pool resynthesis by giving purines could promote the velocity and quality of post-ischemic cardiac recovery (Chapters 10 and 23). It could also provide additional protection against a lethal second ischemic attack. Moreover, myocardial protection or preservation during open-heart surgery or heart transplantation could also gain from well-defined purine administration, or prevention of purine release (Chapter 26).

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### Chapter 2

# **Development of Myocardial Energy Metabolism**

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Age-dependent changes in metabolism, structure and function of the heart may be responsible for differences in tolerance to hypoxia or ischemia. Presumably, the newborn heart has a relatively higher capacity for anaerobic glycolysis and a lower maximal capacity for aerobic ATP-formation than the adult heart. Work performance of the in vivo heart under physiological conditions shows an age-dependent increase. Hypoxic ATP-breakdown rate seems lower in newborn than adult hearts under conditions where anaerobic glycolysis is not higher in newborn than adult hearts. Pathological conditions may influence normal cardiac development and energy metabolism, leading to alterations in the supposedly superior tolerance of the newborn heart to ischemia. During hypoxia and ischemia, the balance between ATPformation and ATP-breakdown determines the eventual outcome. Developmental differences exist in ATP- and adenosine catabolism. These may contribute to agedependent differences in cardiac hemodynamics and coronary flow regulation. Further investigations should show whether and how both formation and breakdown of ATP are altered in newborn versus adult hearts under clinically relevant conditions.

#### Metabolic and Functional Development of the Heart

Birth and the perinatal changes in cardiopulmonary circulation submit the newborn heart to increased work load. Adaptations such as increased cell number follow, as well as changes in intracellular structures and activities of various enzymes. The number and size of mitochondria in cardiac myocytes increase (11). A parallel increase is found in the activity of mitochondrial proteins, related to the oxidative formation of ATP (19). The ratio of mitochondrial over total cardiac protein increases from 0.24 at birth to 0.47 in the adult rat heart. Foetal and neonatal heart has been repeatedly found to have a higher glycogen content and a higher capacity for anaerobic ATP-formation. The rate of anaerobic glycolysis decreases markedly in rats by the day of birth (21,22) with a further steady decline during the first postnatal week. Postnatal lactate dehydrogenase (LDH) and hexokinase activities of the guinea-pig heart decrease in a similar way (4). Increases in LDH-activity in rat heart between 1 and 6 weeks of age have also been reported (3), however. Anaerobic glycolysis is important for cardiac energy production during the hypoxic stress around birth. The increase in fetal rabbit cardiac glycogen content by glucose infusion to the mother, six hours before delivery, causes a better ability of newborn hearts to sustain hypoxic beating (17). In fetal lambs a linear relationship between ATP and glycogen contents exists during hypoxia in utero (23), as well as a correlation between the available glycogen and fetal cardiovascular functioning. Similarly, alterations in fetal heart rate and ECG-patterns, signifying ischemia, are related to ATP and glycogen contents (35).

The above-mentioned postnatal decline in cardiac anaerobic capacity and rise in mitochondrial mass is linked with alterations in fatty acid metabolizing capacity. A decreased carnitine-dependent fatty acid oxidation is found in calf-heart mitochondria as compared to adult bovine mitochondria. Following the suggestion

that palmitoylcarnitine synthesis is rate-limiting in newborn hearts (43), a steady, postnatal increase in palmitoyltransferase activity has been observed in rat- (42) and guinea-pig (4) heart. The general inference from the above (and many other) studies is that the fetal and neonatal heart can presumably synthetize more ATP by anaerobic glycolysis than the adult heart, e.g., during hypoxia (26,27). One needs to bear in mind, however, that anaerobic ATP-formation from glucose would need to have an at least fifteen-fold higher glucose-utilization than oxidative breakdown of glucose to generate a similar amount of ATP. The crucial question, therefore, is how much ATP needs to be generated in hypoxic or ischemic newborn and adult myocardium. The answer is, of course, that the amount of ATP synthetized should preferably equal the amount broken down. The myofibrillar contractile ATP-ases and sodium- and calcium-pumping ATP-ases are the main contributors to ATP-breakdown. Their activity directly relates to the work performance of the heart. A doubling of rat cardiac Ca<sup>2+</sup>-stimulated myofibrillar ATP-ase activity takes place between one and six weeks of age (3) and comparable changes also occur in rabbit heart (32). Moreover, rat cardiac actin, determined as protein-bound ADP, increases with age too (41). Fetal ventricular myocardium contains a higher proportion of non-contractile over contractile mass. The newborn sheep heart presumably contains fewer contractile units or sarcomeres (16). This would predispose to the observed smaller maximal velocity of shortening in isolated muscle of the fetal lamb heart (16) and the altered preload-developed pressure relation in isolated newborn rat hearts (25). Developmental changes in the anatomy and function of sarcoplasmic reticulum and calcium homeostasis are implicated as the direct cause of differences in newborn and adult heart function (5, 15, 32, 38), which in addition can cause differences in ATP-turnover (46).

Does the newborn heart need to synthetize similar amounts of ATP as the adult heart does? It is generally assumed that heart rate is a major determinant of work performance in the developing heart. In rats, the spontaneous heart rate increases during the first ten postnatal days from around 300 to 425 beats/min. Thereafter it gradually decreases in the next two months to around 350. Simultaneously, the resting systolic blood pressure increases over the first 60 postnatal days from around 40 to 110 mmHg, with parallel changes in diastolic blood pressure (from 20 to 80 mmHg, ref. 31). A rough indicator of cardiac work performance (i.e., heart rate times blood pressure) therefore increases nearly three-fold over the first 20 to 30 postnatal days. It appears therefore that the newborn heart has a lower need for ATP-formation than adult heart under similar physiological conditions.

In the next paragraph we will discuss the potential differences in tolerance to hypoxia and ischemia of the newborn and adult heart as reflected in cardiac ATP-contents. From the above, however, it is clear that increased capacity for anaerobic glycolysis will only be the assumed (26,27) relevant factor in tolerance to hypoxia, if the rates of ATP-breakdown are comparable in newborn and adult heart. This is most probably not found under physiological or clinical conditions. In addition it remains to be demonstrated, whether the observed developmental differences in hearts of experimental animals are relevant to human pediatric cardiology. And, if so, on what developmental time scale.

#### Developmental Differences in Cardiac Tolerance to Hypoxia and Ischemia

Supposedly the newborn heart has a greater tolerance to ischemia or hypoxia than the adult heart (26,27) as reflected by better function and ATP-content. Better hypoxic ATP-preservation might result from increased ATP-formation by anaerobic glycolysis as suggested by higher lactate release from hypoxic newborn versus adult heart preparations (26,27). However, the newborn heart seems less tolerant to ischemia as judged from ischemic lactate accumulation (10,45) and/or the time of onset of ischemic contracture (10). Lactate accumulation has been indicated as a



Fig. 1. Energy metabolism in adult and newborn hearts during hypoxia in the presence or absence of glucose. (A,B): Adenine nucleotides and glycogen after 20 min hypoxia plus (A) or without (B) glucose (10 mM). (C,D): Lactate release from adult (circles) and newborn (triangles) hearts during hypoxia plus (C) or without (D) glucose. Newborn hearts have higher glycogen contents (A,B) and lower lactate release (C,D) than adult hearts. Moreover, they have similar (A) or higher (B) ATP-contents. This means that newborn hearts have lower rates of ATP-breakdown during hypoxia than adult hearts. Means  $\pm$  S.E.M. (n = 4).

major factor in determining cardiac function after ischemia and reperfusion (33). Because of the relatively large cytosolic volume in newborn than adult hearts, a larger amount of lactate in newborn heart may reflect a similar increase in the actual concentration of lactate, however. Other factors, which can influence the outcome of ischemia, apart from the degree of coronary flow reduction and species used, are the occurrence of acidosis and calcium-accumulation, the temperature, work-performance, pre-ischemic quality of the heart and probably age. Our studies on isolated, perfused adult and newborn (10-days old) rat hearts show that 15 min of global ischemia at a fixed heart rate leads to a similar decrease in adult and newborn ATP-content (12). Hypothermia during ischemia has a similar protective effect on ATP-loss in ischemic adult and newborn hearts. A second series of experiments (Fig. 1) demonstrates that after 20 min of hypoxia (300 beats/min) in the presence of glucose, adult and newborn hearts display a similar decrease in ATP-content. Adult hearts, however, produce far more lactate and brake down their glycogen content faster than newborn hearts. In the absence of substrate, however, newborn hearts preserve higher ATP-contents than adult hearts, without faster glycogen breakdown. We have to conclude therefore that newborn hearts have a lower rate of hypoxic ATP-breakdown, because we find no indications of faster anaerobic ATP-synthesis. Lower myocardial energy demand of hypoxic newborn than adult heart has also been inferred (46) from studies on rabbit heart ventricular septal preparations. During ischemia in adult heart, glycolysis will eventually be strongly inhibited (34), whereby the onset of inhibition is related to the extent of flow reduction and the duration of ischemia. No comparable studies have been performed in immature hearts yet. How different the conclusions from experiments on age-dependent tolerance to ischemia sometimes may be, the difference between synthesis and breakdown of ATP is a major factor determining the outcome of ischemia in both adult and newborn hearts.

#### **Development of Coronary Vasoactivity**

Contrasting reports have appeared on the development of adenosine-mediated coronary vasodilation. In newborn lamb heart, adenosine is not involved in the metabolic regulation of coronary flow (14). Other authors reach the opposite conclusion for this species (30) and for one-week-old guinea-pig hearts (40). Finally, in rabbit heart the response to infused adenosine increases with age (8). In experiments on isolated ten-day-old and adult rat hearts, we observe large, agedependent differences in the composition of purines released, both during post-ischemic reperfusion (12) and during hypoxia. At similar rates of total purine release, newborn hearts release far less adenosine than adult hearts do (Fig. 2), but similar amounts of inosine. It is not yet clear, whether the cause is more active deamination and/or phosphorylation of adenosine, or a shift from breakdown of AMP to breakdown of IMP in newborn hearts. The catabolism of infused adenosine is very rapid, even in adult rat heart (1). This may greatly influence the adenosine concentration at the site of its vasodilatory receptor. The differences in myocardial adenosine production described here, may influence the sensitivity of the newborn coronary system towards infused adenosine. It may even be the cause of different autoregulatory properties of the newborn heart.

#### Myocardial Reperfusion Damage

Oxygen-radical mediated reperfusion damage may contribute to post-ischemic cardiac dysfunction (20). The enzyme xanthine oxidase is repeatedly implicated as a source of cardiac oxygen-radicals during reperfusion (9). We find the enzyme to be present in adult rat heart (37). However, from both direct measurements of xanthine oxidase activity in homogenates and from the determination of the enzyme's products xanthine and urate in perfusates, we must conclude that xanthine oxidase/dehydrogenase is virtually absent from newborn hearts (2,36). Other sources of oxygen-radicals may well be present in both adult and newborn hearts, however. We

find that during reperfusion after ischemia, newborn hearts release significantly less ATP-catabolites than adult hearts do (2). This strongly suggests a better tolerance of the newborn heart to reperfusion damage. However, young patients, undergoing open-heart surgery for correction of Tetralogy of Fallot, show increased myocardial lipid peroxidation (13) after reperfusion.

#### **Pathological Development of the Heart**

Myocardial ischemia in infancy can be subdivided into several groups (18). First, there is ischemia associated with birth asphysia (hypoxia), cerebrovascular accidents and respiratory distress of the newborn. Secondly, myocardial ischemia can be associated with congenital or acquired abnormalities of coronary arteries. Thirdly, of course there is the group of congenital heart diseases, involving left or right ventricular outflow obstruction. Especially in the latter group, persistent alterations in volume or pressure load cause hypo- or hypertrophic abnormalities. These may alter the sensitivity to ischemic stress. In addition, pediatric cardiac pathology has been observed in a variety of congenital disorders of metabolism (29). Each of these could specifically influence cardiac tolerance to ischemia. Because a large group of pediatric patients with congenital anatomical malformations needs cardiac surgery, insight into the factors, which determine the quality of cardioplegic arrest, is of utmost importance. The potential complications of cardioplegia in infants and young children extend beyond developmental differences in structure, metabolism and function of the heart (7). The following aspects can contribute to different outcome of newborn and adult cardioplegia: The size of the patients (relevant to continuation of hypothermia); the size and anatomy of the heart (collaterals can cause rewarming); the underlying disease (hypertrophy); the properties of the newborn circulation (increased chance of oedema in newborns).

The newborn heart can react to hypoxia and hyperoxia by cell-proliferation and cell-differentiation, respectively. The adult heart appears unable to increase its cell-number, but will react to hypoxia by cellular hypertrophy (24). Hypoxia, moreover,



Fig. 2. Adenosine release from adult and newborn hearts during postischemic reperfusion and hypoxia. At similar rates of total purine release, newborn hearts (triangles) release far less adenosine than adult hearts do (circles), both during hypoxia (closed symbols) or during reperfusion (open symbols).

inhibits the normal age-dependent increase in cardiac mitochondrial protein (19,28). Hypoxia, therefore, probably both distorts and slows down normal cardiac development. Animal-model and patient studies show (44) that in case of ventricular hypertrophy especially the endocardial layer of the hypertrophied ventricle has a decreased high-energy phosphate content. This occurs even before the onset of cardioplegia, after which the difference further increases. In a canine model of congenital cyanotic heart disease, created by surgical induction of an atrialpulmonary shunt and banding of the pulmonary artery, ATP-depletion during cardioplegic arrest is faster than in control animals (39). The full effect is not caused by pulmonary banding alone. After reperfusion, myocardial ATP-content and cardiac function remain depressed in the hypoxemic animals. Cardioplegia in patients proves insufficient, when ischemic time exceeds 85 min. In this study (6), an indirect measure of cardiac ATP (i.e., tissue birefringence) predicts the eventual outcome of surgery. Both low ATP at the onset of cardioplegia and severe ATPdecline during prolonged cardioplegia correlate with negative outcome of surgery as reflected by early death or severely low cardiac output.

#### **Final Conclusions**

Studies on postnatal development of structure, function and metabolism of the heart contribute to the general knowledge of the normal adult and newborn heart. Furthermore, certain aspects of cardiac pathology (e.g., hypertrophy, thyrotoxic heart) should be viewed upon as manifestations of abnormal development. The exact knowledge of normal cardiac development is therefore also a prerequisite for the understanding of pathological development.

We have not discussed differences in catecholamine status and reactivity in adult and newborn hemodynamics nor the possibly different sensitivities to calcium or acidosis. We feel, however, that by looking at the difference between synthesis-rate and breakdown-rate of ATP, both adult and newborn cardiac tolerance to ischemia can be further investigated. Such studies in relevant models and in the clinical setting are needed for pediatric cardiac patients to benefit, as well as for our general understanding of cardiac pathophysiology and development.

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#### Chapter 3

# **Principles of Fuel Metabolism in Heart Muscle**

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The functions of metabolism in the heart are few in number and can be understood without detailed knowledge of elaborate chemical formulas. The primary function of the metabolism of energy-providing substrates is the rapid synthesis of ATP from ADP and inorganic phosphate. Under normal conditions, heart muscle covers its energy needs through the oxidation of glucose, fatty acids, ketone bodies or, to a lesser extent, amino acids. This chapter reviews the mechanisms of metabolic control and fuel selection, as they relate to the function of the heart. Particular emphasis is placed on the interaction between substrates and the control of the citric acid cycle, the final common pathway for all substrates providing reducing equivalents for the respiratory chain.

#### 1. Heart Muscle as Site of Energy Transformation

The purpose of this chapter is to provide a brief outline of cardiac fuel metabolism. To begin with, fuel metabolism cannot be separated from energy metabolism, hence the inclusion of this chapter in a book on Myocardial Energy Metabolism. Secondly, a review on cardiac fuel metabolism seems timely in light of the renewed interest in intermediary metabolism generated by recent techniques using tracer compounds labeled with positron emitting radioisotopes, or by magnetic resonance spectroscopy to assess tissue function and metabolism in the intact heart.

Compared with the vast biochemical literature such a review is necessarily superficial and fragmented. Fortunately, the functions of metabolism in the heart are few in number and can be understood without detailed knowledge of elaborate chemical formulas or reactions. The main purpose of cardiac metabolism is to provide energy for contraction, ion transport, biosynthesis and degradation. The conversion of chemical energy into mechanical energy is termed energy transformation. Like any other living tissue, heart muscle captures and utilizes chemical energy in the form of ATP. The tissue content of ATP is normally around 20 umol/g dry weight. At an  $O_2$  consumption of 4.5 mmol/min/g dry wt. (67) and a P/O ratio of around 3, heart muscle utilizes 1400 umol ATP/min/g dry wt. or 23 umol ATP/s/g dry wt. This calculation shows that under steady-state conditions ATP has to be regenerated as rapidly as it is broken down. The overall concept of this "ATP cycle" is depicted in Fig. 1.

The sources of ATP in heart muscle are oxidative phosphorylation of ADP in the respiratory chain and, to a lesser extent, phosphorylation of ADP either by substrate level phosphorylation in the glycolytic pathway and the citric acid cycle, or by the action of creatine kinase. Since oxidative phosphorylation is quantitatively the main source of ATP and because the key enzymes of oxidative metabolism are located in mitochondria, myocardial cells are well endowed with organelles.

Oxidative phosphorylation of ADP depends on the production of reducing equivalents (protons) and the passage of electrons along the respiratory chain. Since reducing equivalents are produced in the course of substrate catabolism to acetyl-CoA and

subsequent oxidation of acetyl-CoA in the citric acid cycle, it is convenient to group the main energy-providing reactions into three stages (Fig. 2). The first stage comprises all reactions leading to acetyl-CoA, the second stage the oxidation of acetyl-CoA in the citric acid cycle (resulting in liberation of  $CO_2$  and reducing equivalents), and the third stage consists of the flow of the electrons down the respiratory chain leading to release of free energy, which is conserved in the energy-rich phosphate bond of ATP.

#### 2. Some Early Experiments

A major impetus for research in cardiac metabolism came from the work of Richard Bing and his co-workers who measured arterio-venous differences of substrates and products across the heart (see ref. 3 for review).

Using coronary sinus sampling, it was shown that the human heart uses the three foodstuffs carbohydrates, fatty acids and (under certain circumstances also) amino acids to varying degrees for its energy production. The mammalian heart can therefore be likened to an omnivore. Bing's work suggested that the arterial concentration of *carbohydrates* and the presence of insulin affect theire relative myocardial usage. The total aerobic metabolism of *glucose, lactate* and *pynuvate* did, however, not account for the total oxygen consumption of the heart, and it was quickly recognized that oxidation of *fatty acids* and *ketone bodies* made up the considerable balance. Myocardial utilization of fatty acids was particularly high during starvation. While oxidation of ketone bodies accounted for approximately 5% of the total myocardial oxygen consumption in normal individuals, the diabetic dog heart utilized a consistently larger quantity of these fuels (70). The utilization of ketone bodies seemed to be governed by their arterial concentration and the quantity of carbohydrate available (4). Bing et al. (4) further suggested that both the human heart and the dog heart may even extract considerable amounts of *amino acids* from the blood when their plasma levels rise to greater than physiological values. This ability of the heart to utilize various substrates can be regarded as an "important factor of safety" (3) in the maintenance of an adequate fuel supply for an organ of vital importance for the survival of the total organism.

#### **3. Principles of Metabolic Control**

It is a characteristic property of living cells to permit complex chemical reactions to proceed quickly at relatively low temperatures and concentrations. This efficient energy transfer in cells occurs through enzyme-catalyzed reactions and is probably the result of a long process of evolutionary selection (1).



Fig. 1. The cycle of cardiac energy metabolism (ATP cycle). Breakdown of fuels leads to the capture of energy in the form of ATP. Hydrolysis of ATP and ADP and inorganic phosphate  $(P_i)$  provides the energy for contraction, ion transport, as well as synthesis and degradation of biomolecules. Stores of ATP in the tissue are very small and must therefore be continuously replenished by phosphorylation of ADP.

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Activities of enzymes are influenced by temperature, pH, and other factors, such as allosteric activators or inhibitors. The progress of any enzyme-catalyzed reaction can be determined by measuring either product formation or substrate utilization under optimal environmental conditions (Fig. 3). The hyperbolic configuration of enzyme-activity curves shows that the rate of the reaction diminishes with time and reaches its end-point asymptomatically, which is due to a fall in the concentration of substrate as well as a number of other environmental factors.

In most instances substrate availability (as well as product removal) and enzyme activity are the main factors which control the rate of a biochemical reaction. While this is certainly true for the many reactions which are at near-equilibrium, the same does not apply to those enzymes whose activity is modified by factors other than the pathway substrate and which catalyze non-equilibrium reactions. Biochemists identify non-equilibrium or regulatory reations in the first instance by measuring tissue concentrations of products and substrates, by calculating their ratio (mass action ratio) and by comparing this value to the equilibrium constant  $K_{eq}$  of the reaction. If both are similar, the reaction is at near-equilibrium; if they differ significantly, the reaction is at non-equilibrium. The same principle is expressed in the "crossover theorem" by Chance and Williams (9), which simply states that when a pathway is inhibited at a specific reaction, the substrate concentration will increase and the product concentration will decrease. The next step biochemists have taken in the process of identification of a non-equilibrium or regulatory enzyme is testing the enzyme activity in vitro. Such detailed analysis is desirable because regulatory enzymes set the pace for an entire sequence of reactions in a metabolic pathway, as will be discussed below.



Fig. 2. Stages of catabolic reaction in cardiac energy metabolism. Energy-providing substrates are converted to acetyl-CoA, which is, in turn, oxidized in the citric acid cycle. At many stages in their catabolism, the substrates generate reducing equivalents in the form of NADH +  $H^+$  and/or FADH<sub>2</sub>. Transfer of their electrons to the electron carrying chain and their reaction with molecular oxygen leads to synthesis of ATP.

There are two major classes of regulatory enzymes: allosteric (or non-covalently regulated) enzymes and covalently regulated enzymes. The allosteric theory proposes that an enzyme possesses at least two spatially distinct binding sites on the protein molecule, one is termed the catalytic site, the other is termed the regulatory site. A regulator binds to the regulatory site and modulates the activity of the enzyme. Often the product of a reaction or of a metabolic pathway acts as regulator (feedback regulator).

The second class of regulatory enzymes is modulated through interconversion of their active and inactive forms by covalent modification. Examples of covalently modified enzymes are the enzymes of glycogen metabolism and the pyruvate dehydrogenase complex, both of which undergo phosphorylation and dephosphorylation.

#### 4. Control of Metabolic Pathways

The broad objective of the study of enzyme mechanisms in vivo is the understanding of events which link metabolism to function. The full importance of enzymatic reactions can therefore only be understood when they are viewed in the context of a metabolic pathway. A metabolic pathway can be defined as a series of enzymecatalyzed reactions which convert a substrate into a product.

Prevailing ideas on the operational control of metabolic pathways are probably too simplistic when one considers the complex conditions inside the cell. An important concept to be considered is the *concept of flux*, i.e., the flow of matter along a series of reactions. Biochemical maps of a series of reactions provide no more informations than a road map. The real test is the journey itself. According to Newsholme and Leech (42), it is useful to think of a metabolic pathway as a series of reactions initiated and terminated by a flux-generating reaction. The concept of a "pacemaker" reaction for a metabolic pathway was first proposed by Krebs (25). More recently Newsholme and Crabtree (41) have defined a "flux-generating step" as any non-equilibrium reaction saturated with pathway substrate to which all other reactions in the pathway must respond in order to achieve a steady state. As it is understood today, a change in the activity of the regulatory enzyme (or transport step, or work load, for that matter) would cause a change in flux through the pathway as long as the enzyme is saturated with its pathway substrate. Newsholme and Crabtree (41) have termed this process "external regulation" and the control of enzyme activity via changes in the pathway substrate concentration as "internal regulation". The near-equilibrium reactions of the pathway are very sensitive to changes in concentrations of substrate, and a small increase in pathway substrate concentration would stimulate the activity of the enzyme sufficiently to accommodate new rates of flux.



Fig. 3. Progress of an enzyme-catalyzed reaction. Progress of the reaction is in this example monitored by the appearance of a product (solid line) or the utilization of a substrate (broken line) with time (on the abscissa). The reaction assumes the form of a hyperbola. The velocity of the reaction is measured in the initial, linear portion of the curve.

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A traditional approach in studying cardiac metabolism is the analysis of metabolites extracted from rapidly frozen tissue. As discussed earlier, accumulation of metabolites proximal to a metabolic "block" may reveal the presence of a regulatory enzyme. Accumulation of metabolites only suggests, but does not prove, inhibition of flux through a specific step in a pathway. For example, the steady state content of some key metabolites of cardiac energy metabolism may vary considerably depending on the dietary state and substrate supply. It is of note in this regard that there is no correlation between intermediary metabolite levels and cardiac performance. For example, with glucose as the only exogenous substrate, the performance of the heart is excellent, but acetyl-CoA (an important intermediate of oxidative glucose metabolism) is below the detectable range, because the compound is oxidized as fast as it is produced. By contrast, when ketone bodies are the only substrate available to the heart, acetyl-CoA accumulates, but performance and O<sub>2</sub> consumption are low, because of decreased rates of oxidation in the tricarboxylic acid cycle (64). This observation provides an excellent example for the general concept that flux through energy-providing pathways, and not necessarily the steady-state concentrations of metabolites, determines the functional state of the tissue.

#### 5. Metabolism of Energy Providing Substrates

#### A. Carbohydrates

Carbohydrates of major importance as substrates for cardiac energy metabolism are glucose, glycogen, lactate and pyruvate. Glucose is taken up by the heart-muscle cell, phosphorylated and metabolized as glucose 6-phosphate by several distinct enzyme systems to meet the needs of the cell. These systems include: a) degradation via the Embden-Meyerhof pathway (also termed the glycolytic pathway if it entails metabolism of glucose to lactate only), b) conversion of glycogen which serves as an intracellular glucose store, and c) metabolism via the pentose-phosphate pathway which yields ribose and NADPH. The latter pathway is quantitatively of lesser importance in heart muscle than the former two pathways.

In keeping with the scheme presented in Fig. 2, there are three energy-yielding stages of glucose metabolism: the Embden-Meyerhof pathway leading to pyruvate and acetyl-CoA, the citric acid cycle and the respiratory chain. Each stage is regulated by its own set of controls so that overall flux through the pathway proceeds at a rate just sufficient to satisfy the beat-to-beat needs of the heart for the pathway product, ATP. For example, the relative concentrations of ADP in the cytosol control not only the rate of electron transport and oxidative phosphorylation, but also the turnover rate of the citric acid cycle, the rate of pyruvate oxidation and glucose utilization. Thus, whenever hydrolysis of ATP is increased, a whole sequence of enzyme-catalyzed reactions and transport mechanisms is set into motion.

<u>Glucose transport and phosphorylation</u>. Glucose uptake into heart muscle occurs via a stereo-specific passive membrane transport system that displays saturation kinetics and counter transport (35). This system equilibrates intracellular and extracellular concentrations of glucose but does not move sugar against a concentration gradient. The stereo-specificity of the carrier for sugars of the C-1 chain conformation is not matched by the same degree of selectivity.

There are various factors which influence glucose transport by changing either  $K_m$  or  $V_{max}$  of the carrier system. These include hormones, such as insulin, growth hormone, epinephrine and cortisol (for review, see ref. 40), anoxia (34), extracellular glucose concentration and cardiac work (38,39,67). In normal heart muscle, cardiac work, the availability of alternative substrates, as well as plasma insulin and glucose concentrations, act in concert as the most important factors regulating glucose uptake.

When hearts are perfused at very low work loads, insulin nearly doubles glucose utilization and lactate production (34,39,67), and during maximal insulin stimulation net transport exceeds the rate of phosphorylation (34). This finding is in good agreement with recent in vivo studies which have estimated glucose transport with  $[^{3}H]$ -2-deoxyglucose (20).

Phosphorylation of glucose by hexokinase becomes rate limiting for glycolysis at high rates of glucose transport. Rates of glucose phosphorylation measured in vitro are more than twice as high as the maximal rates of glucose utilization by heart at physiological work load and with glucose as the only substrate (41). The intracellular glucose concentration of heart muscle rises with starvation, diabetes, oxidation of fatty acids, ketone bodies or pyruvate, which indicates inhibition of the phosphorylation step. This is most likely due to accumulation of glucose 6phosphate, because of inhibition of the phosphofructokinase reaction (see below).

<u>Glycogen formation and breakdown</u>. Glycogen metabolism is without question the biochemical pathway which has contributed more to the understanding of enzyme regulation and the molecular basis of hormone action than any other known system of metabolic control (10). In this system the first example of the control of enzyme activity by an allosteric regulator (activation of phosphorylase by AMP) was described (13), enzyme regulation by reversible covalent modification (phosphorylation) was discovered (24), and the molecular basis of hormone action was elucidated through the discovery of cAMP (53).

Synthesis and degradation of glycogen occur in two separate pathways. The combined effects of protein phosphorylation and dephosphorylation on glycogen synthase and phosphorylase provide an interlocking system by which homones (such as epinephrine) and mechanical activity (through  $Ca^{2+}$ ) can control the net flux of glucose 1-P into and out of the intracellular glycogen stores. Epinephrine-induced cAMP formation promotes protein phosphorylation and simultaneously inhibits glycogen synthesis while stimulating glycogen breakdown, whereas stimulation of protein dephosphorylation shifts the balance toward glycogen synthesis. Although the interaction of the enzymes of glycogen breakdown and synthesis is more complex than this simple mechanism suggests (for details, see ref. 11), the concept is relatively easy to grasp.

<u>Glycolysis.</u> The catabolism of glucose 6-phosphate in the glycolytic pathway exhibits two major control sites: phosphofructokinase and glyceraldehyde 3-P dehydrogenase.

Allosteric control of phosphofructokinase (PFK) provides for large changes in catalytic activity of the enzyme which consists of six subunits. Phosphofructokinase activity is *increased* by ADP, AMP, Pi, fructose 1,6-bisP (only the alpha anomer), by cAMP, and by ammonium ions. Activity of the enzyme is *decreased* by intracellular acidosis, by ATP, phosphocreatine and citrate. The allosteric control of phosphofructokinase by citrate plays a role in the control of the selection of fuels by decreasing rates of glucose metabolism in the presence of fatty acids, ketone bodies, or pyruvate (51).

The oxidation of glyceraldehyde 3-phosphate to 1,3-diP-glycerate couples reduction of NAD<sup>+</sup> to the formation of a diphosphotriose. It is an energy-conserving step leading to the formation of ATP. The reaction is thought to be at near equilibrium; yet it was found that under conditions of O<sub>2</sub> deprivation or high rates of cardiac work (when PFK becomes strongly activated), glycolysis is controlled further down in the glycolytic pathway (23,55). In vitro the enzyme is strongly inhibited by its products 3-phospho-D-glyceroyl-P and, especially, NADH (33). According to Kobayashi and Neely (23), even in hearts well supplied with oxygen, maximal rates of glucose breakdown are limited by flux through glyceraldehyde 3-P dehydrogenase which is under the restraint of an increasing cytosolic [NADH]/[NAD] as the work load of the heart increases.

<u>Pyruvate Oxidation</u>. In heart muscle pyruvate may either be reduced to lactate (which completes the glycolytic pathway), transaminated to alanine, carboxylated to oxaloacetate or malate, or, most importantly, oxidized to acetyl-CoA. Lactate and most of the alanine are formed in the cytosol by near-equilibrium reactions and both metabolites may be washed out from the cell. In well-oxygenated, working heart muscle, the bulk of pyruvate enters, however, the mitochondrion.

The inner mitochondrial membrane represents a barrier to charged molecules. Specific transport mechanisms are required to enable transport of specific metabolites through the lipid bilayer, and during the past two decades an impressive number of carrier systems have been demonstrated for the transport of both anions and protons (for review, see ref. 29). A specific carrier exists for the transport of pyruvate into the mitochondrial matrix, as it has been most clearly demonstrated with the inhibitor 4-hydroxy-alpha-cyanocinnamate (for review, see ref. 15).

Inside the mitochondrial matrix, pyruvate may be either decarboxylated to acetyl-CoA or carboxylated to oxaloacetate. Although there is a considerable amount of pyruvate carboxylase activity present in heart muscle (59), most of pyruvate is decarboxylated. Oxidative decarboxylation of pyruvate assumes a central position in the regulation of fuel supply to the heart. A system of intricate control mechanisms governs both activation and inactivation of the pyruvate dehydrogenase (PDH) enzyme complex.

The conversion of pyruvate to acetyl-CoA requires the sequential action of three different enzymes: pyruvate decarboxylase, dihydrolipoyl transacetylase, and dihydrolipoyl dehydrogenase. The reaction also requires five different coenzymes or prosthetic groups. These are thiamine pyrophosphate, lipoic acid, CoASH, FAD and NAD. These enzymes and coenzymes are organized into a multienzyme cluster. The overall MW of the PDH complex is about 6 million dalton, i.e., the size of a ribosome. The PDH complex is attached to the inner side of the inner mitochondrial membrane.

PDH is also regulated by covalent modification through a *phosphorylation* and *dephosphorylation* cycle. Multisite phosphorylations provide an indirect means by which the rate and extent of the kinase reactions may regulate reactivation of the PDH complex by the phosphatase. Like most mammalian tissues, heart muscle possesses both active (dephosphorylated) and inactive (phosphorylated) dehydrogenase. The total activity of both together is approximately 30 U/g dry wt. at 30°C (48), of which normally about 20% is in the active form. The relative amount of active PDH increases with an increase in work load. Active PDH may decline to only 1-5% of total PDH during starvation or in alloxan diabetes (22), i.e., when non-carbohydrate substrates become the main fuel for respiration. Both kinase and phosphatae phosphatase are probably active in vivo, and the relative proportion of active PDH must therefore be dependent on the relative activities of kinase and phosphatase as well as the intramitochondrial concentration of the effectors of these enzymes.

Most of the known effectors of the phosphorylation/dephosphorylation cycle are also effectors of the kinase reaction. Of the metabolite pairs ATP/ADP, acetyl-CoA/CoASH, NADH/NAD<sup>+</sup> and lactate/pyruvate, the first component either activates the kinase or serves as substrate, while the second component inhibits the enzyme.  $Ca^{2+}$  and  $Mg^{2+}$  both inhibit the kinase and activate the phosphatase reaction, i.e., lead to activation of PDH. The effects of oxidation of fatty acids or

ketone bodies are likely to be mediated by the increase in acetyl-CoA they induce (22), and the primary effects of the inhibition/inactivation of PDH by fatty acids or ketone bodies is the acetyl-CoA/CoASH ratio (44). Conversely, an increase in cardiac work may inhibit the PDH kinase due to a fall in NADH, acetyl-CoA and ATP, leading to activation of PDH (23).

#### **B.** Fatty Acids

The importance of fatty acids as fuel for respiration and energy metabolism in the heart is easy to understand. First, fatty acids carry considerably more reducing equivalents per g molecular weight than glucose. Secondly, the energy stores of fatty acids in the body in the form of triglycerides (i.e., fatty acids esterified with glycerol) quantitatively by far exceed the energy stores of glucose (i.e., glycogen). Thus, triglycerides constitute the main fuel reserve in mammals. In times of food deprivation, fatty acids are mobilized and serve as the preferred fuel for respiration in most tissues (with the notable exception of nerve tissue and red blood cells).

The mechanism of fatty acid oxidation by beta-oxidation was discovered by Knoop in 1904 long before the glycolytic pathway and the citric acid cycle were known. Despite this early start and despite the importance of fatty acids as fuel for oxidative ATP production in heart muscle and many other tissues, the ratecontrolling steps of fatty acid metabolism are not as well understood as those of carbohydrate metabolism.

The heart can utilize both complex lipids from the blood (14) as well as albuminbound free fatty acids (60). In well-oxygenated heart muscle, the rate of utilization seems to depend chiefly on their plasma concentration and on the work load of the heart (36,45).



Fig. 4. Main pathways of long-chain fatty acid metabolism in heart muscle. Long-chain free fatty acids (FFA) are activated at the outside of the inner mitochondrial membrane and transported across the inner mitochondrial membrane by the carnitine-acyltransferase system. Inside the mitochondrion they undergo beta-oxidation to acetyl-CoA, which is oxidized in the citric acid cycle. FFA activated in the cytosol, but not transported into the mitochondrion, may be esterified with alpha-glycerophosphate to form triglycerides. Heavy lines indicate the main pathway of long-chain fatty acid metabolism in heart muscle. Fatty acids of short chain length do not utilize the carnitine-acyltransferase system and undergo beta-oxidation directly.

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An overview of fatty acid oxidation in heart muscle is shown in Fig. 4. The pathway starts with liberation of fatty acids from triglycerides and/or albumin bound transport of free fatty acids (FFA) in the blood, and it ends with the entry of acetyl-CoA into the citric acid cycle. FFA of any chain length are able to cross the plasma membrane, and are activated on the outer mitochondrial membrane by esterificiation with CoASH to form fatty acyl-CoA. Metabolism of long-chain FFA continues with fatty-acyl transfer to carnitine, transport into the mitochondria in exchange for carnitine, reesterification with CoASH, beta-oxidation and, finally, oxidation of acetyl-CoA. A certain amount of FFA taken up by heart muscle is esterified with glycerol in the cytosol to form triglycerides. Thus, the fatty acyl-CoA formed on the outer mitochondrial membrane has two possible fates, either oxidation via the citric acid cycle in the mitochondria or conversion to triglycerides in the cytosol. Which pathway is taken depends on the rate of transfer of longchain fatty acyl-CoA across the inner mitochondrial membrane and on the rate of esterification in the cytosol. Increased net triglyceride synthesis by the heart has been observed with starvation, diabetes and also with ischemia. It is not known whether this is the result of increased rates of esterification or a decreased rate of lipolysis.

Before the acyl group can be oxidized, it has to be moved from the outer mitochondrial membrane into the mitochondrion. Transfer of long-chain fatty acyl-CoA into mitochondria requires a three-step membrane transport process which is the rate-controlling step for long-chain fatty acid oxidation. The first step of the sequence is the transfer of the acyl group from CoA to carnitine (Fig. 4), catalyzed by the enzyme carnitine acyl-CoA transferase I (CAT I). The carnitine-acylcarnitine translocase system has been partially characterized in isolated heart mitochondria (46) and in the perfused heart (21). An exchange reaction moves acyl-carnitine across the inner mitochondrial membrane in a stoichiometric exchange for carnitine. In the liver, and most likely also in heart muscle, CAT I is the regulatory site for fatty acid oxidation with high concentrations of malonyl-CoA acting as inhibitor of the enzyme during the carbohydrate-fed state (30,31). Malonyl-CoA arises from carboxylation of acetyl-CoA in the cytosol. The presence of this compound in heart muscle has been demonstrated (32). The carnitine-acyl unit traverses through the inner mitochondrial membrane and is transferred to CoASH inside the mitochondrial matrix by a second carnitine acyl-CoA transferase (CAT II) located at the inner surface of the inner mitochondrial membrane.

Once inside the mitochondria, acyl-CoA is committed to oxidation by the system of beta-oxidation. The beta-oxidation pathway is controlled by fluctuations in the concentrations of substrates (acyl-CoA, NAD<sup>+</sup>, and FAD<sup>+</sup>; ref. 7), or, expressed in more physiological terms, by work load and oxygen supply of the heart. It is not yet clear how the rate of oxidation of FFA in the intact heart is related to citric acid cycle activity and the rate of oxidative phosphorylation.

#### C. Ketone Bodies

The ketone bodies, acetoacetate and beta-hydroxybutyrate, are produced by the liver, when supply of FFA is abundant and exceeds the liver's capacity for their complete oxidation in the citric acid cycle. Ketone bodies, like carbohydrates and fatty acids, share in the supply of acetyl-CoA for oxidation in the citric acid cycle and for energy production in heart muscle (72). Compared with the complexity of glucose and long-chain fatty acid metabolism, the metabolism of ketone bodies requires only a few steps, none of which appear to be subject to any regulation.

A unique feature of ketone body metabolism is the activation of acetoacetate to acetoacetyl-CoA by succinyl-CoA, which itself is a citric acid cycle intermediate. A few facts have emerged with respect to the regulation of ketone body metabolism in the heart. The enzymes degrading beta-hydroxybutyrate and acetoaceate to acetyl-

CoA, beta-hydroxybutyrate dehydrogenase, 3-oxoacid-CoA transferase and acetoacetyl-CoA thiolase, are of comparatively high activity in heart muscle (see refs. 2,71). Ketone bodies, when present in relatively high concentrations, spare glucose utilization by the heart. The ability of the heart to oxidize ketone bodies may constitute a buffer against qualitative changes in substrate supply when the relative proportions of oxidizable fuel are shifting as a result of changes in the dietary or hormonal state of the body (56). Even in physiological situations, such as short-term starvation or exercise, ketone body concentrations in the plasma can rise by up to 50-fold (65).

It is of interest to note that ketone bodies, when present as the only exogenous substrate for rat heart, do not sustain the full work output of the heart, and their rate of oxidation in the citric acid cycle is insufficient to meet the energy needs of the heart (64,67). This relative inhibition of the citric acid cycle occurs at the level of 2-oxoglutarate dehydrogenase. It is most likely related to sequestration of the enzyme's cofactor CoASH, which is bound in the form of its thioesters, acetyl-CoA, and as acetoacetyl-CoA, and thus not available for the reaction in the citric acid cycle (64). Whether or not this inhibition of the citric acid cycle is important in clinical situations such as diabetic cardiomyopathy, is a matter of speculation (68).

#### D. Amino Acids

Amino acids are not only the "building blocks" of myocardial proteins, but, through their metabolism, they are also an integral part of myocardial energy metabolism. It cannot be emphasized too strongly that each amino acid is different from the others with respect to control of its metabolism. Many amino acids are either degraded or synthesized by heart muscle (e.g., glutamate, glutamine, aspartate, alanine, leucine) while others are not metabolized at all (e.g., phenylalanine, tyrosine).

In general, transamination is the first step of amino acid degradation in heart muscle. One of the functions of amino-transferases in heart muscle (for comparative data, see ref. 26) is the provision of carbon skeletons for the citric acid cycle. The amino acids aspartate and glutamate also play an important role in the transfer of reducing equivalents across the mitochondrial membrane for oxidation of cytosolic NADH by the mitochondrial electron transport chain (6), as shown in Fig. 5. Current evidence suggests that in heart muscle, and similarly in liver, reducing equivalents are indirectly carried into the mitochondrial compartment by metabolic anions of the malate-aspartate cycle (57,58). In the malate-aspartate cycle, cytosolic NADH, which



Fig. 5. The malate-aspartate shuttle, an important pathway of cytosolic NADH oxidation. Cytosolic NADH arises from the glyceraldehyde 3-P dehydrogenase reaction and can be reoxidized by either the alpha-glycerophosphate dehydrogenase, the malate dehydrogenase or lactate dehydrogenase reaction. The first two reactions are concerned with the transfer of reducing equivalents into the mitochondrial matrix. Quantitatively the most important mechanism for transfer of reducing equivalents is the malate-aspartate shuttle which involves transamination of aspartate and glutamate, as depicted.

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cannot cross the inner mitochondrial membrane, is oxidized by the reduction of oxaloacetate to malate. Malate enters the mitochondrion and is oxidized to oxaloacetate, which is transaminated with glutamate to form 2-oxoglutarate and aspartate. Aspartate and 2-oxoglutarate leave the mitochondrion. Transamination of these metabolites regenerates oxaloacetate and glutamate in the cytosol, the net effect being a transfer of hydrogen ions across the mitochondrial membrane. A second postulated function for the malate-aspartate shuttle is the delivery of intermediates (malate) to the citric acic cycle, acting as a modulator of cycle activity.

Myocardial amino acid metabolism during hypoxia, ischemia and reperfusion has been the subject of a number of studies in the intact and isolated heart muscle (5,66,69). The amino acid alanine was found to be, like lactate, an end product of anaerobic glucose breakdown, arising through transamination of glutamate in the reaction:

This reaction is not the end of the pathway. While alanine leaves the cell, 2oxoglutarate is further metabolized and decarboxylated to succinate via substrate level phosphorylation in the citric acid cycle (63).

#### 6. Competition and Selection of Energy-Providing Fuels

While the heart plays only a minor role in the overall fuel balance of the whole body, it obeys the same rules of metabolic regulation as other tissues do. Specifically, it has been known for a good numner of years that in the fasting animal the energy requirements of the heart are met largely by the oxidation of fatty acids with an respiratory quotient (R.Q.) of 0.7 (18), and that after carbohydrate feeding the heart switches its preference to glucose with an increase of the R.Q. to about 0.9 to 1.0 (19).

It is possible to make a number of assumptions from data available in the literature. As already stated, under aerobic conditions the heart is able to derive energy from a variety of fuel sources such as FFA, ketone bodies, glucose, lactate, and even certain amino acids. Depending on the physiological state of the individual, only plasma glucose levels are relatively constant, whereas levels of lactate, FFA and ketone bodies may vary over a wide range. Thus, it is not surprising that the fuel for aerobic energy production in heart muscle varies with the plasma concentration of individual substrates.

Heart muscle itself is also capable of contributing to the fuels of respiration by releasing both FFA and glucose from their stores of triglycerides and glycogen. The most striking example of endogenous fuel utilization occurs with ischemia, when glycogen is broken down to lactate (see, e.g., ref. 37). Thus, with regard to endogenous fuel consumption, we have the situation that during normoxia and normal coronary flow, exogenous fuels provide the bulk of the fuel for respiration, whereas with ischemia and restricted coronary flow, the endogenous fuel glycogen contributes a large portion of anaerobic energy production through metabolism to pyruvate, lactate and alanine.

Fuel selection by normoxic heart muscle continues to be an intriguing question. The *first factor* which affects the utilization of an individual substrate in a mixture of substrates is its plasma concentration. Glucose concentrations are relatively constant, whereas concentrations of fatty acids, ketone bodies or lactate can fluctuate. Accordingly, FFA have been identified as the preferred fuel for respiration by the heart especially when their plasma concentration was high after an overnight fast (54). Likewise, lactate has been identified as a preferred fuel by the heart when lactate concentrations were high (16).
A second factor which determines selection of fuels by the heart is the ease of access of fuels to the enzymes of oxidative metabolism in the mitochondrion. There are fewer control steps involved in the metabolism of lactate or acetoacetate to acetyl-CoA than in the metabolism of either fatty acids or glucose to acetyl-CoA. Thus, when present in sufficient concentrations, both lactate or ketone bodies may contribute a large amount of acetyl-CoA for oxidation in the citric acid cycle.

A third factor which determines the selection of fuels by the heart is its  $O_2$  supply and the specific P/O ratio (i.e., moles of ADP phosphorylated per mole of  $O_2$ consumed) of fuels. For instance, one mole of glucose, when oxidized, uses 6 moles of  $O_2$  and yields 36 moles of ATP (P/O ratio = 3.0), whereas the oxidation of one mole of palmitate utilizes 23 moles of  $O_2$  and yields 129 moles of ATP (P/O ratio = 2.8). Thus, glucose produces about 8% more ATP per mole of oxygen than do FFA. Although this may not seem a very large difference, an improved P/O ratio may become important for the heart which is extracting very large amounts of oxygen. In situations of limited  $O_2$  reserve, oxidation of glucose can provide a greater yield of ATP. Consequently, the heart may then prefer glucose over FFA as fuel for respiration. Inhibition of the oxidation of a long-chain fatty acid derivative (palmitoylcarnitine) by pyruvate has recently been found in rat-heart mitochondria (8). Suppression of FFA oxidation by glucose and/or pyruvate is different from the situation encountered in true ischemia, where the  $O_2$  supply is completely insufficient to meet the energy needs of cells and rates of anaerobic glycolysis are necessarily increased.

A fourth factor which determines the selection of fuels is the interaction of *intermediary metabolites* arising in the course of degradation of energy-providing substrates with regulatory steps in the pathway of energy conversion of substrates. Such regulatory steps have been described earlier in the discussion of glucose and fatty acid metabolism. For example, the mechanism by which FFA suppress glucose utilization has been delineated in the perfused heart. Addition of fatty acids, ketone bodies or pyruvate cause a major depression in glucose utilization through coordinated inhibition at the four control sites: glucose transport, hexokinase, phosphofructokinase, and pyruvate dehydrogenase by mechanisms already discussed. The opposite question, i.e., how FFA are replaced by glucose as an energy source when the starved organism is refed, has not yet been addressed. In recent experiments with a perfused working heart preparation, capable of sustaining a near-physiological work output in vitro for more than 1 h, glucose (5 mM) suppressed oleate (1 mM) removal by 35% (67). These effects of glucose on FFA metabolism are relatively large and of special interest in light of measurable amounts of the carnitine acyltransferase inhibitor malonyl-CoA in heart muscle from fed animals and its decline in heart muscle from starved animals (32). The exact

The *fifth factor* which determines the selection of fuels is *hormone levels*. Plasma insulin concentration, and probably the insulin/glucagon ratio, play a major role in the fuel preferences of tissues. Insulin regulates glucose entry into muscle, adipose tissue and other cells. Likewise, the rate of adipose tissue lipolysis is under hormonal control. Randle et al. (50) have integrated these observations in the glucose-fatty acids cycle: glucose oxidation is inhibited by FFA when FFA are elevated (low insulin, fasting), and FFA release is inhibited by glucose uptake (high insulin, fed).

#### 7. Control of the Citric Acid Cycle

The function of the citric acid cycle in the heart is to oxidize the acetyl-group of acetyl-CoA to  $CO_2$  in the process of generating reducing equivalents. Reduction of the electron carriers NAD<sup>+</sup> and FAD requires addition of H<sub>2</sub>O, as can be easily seen from the net reaction of the cycle

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$$CH_3 - CO \sim SC_0A + 3 H_2O \longrightarrow 2 CO_2 + 4 [2H] + CoASH,$$

where [2H] represents a pair of reducing equivalents removed to reduce an electron carrier. It is reasonable to assume that the  $H_2O$  added to the cycle arises from reduction of molecular  $O_2$  in the respiratory chain. The production of reducing equivalents from  $H_2O$  is probably the most important function of the citric acid cycle, in addition to the other functions such as the generation of intermediates like citrate (the metabolic signal which regulates phosphofructokinase) and the disposition of the products of carbohydrate, FFA and amino acid metabolism.

There are three non-equilibrium enzymes in the citric acid cycle: citrate synthase, isocitrate dehydrogenase (both the NAD<sup>+</sup>- and the NADP<sup>+</sup>-dependent isozyme) and 2-oxoglutarate dehydrogenase. Under physiological conditions, oxidation of acetyl-CoA in the citric acid cycle is directly related to the rate of ATP production by the oxidative phosphorylation of ADP in the respiratory chain. Not surprisingly, in isolated working rat hearts perfused with glucose as the only substrate, the tissue content of acetyl-CoA is so low that it cannot be measured (64). Hence, the citric acid cycle, as the main source for reducing equivalents, must be under a strict control between supply and demand, because the rate of acetyl-CoA production and oxidation has to respond instantly to a wide range of energy needs.

Control is exercised by the tight relationship between electron transport in the respiratory chain and ATP synthesis. The major determinants of mitochondrial respiratory rate are extramitochondrial [ATP]/[ADP][P<sub>i</sub>], intramitochondrial [NAD<sup>+</sup>]/[NADH] and O<sub>2</sub> tension. If ATP synthesis cannot occur (e.g., when the [ATP]/[ADP] ratio is high), the reducing equivalents NADH and FADH<sub>2</sub> will not be oxidized by the respiratory chain. A very simple mechanism of control of the citric acid cycle is, therefore, that the lack of oxidized cofactors (NAD<sup>+</sup> and FAD) will slow down those reactions of the citric acid cycle which depend on coupled oxidation and reduction. This feedback mechanism prevents unnecessary utilization of fuels when ATP supplies are adequate.

The enzymes of the citric acid cycle are located in the mitochondrial matrix or inner mitochondrial membrane in close proximity to the enzymes of the respiratory chain. However, unlike the respiratory chain or the glycolytic pathway, the citric acid cycle is not a linear pathway. According to Baldwin and Krebs (1), the advantage of a cyclic process is that it can offer opportunities for regulation which a linear pathway cannot provide. Unfortunately, methods of conventional metabolic analysis (such as measuring accumulation of metabolites) frequently fail to elucidate control mechanisms in a cyclic process. Furthermore, understanding of the control of the citric acid cycle is also complicated by lack of knowledge about metabolite concentrations in mitochondria.

Earlier investigators (see, e.g., refs. 27,28) have pointed out that the synthesis of citrate, catalyzed by citrate synthase in the reaction

oxaloacetate + acetyl-CoA + 
$$H_2O \rightarrow$$
 citrate + CoASH,

must be a rate-limiting step, because subsequent metabolites in the citric acid cycle do not normally accumulate. This means that they are removed as rapidly as they are produced or that their rate of removal is limited by their rate of supply. In other words, the citrate synthase reaction is a non-equilibrium reaction which initiates the flux to which other participants in the pathway must respond in order to achieve a steady state. The assumption that citrate synthase is rate limiting has been supported by the discovery of several feedback inhibitors of the enzyme (see ref. 62 for review). Citrate synthase is also inhibited by several products of the pathway which citrate synthase initiates, such as NADH, ATP, or succinyl-CoA. However, for a number of reasons it is unlikely that the in vitro findings also account for the regulation of citrate synthase in the intact, functioning heart muscle. Regulation of citrate synthase by NADH may become clearer when one considers the citric acid cycle as a whole. The overall result of the citric acid cycle (or the oxidation of acetate by  $O_2$ ) is the obligatory phosphorylation of ADP by the respiratory chain.

The key reactions for the cycle cause a reduction of the coenzymes of the respiratory chain. When the relevant products (three NADH<sub>2</sub> and one reduced flavoprotein) enter the respiratory chain,  $H^+$  atoms are transferred from the coenzymes to molecular O<sub>2</sub>, coupled with the synthesis of ATP. Thus, the *primary* end products of the cycle reactions are NADH<sub>2</sub> and reduced flavoprotein, and the *final* product of the cycle and its associated reactions as a whole is ATP.

The obligatory coupling of the oxidation of reduced coenzymes with the phosphorylation of ADP (i.e., synthesis of ATP from ADP and P<sub>i</sub>) means that the [ATP] can indirectly regulate the dehydrogenase steps by controlling the [ADP] and [P<sub>i</sub>]. This type of control would apply to all mitochondrial dehydrogenases; but it would appear that it cannot regulate the initiating step of the cycle, the citrate synthase reaction, because this is not a dehydrogenase reaction. Thus, citrate synthase seems to be the only important enzyme in the cycle not directly regulated by the availability of ADP and P<sub>i</sub> for oxidative phosphorylation.

Certain regulatory mechanisms exist within the cycle when the heart switches from the oxidation of glucose to the oxidation of FFA or ketone bodies. Randle et al. (49) first drew attention to the fact that during the oxidation of acetate plus glucose the cycle transiently operates in two spans, the first span controlled by citrate synthase (and the availability of oxaloacetate), and the second span controlled by 2-oxoglutarate dehydrogenase. This enzyme is a multi-enzyme complex which bears a certain similarity to pyruvate dehydrogenase in its components and reaction sequence. It is also inhibited by its products, succinyl-CoA and NADH (17), and it requires  $Ca^{2+}$ , but, unlike PDH, it is not regulated by a phosphorylationdephosphorylation cycle. Regulation of 2-oxoglutarate dehydrogenase is thought to be of crucial importance in the accumulation of citrate, induced in rat heart by the oxidation of FFA and ketone bodies (52).

The experiments of Randle et al. (49) have shown that flux through citrate synthase is strictly substrate dependent. As more acetyl-CoA becomes available, more oxaloacetate is formed by transamination of aspartate with 2-oxoglutarate. As a result, the tissue content of aspartate falls and the tissue content of glutamate rises. More recent experiments by Hassinen's group have shown that at least part of the oxaloacetate is also derived from carboxylation of pyruvate (43,47).

Regulation of flux through 2-oxoglutarate dehydrogenase is complex and still not well understood. In addition to product inhibition, there are at least two additional factors which need to be taken into account: a) the overall activity of the enzyme, which was found to be directly correlated with flux through the citric acid cycle (12), and b) availability of free CoASH, a cofactor for the reaction (64).

#### 8. Summary and Physiological Implications

The effects of cardiac work on cardiac metabolism are mediated by a feedback system which keeps energy production in step with energy utilization on a beat-tobeat basis. The effects of cardiac work have already been considered in conjunction with individual fuels for respiration. These may be summarized as follows: Within a physiological range, an increase in the work load of the heart leads to: a) an increase in work output, b) an increase in  $O_2$  consumption, c) an increase in citric acid cycle turnover, d) an increase in flux through the Embden-Meyerhoff pathway, and/or e) an increase in beta-oxidation of fatty acids. The integration of this sequence of events is still not completely understood, but it is assumed that the decline in the phosphorylation state of adenine nucleotides, which raises the extramitochondrial [ADP] as a consequence of increased ATP utilization, is the central event.

The phasic mechanical activity of the heart necessitates a continuous supply of nutrients and  $O_2$  for energy production. Heart muscle possesses the ability to choose between a number of different energy-providing substrates circulating in the blood stream, but the mechanisms of fuel selection are not entirely clear as yet. Balance calculations indicate that the energy *value* in the form of reducing equivalents is greater for glucose than for fatty acids, because of the greater P/O ratio of the former.

The selection of fuels rests within a number of regulatory enzymes whose kinetic properties, as discussed earlier, are controlled by factors other than substrate concentrations and product removal. There is good reason to assume that a few regulatory enzymes are responsible for alterations in flux through any metabolic pathway and thus provide for the most economical selection of fuel in a given physiological situation.

To date there are few, if any, examples of in situ measurements of regulatory enzymes relative to flux through the pathway. A rare example of the integrative control of metabolism, neural activity and cardiac function has been reported by Skinner et al. (61). The authors collected multiple freeze-biopsy samples from the myocardium of the conscious pig and demonstrated an increase in activated phosphorylase with repeated stressful situations. The increase declined with time, as the animals became used to the situation. Data like these are, however, few, and it is hoped that more will be learned on the integrative control of cardiac metabolism in the future. It needs to be stressed once more that control of flux through a metabolic pathway cannot be achieved by any single enzyme acting in isolation and that regulation is a shared property of many different intra- and extracellular effectors.

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# Chapter 4

# **Carnitine and Cardiac Energy Supply**

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The purpose of this survey is to briefly consider some metabolic aspects of the human heart. The pathways of lipid utilization for energy production have received special attention. The function of L-carnitine in mitochondrial fatty acid oxidation and its role in myocardial energy production is analysed. Mention is also made of the effects of L-carnitine on myocardial metabolism of patients with coronary artery disease, either at rest or during pacing-induced myocardial ischaemia.

## Introduction

The energy needed for cardiac contraction comes from the breakdown of chemical substances, the substrates. In the heart this process occurs mainly aerobically, i.e., substrates are combined with oxygen to form water, carbon dioxide and energy. The amount of oxygen consumed is often used as a measure of the total energy available. The ultimate fate of the liberated energy is to take the form of heat. However, before this happens completely, part of the energy is used for contraction, a process during which the heart may perform external work, as is usually the case in the intact body.

## Substrate Metabolism of Human Heart

Knowledge of the fuels of the human heart started with the introduction of coronary sinus catherization by Bing and his associates in 1947. The chemical composition of arterial blood entering the heart was compared with that of coronary sinus blood leaving the heart. From such studies it was established that glucose, lactate and fatty acids are the heart's major sources of energy (3,4). It has also been recognized that the myocardium is able to utilize such fuels as pyruvate, acetate, ketone bodies and amino acids, but the normal circulating levels are too low for them to be considered as major sources of energy, even when the external supply is raised (8). The hearts can also use, under certain circumstances, its internal energy stores such as glycogen and lipid (32).

A cruder estimate of the type of substrate used by the heart can be achieved by the respiratory quotient, which is calculated by comparing the rate of oxygen uptake with the rate of production of carbon dioxide. A respiratory quotient near to one implies oxidation of glucose and/or lactate, whereas a lower value implies fatty acid oxidation. Because the myocardial respiratory quotient was frequently low, early workers were alerted to the importance of the role of lipids as the major myocardial fuels (24). However, there is still some controversy about which is the preferred myocardial substrate.

## Substrate Preference

It is often stated that lipids are the preferred myocardial substrates (3,23,33). However, most studies of myocardial utilization of unesterified fatty acids (NEFA) in humans have been undertaken in resting fasting subjects. In this state the circulating NEFA levels are high (over 1000 uEq/L). Thus, they may provide more than 60% of the myocardial energy requirements (2,4,27). But if the energy requirements are satisfied by the supply of carbohydrates, NEFA extraction will be decreased (1). For example, if there is a rise in the arterial lactate, as it occurs during exercise, there is also an increase in the myocardial utilization of lactate (2,4,21,28). Thus, as it is also shown in Fig. 1, myocardial substrate preference is critically dependent on the availability of substrates in the circulation at the moment of the study (12). It is therefore crucial that substrate utilization is studied when all the substrates are available in adequate concentration. Such studies show that in the heart carbohydrates and lipids are concerned predominantly with the provision of ATP for contraction. Oxidation of carbohydrate substrates accounts for up to 35% of oxygen consumption. Extraction of NEFA may account for about 70% of the oxygen consumption in the post-absorptive state. The human heart, like that of the rat, exhibits a predilection for fatty acids as respiratory fuel (3).

As the heart derives most of its energy from oxidative metabolism, enthalpy changes can be taken to be equivalent to the oxygen consumption. The amount of heat liberated per gram of fat oxidized about (9 kcal/g) is more that twice that per gram of carbohydrate about (4 kcal/g). Nevertheless, the enthalpies are quite similar, when expressed per liter of oxygen consumed (fat 4.69 kcal/g  $O_2$ ; carbohydrate 5.05 kcal/L  $O_2$ ). This is because more oxygen is required to oxidize a gram of fat than a gram of carbohydrate.

## Pathways of Substrate Utilization and Energy Production

The overall process of energy production involves the complete oxidation of substrates to carbon dioxide and water (Fig. 2). Energy (as ATP) can be generated in the myocardium by two metabolic pathways, glycolysis (anaerobic) and oxidative phosphorylation (aerobic). Under normal conditions the majority (90%) of ATP is produced by the latter pathway (20).

Before a carbohydrate (such as glucose) can enter the tricarboxylic acid cycle, it must be broken down via the glycolytic pathway. Therefore oxidation of glucose is dependent on the enzymes of anaerobic glycolysis. Lactate, the other major carbohydrate fuel, is dependent on the activity of the lactate dehydrogenase complex.

Although the heart is able to synthesize structural lipids for incorporation into the membranes, the most likely fate of a fatty acid molecule taken up by the heart is oxidation. The first step in fatty acid oxidation is their activation to form acyl-coenzyme A derivatives. The enzymes first concerned with the use of fatty acids



Fig. 1. Relationship between arterial substrate concentration and myocardial arteriovenous difference or extraction of free fatty acids (FFA) and lactate. a = arterial; cs = coronary sinus. Data were obtained in catheterized patients (12).

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are therefore the acyl-coenzyme A synthetases which exist in the heart with different fatty acid specificities and locations (16). Short-chain fatty acids ( $C_8$  or less) are oxidized by heart mitrochondria in the absence of external coenzyme A or of L-carnitine, the role of which is discussed later in this chapter. This is because the relevant acyl-coenzyme A synthetases are in the same compartment (mitochondrial matrix) as the enzymes of beta-oxidation and the citrate cycle. However, for the oxidation of long-chain acids, this activation must be followed by the transfer of long-chain acyl-coenzyme A into the interior of the mitochondria.

It is rather curious that the long-chain fatty acids that are physiologically really important, are converted to their coenzyme A derivatives on the wrong side of the mitochondrial inner membrane, which is impermeable to coenzyme A. This is particularly strange because the activation of fatty acid is accomplished by two distinct enzymes that share the same cytosolic coenzyme A pool, which is very small. One of these enzymes, representing 80% of the total activity, is attached to the mitochondrial outer membrane, and the other to the endoplasmic reticulum (7).

## Function of L-Carnitine in Mitochondrial Fatty Acid Oxidation

Carnitine (L-3-hydroxy-4-trimethylaminobutyric acid) was discovered in 1905 as an abundant constituent of muscle (17). Its metabolic function remained completely obscure for over 50 years. Only in 1959 Fritz (15) showed that carnitine increased long-chain fatty acid oxidation in liver and heart.



Fig. 2. Schematic representation of anaerobic metabolism in the myocardium.

Carnitine is synthetized almost exclusively in the liver, from lysine and methionine (5). Bohmer et al. (6) have found that heart cells in culture possess a specific carnitine uptake mechanism. The plasma concentration of carnitine is about 30 uM; that in the heart is about 100-fold higher. In more recent years, it has been suggested (18) that: a) L-carnitine, the naturally occurring form, is able to stimulate fatty acid oxidation by tissue homogenates; b) fatty acyl esters of carnitine are useful substrates for mitochondria; c) these compounds are formed by enzyme-catalyzed reversible acyl transfer from coenzyme A.

Later it has been recognized that two distinct classes of carnitine acyltransferases exist, normally referred to as carnitine acetyl- and palmitoyltransferases. Little is known about the heart carnitine palmitoyltransferases; however, inhibition experiments with purified liver enzymes have suggested that in the mitochondria they exist in two forms with considerably different properties. One of these transferases, the outer or A enzyme, is accessible to extramitochondrial coenzyme A, while the inner, or B enzyme, is within the inner membrane, and it uses the coenzyme A of the matrix (Fig. 3). The overall steps of free fatty acid oxidation can be summarized as follows (Fig. 3): a) extramitochondrial formation of acylcoenzyme A; b) transfer of the acyl-group to carnitine by the outer transferase; c) delivery of the carnitine ester to the inner transferase; d) transfer of the acylgroup to the matrix coenzyme A; e) beta-oxidation; f) finally, a carnitine acylcarnitine translocase system transfers intramitochondrial carnitine to outside the mitochondria. Experiments with heart mitochondria have demonstrated that penetration of the inner mitochondrial membrane by acyl-carnitine occurs by the acyl-carnitine  $\iff$  carnitine exchangers or translocases. They are analogous to the ADP  $\iff$  ATP exchange.



Fig. 3. Schematic representation of the L-carnitine cycle.

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The process exhibits 1:1 stoichiometry, so that the total carnitine content of mitochondria remains unchanged. This content is about 3 nmol/mg protein in isolated mitochondria, corresponding to a matrix concentration of 2-4 mM, similar to the concentration in the cytostol (25).

#### **Role of L-Carnitine in Myocardial Energy Production**

From the previous discussion, the role of L-carnitine in energy production from fatty acid oxidation is obvious: L-carnitine takes part in the shuttle mechanism whereby long-chain fatty acids are transformed to acyl-carnitine derivatives and transported across the inner mitochondrial membrane. This is impermeable to long-chain fatty acids and to their coenzyme A esters. Once across the membrane, the acyl-carnitine is reconverted to carnitine and to acyl-coenzyme A, which undergoes beta-oxidation. This role of L-carnitine is essential in heart free fatty acid metabolism. Interestingly, the myopathic-type of carnitine deficiency is associated with cardiomyopathy as well as specific carnitine deficiency in the heart (10). In such cases FFA have always been shown to accumulate in the myocardium, with impaired long-chain fatty acid oxidation by muscle homogenates. These alterations can be corrected by exogenous administration of L-carnitine.

From Fig. 4 it also appears that L-carnitine might have a role not only in energy production, but also, indirectly, in energy delivery from the mitochondria to the cytosol. In fact long-chain acyl-coenzyme A is known to inhibit mitochondrial adenine-nucleotide translocase (26,29-31), the enzyme responsible for the one to one exchange of ATP and ADP (19). There is evidence that one of the most important effects of L-carnitine administration is the reduction of long-chain acyl-coenzyme A in heart tissue, resulting in a reduction of the inhibition of the adenine-nucleotide translocase. In this way, L-carnitine allows ATP to be formed from long-chain acyl-coenzyme A and transported from the mitochondria to the cytosol.

It is often assumed that L-carnitine has also a role to permit the import of acetylcoenzyme A by mitochondria as with the long-chain derivatives. However, doubts are threefold: a) acetyl-coenzyme A is generated within the mitochondrial matrix; b) isolated mitochondria scarsely use external acetyl-coenzyme A, even in the presence of carnitine; c) isolated mitochondria apparently contain only one form of carnitine acetyltransferase (9).



Fig. 4. Schematic representation of the effect of L-carnitine on adenine-nucleotide translocase (ANT).

#### Effects of L-Carnitine on Myocardial Metabolism of Patients at Rest and during Pacing-Induced Ischaemia

We have investigated the influence of L-carnitine, the necessary cofactor for mitochondrial transport of long-chain fatty acids (11-14), on myocardial carbohydrate and lipid metabolism in 16 patients with coronary artery disease. The patients were also subjected to two periods of rapid coronary sinus pacing (Fig. 5). They received a central venous infusion of L-carnitine (40 mg/kg) in 50 ml at 200 mg/min prior to the second pacing period. The two pacing periods were separated by a 45-min interval (12). The haemodynamic data at rest, reported in Table 1, show that acute administration of L-carnitine did not significantly change the parameters



Fig. 5. Effect of L-carnitine on myocardial uptake of FFA, glucose and lactate of 16 patients with coronary artery disease, either at rest or during pacing. Data represent the mean  $\pm$  S.E. L-carnitine significantly (P<0.001, analysis of variance) modified the behaviour of myocardial uptake of glucose and lactate during the second pacing period, when lactate and glucose production was converted into extraction.

measured, including total coronary flow. In addition L-carnitine failed to modify the changes in haemodynamic parameters induced by atrial pacing, except for the left ventricular end-diastolic pressure which was reduced from  $22.1 \pm 8.2$  to  $16.7 \pm 48$  mmHg (P<0.05, ref. 13).

Figure 5 shows that the same concentration of L-carnitine induced important changes in myocardial substrate uptake, either at rest or after pacing. At rest, L-carnitine induced a small increase in myocardial uptake of free fatty acids (FFA), concomitant with a decrease of glucose uptake and no major changes in lactate uptake. During pacing, lacatate production was converted to extraction. An increase in the uptake of FFA and of glucose, though small, was demonstrated. All these alterations are due to the metabolic properties of L-carnitine, as coronary flow during atrial pacing was unchanged.

There is no simple explanation for these effects of L-carnitine. During ischaemia, the reduced availability of oxygen leads to a decrease in mitochondrial electron transport, which in turn causes an accumulation of NADH and long-chain acyl-coenzyme A. An increased NADH/NAD ratio and acyl-coenzyme A/coenzyme A ratio may inhibit the activity of pyruvate dehydrogenase (PDH), the enzyme which regulates the entry of pyruvate into the citric acid cycle. Under these conditions pyruvate is preferentially converted to lactate with consequent lactate production. L-carnitine can reduce lactate production under ischaemic conditions either by activating PDH directly or by provoking a decrease in the acetyl-coenzyme A/coenzyme A ratio, as a consequence of acetyl removal from coenzyme A, mediated by carnitine coenzyme A acetyltransferase. Therefore, the utilization of pyruvate and consequently of lactate in the oxidative pathway, induced by L-carnitine, may be attributed to enhanced PDH activity, rather than to an increase in oxygen availability induced by L-carnitine.

Another possible explanation for the effects of L-carnitine on lactate metabolism is an indirect effect on phosphofructokinase activity (PFK), the enzyme regulating the rate of anaerobic glycolysis. The activity of this enzyme is inhibited by a high cytosolic ATP concentration, whilst it is stimulated by low cytosolic ATP concentration. Under ischaemic conditions the ATP concentration in the cytosol decreases either as a result of reduced oxidative metabolism or because the activity of adenine-nucleotide translocase is inhibited by long-chain acyl-coenzyme A. This causes sequestration of ATP into the mitochondrial matrix with a decrease of cytosolic ATP and a consequent stimulation of PFK. By removing long-chain acylcoenzyme A, carnitine can also prevent its inhibitory action on adenine-nucleotide translocase, thus improving the ATP transfer to the cytosolic compartment. The increased cytosolic ATP levels might decrease PFK activity and lactate production.

Variable	Before	After L-carnitine	
Heart rate (beats/min)	79 ± 4	76 ± 3	
Mean aortic systolic pressure (mmHg)	$146 \pm 4$	$144 \pm 6$	
Mean aortic diastolic pressure (mmHg)	76 ± 6	73 ± 2	
Pulmonary artery pressure (mmHg)	$18 \pm 1$	$18.2 \pm 3$	
Cardiac output (litre/min)	$5.9 \pm 0.7$	$5.9 \pm 0.6$	
Coronary sinus blood flow (ml/min)	127 ± 14	129 ± 12	

 Table 1. Effect of L-carnitine on haemodynamic variables in cardiac patients at rest

See legend to Fig. 5. No statistically significant differences were observed.

It is possible that the increased PDH and the decreased PFK activity are concurrently involved in the reduction of lactate formation induced by L-carnitine administration. These data show the complexity of lactate metabolism that can be positively affected by interventions like with L-carnitine, a compound without haemodynamic effects, acting primarily on lipid metabolism.

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# Chapter 5

# Adenine Nucleotides, Purine Metabolism and Myocardial Function

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During cardiac contraction there is a very rapid breakdown and synthesis of the high-energy phosphates ATP and creatine phosphate. Whenever a disbalance exists between synthesis and breakdown of ATP a series of enzymes will act to catabolize AMP to adenosine, inosine, (hypo)xanthine and urate which will leave the heart. This will eventually result in decreased adenine nucleotide and ATP levels. A number of observations are discussed in order to relate the function of the heart and the metabolism of ATP and its catabolites. Good evidence is still lacking to conclude that ATP-content and heart function are either directly related or non-related. Major problems appear to be adequate definition of heart function and the fact that the ATP-catabolite adenosine exerts several modulatory effects on heart function. For these and other reasons further studies on purine (nucleotide) metabolism and heart function are needed.

## Introduction

The continuous pumping of blood by the heart requires a large amount of (bio)chemical energy in the form of adenosine triphosphate (ATP) in order to drive the relaxation of the myofibrillar contractile proteins, to maintain cationic homeostasis and to activate intermediary metabolism and synthesis of high molecular structural and functional cell components. The normal ATP-content of cardiac tissue falls around 20-30 umol per g dry wt. for most mammalian species studied. In addition the heart will contain between 3 and 5 umol ADP per g dry wt., most of which is probably protein-bound, and an even smaller amount of AMP (0.3-1.0 umol/g dry wt.). In well-oxygenated hearts ATP therefore makes up 80 to 90% of total adenine nucleotides, if we do not include the nicotinamide adenine dinucleotides. This apparently constant amount of cardiac ATP is in fact the result of a continuous, fast breakdown and synthesis of ATP. The latter has been estimated between 150 and 600 umoles of ATP synthetized per g dry wt. per minute in isolated rat hearts (28). This means that between 5 and 10% of total ATP can be used (and resynthetized) during a single contraction. Only oxidative (oxygen dependent) phosphorylation in the cardiac mitochondria is able to provide such a high ATP-synthesis rate. Apart from ATP, the heart contains relatively large amounts (25-40 umol/g dry wt.) of another high-energy phosphate, i.e., creatine phosphate, which is also continually turned over to creatine, with formation of ATP from ADP, the first-formed ATP-catabolite. When the formation of ATP is inhibited, either by lack of oxygen, hypoxia, poisoning of oxidative phosphorylation, or ischemia, the ATP-content in the heart will decline, ADP and AMP contents will rise, and AMP will be broken down. A long chain of enzymatically mediated events will lead to the formation of adenosine, inosine, hypoxanthine and, occasionally, xanthine and urate (1-3,17). These purine nucleosides (adenosine, inosine) and oxypurines can pass the cell membrane relatively easy and will be washed out off the heart. Fig. 1 shows the time-course of the ATP-breakdown cascade in autolyzing rat heart as determined by Gerlach and co-workers (13,15) more than 20 years ago.

If reoxygenation occurs after a period of ischemia and adequate oxidative metabolism is restored, the heart will be left with a decreased amount of adenine nucleotides. We suggest that adequate reperfusion of non-irreversibly damaged hearts will result in ADP- and AMP-contents which are no longer increased. If still increased, however, AMP-catabolism and purine release will rapidly lead to further adenine nucleotide loss and ATP-decrease, which, beyond a certain level, is incompatible with adequate cellular function and will be associated with cell death (26). During hypoxia, increased work-load or hypoperfusion, the inadequate oxygen supply to the heart can be increased by coronary vasodilation, which is at least partially mediated by the ATP-catabolite adenosine (7). The supply-demand ratio for oxygen, effected by the balance between formation and breakdown of ATP, can thus be considered as the trigger for adenosine-mediated coronary vasodilation (2,38). From the above considerations alone, it is obvious that there exists an intricate and complex relationship between myocardial ATP-content and turnover, purine metabolism and heart function, which will be further discussed in the following paragraphs.

## **ATP-Content and Myocardial Function**

During adequate supply of oxygen and substrates, the heart is capable of increasing its work-performance and ATP-turnover rate several fold. This will not be accompanied by major changes in myocardial ATP-content. In addition, when cardiac



Fig. 1. Time-course of adenine nucleotide degradation (A) and accumulation of purine nucleosides and oxypurines (B) in autolyzing rat heart. The data were redrawn from Deuticke and Gerlach (13). Abbreviations: INO = inosine, ADO = adenosine, HYP = hypoxanthine, XAN = xanthine. Note the very fast ATP-decline over the first 20 min and the simultaneous rapid rise in AMP, adenosine and inosine.

work-performance is fully arrested in the presence of oxygen and substrate (e.g., by giving high doses of calcium antagonists or providing zero calcium) again normal ATP-values will be found. Superficially, this would appear to contradict any relation at all between ATP-content and heart function. Still, it has been found that myocardial ATP-content during hypoxia of varying degrees of severity correlates with myocardial function (see Fig. 2). Other studies, however, failed to observe significant decline in ATP-contents during hypoxia, while cardiac function was already severely depressed (8). The latter NMR-study did show, however, a correlation between ATP-turnover rate via the creatine kinase reaction and cardiac function, expressed as the rate-pressure product. It should be noted that NMR measurements of ATP-content are not very sensitive.

Decreased myocardial ATP-contents after short and reversible ischemia in vivo are restored very slowly (10,23,35,40). A similar slow restoration is found for the decreased post-ischemic function (10,27). Myocardial ATP-content after reperfusion is directly and sometimes linearly (20) related to the ATP-content at the end of ischemia. During hypoxia, ischemia or during inadequate reperfusion, metabolic alterations, other than decreased ATP-content, are thought to contribute to depression of function or to the alteration of the ATP-function relationship. Acidosis (lactate accumulation) or intracellular build-up of calcium or phosphate have been suggested as causative factors, together with fatty acids or fatty acid derivatives. In general it can be expected that factors which inhibit ATP production rate, will lead to larger disturbances of ATP-content than factors which inhibit ATP-consumption, i.e., contractile function.

Indirect evidence for the importance of cardiac ATP-content in maintaining cardiac function comes from studies on the protective effects of calcium-antagonists against ischemic damage. Calcium antagonists (diltiazem, nifedipine, nisoldipine, bepridil) have a dose-dependent negative inotropic action on the heart. When given before the onset of ischemia, there is a dose-dependent inhibition on ischemic ATP-breakdown as reflected by the decreased release of ATP-catabolites (purines) during ischemia (11,16,19). Work performance before the onset of low-flow ischemia is directly related to purine release during ischemia (12). Under these conditions (low-flow ischemia, glucose as substrate), it is most probably the formation of ATP which is restricted. Therefore reduction of ATP-breakdown is the palliative measure of choice.



Fig. 2. Relation between ATP-content and contractility in hypoxic rat heart. ATP-contents were determined in isolated rat hearts, subjected to varying degrees of hypoxia ( $\bullet$ ) or ischemia (o). Myocardial function is given as percentage of prehypoxic function. Apex displacement is used as a measure of contractility. Data were redrawn from Stam and De Jong (39).

In a number of studies, good correlations have been observed between ATP-content and myocardial function after ischemia and reperfusion (22,24,25,30,34,41). The actual curves, however, which describe the ATP-function relationship (Fig. 3) are different in these studies. After pre-ischemic conditioning of hearts by 10 min anoxia, Neely and Grotyohann (31) found a very steep relationship, i.e., fast decline of function over a short range of ATP-contents. They concluded that there is no relation between ATP-content and heart function. Whether AMP and ADP contents returned to normal levels in these anoxia-pretreated, reperfused hearts, was not reported (31). Non pretreated, reperfused hearts in the same study still had enormously elevated AMP-levels. Because evidence has been gathered (3,9) that adenine nucleotide breakdown to purines is regulated by cytosolic AMP-content, hearts with strongly elevated AMP-contents are bound to loose adenine nucleotides (and thus ATP) at a fast rate. Hearts of cardiomyopathic hamsters contain severely depressed amounts of ATP at the moment when signs of heart-failure in vivo are apparent. The protective effect of high oral doses of calcium antagonists on these hearts coincides with the finding of normalized ATP-values (42).

The discrepancies observed in various reports on the correlation between ATP and function (Fig. 3) could be related to the manner in which cardiac function is determined or expressed. For instance, developed tension or pressure of the heart is directly related to the applied or existing preload. It is repeatedly found that postischemic, reperfused hearts exhibit increased preload (beginning contracture?). Function determined at this increased preload should be expressed in relation to the control function at similarly high preload, and not, as is often done, in relation to function (developed pressure) at a lower preload. It has on occasion been reported that developed pressure of isolated hearts during early post-ischemic reperfusion was actually higher than the pre-ischemic control value (29). In addition, changes in heart rate may also cause changes in the inotropic status of the heart. In general, it proves extremely difficult to obtain relevant measures of cardiac function (14). We feel that most controversies on the relationship between heart function and metabolic parameters can be reduced to problems related to adequate determination of function or inotropic status of the hearts investigated. Another factor which tends to obscure good interpretation of data is the fact that differences may exist in both ATP-content and function in the outer and inner layers of the ischemic and reperfused heart (33). NMR-measurements and biochemical determinations of ATP in whole heart homogenates will be inadequate if such differences exist. We conclude that it is not yet fully clear how the dynamic relationship between ATP-content or ATP-turnover on the one hand and contractile activity or contractile potential on the other is regulated. A major contribution to the regulatory dynamics appears to be effected through operation of the creatine kinase reaction. In this respect it should be considered that gradients of ATP and ADP may exist between mitochondria and the site of the myofibrillar, contractile ATP-ase, with the creatine kinase equilibrium functioning to lessen the steepness of ATP- and ADP-gradients. The fact that the flux through creatine kinase is at least severalfold higher than the ATP-formation rate could point into that direction. Evidence for the existence of ATP-gradients has been found in hypoxic liver cells (5), which have far lower rates of ATP-turnover than the heart.

It is often argued that the cellular ATP-concentration cannot directly influence contractility, because the  $K_m$  of the contractile ATP-ases will always be far lower than estimated ATP-concentrations. It should be noted, however, that ordinary Michaelis-Menten kinetics do not apply when enzyme concentrations are extremely high, such as at the myofibrillar ATP-ase sites. Again the possible existence of ATP-gradients could strongly influence the actual ATP-concentration at the ATP-ase site.

Myocardial ATP-contents in physiologically normal hearts tend to be constant and depleted ATP-contents will be increased, albeit slowly, during reperfusion (10,23,35,40). It appears therefore that normal ATP-content is a prerequisite for normal cardiac metabolism and function. ATP is a known modulator and substrate for a host of cardiac enzymes (e.g., kinases, ATP-ases), which execute a number of essential reactions. The precise sites of action, which will be critically influenced by a decreased cytosolic or mitochondrial ATP-concentration, are therefore very difficult to identify. It would be interesting in this respect to investigate the metabolic properties of non-irreversibly damaged, but partially ATP-depleted hearts at various work loads.

One important implication of a direct relationship between ATP-content, heart function and metabolism may be that an already ATP-depleted heart is bound to completely loose its function or enter a stage of irreversible damage, if ATP-levels have not been restored in the inter-ischemic period. Studies have been performed (18,36), in which short periods of repeated ischemia/reperfusion were compared with a long period of ischemia of similar total duration. It appeared, however, that short-during ischemia and reperfusion might have a protective effect on ATP-decline during a next ischemic period (18,36). It is not clear, whether the observed decreased heart function, the observed "overshoot" of creatine phosphate or the combination of both provides the temporarily protection against ATP-decline. It is tempting to conclude that creatine phosphate overshoot reflects an intact capability



Fig. 3. Myocardial ATP-content and function in reperfused isolated rat heart preparations. Data were taken from publications by different authors. (----) ref. 22; (---) ref. 31; (----) ref. 30; (xxxx) ref. 41.

of the heart to synthetize but not to use ATP (21). Finally, there is no solid evidence that severely ATP-depleted hearts will continue to perform in the normal to maximal range of work, which can be observed in non ATP-depleted hearts. In the next paragraph we shall further elaborate on the effects of ATP-catabolites, with the emphasis on adenosine, on regulation of cardiac function.

#### **Purine Metabolism and Function of the Heart**

Adenosine release from the heart is increased, whenever a disbalance exists between formation and breakdown of ATP. This can be either caused by increased work-load, decreased oxygen supply, mitochondrial poisoning, catecholamine stress or substrateinduced (acetate) breakdown of ATP to the direct adenosine precursor AMP (1,3). The physiological effects of adenosine may be looked upon as controlling the balance between supply and demand of oxygen or energy (32,38). Adenosine is a coronary vasodilator, also in man, and many coronary vasodilatory drugs (dipyridamole, dilazep, hexobendine, papaverine) presumably act by enhancing adenosine effects through interference with the normally fast transport and catabolism of adenosine.

In addition, adenosine reduces heart rate and slows conduction through the atrioventricular node (6) and anti-catecholaminergic effects of adenosine have been noticed (37). Several beneficial effects on the ischemic heart have also been reported for the adenosine catabolite inosine (4). Both vasodilation and positive inotropic effects by inosine have been noticed in animal and human studies (see Chapter 26). Inosine effects may well be caused by modification or attenuation of adenosine effects or of the availability of adenosine at the site of its receptors. Both adenosine (via adenosine kinase) and inosine can be reincorporated in the myocardial adenine nucleotide pool. Inosine is first catabolized to hypoxanthine before incorporation. A number of studies have indicated the feasibility of increased adenine nucleotide formation in the post-ischemic heart by giving purine compounds (adenosine, inosine, hypoxanthine, adenine, 5-amino-4-imidazolecarboxamide riboside; see Chapters 1,10,26). No consensus of opinion exists as yet, however, whether this will actually result in increased recovery of cardiac function (18). Enhanced ATPrestoration by 5-amino-4-imidazolecarboxamide riboside coincides with decreased cardiac function for which no adequate explanation has as yet been provided. As outlined in a previous paragraph, this might well be caused by the difficulties in determining the relevant indices of cardiac function. The study of the intricate relationship between myocardial purine metabolism, function and general metabolism of the heart deserves much further attention. Attempts to speed up restoration of post-ischemic ATP-content might be worthwhile, even if ATP-concentration is not directly related to cardiac function. Restoration of ATP-regulated metabolism of the post-ischemic heart remains a goal worthy of pursuit.

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# Chapter 6

# **On the Role of Myocardial AMP-Deaminase**

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In this chapter the AMP-deaminase activity in hearts of different species is compared with that in skeletal muscle. Some features of the well-characterized skeletal muscle enzyme, like binding to myosin filaments and deficiencies in humans, are described. Experimental models to study the role of cardiac AMP-deaminase, i.e., isolated enzyme, cardiomyocytes and isolated perfused heart, are discussed. Substrate saturation kinetics and regulatory effects of nucleotides, phosphate, and activated fatty acids are summarized; their significance for in vivo regulation considered. In addition the behavior of AMP-deaminase under various adenylate energy charges and its implications for the regulatory of enzyme in ischemic tissue is examined. Myocardial AMP-deaminase is shown to be a tissue-specific isoenzyme, with different kinetic and regulatory forms. Developmental changes of AMP-deaminase and other enzymes of adenylate metabolism enzymes in myocardial cells are reviewed. The flux through the pathway catalyzed by AMP-deaminase in normoxic and ischemic heart is analyzed. A probable function of the reaction - preservation of purine nucleotides inside the myocardial cell - is proposed.

# Introduction

Adenosine 5'-monophosphate is the first Lonenergetic catabolite of ATP-breakdown in contracting muscle. It is also one of the precursor metabolites for the restoration of high-energy phosphates. The large, time-dependent changes of AMP levels in ischemic/reperfused heart (58) suggest an efficiently regulated system for its transformation. The adenylate kinase equilibrium is responsible for AMP-conversion to the di- and triphosphate level. AMP can be either dephosphorylated to adenosine or deaminated to IMP. The former reaction is involved in the main stream of adenine nucleotide catabolism in the heart. It produces physiologically active adenosine (see Chapters 8 and 9). The parallel hydrolysis of the 6-amino group attached to adenine ring in AMP gives IMP. Independently of further degradation, IMP is also the common point between de novo purine nucleotide synthesis and hypoxanthine salvage. Both reactions are part of ATP and GTP biosynthesis, thus being of undoubted importance for energy-metabolite replenishment. The reamination of IMP to AMP, an energy-dependent two-step process, is involved in AMP deamination in the purine nucleotide cycle (37).

AMP-deaminase (AMP-aminohydrolase, AMP-D; EC 3.5.4.6) catalyzes the hydrolysis of AMP to IMP and ammonia. It is found in the majority of cells and organs of all species studied so far (79). The reaction is irreversible. The enzyme isolated from

vertebrates is highly specific towards 5'-AMP, with only slight reactivity with 2'deoxy-5'-AMP (61). AMP-deaminase is exclusively localized in the cytosolic compartment. Only one report indicates also its location in the mitochondria (22).

The physiological role of AMP-deaminase is still unclear, despite its general occurrence. This holds even for skeletal muscle, where its activity is one to two orders of magnitude higher than in other tissues, including heart muscle (52). Nevertheless, the activity changes during myoblast differentiation (40) as well as the metabolite flux through the AMP-deamination pathway (see, e.g., refs. 1,2,39) indicate the significance of AMP-deaminase in heart purine nucleotide catabolism. This contribution may be lower than in skeletal muscle. The aim of this chapter is to discuss these potencies and to suggest a physiological role of AMP-deaminase in heart tissue.

## Skeletal and Heart Muscle AMP-Deaminase

#### **Tissue** Activities

Purzycka (52) proved AMP-deaminase to be present in rat-heart muscle. This author revised the hypothesis of Conway and Cooke (13): compulsory dephosphorylation prior to deamination of AMP in heart ventricle, auricle being an exception in that sequence of adenylic acid breakdown. Nakatsu and Drummond (41) compared the activity of adenylate deaminase in heart of several species. They ranked them from nondetectable in dog heart, through rabbit and pigeon heart with 0.75 U/g wet wt., rat heart with 1.8 U/g, to the exceptionally high activity of 7 U/g in turtle heart. The activities of heart AMP-deaminase reported by other authors varied from 0.15 U/g wet wt. (rabbit, ref. 74), through about 0.75 U/g (man, ref. 47), to about 2.4 U/g (rat, ref. 76), depending also on the assay conditions.

Comparative studies revealed a wide spectrum of skeletal muscle AMP-D to heart AMP-D activity ratios in tissue extracts. These extend from 26 (pig, ref. 23), through 55 (man, ref. 47), to about 260 (rabbit heart and white skeletal muscle, ref. 45). If one disregards the assay conditions, the activity of heart AMP-deaminase is about two orders of magnitude lower than that of skeletal muscle. However, the enzyme activity in many tissues could be either overestimated due to superimposed activity of both 5'-nucleotidase and adenosine deaminase or underestimated because of nonoptimal conditions. The former can be avoided by the use of inhibitors. The enzyme activity in heart extracts is not inhibited by endogenous low-molecular compounds, but it is in homogenized skeletal muscle (23).

AMP-deaminase activity is much more pronounced in white type muscles in rabbit; it exceeds 6-10 times the activity in red muscles (55). A histochemical technique applied to hamster skeletal muscle demonstrated the strongest staining for the enzyme in the gastrocnemius fast, glycolytic fibers, and only weak staining in the slow, oxidative fibers (21). Thus well-oxygenated muscles (red skeletal muscle and heart, with oxidative energy production prevailing over glycolysis) clearly contain a lower capacity to deaminate AMP directly than the glycolysis-dependent muscles do.

#### Binding to Myosin Filament

Essential in the physiological role of AMP-deaminase catalysis could be enzyme regulation during muscle contraction. The formation of complexes between proteins

of the contractile apparatus and the enzymes involved in muscle energy formation is well-known [e.g., creatine kinase (77), adenylate kinase (57)]. The interaction which exists between AMP-deaminase and myofibrils is sufficiently strong ( $K_d = 10^{-10}$  M) to get more than 99% of the binding sites of the enzyme occupied in vivo (14). The binding is remarkably increased in skeletal muscle contracting by electrical stimulation of the sciatic nerve (66). This treatment does not change the total enzyme activity. However, it increases the muscle ammonia and IMP content. Phosphate (<10 mM) and GTP, which are inhibitors of AMP-deaminase, abolish the binding completely (44).

The purified enzyme binds to rat-muscle myosin in a molar ratio 1:3, mainly with the light meromyosin portion of the myosine molecule (67). However, the exclusive binding with the heavy meromyosin part or with subfragment 2 has also been reported for the rabbit enzyme (3). Myosin-bound AMP-deaminase is more sensitive to regulation by nucleotides than is the free enzyme (68). On the other hand, adenylosuccinate synthetase from rat muscle, the second enzyme of the purine nucleotide cycle, interacts with F-actin. This is not observed with AMP-deaminase (48). During contraction, when myosin (to which AMP-deaminase binds) slides along actin (to which adenylosuccinate synthetase binds), the cyclic operation of the purine nucleotide cycle could be triggered (67). Contradictory results have been published about the isozymic specificity of deaminase binding with myosin. Slight interference exists between rat muscle myosin and cardiac AMP-deaminase on the one hand (67). Efficient binding of heart enzyme to myosin is reported on the other (44). Both heart and skeletal muscle AMP-D bind by about 70% in that case (44).

The physiological role of the interaction adenylate deaminase-thick filament is unclear. The extent of interaction could be regulated by pH and phosphate (5). Transitions could occur in the range measured for exercising skeletal or ischemic heart muscle. The association could vary with the exercise-recovery cycle or with ischemic-postischemic changes in the heart. Thus the process may have significance in the metabolism of contractile tissues.

## **Enzyme** Deficiencies

Myoadenylate deaminase deficiency is the most common of the known enzyme defects in muscle (for review, see ref. 17). Two forms occur, one being a primary type inherited as a complete gene block. The other is a secondary form consequent to muscle damage from other diseases. In the last form, the residual activity is 1-10% of normal; the other muscle enzymes are also depleted. Neuromuscular dysfunctions are more often associated with the secondary form. The complete lack of skeletal muscle AMP-D isozyme is accompanied by normal AMP-D levels in other tissues.

Among many cases, two related individuals have been described with dilated cardiomyopathy combined with skeletal myopathy, characterized by type I fiber atrophy (62). The metabolic consequences of AMP-D deficiency are mostly treated as connected with the disruption of the purine nucleotide cycle and unability of the muscle to restore high-energy phosphates deposit from adenylate catabolites after exercise (59) or turning off the one from the anaplerotic processes involved (63).

#### Models to Study Cardiac AMP-Deaminase

Different experimental models addressed the question whether the low activity of cardiac AMP-deaminase correlates with its minor role in heart adenylate catabolism. It is of particular interest to compare the properties of isolated AMP-deaminase with those of AMP-D in situ. Many authors isolated and characterized cardiac AMP-deaminase from various species. All purification procedures include a very efficacious phosphocellulose chromatographic step, first used by Smiley et al. (72) for rabbit skeletal muscle enzyme. The specific activities of the obtained highly purified AMP-D preparations were: 30 [pig (23)], 37 [duck (50)], 40-50 [rat (42)] and 30-50 U/mg protein [cow (Skladanowski, unpublished data)]. Only one was apparently homogeneous according to the criterion of SDS-electrophoresis (23). The molecular weights were estimated only for beef- and duck-heart AMP-deaminase. The former appeared to be a tetrameric structure with mol. wt. 161,000 D (71). The duck-heart enzyme contained two homotetrameric protein molecules of different isozymes, mol. wt. 190,000 and 240,000 D (50).

Isolated cardiomyocytes allow us to estimate AMP-deaminase's share in the whole adenylate metabolism. Preparations of isolated adult ventricular cells are now available in which the morphological and metabolic features of intact heart tissues are well preserved (e.g., refs. 11,16,25,33, and Chapter 15). Such myocyte preparations offer a convenient way to study nucleotide synthesis and degradation under conditions existing in the myocardial cell. In addition, we can look at the intracellular compartmentation of various components. Myocytes lose ATP and adenylates as a function of time during anaerobic incubation in the absence of glucose (25), the presence of uncouplers and glycolysis inhibitors (20), or the presence of 2-deoxyglucose and oligomycin (39). Measurement of ATP-catabolites in a cell homogenate, which can be done by HPLC, may reveal AMP-deaminase activity in situ.

The activities of various pathways may be gauged in intact cells of primary cardiomyocyte culture under physiological conditions, employing labelled precursors (80). This approach allows us to design a model of synthesis - degradation pathways and estimate the importance of different reactions, for example those catalyzed by AMP-deaminase. Its activity estimated in cardiomyocyte extract equals to 0.078 U/mg protein. It is of the order found in human fibroblasts and 8-fold lower than in skeletal muscle myotubes (80). Developmental changes of cardiac AMP-deaminase could be observed with embryonic heart cells (myoblasts), differentiated to myocytes, using selective cultivation conditions (40). The extensive change in the amount of AMP-deaminase, occurring during cell transformation, could be a promising approach to estimate its role in this kind of cell.

Important conclusions on the role of AMP-deamination in the myocardium have been derived from experiments with perfused isolated heart. In the Langendorff preparation (36), most commonly used, decreased perfusion or changed gas composition (from  $O_2$  to  $N_2$ ) initiates net ATP-catabolism. Catabolic enzymes could be activated. Tissue IMP changes during ischemia could indicate enhanced AMP-D activity, the sole pathway able to produce IMP during a relatively short time. Perfusion with substrates for ATP-resynthesis (adenine, adenosine, inosine, hypoxanthine, etc.) could establish AMP-deaminase's share in anabolic routes (see Chapter 26). Additionally, inclusion of enzyme inhibitors [like erythro-9-(2-hydroxy-3-nonyl)adenine (EHNA) for adenosine deaminase] and analysis of the degradation

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products in the perfusate allow us to revaluate two deamination steps existing in adenylate metabolism: adenosine and AMP-deamination. A specific, membranepermeable AMP-deaminase inhibitor is not available. Therefore, direct insight in the AMP-D activity in the whole organ is impossible.

Preliminary data in human-heart biopsies, collected during open-heart surgery and analyzed with HPLC, demonstrate distinct IMP production, which correlates with decreased ATP-levels (Smolenski & Skladanowski, unpublished results).

## **Kinetic Properties and Regulation of Cardiac AMP-Deaminase**

Substrate saturation kinetics of AMP-D under optimal conditions of ionic strength and pH ( $\geq 100$  mM KCl, pH 6.5) reveals a slightly sigmoidal curve, suggesting allosteric regulation. However, the Hill coefficient is reported sometimes as close to 1 (4,23), whereas the majority of values obtained are higher: 1.3 [rat (30)], 1.3 [duck (51)], 1.33 [man (32)], 1.7 [pigeon (29)], 2.0 [cow (70)], 2.35 [pig (53)]. Cooperativity between AMP-binding sites presumably exists, as the calculated number of functional protomers according to the model of Monod et al. is 2 (29, 30).

The half-saturation constant for heart AMP-D under optimal K<sup>+</sup> (or Na<sup>+</sup>) concentrations and in the absence of other effectors is higher than that for skeletal muscle AMP-D (for exception, see ref. 42). It equals to: 1.86 mM [duck (51)], 2.11 mM [hen (31)], 2.78 mM [pigeon (29)], 3.04 mM [rat (30)], 3.1 mM [rabbit (4)], 3.8 mM [pig (23)], 3.9 mM [cow (70)], and 5.0 mM [pig (53)].

Purified myocardial AMP-deaminase is regulated by nucleotides, inorganic and organic phosphates, activated fatty acids, and alkali metal ions. The isolated enzyme is unstable and practically inactive in the absence of metal cations (Skladanowski, unpublished results). In the presence of at least 100 mM K<sup>+</sup> or Na<sup>+</sup>, confirmed as saturating, AMP-D from beef heart is regulated by nucleotides and phosphate (Skladanowski, unpublished data) in a way corresponding to a "K<sub>M</sub>-type" of enzyme, whereas a mixed "K<sub>M</sub>, V<sub>max</sub>-type" of regulation is observed at low ion concentrations (70).

# Activation by ATP and ADP

Subphysiological concentrations of ATP (0.5 mM) and near-physiological concentrations of ADP (0.5 - 1.5 mM) are able to activate maximally the majority of heart AMP-D preparations studied so far. The activation constants for ATP and ADP, calculated for rat-heart AMP-D at 0.75 mM AMP concentration, were 0.037 and 0.095 mM, respectively (30).  $K_{0.5}$  was decreased from 3.04 to about 0.6 mM in the presence of 0.5 mM ATP, and to about 1 mM in the presence of 1.5 mM ADP (30). Rabbit-heart AMP-D was activated up to 300% by 0.05 mM ATP at 0.1 mM AMP concentration, but only to 30% by 0.05 mM ADP (4). ATP-Mg<sup>2+</sup> was shown to be an even stronger activator (4); at a physiological Mg<sup>2+</sup> concentration, a part of ATP exists as a magnesium complex (19). ATP and ADP effects are further enhanced (about 100% with ATP, and 50% with ADP) in the presence of artificial phospholipid membranes (53). This is not true for skeletal muscle AMP-deaminase. Chung and Bridger (12) reported a stronger stimulation of rabbit-heart AMP-D by ADP than by ATP. This dissimilarity is probably caused by a chelating effect of citrate.

#### Inhibition by Phosphate and GTP

Inorganic phosphate, phosphocreatine and several other organic phosphate compounds inhibit the cardiac enzyme analogous to skeletal muscle AMP-deaminase.  $I_{s0}$  values are calculated to be about 0.5 mM and 2.0 mM for phosphocreatine and  $P_i$  inhibition of rabbit-heart AMP-D at 0.1 mM AMP, respectively (4). The binding of the phosphate to the enzyme molecule shifts the substrate saturation curve to a distinct sigmoid (4), but ATP reverses this effect instantly (28).

GTP is known as the most effective inhibitor of AMP-D from skeletal muscle (7). However, its effect on rat-heart AMP-deaminase is more complex. Activation is even observed at low GTP concentrations (30). Rabbit-heart AMP-D is inhibited by about 25% at 0.1 mM AMP in the presence of 5 uM GTP. The activation by micromolar concentrations of ADP and ATP is abolished (4). The inhibition of duck-heart AMP-D by various GTP concentrations is biphasic. It is probably caused by a different sensitivity of the two existing isoforms (51).

## Inhibition by Activated Fatty Acids

Palmitoyl-CoA and stearyl-CoA strongly inhibit the activity of AMP-deaminase from beef (69), hen (31) and pigeon (29) heart at concentrations as low as 5 uM. Both acyl-CoA's inhibit well below their critical micelle concentrations (73). Many enzymes are inhibited by activated fatty acids (78). For some the effect is due to ligand-induced irreversible dissociation of polymeric enzymes (34). The inhibition of heart AMP-D could be of physiological importance during ischemia when an increase in activated fatty acids level in the tissue is observed (6).

### Regulation by Adenylate Energy Charge

The response of AMP-deaminase to variations in adenylate energy charge ([ATP] + 0.5[ADP])/([ATP] + [ADP] + [AMP]) reflects best its behavior in a dynamically changing intracellular environment. The activities of pig-heart AMP-D (23) and the beef-heart enzyme (Skladanowski, unpublished results) are suppressed at energy charge values in the physiological range (0.8 - 0.9). A decrease of this parameter, occurring during ischemia (56), could deinhibit AMP-D and be responsible for IMP accumulation in myocardium (refs. 20,26; Smolenski & Skladanowski, unpublished data). On the other hand, rabbit-heart AMP-D (74) and the rat-heart enzyme (Skladanowski, unpublished results) are unaffected by energy charges below 0.9. In contrast, the skeletal muscle enzyme increases gradually in activity, when the energy charge decreases from 1.0 to 0.0 (74). Rabbit- and rat-heart AMP-deaminase always seem to be in the activated state.

#### Is Cardiac AMP-Deaminase regulated in Vivo?

According to Burger and Lowenstein (10), AMP deamination dominates AMP dephosphorylation due to the cardiac ATP level. However, during accelerated adenylate catabolism, the heart produces considerable amounts of adenosine, and little IMP (58).

The activation of cardiac AMP-D by ATP seems to be insignificant for in vivo regulation. We disregard the question how 5'-nucleotidase, responsible for adenosine production, is affected by ATP (27,38). The fall of ATP-level during ischemia may

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not exceed 60% (26). Even then AMP-D remains saturated by ATP. Inhibitory effects by GTP as well as  $P_i$  are most likely permanently suppressed by ATP-activation (4). Therefore, an ischemia-induced  $P_i$ -increase (24) probably does not play a significant role in heart AMP deamination. This thesis is confirmed by the fact that 2 mM  $P_i$  does not change hen-heart AMP-D substrate saturation kinetics in the presence of 5 mM ATP (28). ADP is unlikely to regulate AMP-D activity in the muscle cell, because it preferentially binds to actin filaments.

The key role in AMP-D regulation in heart therefore seems to be played by the level and accessibility of substrate. The free AMP concentration in guinea-pig heart is well below 1 uM (8). This has implications for both cytosolic 5'-nucleotidase and adenylate deaminase. Since the  $K_{0.5}$  for AMP of each enzyme appears to be considerable higher than its free cytosolic concentration, the catalytic properties of either enzyme are unlikely to be rate-controlling. As mentioned before, AMP dephosphorylation prevails of deamination. Substrate availability for these potentially competing enzymes is therefore probably limited by intracellular compartmentation.

In conclusion none of the properties of heart AMP-D seems to be more significant than the other for in vivo regulation of this enzyme. More complex experiments on the cumulative effects of effectors with micromolar AMP concentrations could settle this problem.

#### **Development of Heart AMP-Deaminase**

Changes in properties of skeletal muscle AMP-D during growth are due to isozymic pattern differentiation (61). According to Ogasawara et al. (43), rat-heart tissue possesses one type of AMP-deaminase isozyme (denoted C). This belongs to the isozyme family: homoprotomeric A (skeletal muscle), B (liver), C (heart), and heteroprotomeric hybrids composed of the B and C type subunits. On the other hand, three types of isozymes are found in human heart. Two have the properties of the erythrocytic forms  $E_1$  and  $E_2$ , and one those of the hepatic form L (47). Two forms differing in kinetic and regulatory properties, have also been detected in beef heart (Skladanowski, unpublished results).

Developmental changes occur in the amount of isozymes in rat heart, with a shift from five isoenzymes to only one adult form (46). In most tissues (including heart), AMP-D activities are higher in mature cells or organs than in the early postnatal stage (46). Chicken-heart AMP-D activity increases about four times between 18 days of postnatal development and adulthood (54).

Striking differences in adenylate metabolism are encountered in rat-heart myoblasts, differentiating to myocytes. The 14-fold increase in AMP-D activity correlates well with a jump in creatine kinase activity (40). These and other enzyme activity changes during development are presented in Table 1.

AMP-deaminase may be important for the conservation of purine nucleotides through the operation of purine nucleotide cycle. This also holds for 5'-nucleotidase, operating through vasodilation by adenosine (or inosine). The ratio AMP-D : 5'nucleotidase changes during differentiation from 0.11 to 0.48. This indicates that, in rat-heart myocytes, the interconversion of purine nucleotides is mainly from AMP to IMP, not inversely (40). The same is true for rat skeletal muscle cells (81). The increase of enzymes responsible for ATP sparing and purine ring conservation in myocytes correlates temporarily with an increase of the ATP demand of and use by contracting myocytes.

# Heart AMP-Deaminase and AMP Catabolism

During oxygen deficiency cytosolic ADP cannot be rephosphorylated by creatine phosphate. ADP is then used more effectively by myokinase than by creatine kinase and mitochondrial oxidative phosphorylation. Judged from AMP-binding equilibria constants, AMP levels increase from about 100 nM during normoxia to about 800 nM in ischemic hearts (8). This change occurs in a strikingly lower range than that suggested by total tissue AMP analysis: 100-200 uM (9). The change in free AMP is several fold larger, and thus could be a better stimulus for cytoplasmic enzymes, converting AMP. Under these conditions AMP becomes available for adenylate deaminase and 5'-nucleotidase, either for further degradation to IMP or to the coronary vasodilator adenosine, respectively. In fact, several studies demonstrate a rise in myocardial IMP level during ischemia (15,26,49) and especially during anoxia (18). Also in cardiomyocytes, subjected to conditions stimulating ATP catabolism, IMP distinctly increases (20,39). However, the superiority of the AMP  $\Rightarrow$  adenosine route over that of AMP  $\Rightarrow$  IMP in ischemic heart adenylate catabolism is not questioned. It cannot be explained by kinetic differences between AMP-D and cytosolic 5'-nucleotidase, because these enzymes, isolated from rat heart (27,30), have very similar properties. Nevertheless, the total enzyme activity of 5'nucleotidase is about 15 times higher than that of AMP-deaminase (60).

The question remains: What are the further fates of accumulated IMP? Part is most likely hydrolyzed to inosine, transported from the myocytes to the endothelial cells, and degraded to uric acid. The rest establishes a pool of nondiffusable nucleotides, ready for recovery of depleted adenylates by the purine nucleotide cycle. The presence and role of this cycle in heart musle is still controversial (75). Time-delayed variation in flux through different parts of the cycle – AMP-deaminase during

Enzyme	Increase	Reference	
Creatine kinase	10-12 <b>v</b>	(35.40)	
AMP-deaminase	10-12x 14x	(40)	
Adenosine deaminase	1.4x	(40)	
5'-Nucleotidase	3.1x	(40)	
Purine nucleoside			
phosphorylase	1.7x	(40)	
Myokinase	ca. 3x	(35)	
Xanthine oxidase	manifold	(64)	
Adenosine kinase	1.4x	(40)	
HGPRT	1.7x	(40)	
APRT	1.6x	(40)	

Table 1. Changes of enzyme activities in heart adenylate metabolism during cell differentiation and development

ischemia and adenylosuccinate synthetase with adenylosuccinase during recovery – could be an explanation. The size of the remaining IMP pool must depend on the rate of cytosolic dephosphorylation by 5'-nucleotidase. This reaction is inhibited by increased  $P_i$  and by decreased energy charge (27). Thus the preservation of IMP would be better during more severe ischemia.

The importance of AMP-deamination in AMP catabolism during either normoxia or ischemia is indirectly confirmed by the use of EHNA, a strong adenosine deaminase inhibitor, during rat-heart perfusion with adenosine (2), or 5'-amino-5'-deoxyadenosine (an adenosine kinase inhibitor) combined with EHNA during guineapig heart perfusion (65). EHNA increases the proportion of adenosine released, without changing total purine release in normoxia from 7 (without inhibitor) to 18%, and from 15 (without inhibitor) to 60% of total purine release during ischemia. It suggests that purine formation through AMP  $\Rightarrow$  IMP  $\Rightarrow$  inosine, bypassing the inhibited step is significant under both conditions (2). The decrease of intramyocardial adenosine without change of total purine release after homocysteine infusion during ischemia also indicates that purine formation could occur partly via IMP under ischemic conditions (1).

#### **Concluding Remarks**

1. Myocardium contains AMP-deaminase activity which is about two orders of magnitude lower than that in skeletal muscle. The enzyme content rises sharply in the myocardial cell during development, together with other enzymes responsible for ATP maintenance.

2. Isolated AMP-deaminase is activated by ATP, ADP, K<sup>+</sup> or Na<sup>+</sup> ions, and inhibited by GTP, P<sub>i</sub>, activated fatty acids. The activation by ATP is enhanced by phospholipid bilayers. All nucleotide effects are modulated by  $Mg^{2+}$ . The reaction rate in vivo is mainly regulated by availability of the substrate AMP, which is normally present in submicromolar amounts. Changes of enzyme effectors are probably insignificant for in vivo regulation of AMP-deaminase in view of the prevailing ATP activation.

3. AMP deamination is responsible for a considerable part of the purine catabolites released from ischemic heart and probably for the entire purine efflux during normoxia.

4. The postulated physiological role for cardiac AMP-deaminase is preservation of the purine ring inside myocytes (as nondiffusable IMP) during increased ATP catabolism. Salvaged IMP could later be the source for fast adenylate pool restoration.

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# Chapter 7

# Redox Manipulation of Free Cardiac Adenylates and Purine Nucleoside Release

## **Reciprocity between Cytosolic Phosphorylation Potential and Reduction-Oxidation State or Free AMP in Perfused Working Heart**

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The thermodynamic equilibrium relations between free adenylates and redox state in the cytosol were reviewed as applicable to perfused heart. Studies with isolated working guinea-pig hearts examined cytosolic redox state, phosphorylation potential, and the free concentration of AMP ([AMP]) in relation to the coronary venous output of adenylate catabolites. Excess pyruvate (20 mM), relative to 20 mM lactate, produced increases in the phosphorylation potential and total cardiac AMP content, but inorganic phosphate content, [ADP], [AMP], and purine nucleoside release were decreased. Conversely, excess lactate induced the releases of inosine and large amounts of xanthine, the reduced precursor of uric acid. When 15 mM glucose was the sole substrate, lactate output and adenosine plus inosine release changed concordantly during adrenergic stimulation. During low-flow ischemia with adrenergic stimulation, output of purines showed a large but transient increase, was mainly (67% - 87%) in the form of adenosine plus inosine, and was attenuated by cytosolic oxidation due to intracoronary administration of 5 mM pyruvate. Uric acid output accounted for 40% - 50% of total purine release during normoxia, but for not more than 10% during early ischemia. It was concluded that, in perfused working heart, 1) the cytosolic oxidation by pyruvate can increase myocardial energy state, apparently in accordance with a reciprocal relation between the cytosolic [NADH]/[NAD+] ratio and the phosphorylation potential, 2) a high phosphorylation potential is associated with very low, i.e., submicromolar [AMP] and minimum adenylate catabolite output, and 3) the cytosolic phosphorylation potential may exert control over free [AMP], which appears to function as a major determinant of 5'-nucleotidase flux.

#### **Introduction and Overview**

Previously we demonstrated a quantitative association between the rate of myocardial purine nucleoside release and the calculated concentration of free AMP ([AMP]) in the cytosol (6,10,11). Because [AMP], like the free [ADP]/[ATP] ratio, was highly responsive to physiologic changes in myocardial energy balance and/or utilization, free cytosolic AMP was regarded as an integral component of the free extramitochondrial adenylate system. Established characteristics of the free cytosolic adenylate system in working heart are 1) adaptability of the cytosolic phosphorylation potential (ATP potential) to the acute myocardial energy balance over a wide range of workloads, 2) an apparent linear relationship between free cytosolic [ADP] and myocardial oxygen consumption (mitochondrial respiration) in the physiological range, with [ADP] between 10 and 100 uM, and 3) strict compartmentation of AMP between cytosol and mitochondrial matrix, with [AMP] in the upper submicromolar range being highly responsive to the [ADP]/[ATP] ratio. Since, moreover, the [ADP]/[ATP] ratio is a major determinant of the cytosolic

phosphorylation potential ([ATP]/([ADP]  $\cdot$  [P<sub>i</sub>]), myocardial energy state and [AMP] are reciprocally related. Net release of adenosine plus inosine appeared to correlate more closely with [AMP] than with total myocardial AMP content (6,10), supporting the possibility that 5'-nucleotidase flux is controlled by the availability of cytosolic AMP. Studies in skeletal muscle, liver, brain and erythrocytes have suggested that cytosolic redox state and ATP potential are, in principle, reciprocally related (27,54). It seems therefore logical to assume that cytosolic reduction potential and [AMP] may change concordantly. This raised the possibility that myocardial release of adenylate catabolites (purine nucleosides and bases) might be responsive to the cytosolic redox state per se, with the phosphorylation potential functioning as the intermediary metabolite mediator system. In fact, the coronary effluent lactate/ pyruvate concentration ratio and the rate of myocardial purine nucleoside output were directly related (6).

In the present study we examined further the relationships between cytosolic oxidation-reduction state, ATP phosphorylation potential, free cytosolic [AMP], and net release of purine nucleosides and bases, using the isolated physiologically performing guinea-pig heart as the experimental model. The cytosolic redox state was externally manipulated by infusion of excess amounts of redox substrates (lactate, pyruvate) under the conditions of normoxia and global, low-flow ischemia with residual oxygen consumption.

#### Theoretical background

Creatine kinase and myokinase are powerful enzymes in skeletal muscle, heart and brain, i.e., in tissues with potentially high ATP turnover rates (36,38). The enzyme couple is strategically located in the extramitochondrial space (mitochondrial intermembrane + cytosolic compartments), but is absent from the intramitochondrial matrix (23,38,46). The kinase equilibria are only responsive to the thermodynamic ("free") concentrations of ATP, ADP, and AMP ([ATP], [ADP], [AMP]) or to the cytosolic [ATP]/[ADP] ratio. Since the bulk of myocardial ATP is located in the extramitochondrial compartment (10,20,52) and because actomyosin only insignificantly binds ATP (40), the chemically determined overall myocardial ATP content has been taken as a reasonable approximation of [ATP]. Current estimates based on total tissue analysis, non-aqueous or digitonin fractionation techniques place cytosolic [ATP] in the 7 to 12 mM range (10,20,52). These concentrations are principally found in hearts or cardiomyocytes with "energized" mitochondria under respiratory control, since energization of the inner mitochondrial membrane produces a gradient for the [ATP]/[ADP] ratio (higher outside) across this membrane (22,25,26,52). The subcellular distribution of ADP is complex, because actin and also myosin tightly bind ADP (21,40,48) and because the energized ADP-ATP translocase of the inner mitochondrial membrane exchanges extramitochondrial ADP in preference for intramitochondrial ATP (22,25,26). Thus, ADP rephosphorylation by creatine kinase, binding by actomyosin, and the vectorial ADP-ATP translocase during mitochondrial respiratory control act in concert to reduce the extramitochondrial [ADP] to a physiologically permissible minimum. In fact, the creatine kinase equilibrium predicts [ADP] in the 10 to 100 uM range as a function of the physiological state of the heart (6,10).

Free ADP concentrations in the low submillimolar range are consistent with the absence of a definitive ADP signal in <sup>31</sup>P-NMR studies in, e.g., heart and skeletal muscle. Consequently, relatively high values between 10,000 - 100,000 M<sup>-1</sup> of the cytosolic phosphorylation potential are obtained. The value of  $[ATP]/([ADP] \cdot [P_i])$ , combined with the standard free energy of ATP hydrolysis  $[\Delta G_{ATP}^{\circ} = -32.35 \text{ kJ/mol}$  under conditions prevailing in vivo (54)], determines the actual free energy change during ATP hydrolysis in the cytosol:

(Eq. 1)  $\Delta G_{ATP} = \Delta G_{ATP}^{\circ} + RT \ln([ADP] \cdot [P_i])/[ATP]$ 

## in which R = gas constant, 8.314 J/°K/mol, and T = absolute temperature, ° Kelvin.

In pressure-overloaded, isolated working guinea-pig heart, the cytosolic phosphorylation potential can fall to near 7,000 M<sup>-1</sup> yielding  $\Delta G_{ATP} = -55.17$  kJ/mol (13.18 kcal/mol); on the other hand, in non-working, empty, beating Langendorff heart, the ATP potential is near or higher than 80,000 M<sup>-1</sup> [ $\Delta G_{ATP} \geq -61.45$  kJ/mol (14.68 kcal/mol), ref. 10]. The free energy change of cytosolic ATP hydrolysis may be compared with the 52.7 kcal liberated by the mitochondrial respiration chain when a pair of electron passes from intramitochondrial NADH to oxygen to yield H<sub>2</sub>O (30). Assuming a coupling ratio of 3 ATP per oxygen atom reduced, the energy stored in cytosolic ATP varies between 39.54 kcal (75% efficiency) at high workloads to 44.04 kcal (83.5% efficiency) at low workloads. The reason for the apparent decrease in cytosolic energy storage efficiency at high workloads (i.e., high rates of ATP turnover) is not clear, but may be related in part to the fact that energy is obligatorily expended when intramitochondrial ATP (with a relatively low energy potential) must be exported at increased rates into the cytosol, which has a relatively higher phosphorylation potential (22,25,26,52).

Free cytosolic ADP cannot be directly measured by current invasive techniques (acid extraction, sonication plus density gradient fractionation, digitonin cell membrane lysis). However, division of the chemically measured cellular ATP concentration by the value of cytosolic [ATP]/[ADP] (derived from the creatine kinase equilibrium equation), yields the mean cytosolic [ADP]. This approach appears to be justified, since most of guinea-pig cardiac creatine kinase (about 89%) may be located in the cytosol (37). There appears to be no substantial ATP concentration gradient between cytosol and nucleus (50) and the intermembrane space of mitochondria is freely permeable to cytosolic adenylates, protons, and creatine compounds. However, the [ADP] obtained by equilibrium equations ignores the possible existence of microenvironmental diffusion gradients (e.g., ref. 34). Possible differences in the chemical activity of water in the various subparenchymal and subcellular compartments are also not taken into consideration.

The myokinase reaction, like the creatine kinase reaction, is reversible. In skeletal muscle, heart, and brain, myokinase activities are at least as high as those of creatine kinase (38). Coronary endothelium also contains creatine phosphate (and hence creatine kinase) and myokinase in substantial quantities (35). The myokinase equilibrium constant is dependent on the hydrogen ion concentration and free Mg<sup>2+</sup> ([Mg<sup>2+</sup>]) in the cytosol. Thus, under conditions of low constant metabolic throughput (relative to the total potential activity) and constant metabolite levels for a sufficiently extended period (metabolic steady state), the myokinase reaction seems likely to reach the thermodynamic near-equilibrium state:

(Eq. 2) [AMP] =  $K_{mvk} \cdot [ADP]^2/[ATP]$ 

 $K_{mvk} = 1.12$  at pH 7.2,  $[Mg^{2+}] = 0.9$  mM (29).

Even in systems with rather low myokinase activities, e.g., erythrocytes and liver, the enzyme probably operates at near-equilibrium (54). Moreover, it has been argued (49,54) that in brain, where activities of myokinase and creatine kinase are about 30% of those in heart (38), the enzymes are nevertheless sufficiently powerful to catalyze near-equilibrium reactions under appropriate conditions. In isolated working guinea-pig heart perfused for prolonged periods at constant pre- and afterloads, whether examined under the conditions of low, normal or high-energy outputs or during low-flow ischemia, we recently found essentially the same square root-dependencies between thermodynamic [AMP] (myokinase equation) and cytosolic [ADP]/[ATP] (creatine kinase equation). These findings suggest near-equilibrium catalyses by the kinase couple in perfused heart under various steady-state physiological and pathophysiological conditions (6).

Since H<sup>+</sup> is an important reactant of the creatine kinase and because this enzyme is  $Mg^{2+}$ -dependent, the value of the expression  $[H^+]/K_{cpk}$ , an essential term in the creatine kinase equation, is strongly H<sup>+</sup>- and Mg<sup>2+</sup>-dependent (29):

(Eq. 3)  $[ATP]/[ADP] = [CrP]/[Cr] \cdot [H^+]/K_{cpk}$ (Eq. 4a)  $log([H^+]/K_{cpk}) = -0.87 \cdot pH + 8.31$ ; free  $[Mg^{2+}] = 0.9 \text{ mM}$ (Eq. 4b)  $log([H^+]/K_{cpk}) = -0.86 \cdot pH + 8.30$ ; free  $[Mg^{2+}] = 3.6 \text{ mM}$ 

in which CrP = creatine phosphate, Cr = creatine.

It is thus critical to obtain independent values for cytosolic  $[H^+]$  and  $[Mg^{2+}]$ , in order to strengthen confidence in data and conclusions derived from myokinase and creatine kinase equilibrium calculations. A number of recent physico-chemical, cell biological and <sup>31</sup>P-NMR studies indicate that the free Mg<sup>2+</sup> concentration in cell water is relatively constant at 0.5 to 1.0 mM in mammalian organs, including heart (15,19,32,43,55). Bicarbonate and CO<sub>2</sub> measurements in tissues and perfusing media as well as <sup>31</sup>P-NMR detectable phosphate chemical shifts indicate intracellular pH values near 7.2 in well-perfused and oxygenated tissues (10,15,19,32,54). Using the Henderson-Hasselbalch equation in combination with measured tissue bicarbonate and perfusion fluid [CO<sub>2</sub>] we obtained the following operational equation for calculation of intracellular pH in isolated working guinea-pig hearts (6,10):

(Eq. 5)  $pH_i = 7.524 \cdot e^{(-0.0008786 \cdot pCO_2)}, n = 20, r = 0.959$ 

in which  $pCO_2$  = coronary effluent  $pCO_2$  (mmHg) in hearts perfused with a Krebs-Henseleit buffer, pH 7.4 (25 mM bicarbonate), oxygenated with  $O_2/CO_2$  = 95/5.

Three powerful cytosolic enzymes, glyceraldehyde-3-phosphate dehydrogenase (GAPDH), 3-phosphoglycerate kinase (PGK), and lactate dehydrogenase (LDH) (41), associate the cytosolic ATP potential with the overall cytosolic oxidation-reduction state, if it is assumed that GAPDH and LDH share free cytosolic NAD<sup>+</sup>(H) as a common coenzyme. In theory cytosolic reduction state and ATP potential are reciprocally related according to the following two relationships (27,54):

(Eq. 6)  $[ATP]/([ADP] \cdot [P_i]) = [GAP]/[3-PG] \cdot [Pyr]/[Lac] \cdot K$ 

in which  $K = K_{gapdh} \cdot K_{pgk}/K_{ldh}$ ; GAP = glyceraldehyde-3-phosphate, 3-PG = 3-phosphoglycerate, Pyr = pyruvate, Lac = lactate; and

(Eq. 7) [Pyr]/[Lac] =  $K_{idh} \cdot [NAD^+]/([NADH] \cdot [H^+])$ .

Since the [ADP]/[ATP] ratio appears to serve a dual role as the major determinant of the value of the cytosolic phosphorylation potential (Eq. 3 divided by  $[P_i]$ ) and of [AMP] (Eq. 2), it seems logical to suggest that [AMP] and the free cytosolic [NADH]  $\cdot$  [H<sup>+</sup>]/[NAD<sup>+</sup>] ratio are directly related, provided induced changes in [Pyr]/[Lac] are not compensated by those in [GAP]/[3-PG]. The complexity of the interactions between the cytosolic NAD<sup>+</sup>(H)-dependent dehydrogenases, phosphoglycerate kinase, and myokinase is increased by the fact that the equilibrium constants of both kinases, especially  $K_{pgk}$ , are highly sensitive to [Mg<sup>2+</sup>] and ionic strength (14,29).

Hearts were isolated from male albino guinea pigs (350-600 g body wt.) fed a standard chow diet. Isolation, preparation, non-recirculating hemoglobin-free perfusion and physiologic performance characteristics of the isolated, perfused working hearts have been detailed elsewhere (7,9). Hemodynamic parameters [spontaneous heart rate (HR), coronary venous outflow and aortic flow, mean left ventricular filling pressure ( $P_v$ ), and phasic and mean aortic pressure ( $P_a$ )] were continuously monitored using an 8-channel direct recorder (Hewlett Packard). Left continuously monitored using an o-channer uncer recorder (newlett rackard). Lett ventricular pressure work per min was judged from the heart rate-pressure product expressed as HR  $\Delta P$  ( $\Delta P = P_a \cdot P_v$ ) (e.g., refs. 7,12). All hearts were perfused at constant filling pressures between 12 and 16 cmH<sub>2</sub>O. Perfusion medium was a Krebs-Henseleit bicarbonate buffer (28), pH 7.40-7.45, 38°C, 294 ± 0.4 mOsm (n = 29), oxygenated with 95%  $O_2$ :5%  $\dot{CO}_2$ ; the medium was modified to contain free calcium in the physiologic concentration range between 1.5 - 2.5 mEq/L. The perfusion fluid was supplemented with 5 mM glucose (+ 5 U/L bovine insulin) as the sole energy substrate or in combination with pyruvate or L-lactate as described in the table and Fig. 1. When additional excess Na-pyruvate or Na-lactate were used the NaCl concentration of the medium was reduced accordingly to maintain osmolarity constant. L-Norepinephrine (NE) stock solution (500 uM), stabilized by 2 mg/ml sodium metabisulfite (Winthrop-Breon, New York, NY), was infused directly into the left atrium of the working hearts, yielding arterial perfusate NÉ concentrations between 0.2-1.1 uM; these NE levels represented near-maximum effective NE doses in the isolated perfused guinea-pig heart (7). Water for perfusion media and analytical tests was freshly prepared from a low-output quartz doubledistillation system. To remove bacteria and particulate impurities all solutions and media including those for enzymatic analyses and HPLC were filtered (Millipore filter, 0.22 um) immediately prior to use.

Myocardial  $O_2$  uptake was calculated from the continuously monitored arterio-venous  $O_2$  concentration difference (Clark-type oxygen electrode) and coronary venous effluent rate. Arterial and coronary venous p $O_2$ , p $CO_2$ , and pH were measured as previously described (10,11). Intracellular pH was estimated from measured coronary venous p $CO_2$ , using Eq. 5 (6).

To measure the volume of the extracellular space, [<sup>14</sup>C]carboxylate-inulin (15 mg/L<sup>i</sup> sp. act. =  $1.5 \cdot 10^6$  dpm/mg) was infused for 6 min prior to stop-freezing the hearts. Radioactivity was measured by liquid scintillation counting using a standard cocktail (9). When experiments were terminated by freeze-fixation of the working hearts, a Wollenberger clamp at the temperature of liquid N<sub>2</sub> was used. Myocardial metabolites were measured in neutralized perchloric acid tissue extracts, using the following standardized extraction procedure. Frozen tissue was pulverized in a mortar under liquid  $N_2$ ; 1 g pulverized myocardium at the temperature of liquid  $N_2$  was added to 4 ml ice cold 0.3 N perchloric acid; this ensured slow acidic thawing of tissue powder at temperatures below 0°C. The crude ice cold homogenate was stirred (660 rpm) for 1 min using a precooled (0°C) Teflon rod of a diameter 2 mm smaller than that of the plastic test tube (inner diameter = 14 mm); during stirring the test tube was placed in an ice/water mixture. Subsequent centrifugation (12,000 g, Sorvall RC-5) at  $0 - 4^{\circ}$ C for 10 min was followed by gentle pH-controlled neutralization of the supernatant fluid to a slightly acidic pH (5.8 to 6.3), using small aliquots of refrigerated (4°C) 1 N and 0.1 N KOH, respectively. After 30 minstanding in ice, potassium perchlorate was removed by centrifugation (12,000 g, 10 min, 0-4°C). Aliquots of the slightly acidic extracts were immediately assayed for cardiac metabolites (enzymatic analysis). Reextraction (n = 8) of the homogenate pellet for the key metabolites ATP, ADP, AMP, creatine phosphate, creatine, inorganic phosphate, pyruvate, and dihydroxyacetone phosphate indicated that the extraction procedure yielded the following percentage recoveries, respectively: 92.7, 89.5, 83.7, 92.8, 95.1, 95.1, 82.7, 89.8. Data presented were not corrected for

measured recoveries. Most of the enzymatic tests were done in the dual beam/dual wavelength mode (measuring wavelength = 340 nm, reference wavelength = 385 nm, millimolar absorption coefficient =  $6.097 \text{ cm}^2 \cdot \text{umol}^{-1}$ , Perkin-Elmer spectrophotometer model 556). Intracellular contents of the perfusion medium metabolites (pyruvate, lactate, inorganic phosphate) were calculated as the difference between respective total myocardial contents and extracellular amounts using the [<sup>14</sup>C]inulin distribution data. The mean extracellular metabolite concentration was taken as (arterial plus venous metabolite concentration)/2. The [<sup>14</sup>C]inulin space in freeze-clamped hearts was 0.57 ± 0.01 ml/g wet wt. (n = 16 mean ± S.E.M.). The intracellular solvent space was taken as the difference between total tissue water and [<sup>14</sup>C]inulin space. Intracellular water was not corrected for mitochondrial water. Perfused myocardium wet weight/dry weight ratio after freeze-clamping was 9.6 ± 0.2 (n = 44).

Coronary venous purine nucleosides (adenosine, inosine) and other natural purines (hypoxanthine, xanthine, uric acid) were separated using reverse phase HPLC as described in detail previously (10,12). Purines were identified employing known retention times in combination with measured absorbance characteristics at four different wavelengths (254, 263, 273, 293 nm) using a multiwavelength detector (Waters model 490). Quantitation of the purines was accomplished by comparisons with calibrated standards. Standards and experimental samples were kept at about  $4^{\circ}$ C prior to separation by HPLC using a refrigerated automatic injector system (Waters WISP model 710B). Repetitive injections of standards showed that all purines were stable for more than 24 h under these conditions.

Data were statistically analyzed using Student's t-test for unpaired samples. For multiple comparisons Duncan's variance analysis was performed.

#### **Results and Discussion**

#### Effects of Excess Pyruvate vs. Lactate on Cytosolic Adenylates and Purine Nucleoside plus Xanthine Output

The oxidation-reduction state of cytosolic redox couples (e.g.,  $[NAD^+]/([NADH] - [H^+]))$  was altered by perfusing the isolated working guinea-pig hearts with excess pyruvate (20 mM) or excess lactate (20 mM) in the presence of 5 mM glucose (plus insulin). Table 1 shows that, relative to lactate, pyruvate effected respective increases in total cardiac AMP content, the [CrP]/[Cr] ratio, the phosphorylation states of creatine phosphate and ATP, and the cytosolic [ATP]/[ADP] ratio; intracellular  $[H^+]$  and the total myocardial contents of creatine phosphate and ATP were not appreciably different between the two conditions. Similarly, the heart rate-pressure product was virtually identical with either pyruvate or lactate, but myocardial oxygen uptake tended to increase slightly in presence of lactate; this was expected from the increase in [ADP] from about 25 to 41 uM under the same conditions (6,10). In accordance with the small increase in the [CrP]/[Cr] ratio, pyruvate also ef-fected a small but definitive decrease in the cellular inorganic phosphate concentration.

The AMP content was increased almost 8-fold by excess pyruvate relative to lactate. On the other hand, calculated [AMP] fell 51% (227 vs. 108 nM, Table 1). The coronary venous concentration of adenosine was below the detection limit of 2 picomoles per 200 uL (10 nM) in the presence of pyruvate or lactate. Nevertheless, in agreement with our concept (6,10,11) that it is the cytosolic [AMP], not intramitochondrial [AMP], which functions as the common precursor of released adenosine plus inosine, coronary venous adenosine plus inosine release was minimal in the presence of pyruvate ( $\leq 2$  nmol/min per g dry wt.), but was increased to 8.35 nmol/min per g dry wt. in the presence of lactate (Table 1).

Free Cardiac Adenylates and Purine Nucleosides

The lactate-perfused hearts released additionally very large quantities of xanthine  $(44.6 \pm 8.4 \text{ nmol/min} \text{ per g dry wt.})$ , the immediate precursor of uric acid; released hypoxanthine and uric acid, however, remained below the detection limit. Thus, inosine and xanthine were the major purines released from the redox-manipulated working guinea-pig hearts. Myocardial xanthine output was apparently dependent on a high cytosolic reduction potential, indicating that the native form of cardiac xanthine oxidase functions as an NAD<sup>+</sup>-dependent dehydrogenase (3). The xanthine dehydrogenase/oxidase reaction was presumably feed-back inhibited by NADH accumulated in the cytoplasm, resulting in the output of xanthine instead of uric acid. Since xanthine oxidase is a marker enzyme for coronary capillaries (24), the xanthine of the coronary outflow tract most likely originated in the coronary capillary endothelium. Furthermore, in accordance with Eqs. 2, 6 and 7, the large output of xanthine probably reflected a fall in the endothelial phosphorylation potential in association with an increase in [AMP]. In fact, myokinase activity is about 10 times higher than that of adenosine deaminase in the coronary endothelium of the guinea pig (35).

Measured concentrations of reduced intracellular metabolites (lactate,  $0.45 \pm 0.11$  mM; alpha-glycerophosphate,  $0.02 \pm 0.004$  mM) were quite low in the 20 mM pyruvate-perfused hearts, but greatly increased (12.3 and 5.4 mM, respectively) in presence of 20 mM lactate. Similarly, the cellular concentration of alanine, which is likely in near-equilibrium with cytosolic pyruvate via alanine aminotransferase (e.g., ref. 42), was 13-fold higher with pyruvate (7.8  $\pm$  0.8 mM) than with lactate.

Variable	Addition to 5 mM glucose + 5 U/L insulin		
	20 mM pyruvate	20 mM lactate	
pH <sub>i</sub>	7.224	7.235	
ATP content, umol/g	$24.6 \pm 0.6$	$23.9 \pm 0.7$	
AMP content, umol/g	$1.14 \pm 0.13$	$0.15 \pm 0.05^*$	
CrP content, umol/g	$50 \pm 3$	$46 \pm 3$	
[CrP]/[Cr]	$2.30 \pm 0.10$	$1.74 \pm 0.10^*$	
[P <sub>i</sub> ], mM	$4.6 \pm 0.3$	$5.8 \pm 0.3^{\#}$	
$[CrP]/([Cr][P_i]), mM^{-1}$	$0.52 \pm 0.04$	$0.280 \pm 0.017^*$	
$[ATP]/([ADP][P_i]), M^{-1}$	57,276	31.080	
[ATP]/[ADP]	254	193	
[ADP], uM	24	39	
[AMP], nM	108	227	
V(AR + INO), nmol/min · g			
measured	< 2.0	$8.4 \pm 0.4$	
predicted from Eq. 8	2.89	8.75	
HR $\Delta P$ , 10 <sup>-3</sup> cmH <sub>2</sub> O/min	$20.56 \pm 0.02$	$19.91 \pm 0.02^*$	
MVO <sub>2</sub> , umol/min · g	45.1 ± 1.6	$50 \pm 4$	
$H_2O_{ic}, ml/g$	$4.1 \pm 0.3$	$3.18 \pm 0.08 \#$	

Table 1. Effects of excess pyruvate vs. excess lactate on myocardial adenylates, free ADP, AMP, and nucleoside release

Values are means  $\pm$  S.E., n = 5-7. Working hearts perfused for 20 min at  $P_v = 16 \text{ cmH}_2\text{O}$  and  $P_a = 100 \text{ cmH}_2\text{O}$ . Where applicable, data are expressed per g dry weight; [] denotes intracellular thermodynamic concentration. pH<sub>i</sub>: Intracellular pH calculated from coronary venous pCO<sub>2</sub> according to Eq. 5. V(AR + INO): Coronary venous output of adenosine plus inosine. MVO<sub>2</sub>: Myocardial oxygen consumption. H<sub>2</sub>O<sub>ic</sub>: Intracellular water space. Intracellular free [Mg<sup>2+</sup>] was assumed near 1 mM (Eq. 4a). K<sub>myk</sub> = 1.12 (Eq. 2). Total cardiac AMP was 279 and 47 uM in the presence of 20 mM pyruvate and 20 mM lactate, respectively. Consequently, free [AMP] was less than 1% of total AMP (10). # p<0.05, \* p<0.01.

Assuming that pyruvate distributed across the cell membrane in a manner similar to that of lactate (both metabolites are 3-carbon monocarboxylates and highly dissociated at physiological pH) the calculated intracellular [Pyr]/[Lac] ratio was about 28 in presence of excess pyruvate, but only 0.007 in presence of excess lactate, i.e., about 4000-fold lower. Clearly pyruvate, relative to lactate, oxidized the cytosolic NAD<sup>+</sup>-NADH pool, and this imposed oxidation was associated with a significant increase in the cytosolic phosphorylation potential (Table 1). The data are consistent with the previously reported association between coronary effluent [Lac]/[Pyr] ratio and purine nucleoside release in glucose-perfused heart (6) and also with the concordant changes in lactate output and purine nucleoside release during adrenergic stimulation (see Fig. 2). On the other hand, the present reciprocity between cardiac redox and energy state apparently disagrees with the stability of the phosphorylation potential in the liver of ethanol-cyanamide treated rats (15). Here the [Lac]/[Pyr] ratio increased only about 3-fold, not several orders



Fig. 1. Effects of external redox manipulations on coronary venous release of adenylate catabolites from isolated working guinea-pig hearts under conditions of normoxia and NE-ischemia. Data are means  $\pm$  S.E., n = 6-8. NE: norepinephrine; AR: adenosine; INO: inosine; HX: hypoxanthine; X: xanthine; U<sup>-</sup>: uric acid. HR  $\Delta P$ : normoxia, ~ 19,000 cmH<sub>2</sub>O/min; during ischemia, HR  $\Delta P$  fell progressively from about 4,500 cmH<sub>2</sub>O/min at 2 min to zero at 25-45 min, and did not appreciably differ among the three conditions tested. Variance analysis was done after square-root transformation of raw data. Total purine release: \*p<0.05 vs. 5 mM glucose; (+)p<0.05 vs. 5 mM glucose; + 5 mM lactate. Adenosine plus inosine release: (\*)p<0.05 vs. 5 mM glucose; (+)p<0.05 vs. 5 mM glucose; + 5 mM lactate.

of magnitude; moreover, the GAP/3-PG couple can possibly compensate changes in [Lac]/[Pyr] more effectively in rat liver in vivo than in isolated working guinea-pig heart.

Of special interest was the fact that pyruvate induced a large increase in AMP content, but a decrease rather than increase in inosine release. Even if one accepts the widely held view that most of cardiac AMP is located in the extramitochondrial compartment, the pyruvate-dependent decrease in inosine release in presence of increased AMP content is difficult to explain. According to Table 1 pyruvate did not appreciably increase the already very high concentration of the putative 5'-nucleotidase inhibitor creatine phosphate (45). Pyruvate decreased (19%) the concentration of ATP slightly from about 7.5 to 6.1 mM. Allosteric ATP inhibition, not activation (13), of cytosol 5'-nucleotidase has been discussed for guinea-pig cardiac muscle (16). The minor decrease in myocardial ATP level would be expected to de-inhibit cytosolic 5'-nucleotidase in vivo, but measured nucleoside release was decreased instead of increased (Table 1).

Alternatively, the present AMP concentration and purine nucleoside release data can be discussed in terms of the recently proposed AMP-availability model (6,10,39); accordingly, the free cytosolic ATP-ADP/myokinase system can "downregulate" [AMP] into the submicromolar concentration range, effectively reducing cytosolic [AMP] to levels several orders of magnitude lower than  $K_{mAMP}$  values of 5'nucleotidase and adenylate deaminase (13,16,47). Under such conditions cytosolic 5'nucleotidase and adenylate deaminase fluxes would be controlled, at least in part, by [AMP]. Independent evidence for normally extremely low cytosolic AMP levels comes from digitonin fractionation of isolated rat cardiomyocytes (20); here the cytosolic fraction did not contain measurable amounts of AMP under the conditions of nor-



Fig. 2. Correlation between purine nucleoside release and lactate release in isolated working guineapig heart stimulated by norepinephrine (NE) in the presence of 1.5 mEq/L Ca<sup>2+</sup> under conditions of normoxia.  $P_v = 12 \text{ cmH}_2\text{O}$ ;  $P_a = 90 \text{ cmH}_2\text{O}$ . Data are means  $\pm$  S.E., n = 6. MVO<sub>2</sub>: myocardial oxygen consumption;  $R_{cor}$ : coronary resistance. An intraventricular needle was introduced through the cardiac apex to measure ventricular dP/dt<sub>max</sub>: control = 1250  $\pm$  85 mmHg/s; NE = 2833  $\pm$  228 mmHg/s.

moxia; cytosolic AMP became measurable only during anoxia. Furthermore, results from two independent series of experiments in working guinea-pig hearts, one used to determine the relationship between  $MVO_2$  and calculated [AMP] (6), the other to determine the relationship between  $MVO_2$  and adenosine plus inosine release (6,10), allowed us to derive an operational formula that characterized the steady-state correlation between [AMP] and measured adenosine plus inosine release:

# (Eq. 8) $V(AR + INO) = 0.0027 [AMP]^{1.49}$ (ref. 6)

where V(AR + INO) = adenosine plus inosine release (nmol/min per g dry wt.) and [AMP] designates free cytosolic AMP concentrations (nM). Measured and predicted V(AR + INO), using the [AMP] data from Table 1, show reasonable agreements (Table 1), supporting the concept that cytosolic AMP availability may normally be an important determinant of the rate of purine nucleoside formation and release in working guinea-pig heart. It should be emphasized that Eq. 8 was derived from predominantly myocytic, not endothelial parameters (MVO<sub>2</sub>, ATP content, [CrP]/[Cr], [ATP]/[ADP], intracellular pH), i.e., [AMP] refers to free AMP in the myocyte cytosol.

It is clear that most (although less than 100%) of cardiac LDH, GAPDH and PGK is located in the cytoplasm of the myocytes, which make up about 85% of the myocardial wet weight (31). It thus seems reasonable to assume that most of the present effect of pyruvate/lactate infusion on inosine release originated in the free cytosolic AMP of the myocytes. The possible inosine release from endothelial cells, which comprise only about 2% of myocardial wet weight (51), is difficult to assess under the present conditions. However, dipyridamole blocked coronary venous uric acid output without inhibiting inosine release (4), and isolated heart cells release purine nucleosides in appreciable amounts (17,20,33). Taken with these reports, the present response of inosine release to cytosolic redox alterations further supports the AMP-availability model for the working heart. It is then not surprising that adenosine plus inosine release predicted from Eq. 8 and the measured adenosine plus inosine release were in reasonable agreement, both during the present redox interventions (Table 1) and during steady state low-flow ischemia (6). However, a potential complication is the fact that pyruvate can accelerate while lactate appears to decelerate hypoxanthine salvage in cardiomyocytes in culture (44). Such pyruvate stimulation of the salvage pathway via direct oxidation of cytosolic NADH cannot be ruled out with certainty under the present conditions.

In theory, the present coronary venous inosine fraction could also have been derived from cytosolic IMP generated from AMP by adenylate deaminase. However, several observations argue against this interpretation: 1) most of cardiac IMP appears to be located in the mitochondria rather than in the cytosol under the conditions of normoxia (20); 2) the thermodynamic AMP concentration is several orders of magnitude below the  $K_{mAMP}$  of adenylate deaminase; 3) guinea-pig myocardium does not appear to contain appreciable quantities of adenylate deaminase (16); and 4) the soluble 5'-nucleotidase has a higher affinity towards AMP than IMP (13,16).

Cytoplasmic 5'-nucleotidase accounts for only 3-17% of total 5'-nucleotidase activity in guinea-pig myocardium (16,47). The enzyme appears to be subcompartmented between the free cytosol and lysosomes (2,13). Nevertheless, several reports strongly support intracellular formation of adenosine (17,33,47,56). Uncertainty prevails regarding the subparenchymal site(s) of adenosine deaminase (16,47). Up to 30% of total cardiac adenosine deaminase may be localized in the endothelial compartment with only a small fraction (about 5%) in the myocytes (16). However, dipyridamole as an inhibitor of the endothelial nucleoside transporter did not appreciably block coronary venous inosine output (4); this seems to imply that inosine export from extramyocytic elements is of only minor importance in the beating heart. Free Cardiac Adenylates and Purine Nucleosides

Consequently, the presently released inosine fraction could actually reflect, at least in part, myocytic adenosine deaminase activity. Although the possibility has been raised that adenosine is exclusively derived from the endothelium in the resting normoxic heart (35), dipyridamole greatly increased (not decreased) cardiac adenosine output (about 20-fold), whereas uric acid release was nearly completely blocked (4). These data obviously imply that the beating myocytes can generate free interstitial adenosine which is readily taken up by and degraded within endothelial structures. However, this adenosine is also available for washout via the coronary outflow tract (1,4). Alternatively, based on studies with special isolated myocyte and endothelial preparations, it was recently emphasized (16) that guinea-pig myocytes normally release only minimum, if any, amounts of free adenosine and inosine. Instead myocytic AMP and other adenylates are released to undergo subsequent dephosphorylation by ecto-nucleotidases of various subparenchymal origins (18,53).

#### Effects of Excess Pyruvate vs. Lactate on Coronary Venous Purine Release during Global Low-Flow Ischemia

During states of acute myocardial oxygen- or flow-deficiency, ATP breakdown is most pronounced in the working myocytes; it is this cell type which by far predominates in cardiac parenchyma and which, due to its hydraulic workload, necessarily incurs the greatest immediate energy deficit during ischemia conditions. It is not known whether other formed elements of the myocardium (endothelial and epicardial cells, interstitial fibroblasts, leukocytes, pericytes, smooth muscle cells) develop energy deficiencies during ischemic or hypoxic perfusion. However, it appears reasonable to assume that, relative to the working myocytes, the extent and time course of any extramyocytic energy imbalance is likely to be minor and delayed. Obviously, cardiac ischemia causes ATP degradation, accumulations of lactate, NADH, and free AMP, and intracellular acidification mainly in the myocyte.

If released adenosine plus inosine are primarily derived from free cytosolic AMP of the myocytes, especially during states of acute myocardial ischemia, and if [AMP] is actually under the control, at least in part, of the cytosolic phosphorylation potential and hence linked to the NAD<sup>+</sup>-NADH oxidation-reduction state (Table 1), ischemic purine nucleoside release would be expected to respond accordingly to an imposed (i.e., "clamped") redox state in the form of a constant arterial [Pyr]/[Lac] ratio.

We tested this rationale in a special low-flow, globally ischemic, isolated working guinea-pig heart in which coronary hypoperfusion was combined with maximum norepinephrine stimulation ("NE-ischemia") to produce a substantial rather than mild energy deficit of the myocytes. Hearts received as substrate 5 mM glucose or 5 mM glucose plus 5 mM lactate or 5 mM glucose plus 5 mM lactate plus 5 mM pyruvate. Such excess concentrations of pyruvate or lactate substantially elevate the intracellular [Pyr] and/or [Lac]. We also showed that pyruvate, when applied in a concentration of 2 mM in the presence of 5 mM lactate is oxidized only during normoxia, but released during states of oxygen deficiency (12). In addition, under low-flow ischemic conditions, external 2 mM pyruvate produces a large increase (about 12-fold) in cellular [Pyr] in association with a relative oxidation of the cytosolic NADH-NAD<sup>+</sup> system as evidenced by increased cellular [Pyr]/[Lac] or [dihydroxyacetone phosphate]/[alpha-glycerophosphate] ratios (data not shown). Consequently, experimental clamping of the external pyruvate-lactate oxidation-reduction state in the globally ischemic working heart.

Figure 1 shows that those ischemic working hearts which received glucose alone or in combination with lactate released larger amounts of purines during the first 14 min of the NE-ischemia, as compared to those hearts which received excess pyruvate (arterial [Pyr]/[Lac] = 1.0). The data also show that the major portion (75-80%) of the ischemically released purines was in the form of nucleosides (adenosine plus inosine) and the remainder in the form of purine bases (xanthine compounds) plus uric acid. The decrease in purine release in the presence of the externally fixed [Pyr]/[Lac] ratio was not due to differences in residual ischemic cardiac pressure work, since the heart rate - pressure products were comparable in the absence and presence of pyruvate and/or lactate (see legend to Fig. 1). In the steady state after about 45 min NE-ischemia, purine release from the pyruvate- plus lactate-perfused hearts was only marginally smaller than that from hearts receiving glucose plus lactate. These findings thus indicated that external "clamping" or buffering of the cytosolic [NADH]/[NAD<sup>+</sup>] ratio can significantly attenuate purine release, especially during short-term non-steady state ischemia conditions. Figure 1 also shows that ischemic stress, if extended for 45 min or more, is associated with an only transitory excess release in adenylate catabolites, i.e., only during early ischemia. A similar time-dependence of adenosine plus inosine release was recently reported in high-flow hypoxic working hearts (12).

The present data show unequivocally that the purine nucleosides, not the free purine bases plus uric acid, are the major adenylate catabolites during severe acute ischemic stress of the perfused working heart. Uric acid accounted for only 4% to 10% of net purine release during the first 14 min of the NE-ischemia (Fig. 1). However, uric acid output accounted for 41% to 52% of total purine release during normoxia, although after 45 min NE-ischemia only 9% to 19% of total purine release was in the form of uric acid. The situation is different in some Langendorff hearts (1,4); here uric acid release comprises more than 70% of total purine output during normoxia and also during 1 min reperfusion after a brief period (30 s) of zero-flow ischemia (1). On the other hand, the Langendorff heart, when tested under highflow anoxia, releases adenosine plus inosine, not the free purine bases plus uric acid, as the major adenylate catabolites (1,4). Apparently, under conditions with minimum myocyte energy deficit, much of the relatively small amount of adenylate catabolite(s) can still be fully converted to uric acid due to the active purine metabolism within the endothelium. However, when the myocyte energy deficit heatometabolism compared and a plus income formation appears to every helm the becomes severe, cardiac adenosine plus inosine formation appears to overwhelm the metabolic capacity of the endothelium resulting in an increased appearance of nucleosides in the coronary effluent. Ultimately the interstitial nucleoside concentration may rise to nearly saturate binding, transport, and receptor sites, which would lead to an increase in the intracoronary ratio of [free nucleosides]/[free bases plus uric acid] with the uric acid output reaching a maximum (e.g., Fig. 1 and ref. 1). In fact, myocardial output of free adenosine can experimentally be increased at the expense of uric acid release when dipyridamole is applied as an adenosine transport blocker of the endothelium (4). A finite purine transport and degradation capacity of the endothelium is also consistent with Eq. 8, according to which adenosine plus inosine release increases more than proportionally with free cytosolic [AMP].

Figure 2 shows that adenosine plus inosine output and lactate release change concordantly in working hearts stimulated by a high dose of norepinephrine under the conditions of normoxia. Since these hearts received 15 mM glucose as the sole substrate, lactate release was a qualitative index of the cytosolic [NADH]/[NAD<sup>+</sup>] ratio (5,6). In other words, cytosolic reduction potential of the working myocytes and coronary venous purine nucleoside output changed in parallel (6) when the phosphorylation potential decreased due to a physiologic adrenergic stimulus (10). In fact, according to Eqs. 6 and 7, the increased lactate output reflected an increase in the cytosolic [NADH]/[NAD<sup>+</sup>] ratio in association with a fall in the phosphorylation potential. Consequently, [AMP] was probably increased during

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adrenergic stimulation due to the fall in the phosphorylation potential (6,10), most likely stimulating 5'-nucleotidase flux according to the AMP-availability model. Since adenosine and inosine appeared in the coronary outflow tract in increasing amounts, a significant fraction of myocytic adenosine plus inosine probably bypassed endothelial nucleoside transport sites. This argument is valid if one assumes that adrenergic stimulation does not appreciably alter the energy balance of the extramyocytic elements of the working heart. Indeed, increased oxygen consumption (a predominantly myocytic parameter) is associated with increased adenosine plus inosine release, regardless of whether oxidative metabolism is stimulated by, e.g., acetate (1,57) or by infusion of adrenergic agonists (6,10). Since increased oxygen consumption of perfused heart is normally associated with a fall in cytosolic phosphorylation potential (see ref. 10 and refs. therein), it appears likely that [AMP] was increased under the same conditions (6,10).

It should finally be pointed out that such energy-linked adaptations in the free concentrations of cytosolic adenylates and coronary purine nucleoside output do not necessarily identify the site(s) of primary AMP degradation. A functional coupling between cytosolic AMP degradation at the inner surface of the sarcolemma followed by vectorial adenosine export (18) is also not necessarily excluded. It appears to be important to develop an experimentally testable hypothesis to differentiate between intra- and extracellular sites of endogenous AMP metabolism in the physiologically performing heart.

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# Chapter 8

# **Regulation of Myocardial Perfusion by Purine Metabolites**

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This review of the roles of cardiac purines in the regulation of myocardial perfusion begins with brief descriptions of the general characteristics of models of the metabolic control of tissue blood flow and of cellular energy potential. The body of the review, an analysis of the evidence for the participation of adenosine in coronary regulation, is divided into four sections: 1) the enzymatic synthesis of adenosine and the coupling of production to cardiac energy state; 2) the regulation of adenosine concentration at the coronary resistance vessels; 3) the mechanism of adenosine coronary vasodilation; and 4) the relationship of adenosine to other vasodilatory metabolites and mechanisms that might participate in coronary flow regulation. Two recent reviews provide additional information about the role of adenosine in the metabolic control of coronary flow (43,57).

# The Metabolic Control of Coronary Perfusion

The coronary circulation is a vascular bed primarily under metabolic control, that is, one in which tissue perfusion varies according to changes in tissue energy state. The notion that energy production depends on the supply of some rate-limiting metabolite is implicit in such a definition, as is the concept of a cellular energy potential. The tight coupling of coronary flow to cardiac metabolism (4) stems from attributes of both the cardiac striated and coronary smooth muscle cells. Working cardiocytes rely almost entirely on aerobic metabolism for the large amounts of energy needed to support their continuous activity. Because the blood concentrations of energy-yielding substrates such as glucose or organic acids are relatively high and the heart is unselective as to the source of reducing equivalents, oxygen is the limiting substrate; cardiac performance begins to deteriorate within a few heartbeats after the interruption of oxygen delivery by coronary occlusion (44). Coronary vascular resistance is high relative to the rate of substrate usage, the results of some poorly understood influence on - or perhaps an intrinsic property of coronary smooth muscle cells. Consequently, the transcoronary extraction of oxygen is high, typically 60-80 percent under basal conditions. Since the potential for a further increase in oxygen extraction is limited, a change in coronary flow is the primary response to the change in the rate of cardiac energy usage.

Models of metabolic coronary vasoregulation have three fundamental characteristics. The first is a vasodilatory chemical whose concentration is coupled to cardiac energy state. The same working cardiocytes that generate the demand for energy-yielding substrates also produce the vasoregulatory metabolite; thus, flow is said to be under local control. The second characteristic is a mechanism whereby the endogenous vasoregulator influences the contractile state of the smooth muscle cells in the coronary resistance vessels. A third characteristic of such models is that control lies in the direction of vasodilation. Although vasoconstriction may determine perfusion rate under pathological control of coronary flow consists of

dynamic metabolic vasodilation superimposed on a tonic constriction of unknown origin.

#### **Cardiac Energy State**

The energy state of a cell is a chemical potential that reflects the balance between energy generated through cellular metabolism and the consumption of that energy by cellular functions such as contraction, maintenance of a membrane potential, secretion and so forth. In "aerobic" organs such as the heart, energy metabolism consists of two sequential steps. The first is the conversion into reducing equivalents, mainly NADH, of the energy in the electrons liberated by oxidation of substrate. The second step is the orderly transfer of these electrons into mitochondria and conservation of the energy as the ATP generated by oxidative phosphorylation. Either of these steps can be the basis for a formal, thermodynamically correct statement of the cellular energy state. One is the redox potential, [NADH][H<sup>+</sup>]/[NAD<sup>+</sup>], which is based on the concentrations of reducing equivalents. This potential is not uniform throughout the cell, mitochondria being more "reduced" than the cytoplasm (61). However, the cytosolic redox potential is still a useful indicator of energy state and, because it is coupled to the reaction catalyzed by lactic dehydrogenase, can be judged by the lactate/pyruvate ratio. The phosphorylation state of ATP in the cytosol is another indicator of cellular energy state (60). The two steps in glycolysis that are catalyzed by glyceraldehyde phosphate dehydrogenase and phosphoglycerate kinase couple the ATP potential, defined as [ATP]<sub>free</sub>/[ADP]<sub>free</sub>, to the redox potential (Fig. 1):

GAPDH  $GAP + NAD^{+} + Pi \xrightarrow{\text{GAPDH}} NADH + H^{+} + 1,3 \cdot DPG$   $1,3 \cdot DPG + ADP \xrightarrow{\text{PGK}} 3 \cdot PG + ATP$   $\frac{[ATP]_{free}}{[ADP]_{free} [Pi]} = \frac{[GAP]}{[3 \cdot PG]} \cdot \frac{[NAD^{+}]}{[NADH] [H^{+}]} \cdot K_{GADPH} \cdot K_{PGK}$ 

Fig. 1. Coupling of ATP potential to redox potential. 1,3-DPG = 1,3-diphosphoglycerate; GAP = glyceraldehyde-3-phosphate; GAPDH = glyceraldehyde phosphate dehydrogenase; 3-PG = 3-phosphoglycerate; PGK = phosphoglycerate kinase.

The subscript "free" in each term of the ATP potential emphasizes that the concentrations of these reactants refer to those in solution in cytosolic water rather than the total content in tissue. The ADP pool of heart muscle is highly compartmentalized, mainly through binding to actomyosin (49); ADP thus sequestered cannot be in equilibrium with ATP and Pi, which exist mainly in free solution, so calculations based on total tissue ADP are meaningless.

Although the redox and the ATP potentials are equivalent as indices of energy state, the ATP potential is perhaps preferable because ATP is the immediate source of energy for cellular function. At present it is not possible to measure  $[ADP]_{free}$  directly; however, the creatine phosphate potential, [CrP]/[Cr][Pi], serves as an index of the ATP potential in much the same way as the lactate/pyruvate ratio serves as an index of redox potential.

## Production of Adenosine and Coupling to Cardiac Energy State

The hydrolysis of either of two substrates could account for adenosine production in heart muscle, that of AMP by 5'-nucleotidase or that of S-adenosylhomocysteine by S-adenosylhomocysteine hydrolase (Fig. 2):

 $AMP + H_2O \xrightarrow{5'-NP} Pi + Ado$ 

AdoHcy +  $H_2O_{4}$  Ado + Hcy

Fig. 2. Two pathways of adenosine production in heart muscle. Ado = adenosine; AdoHcy = S-adenosylhomocysteine; Hcy = homocysteine; 5'-NP = 5'-nucleotidase; SAH = S-adenosylhomocysteine hydrolase.

## **Cardiac 5'-Nucleotidases**

Studies employing enzyme histochemistry and cell fractionation show that AMP phosphohydrolase activity resides in both the sarcolemma and the cytosol of cardiac muscle cells (11,39,52). Several lines of evidence exclude the sarcolemmal enzyme as the source of adenosine for coronary vasoregulation. This 5'-nucleotidase is an *ecto* enzyme, that is, its catalytic site is located on the external surface of the cardiac cell (18). There has never been a satisfactory explanation for how its substrate, AMP in the interior of the cell, might traverse the plasma membrane to gain access to the site of hydrolysis. ADP is a powerful inhibitor of the *ecto* nucleotidase (9); accordingly, it is not possible to reconcile the accelerated rate of adenosine production during hypoxia (2,19) and ischemia (42) with the rise in ADP levels that occur under such conditions and which should inhibit the enzyme. The most convincing evidence, however, is from studies showing that the rate of adenosine unaffected by essentially complete inhibition of the *ecto* nucleotidase by specific antibodies, by nucleoside phosphonates or by lectins such as concanavalin A (40,62).

New information about the properties of the 5'-nucleotidase in the cardiac cytosol (11,41) and about the compartmentalization of the AMP pool of heart muscle (7,8,56) support a plausible model of adenosine production coupled to cardiac energy state (43). AMP is the preferred substrate of the cytosolic nucleotidase of rabbit heart. ATP, at the low millimolar concentrations that occur within a cell, fully activates the enzyme by increasing its affinity for substrate, the  $K_{app}$  for AMP, averaging 2.5 mM (11,41). ADP also activates the nucleotidase when the AMP concentration is low. Purification of the enzyme abolishes regulation by ATP, evidence that the enzyme in unfractionated extracts of ventricular muscle from rabbit or guinea pig is about 300 umole/min per g wet wt., more than enough to account for adenosine production at the accelerated rates, up to 1 umole/min per g wet wt., that have been measured in ischemic heart muscle (41).

The model posits that ATP potential controls adenosine production by regulating the availability of substrate, rather than by modifying the catalytic properties of the cytosolic 5'-nucleotidase. As pointed out above, the cytosolic concentration of ATP is sufficient to fully activate the enzyme and, owing to the buffering effect of the creatine phosphate system, it changes only modestly even under such extreme con-

ditions as hypoxia. Accordingly, the enzyme probably remains fully activated at all times and adenosine production varies according to substrate availability. Very nearly the entire cardiac AMP pool is intramitochondrial (8,56); the small fraction in the cytosol probably arises from the dismutation of ADP catalyzed by myokinase (Fig. 3). Cytosolic AMP concentrations are in the submicromolar range, vary as an inverse function of the ATP potential and, according to the myokinase equilibrium, in proportion to the square root of [ADP]<sub>free</sub>, between about 50 and 750 nM (7). Because such substrate concentrations are three to four orders of magnitude below the  $K_{app}$  of the activated cytosolic nucleotidase, any change in AMP level produces a proportional change in the rate of adenosine production.

#### S-Adenosylhomocysteine Hydrolase (SAH)

SAH has one clearly established function, the degradation of the Sadenosylhomocysteine that results from and is a very potent competitve inhibitor of the S-adenosylmethionine-dependent regulation of such diverse substrates as biogenic amines, proteins and membrane phospholipids (10). In most eukaryotic cells the hydrolase also reversibly binds adenosine (24,58). Adenosine sequestered in this intracellular compartment accounts for very nearly the entire cardiac pool of 1-2 nmole/g wet wt. (46), but whether it has a physiological role is unknown. Heart muscle also contains a pool of S-adenosylhomocysteine, somewhat smaller than the adenosine pool, about 0.2-0.6 nmole/g wet wt. Studies of the turnover of this pool in isolated, perfused rodent hearts suggest that adenosine formation from Sadenosylhomocysteine is somewhat less than 1 nmole/min per g under normoxic conditions and less than twice that rate during hypoxia. Since adenosine formation during hypoxia is so much greater, the transmethylation pathway may be a significant source of adenosine only under basal conditions (1,33).

One encounters important problems in trying to account for the metabolic control of coronary flow in terms of the SAH model. First, the enzyme does not appear to be under the control of cardiac energy state. SAH contains NAD<sup>+</sup> (47), but it is very tightly bound and there is no evidence for coupling to the cytosolic pyridine nucleotides that determine redox state. Second, the poise of the equilibrium catalyzed by SAH lies strongly in the direction of synthesis, and even in the absence of homocysteine a high affinity for adenosine promotes retention of the nucleoside in the SAH  $\cdot$  adenosine complex (58). A physiological stimulus to the dissociation of this complex is not known. Trapping adenosine by a second enzymat-



Fig. 3. Coupling of adenosine production to the ATP potential. Abbreviations are: Ado, adenosine; MK, myokinase; and 5'NP, 5'-nucleotidase. The (+) denotes ATP activation of the cytosolic 5'-nucleotidase. See text for discussion.

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ic reaction is, of course, one way to effect dissociation of the SAH  $\cdot$  adenosine complex, but the net result is not the production of adenosine but rather the product of the trapping reaction. Third, since the adenosine produced from transmethylation may have physiological significance only under basal conditions, its significance diminishes further in light of reports that basal coronary resistance is insensitive to either the destruction of adenosine receptors by dialkylxanthines, evidence that adenosine plays no role in setting basal coronary tone (16,20,22,28,45,53,54).

# **Regulation of Cardiac Interstitial Adenosine Concentration**

Since the coronary resistance vessels lie in the interstitial space of the heart, the concentration of adenosine in this anatomical compartment is the independent variable in those tests of the adenosine hypothesis that examine the dose-response relationship between adenosine and coronary resistance. In terms of mass balance, this concentration is the resultant of adenosine release from the cells that constitute the boundary of the interstitial space, the re-uptake of adenosine by these same cells, washout into the coronary venous drainage as well as relatively minor losses via the cardiac lymphatics and into the pericardial space. A large literature documents the movement of adenosine in either direction across cellular membranes by a combination of carrier-mediated and simple diffusion. The cardiac interstitium is not accessible to direct sampling of labile metabolites such as adenosine, a fact that greatly complicates studies of the control of adenosine concentration in this compartment. To appear in the coronary venous effluent, adenosine must first pass the endothelial barrier, the cells of which have an extremely active purine metabolism (39,48). Endothelial cells in culture not only take up adenosine avidly, but also release significant amounts of this and other purines (29,38). Perhaps as much as one-sixth of the cardiac purine efflux originates in the coronary endothelium rather than in cardiac myocytes (13). Likewise, the adenosine appearing in an epicardial transudate (17,23) or in a pericardial superfusate (21,51) will have passed through a layer of epicardial mesothelial cells. Although cardiac lymphatics drain the interstitium directly, the flow rate through these channels is so slow that the adenosine concentration in lymph probably reflects metabolism by the lymphatic endothelium rather than that in the interstitium.

## **Coronary Adenosine Receptor and Coupling to Effectors**

The identity of the coronary artery adenosine receptor is fairly certain but that of the systems(s) through which receptor activation leads to coronary relaxation is not. Low affinity A2 (or Ra) receptors that stimulate adenylate cyclase appear to mediate coronary vasodilation by adenosine. The several lines of evidence supporting such an identification are: 1) the coronary receptor is located on the myocyte surface; 2) alkylxanthines such as theophylline competitively antagonize adenosine coronary vasodilation; 3) forskolin potentiates adenosine vasodilation; 4) the coronary vasoactivity ranking of receptor-selective analogues resembles that of A2 receptors; and 5) adenosine stimulates cyclic AMP accumulation and relaxation of coronary artery strips and in suspensions of coronary microvessels (43). The functional coupling of the coronary adenosine receptor to adenylate cyclase implies that receptor activation initiates protein phosphorylation by cyclic AMP-dependent protein kinase (A-kinase), but the identity of the substrate(s) is uncertain. A kinase can phosphorylate and thereby inactivate myosin light chain kinase (14,55); other possible substrates that could influence smooth muscle contractile state are proteins in the sarcolemma (26) or in intracellular organelles that sequester the Ca<sup>2+</sup>- supporting smooth muscle contraction (59).

Two recent reports raise the possibility that adenosine causes coronary relaxation through receptors other than the A2 adenosine receptor and/or effectors other than

A-kinase. The first paper (15) describes the inhibition of the phosphatidylinositol kinase (PI-kinase) of aortic smooth muscle by adenosine and several analogues modified in the ribose, but not the purine, that compete with ATP for binding to the catalytic site. The  $K_i$  of 5'-chloro-5'-deoxyadenosine, a metabolically inert analogue, is much lower than the  $K_m$  for ATP. Cell fractionation studies show that PI-kinase purifies along with markers of sarcoplasmic reticulum, suggesting that inhibiting the formation of a precursor of phosphoinositol-4,5-diphosphate, a Ca<sup>2+</sup> ionophore, could prevent the release of Ca<sup>2+</sup> from intracellular stores (6). The inhibition of coronary smooth muscle PI-kinase by adenosine could well be physiologically important, but it is not a mechanism for the *metabolic* control of coronary perfusion, because adenosine production occurs inside the coronary myocyte and thus is not coupled to the cytosolic ATP potential of the working cardiocytes.

The second report describes evidence that A1 receptors coupled to guanylate cyclase mediate adenosine vasodilation, a model exactly opposite the conventional model of A2 receptors coupled to adenylate cyclase. In aortic smooth muscle cells in culture, adenosine and two analogues, N<sup>6</sup>-cyclohexyladenosine (CHA) and N-ethyl adenosine-5'-uronamide (NECA), at concentrations between 10<sup>-9</sup> and 10<sup>-4</sup> M, promote the concentration-dependent 2- to 2.5-fold accumulation of cyclic GMP. Activity in the nanomolar range, an agonist potency ranking of CHA > NECA > adenosine, and antagonism by theophylline suggest that A1 adenosine receptors mediate this response. Parallel studies of the guanylate cyclases of cultured aortic smooth muscle cells and of freshly isolated media of rat aorta and also cow coronary artery show that the three adenosines stimulate the particulate, but not the soluble, guanylate cyclases of these vascular muscles (30). This report is intriguing, because it suggest that adenosine might act through the same cyclic GMP-activated protein phosphorylation cascade as, for example, the endothelium-dependent vasodilators (50). Interpretation is not straightforward, however, for these observations do not correspond to the intact heart, where the coronary vasoactivity of NECA is two orders of magnitude greater than that of CHA (31).

## **Relationship of Adenosine to other Endogenous Vasodilators**

It is doubtful that adenosine is the sole mediator in the metabolic control of coronary blood flow. Models in which there is a single mediator of flow posit an unique quantitative relationship between agonist concentration and coronary flow, that is, one that is identical under widely differing experimental conditions (3). A number of studies purport to show a correlation between coronary flow and adenosine concentration, as reflected by one or another of the indirect indices of adenosine concentration in the interstitial space of the heart (5). While this sort of evidence is certainly consistent with the participation of adenosine in flow regulation, it is not robust enough to establish adenosine as the sole mediator at the exclusion of other endogenous regulatory substances. For want of a technique for d'ect measurements of adenosine in the interstitial space to serve as a Gold Standard, the validity of all the indirect methods is uncertain and the inferences drawn from such data are necessarily conditional. Further, random errors in the estimates of adenosine concentration are large enough to obscure significant differences in the relationship between flow and concentration under different experimental conditions.

Two additional lines of evidence argue more directly against an unique relationship between coronary flow and interstitial adenosine concentration. The first line consists of two studies of adenosine release into the coronary effluent. Owing to adenosine release from coronary endothelial cells (38), the adenosine concentration in the coronary veins only reflects directional changes in interstitial concentration. Even so, the experiments suggest, respectively, that adenosine production may depend on the nature of the stimulus and that adenosine may only initiate Myocardial Perfusion and Purine Metabolites

hyperemic responses, which are then sustained by other agents or mechanisms. In open-chest dog preparations, raising coronary flow by means of beta-adrenergic stimulation elicits adenosine release, whereas electrical pacing to the same level of cardiac performance and coronary flow does not cause adenosine release (34). In isolated guinea-pig hearts, the infusion of norepinephrine or perfusion with a highcalcium buffer raises oxygen consumption and coronary flow to new steady-state levels that are stable over many minutes. These interventions also evoke adenosine release, but this is only transitory, reaching a peak within a minute and declining steadily thereafter toward control (12). To the extent that coronary venous adenosine reflects the interstitial concentration, the changing dose-response relationship between adenosine and coronary flow suggests the participation of other vasodilators.

The second line of evidence against an unique dependance of coronary flow on adenosine concentration comes from experiments employing specific adenosine receptor blockade by dialkylxanthines such as theophylline or the destruction of endogenous adenosine by intracoronary adenosine deaminase. These interventions reduce but do not abolish the reactive hyperemia response to ischemia (53), to beta-adrenergic stimulation (45,27), or to hypoxia (36), but have no effect on basal coronary flow or on the autoregulatory response to a change in coronary perfusion pressure (16,20,22).

One can trace a causal relationship between the release of two endogenous vasodilators, carbon dioxide and potassium, and the cytosolic ATP potential. The ATP potential controls the sarcolemmal ATPase that mediates the exchange of Na<sup>+</sup> for K<sup>+</sup> and, consequently, of K<sup>+</sup> release. So, too, the ATP potential regulates the rate of the oxidative substrate metabolism that generates carbon dioxide. Like adenosine, however, neither potassium nor carbon dioxide alone seems able to account for metabolic coronary flow control. Potassium lacks the potency to account for more than a fraction of coronary flow responses (37). The relationship between carbon dioxide production and oxidative metabolism, as defined by the respiratory quotient, varies according to the type of substrate. In other words, there is not an unique relationship between vasodilator release and cardiac energy state. It might well be that adenosine, potassium and carbon dioxide act in concert to control coronary flow, just as these three metabolites regulate cerebral blood flow (32). To complicate matters still further, the discovery that a high coronary flow rate by itself can cause endothelium-dependent vasodilation (35) suggests that even hyperemic responses that begin as metabolically-linked vasodilation might be sustained, in part, by non-metabolic, flow-linked vasodilation (25).

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# Chapter 9

# **Cardiac Electrophysiological Effects of Adenosine**

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Adenosine, in addition to its well-established vasodilator actions, also modulates a) pacemaker activity, b) atrioventricular (AV) transmission, c) atrial contractility, and d) the myocardial actions of catecholamines. Based on pharmacological response and mechanism of action, the above cardiac effects of adenosine can be tentatively subdivided into two categories, i.e., non-cAMP-mediated effects which appear to be due to an activation of  $K^+$  channels, and cAMP-dependent effects characterized by a modulation of calcium inward current. In working ventricular myocardium, only the cAMP-dependent mechanism appears to be operative, whereas in the atrial myocardium and specialized tissues [e.g., sinoatrial (SA) and AV node], both the cAMP-independent and cAMP-dependent mechanisms are operative. In the SA node, adenosine causes sinus slowing and a shift in location of the primary pacemaker site. In atrial myocytes, adenosine depresses contractility and shortens the action potential; in the AV node, it depresses the nodal (N-cell) action potential which is accompanied by prolongation of AV conduction time. In ventricular myocardium, adenosine antagonizes the stimulatory effect of agents that stimulate adenylate cyclase activity but has no effect in their absence. In summary, adenosine can affect cardiac function at several levels by different mechanisms and may have greater physiological importance than previously recognized.

# Introduction

In addition to its well-established vasodilator effects, the nucleoside adenosine has many other cardiac actions. Adenosine has a depressant effect on a) sinoatrial (SA) and atrioventricular (AV) node function (i.e., negative chronotropic and dromotropic action, respectively), b) atrial contractility (negative inotropic effect), and c) the cardiac stimulatory actions of catecholamines. Based on pharmacological response and mechanism of action, the above cardiac effects of adenosine can be tentatively subdivided into two categories: cAMP-independent and cAMP-dependent (Fig. 1). The cAMP-independent effects appear to be due to an increase in K<sup>+</sup>-conductance (e.g., hyperpolarization of SA-node cells). On the other hand, the cAMP-dependent effects appear to be the result of an inhibition of adenylate cyclase. They are characterized by an attenuation of the catecholamine-enhanced calcium inward current (e.g., antagonism of the positive inotropic and arrhythmogenic effect of isoproterenol). In working ventricular myocardium, only the cAMP-dependent mechanism appears to be operative, whereas in the atrial myocardium and specialized tissues (e.g., SA and AV node), both the direct and indirect mechanisms are operative (Fig. 1).

Current evidence indicates that all of the aforementioned actions of adenosine are a) mediated by extracellular receptors - subtype  $A_1$  (or  $R_1$ ), b) potentiated by nucleoside transport blockers (dipyridamole) and adenosine deaminase inhibitors, e.g., erythro-6-amino-9-(2-hydroxy-3-nonyl)purine, and c) antagonized competitively by alkylxanthines (such as theophylline) and abolished by adenosine deaminase. In summary, the adenosine system consists of three components: a receptor, a mechanism for production of adenosine, and a mechanism for its removal. The intent of this review is to describe the more general features of the effects of adenosine on the cardiac functions listed above. In addition, the present state of knowledge regarding the mechanism of action of adenosine in various cardiac cell types, as well as the possible role of adenosine as a modulator of certain rhythm disturbances associated with myocardial hypoxia/ischemia will be discussed.

## I. Chronotropic Effect (SA Node)

Adenosine, ATP and adenine derivatives (e.g., ADP, AMP) exert a negative chronotropic effect on the sinoatrial node. This effect of the nucleoside has been demonstrated in a variety of isolated preparations from mammalian (7,13,51,56,57) and non-mammalian species (25) as well as in intact anesthetized (7,22,34,45,54) and awake animals (29). Recently, sinus slowing and arrest was shown in humans after intravenous injection of adenosine (19). As illustrated in Fig. 2, adenosine, in addition to slowing sinus rate, causes pacemaker shift, i.e., in the presence of adenosine the earliest site of activation shifts from the SA node to the right atria (56). At high concentrations adenosine also causes sinoatrial exit block (56). Similar to its many other actions, the depressant effect of adenosine in pacemakers (SA, atrial or ventricular) can be antagonized by methylxanthines (23,28,45) and potentiated by nucleoside transport inhibitors (13,25,37,45,56). As shown in Fig. 3, the negative chronotropic effect of adenosine in atrial and His-bundle pacemakers are antagonized by aminophylline (Fig. 3, panel A) but enhanced by dipyridamole (panel B). From this figure it can be seen, as has been reported previously (7,51), that in the guinea-pig heart the His-bundle pacemaker is more sensitive to adenosine than atrial pacemaker, i.e.,  $IC_{50}$  of 1 x 10-5 and 2 x 10-4 M, respectively.

Adenosine-induced sinus slowing in frog sinus venosus (25) and isolated rabbit SA node (56) is associated with a hyperpolarization to about the K<sup>+</sup>-equilibrium potential ( $E_{\rm K}$ ). Like the sinus slowing, the adenosine-induced hyperpolarization can be reversed by aminophylline (57). It is worth noting that the magnitude of hyperpolarization caused by adenosine correlates with the degree of slowing (57), i.e., the greater the hyperpolarization the greater the sinus slowing. Since hyperpolarization of pacemaker cells causes slowing of pacemaker firing (11), the above findings suggest that the mechanism of the adenosine-induced slowing is the same as the hyperpolarization. Because the hyperpolarization caused by adenosine is



Fig. 1. Cardiac actions of adenosine. Mechanisms involved in the cAMP-independent and cAMPdependent actions of adenosine on sinoatrial (SA), atrioventricular (AV) node, atrial and ventricular myocardium. The indirect actions of adenosine, which can be demonstrated only when cAMP levels have been elevated, seem to be mediated by an inhibitory effect on the adenylate cyclase. See text for discussion. g = conductance.

neither blocked by ouabain nor low concentrations (1-3 mM) of cesium chloride, it is unlikely that the adenosine-induced hyperpolarization is due to an effect on Na<sup>+</sup>/K<sup>+</sup> pump and/or i<sub>f</sub>, i.e., the time- and voltage-dependent inward current activated by hyperpolarization (-80 to -120 mV; ref. 57).

Recent studies (59) of the macroscopic and microscopic currents in single isolated SA-node cells have confirmed as well as extended the results obtained in the multicellular preparations. Adenosine caused a dose-dependent slowing of spontaneous rate and hyperpolarization in isolated nodal cells. In whole-cell voltage clamp experiments, adenosine caused an increase in background K<sup>+</sup> current, which was dose-dependent. Adenosine alone had no significant effect on the control (basal) calcium current  $i_{Ca}$  nor the hyperpolarization-activated inward current  $i_{f}$ . However, adenosine did attenuate isoproterenol-enhanced  $i_{Ca}$  and  $i_{f}$ . In studies of microscopic current in SA nodal cells, adenosine-activated channels had singel-channel conductance of  $8 \pm 2$  pA and showed significant inward rectification. The activation of the K<sup>+</sup> channel by adenosine, as has been shown in atrial cells (33,38), involves a GTP-dependent coupling protein, e.g.,  $G_{o}$ , mediating the coupling of the receptor and K<sup>+</sup> channel. The K<sup>+</sup> channel is thought to be independent of cAMP. Thus, in the modulation of pacemaker rate, adenosine activates K<sup>+</sup> channels and, in addition, during increased adrenergic tone, affects  $i_{Ca}$  and  $i_{f}$  through a cAMP-dependent mechanism.



Fig. 2. Adenosine (Ado)-induced sinus slowing and shift in pacemaker site in SA-node preparation. First two rows show same recording at two different sweep speeds. Ladder diagram depicts relationship between activation of SA node with respects to activation of crista terminalis during each condition. Impalement was maintained throughout control, adenosine and washout. A: Control. Cycle length = 460 milliseconds, activation of SAN precedes that of  $CT_1$  and  $CT_2$  by 44 ms. B: Adenosine, 1 x  $10^4$  M. Cycle length = 540 ms. Note change in configuration of SA-node action potential and that activation of  $CT_1$  and  $CT_2$  precedes that of SAN by 24 ms. C: Wash (3 min). Cycle length = 465 ms, SAN again leads  $CT_1$  and  $CT_2$  by 46 ms. SAN = intracellular recording from leading pacemaker cell.  $CT_1$  and  $CT_2$  = extracellular recording from upper and lower crista terminalis, respectively. Reproduced from ref. 56, with permission.

#### II. Dromotropic Effect (AV Node)

Since 1929, adenosine and related compounds have been known to cause heart block (22). In the 1970s and through the 1980s, numerous studies confirmed the observations that adenosine and adenine nucleotides (e.g., ATP) depress impulse conduction through the AV node in various preparations and species including humans (4,5,14,19,50,53). Although the depressant effect of adenosine in the AV node is a reproducible finding, the magnitude of the slowing of AV-node conduction varies greatly with the species and preparation (7,14,53). This negative dromotropic effect of adenosine is due to activation of a cell surface receptor that can be competitively blocked by alkylxanthines (2). Three key observations support the hypothesis that adenosine action in the AV node is mediated by a specific extracellular receptor, i.e., a) the effect of adenosine is enhanced by nucleoside transport inhibitors such as dipyridamole (2,50), b) adenosine analogs, too large to permeate the cell membrane, cause a similar or greater effect than does free adenosine (2), and c) the alkylxanthine 8-(p-sulfophenyl)theophylline, which has a polar subgroup that limits its cell permeability (28), is more potent than theophylline) to antagonize adenosine. Furthermore, the adenosine P-site analog 2'-deoxyadenosine has no dromotropic effect (2). Despite a large number of



Fig. 3. Attenuation (Panel A) and potentiation (Panel B) of the negative chronotropic effect of adenosine (ADO) on atrial and His-bundle rate by 30 uM aminophylline (AMINO) and 5 uM dipyridamole (DPM), respectively. The control (i.e., absence of adenosine) atrial and His-bundle rates were  $190 \pm 12$  and  $90 \pm 5$  beats/min, respectively. Numbers in parentheses indicate number of guineapig hearts studied. \*Significantly different from ADO alone (P<0.05).

reports demonstrating the negative dromotropic effect of adenosine, few or no attempts have been made to elucidate the mechanism and specific site of adenosine action responsible for its negative dromotropic effect. Previous studies (2,5), including those in human subjects (8,19), have shown that the AV block caused by adenosine is due to an increase in conduction time between the atrium and His bundle (i.e., an increase of the AH interval), whereas the His-bundle to ventricle (H-V interval) conduction time is not affected (2,5,53). In a more recent study, it was found that the electrophysiological effects of adenosine on the cells of the AVnode region are concentration-dependent and varied according to the cell type (15). As shown in Fig. 4, adenosine causes an overall depression (i.e., amplitude, duration and rate of rise) of the nodal (N) cell action potential. In comparison, in atrial and atrial-nodal (AN) cells the only significant effect of adenosine is a shortening of the action potential. Thus, the depression and eventual abolition of the action potential of N-cells identifies the N-zone of the AV node as the site of adenosineinduced AV block (15). In contrast, the nucleoside had no effect on the action potentials recorded from nodal-His (NH), His-bundle (HB) and ventricular (V) cells (15), as expected from results which showed a lack of effect on H-V interval.

Little is known regarding the cellular mechanism by which adenosine depresses the N-cell action potential and hence impairs AV conduction. The depression of the N-cell action potential could be explained by a) an increase in K<sup>+</sup>-conductance as is the case of SA-node cells and atrial myocytes, b) a decrease in calcium inward current ( $i_{Ca}$ ), or c) both. As mentioned previously, it should be noted that adenosine does not decrease  $i_{Ca}$  in sinoatrial-node cells in the absence of catecholamine stimulation. Furthermore, adenosine has been shown to increase K<sup>+</sup>-conductance not only in sinoatrial-node cells (59) and atrial myocytes (4), but also in cultured striatal neurons (52), Xenopus oocytes (42), and smooth muscle from urinary bladder (G. Isenberg, personal communication). Thus, it would not be surprising if the mechanism by which adenosine causes AV block is due to an increase in K<sup>+</sup>-conductance. Additional support comes from data showing that acetylcholine (Ach), which in the heart has similar effects as adenosine, increases single channel K<sup>+</sup>-currents in both isolated SA- and AV-node cells (48,59). Furthermore, studies in isolated hearts from animals treated with pertussis toxin show an attenuation of the adenosine-induced AH prolongation (60), similar to the effects in SA-node and atrial tissue. This implies a coupling of adenosine receptor and G-protein to the effector. Although indirect, this latter finding further suggests that the action of adenosine in the N-cells may also be due to an increase in K<sup>+</sup>-conductance.



Fig. 4. Effect of adenosine on the action potential of a nodal (N) cell from a small  $(1 \text{ mm}^2)$  AVnode preparation of a guinea-pig heart. Traces under each action potential are dV/dt. Panel A: Control; Panel B: Overall depression 2 min after addition of adenosine; and C: Five min after washout of adenosine the action potential returned to control. Preparation was paced at a rate of 3 Hz. All records were obtained from a single impalement. Time and voltage calibrations in panel C apply to all panels.

#### **III. Electrophysiological and Inotropic Effect (Atrium and Ventricle)**

In this section, the electrophysiological and inotropic effects of adenosine on atrial and ventricular myocardium are described. Also discussed are the differences in response and mechanism of action of adenosine in atrial and ventricular tissue.

#### A. Atrial Myocardium

Since the early study of Drury and Szent-Györgyi (22), the negative inotropic effect of adenosine in atrial tissue has been confirmed in various preparations and species (12,18,20,24,43,47). In contrast to the effect on ventricular tissue, in atrial myocardium adenosine and its analogs exert a concentration-dependent negative inotropic effect that is independent of prior catecholamine stimulation, i.e., adenosine depresses basal contractility. Furthermore, this direct negative inotropic effect of adenosine seems to be independent of a decrease in cellular cAMP levels



Fig. 5. Schematic of a model to describe the cAMP-independent and cAMP-dependent actions of adenosine. In A, both the cAMP-dependent and -independent actions are shown, both being present in atria, SA node and AV node. The cAMP-dependent action of adenosine is thought to result in an increase in K<sup>+</sup>-conductance ( $g_K$ ), whereas the cAMP-independent action involves the inhibition of adenylate cyclase and hence a decrease in cAMP levels. In B, as occurs in ventricular tissue, only the cAMP-dependent action is shown. The adenosine receptors mediating both the cAMP-dependent and -independent effects are of the subtype A<sub>1</sub>, i.e., both the R<sub>i</sub> (inhibitory on adenylate cyclase) and R<sup>\*</sup> (increasing K<sup>+</sup>-conductance) are the same receptor types termed A<sub>1</sub>. G<sub>s</sub>, G<sub>i</sub> and G<sub>o</sub> represent guanine-nucleotide coupling proteins, and C is the catalytic subunit of adenylate cyclase. See text for discussion. PDE = phosphodiesterase.

(12,23). Thus, like in the SA and AV node, in atrial tissue adenosine has a direct depressant effect (Fig. 1). The negative inotropic effect of adenosine in atrial myocardium is accompanied by a shortening of the action potential. It is conceivable that this shortening of the atrial action potential (4,12,30,35,36) indirectly limits calcium entry and, hence, accounts for the decrease in contractility (12,35). The marked abbreviation of the action potential and attenuation of plateau can be largely accounted for by an adenosine-induced increase in steady-state outward potassium currents (4,35). In support of this hypothesis, adenosine has been also shown to increase  $^{42}$ K efflux in atrial trabeculae (35) and sinus venosus (31). It is worth noting that these actions of adenosine in atrial tissue are similar to those of Ach, but due to activation of specific adenosine receptors that are blocked by theophylline and not atropine. Furthermore, it appears that both Ach and adenosine have a common mechanism of action, i.e., activation of a potassium conductance (4). Recent patch-clamp studies in atrial myocytes indicate that adenosine activates the same population of K<sup>+</sup>-channels as does Ach (38), but the activation of the channels is mediated by different receptors and both receptors are coupled to the K<sup>+</sup>-channel by a GTP-dependent protein.

Pharmacological and ligand binding studies indicate that the above effects of adenosine are mediated by the extracellular  $A_1$ -receptor (12,24,41). As discussed below and depicted in Fig. 5, the adenosine antagonism of the effects of isoproterenol in ventricular muscle is mediated by  $A_1$ -receptors coupled to adenylate cyclase, whereas in atrial tissue this receptor seems to be coupled to a K<sup>+</sup>-channel. Whether this atrial  $A_1$ -receptor is coupled to K<sup>+</sup>-channels by a guanine regulatory protein (G), as has been suggested for Ach receptors in atrial tissue (46), is not known. It has also been suggested that the A<sub>1</sub>-receptors that mediate the direct action of adenosine in atrial tissue, are also coupled to adenylate cyclase and the negative inotropic effect of adenosine is mediated by cAMP (40). However, much of the current evidence does not support such a hypothesis (19,12,23). It has yet to be determined whether the adenosine receptors that mediate the cAMP-independent and cAMP-dependent actions of adenosine in the atria are the same, i.e., the receptor coupled to a K<sup>+</sup>-channel and the receptor linked to adenylate cyclase are the same. However, that both induce K<sup>+</sup>-channel activation and adenylate cyclase inhibition can be clearly demonstrated. Regardless of the molecular basis of the adenosine action, the adenosine-induced increase in  $i_{\kappa}$  is a major mechanism in the action of this nucleoside in atrial myocardium.



Fig. 6. Example of the anti-adrenergic effect of adenosine in an isolated guinea-pig heart. In the first panel, a bolus (pressure injected, 0.6 s pulse) of 1 uM isoproterenol (ISO) caused a 175% and 285% increase in left ventricular pressure (LVP) and dP/dt, respectively. As shown in the middle panel, in the presence of 5 uM adenosine (ADO), ISO caused only a 32% and 45% increase in LVP and dP/dt, respectively. Note that ADO alone had no direct effect on LVP or dP/dt and only attenuated the ISO-induced increase in LVP and dP/dt. After washout of adenosine, third panel, the response to ISO was the same as during control.

#### B. Ventricular Myocardium ("Anti-Adrenergic" Effect)

Evidence has accumulated that adenosine modulates some of the myocardial actions of catecholamines (e.g., inotropic and arrhythmogenic; refs. 1,6,21,49). This modulatory role of adenosine on the cardiac effects of catecholamines occurs at two levels: a) pre-junctionally at the level of adrenergic nerve terminals (27,55), and b) post-junctionally at the level of cardiac myocytes (3,32).

In contrast to atrial myocardium, the negative inotropic action of adenosine in ventricular muscle can only be demonstrated when the adenylate cyclase activity has been previously stimulated above basal levels. As illustrated in Fig. 6, adenosine alone does not exert a negative inotropic effect, but it significantly attenuates the positive inotropic effect of isoproterenol. In fact, adenosine attenuates the positive inotropic action of forskolin, amrinone or histamine, agents known to cause an increase in cellular cyclic AMP levels (1,10,58). Similarly, adenosine and its non-metabolizable analogs have also been shown to antagonize the electrophysiologic effects of catecholamines and forskolin (3,32,58). For instance, in isolated guinea-pig and bovine-ventricular myocytes, adenosine antagonizes the isoproterenol- and forskolin-induced a) prolongation of the action potential, b) displacement of the plateau to more positive potentials, c) depolarizing afterpotentials, and d) triggered activity (3,58). These actions of adenosine, i.e., antagonism of the inotropic and electrophysiological effects of isoproterenol and forskolin, can be explained by the ability of the nucleoside to attenuate the isoproterenol- and forskolin-induced increase in calcium inward current ( $i_{Ca}$ ; ref. 32).

There is evidence to indicate that the adenosine antagonism of the aforementioned actions of catecholamines, as well as forskolin and histamine, is due to the ability of the nucleoside to antagonize the increased accumulation of cAMP caused by these substances (Fig. 7). In keeping with these findings is the observation that adenosine does not attenuate either the positive inotropic or electrophysiological effects of dibutyryl-cAMP, i.e., the site of action of adenosine is likely to be due to attenuation of adenylate cyclase activity (47,58). Based on biochemical and pharmacological studies, it appears that the adenosine receptor responsible for this inhibitory action is the  $A_1$ -receptor as identified in other preparations (40,41). As



Fig. 7. Inhibition by 0.5 uM N<sup>6</sup>-R-phenylisopropyladenosine (L-PIA) of forskolin-induced cyclic AMP accumulation in dispersed guinea-pig ventricular myocytes. Basal levels of cAMP were  $4.53 \pm 0.64$  pmol/mg protein. Note that L-PIA alone had no effect on basal cAMP levels, but it significantly (P <0.05) attenuated the forskolin-induced increase in cAMP. Reproduced from ref. 59, with permission.

illustrated in Fig. 5, in ventricular cells this receptor is coupled to the catalytic subunit of adenylate cyclase via the guanine regulatory protein ( $G_i$ ). Likewise, in the atria the receptor that mediates the indirect action ("anti-adrenergic") seems also to be linked to the adenylate cyclase via  $G_i$  (40). In support of the working model depicted in Fig. 5 is the observation that pertussis toxin, known to cause ADP-ribosylation of  $G_i$ , and, hence, uncouple the inhibitory receptors (e.g., muscarinic and adenosine- $A_1$ ) from adenylate cyclase, significantly diminishes the ability of N<sup>6</sup>-methyladenosine to antagonize the isoproterenol-induced cAMP accumulation (26). Thus, the susceptibility of the adenosine-mediated inhibition of adenylate cyclase activity to pertussis-toxin treatment provides further evidence that the adenosine  $A_1$ -receptors in ventricular myocytes are located on the cell-surface membrane and are linked to adenylate cyclase.

It should be also noted that there are a few reports suggesting the mechanism of action of adenosine in ventricular myocardium to be independent of adenylate cyclase (9,10,12). The reasons for the contradictory results are not clear, but in part may be explained by experimental technique.

#### **IV. The Physiological Role of Adenosine**

Adenosine is normally formed and released by myocardial cells into the intracellular space. Whether the amount of adenosine that accumulates in the interstitium under normal physiological conditions (i.e., well-oxygenated heart) is sufficient to affect any of the cardiac functions discussed in the previous sections remains uncertain. Although the components and their interactions of the adenosine system in the heart is becoming better understood, there are still many aspects to be elucidated. For example, the true interstitial adenosine concentration under various physiological conditions is still not known. The lack of potent and specific adenosine antagonists has hindered the characterization of adenosine receptors, as well as its physiological roles. Although recent discoveries of selective and very potent adenosine receptor antagonists are beginning to appear (16,17), the full applications and interpretation of the results with these substances remain to be clarified. Likewise, nucleoside transport blockers such as dipyridamole have multiple actions and a debate exists whether they block both uptake as well as release of adenosine. Thus, more potent and specific tools that modify the various components of the adenosine system are needed before one can more precisely define the physiological role of the nucleoside.

Despite some lack of understanding the details of the process, in recent years increasing evidence implicating adenosine as a biochemical mediator of sinus slowing and AV-conduction disturbances due to hypoxia and ischemia has accumulated (2,5,44). The observation that the effects of hypoxia and ischemia on sinus- and AV-node function can be modified in a more or less predictable manner by modulators of the actions of adenosine supports the hypothesis that the chronotropic and dromotropic effects of hypoxia and ischemia are at least in part mediated by adenosine released from oxygen-deprived myocardial cells (2,5,44). Furthermore, since adenosine has been shown to have negative chronotropic, dromotropic and anti-adrenergic effects in the human heart (19,39), the recognition and understanding of its actions may provide new insight into the etiology and treatment of certain heart diseases.

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# Chapter 10

# Acceleration of Adenine Nucleotide Biosynthesis after Ischemic Insult

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The possibility was examined that it is the slow rate of cardiac adenine nucleotide biosynthesis that may be responsible for the retarded metabolic recovery of the myocardium during the reperfusion period following a brief ischemic insult. The approach that was used is based on the fact that ribose expands the available pool of 5-phosphoribosyl-1-pyrophosphate and thus leads to the stimulation of the bio-synthesis of adenine nucleotides in heart and skeletal muscle. The effect of this stimulation on the cardiac ATP pool was then studied. Two experimental models of myocardial ischemia and reperfusion were examined in rats: recovery from multiple asphyxic periods and recovery from temporary regional ischemia. The spontaneous increase in adenine nucleotide biosynthesis that occurred in the myocardium during the reperfusion period of regional ischemia, the restoration of the ATP pool was achieved much earlier when ribose had been applied. It can therefore be concluded that the slow adenine nucleotide biosynthesis determines the speed of ATP repletion, and that ribose accelerates this metabolic recovery process. It is shown that ribose fulfills the essential criteria that should be met by a suitable cardioprotective agent.

### **The Reperfusion Problem**

Since the introduction of nonsurgical recanalization of occluded coronary arteries by means of intracoronary thrombolysis (22) and percutaneous transluminal coronary angioplasty (10), reperfusion of the previously ischemic myocardium has received much attention (2,3). Although the restoration of normal blood flow is the adequate therapeutic intervention to prevent the further development of ischemia and ultimately infarction, there are several risks involved. Reperfusion leads to the washout of adenine nucleotide degradation products that have accumulated during ischemia (8). They can therefore not be utilized to a full extent for the resynthesis of adenine nucleotides so that the previously ischemic myocardium is dependent predominantly on the de novo synthesis process (29). A further risk is the readmission of calcium which may ultimately culminate in the calcium paradox, i.e., tissue distruption, enzyme release, development of contracture and marked reduction in high-energy phosphates (31). Concomitantly, cell volume regulation may also be disturbed resulting in swelling of myocytes and endothelial cells which may contribute to vascular compression, a phenomenon termed "no-reflow" (16). Leukocytes usually migrate into the ischemic myocardium from the adjacent tissue. During reperfusion, this migration is accelerated, particularly when the vascular endothelium has been injured. That leukocytes may be involved in aggravating myocardial necrosis has been suggested, since leukocyte depletion achieved by administration of neutrophil antiserum has been shown to result in reduction of infarct size in dogs (23). Neutrophils may also contribute to the production of oxygen free radicals. It has been reported there is a burst of oxygen radical generation during reperfusion (33), that organic spin trap agents can reduce reperfusion-induced arrhythmias (13), and that superoxide dismutase and catalase protect the jeopardized myocardium (15). However, there are also studies with negative results (18,24). Another concern about reperfusion is that hemorrhage may occur and lead to the extension of myocardial necrosis.

Of particular experimental interest are the findings that the myocardial ATP content remains low for days during reperfusion after a brief ischemic period (4,21,27), and that regional heart function is impaired for a prolonged period (14). This phenomenon has been termed "stunned myocardium" (2). Several mechanisms have been proposed to be responsible for this phenomenon. First, there may be a depressed Ca<sup>++</sup>activation of contraction and a shift in Ca<sup>++</sup>-sensitivity (17). Furthermore, oxygen



Fig. 1. Schematic presentation of the pentose phosphate pathway. The reactions of the oxidative branch are shown in detail in the center. G-1-P: Glucose-1-phosphate; G-6-P: Glucose-6-phosphate; G-6-PDH: Glucose-6-phosphate dehydrogenase; 6-PGL: 6-Phosphogluconalatione; 6-P-G: 6-Phosphogluconate; 6-PGDH: 6-Phosphogluconate dehydrogenase; Ru-5-P: Ribulose-5-phosphate; PRPP: 5-phosphoribosyl-1-pyrophosphate; F-6-P: Fructose-6-phosphate; GAP: Glyceraldehyde-3-phosphate; NADP<sup>+</sup>: Nicotinamide adenine dinucleotide phosphate; GSSG: Oxidized glutathione; GSH: Reduced glutathione.

Adenine Nucleotide Biosynthesis

free radicals may be responsible, since superoxide dismutase and catalase improved segment shortening of the previously ischemic myocardium (20). In addition, there seems to be a disruption of myofibrillar energy use due to a decreased creatine kinase activity associated with the myofibril and to limitation of substrate necessary for maximal creatine kinase activity (9). Finally, the recovery of ATP seems to be limited by an insufficient increase in adenine nucleotide biosynthesis (29). In this contribution, it will be examined whether the restoration of the ATP pool during reperfusion can be accelerated by a metabolic intervention directed at stimulating adenine nucleotide biosynthesis. This can be done with ribose.

# Mode of Action of Ribose

To understand the ribose effects, one has to examine the oxidative pentose phosphate pathway. Glucose-6-phosphate originating from glycogenolysis or from glucose taken up by the cell is metabolized mainly via glycolysis (Fig. 1, right hand side).



Fig. 2. Activities of glucose-6-phosphate dehydrogenase (upper panel), the available pool of 5-phosphoribosyl-1-pyrophosphate measured as the incorporation of <sup>14</sup>C-adenine into adenine nucleotides after 15 minutes of in vivo exposure (middle panel), and rates of adenine nucleotide biosynthesis (bottom panel) in various rat organs. Mean values  $\pm$  SEM; number of experiments in parentheses; middle panel, n = 2.

An alternative possibility is the processing via the oxidative pentose phosphate pathway (central portion in Fig. 1) of which glucose-6-phosphate dehydrogenase (G-6-PDH) is the first and rate-limiting enzyme (6). After a further step catalyzed by lactonase, 6-phosphogluconate is converted to ribulose-5-phosphate by 6-phosphogluconate dehydrogenase. From ribulose-5-phosphate originates ribose-5-phosphate, the immediate precursor for 5-phosphoribosyl-1-pyrophosphate (PRPP). This substrate is essential both for pyrimidine and purine nucleotide biosynthesis as well as for the salvage pathway.

The oxidative pentose phosphate pathway has many important functions. It provides reducing equivalents in the form of NADPH. In this way, oxidized glutathione (GSSG) can be transformed to reduced glutathione. In the myocardium, an increase in GSSG may indicate oxidative stress such as during post-ischemic reperfusion. Moreover, NADPH is needed for the synthesis of fatty acids, particularly in liver, mammary gland and adrenal cortex. There are connections to glycolysis via the transaldolase and transketolase reactions, which have not been investigated exten-



Fig. 3. Effect of ribose on the available pool of 5-phosphoribosyl-1-pyrophosphate (upper panel) and on the rate of adenine nucleotide biosynthesis (bottom panel) in various rat organs. Mean values  $\pm$  SEM; number of experiments in parentheses.

sively in the myocardium. The most important function in the present context is the generation of ribose-5-phosphate for the synthesis of PRPP.

To assess the capacity of the oxidative pentose phosphate pathway in various rat organs, the activity of G-6-PDH, the available pool of PRPP and the rate of adenine nucleotide biosynthesis were determined. Fig. 2 shows that the highest activity of G-6-PDH, the greatest PRPP pool and the fastest rate of adenine nucleotide biosynthesis were found in kidney, followed by liver, heart and skeletal muscle. From these results it appears that the capacity of the oxidative pentose phosphate pathway is very small in muscular organs. The consequence of this is that the PRPP pool is minute and that adenine nucleotide biosynthesis is small in the heart and not measureable at all in the resting skeletal muscle.

The limitation of the PRPP pool can be overcome by bypassing the critical step in the oxidative pentose phosphate pathway with ribose (Fig. 1). It was thereforeexamined which effect ribose has on the available pool of PRPP and on the rate of adenine nucleotide biosynsthesis in these organs (Fig. 3). Ribose had no effect in kidney and liver. In heart and skeletal muscle, however, there was a threefold increase. Thus, ribose had a stimulating effect only in muscular organs in which the activity of G-6-PDH is very low. This concept has been confirmed in studies on the isolated perfused rat heart (12) and on mature rat cardiac myocytes (5). This investigation and other studies demonstrated that the metabolic influence of ribose is specific for muscle cells, so endothelial or interstitial cells seem not to contribute to the results obtained in the heart in vivo (26). Since ribose had such a marked stimulating effect already in the normal heart, it was of interest to examine whether it might enhance cardiac adenine nucleotide biosynthesis during reperfusion following an ischemic insult, and whether this would then result in changes in the adenine nucleotide content.



Fig. 4. Effect of successive periods of asphysia (horizontal black bars) on heart rate (HR), on the maximal rate of rise in left ventricular pressure  $(dP/dt_{max})$ , and on left ventricular pressure (LVP) in rats anesthetized with ether and ventilated artificially using a Rodent respirator.

# **Experimental Models to Study Post-Ischemic Recovery**

Two experimental models were used to induce myocardial ischemia. In the first, successive periods of total myocardial ischemia were achieved by intermittent asphysia of 1-min duration in rats induced by interruption of the artificial respiration. Each asphysic period was characterized by a decline in heart rate, LV dp/dt<sub>max</sub> and in left ventricular systolic pressure, while diastolic pressure was elevated (Fig. 4). The decrease in left ventricular systolic pressure was more pronounced with each successive asphysic period so coronary artery perfusion must have been severely impaired. As a result, the ATP content was reduced from 4.4  $\pm$  0.008 umol/g wet wt. (n = 25) to 3.7  $\pm$  0.3 umol/g (n = 6; P<0.05) at the end of the last asphysic period (25).

In the second model, regional myocardial ischemia was induced in rats by ligation of the descending branch of the left coronary artery for a brief period (1). Thereafter, reperfusion was allowed for 24 h (27). There was a progressive decline in the adenine nucleotide content depending on the duration of the ischemic period (Fig. 5). For a further detailed analysis of adenine nucleotide metabolism, an ischemic period of 15 min was chosen.

### **Ribose and Adenine Nucleotide Metabolism during Reperfusion**

During the first hour of recovery from multiple asphysic periods, there was an increase in cardiac adenine nucleotide biosynthesis, when no intervention was made. When ribose was administered as continuous i.v. infusion, there was a further stimulation of this biosynthetic process in a dose-dependent fashion (Fig. 6). However, in this model it could not be examined whether this increase might have an effect on



Fig. 5. Adenine nucleotide content (umol/g wet wt.) in rat hearts after 24 hours of reperfusion following a 10, 15 or 20 min period of regional myocardial ischemia. Mean values  $\pm$  SEM; number of experiments in parentheses.

the actual ATP content in the myocardium, since the recovery period could not be extended for a longer period of time, because of the artifical respiration and the ether anesthesia that was used in this experimental series (25).

Essentially similar results were obtained during the reperfusion period following a 15-minutes' regional ischemia (Fig. 7). After 5 h of recovery there was a spontaneous enhancement of adenine nucleotide biosynthesis which became more pronounced when ribose had been applied. This ribose-induced additional increase was of such an extent that it should have an effect on the ATP content. Fig. 8 shows the decline of the ATP pool at the end of the 15-minutes' period of regional ischemia. Without any intervention there was a spontaneous increase after 5 h of postischemic recovery, and ribose had no effect at this time. However, after 12 h, the ATP pool remained unchanged compared to the 5-h value in the untreated hearts, but had become normalized when ribose had been infused. Also after 24 h, the ATP in the untreated group was reduced to the same extent as after 12 h, and was normal with ribose treatment. Thus, the time required to reach complete ATP recovery was markedly shortened by ribose. Studies on isolated perfused working rat hearts (19) and on dog hearts in vivo (7,11) have shown that also the recovery of function was improved with ribose had a cardioprotective effect.

#### A Suitable Substrate to Improve Post-Ischemic Recovery

Since there are many competing substrates for metabolic intervention, it may be useful to have some criteria for the selection of the most suitable one. Such criteria include:

- 1. Its mechanism of action on cardiac metabolism must be known.
- 2. There should be a marked and long-term effect on myocardial metabolism in several pathophysiological situations.



Fig. 6. Rates of adenine nucleotide biosynthesis in rat hearts at the end of 60 minutes of recovery from 5 asphysic periods of 4.5-minutes' total duration (see Fig. 4), and the effect of ribose. Mean values  $\pm$  SEM; number of experiments in parentheses.

- 3. The beneficial effects should outweigh the possible risks of adverse side effects.
- 4. It should be effective in man.
- 5. It must not have hemodynamic or vasoactive properties.
- 6. Its combination with conventional therapy must be possible.

Ribose fulfills all these criteria. As to its possible effects in man, it is worth mentioning that the activity of glucose-6-phosphate dehydrogenase, the first and regulating enzyme of the oxidative pentose phosphate pathway, is even lower in the human heart than its activity in the rat or guinea-pig heart in which the cardioprotective influence of ribose has been demonstrated (28). That ribose is effective in man is also suggested by the therapeutic effectiveness of ribose in a patient with myoadenylate deaminase deficiency in skeletal musle (32). Of great advantage is the fact that ribose has no hemodynamic influence of its own and that it does not affect the functional effects of calcium antagonists and  $\beta$ -receptor blockers (30). Moreover, even when heart function is severely depressed by negative inotropic drugs, the metabolic stimulation of ribose is still preserved (30).

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Fig. 7. Rates of adenine nucleotide biosynthesis after 5 hours of recovery from a 15-minutes' period of regional myocardial ischemia in rats with continuous i.v. infusion of 0.9% NaCl or ribose (200 mg/kg/h). Mean values  $\pm$  SEM; number of experiments in parentheses.

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Fig. 8. Alterations in the cardiac ATP pool (umol/g wet wt.) at the end of temporary regional ischemia induced by ligation for 15 minutes of the descending branch of the left coronary artery in rats and during the subsequent post-ischemic recovery period. Infusion of 0.9% NaCl (hatched bars) or ribose (stippled bars) at a dose of 200 mg/kg/h was started at the end of the ischemic period and maintained throughout the repective recovery periods indicated on the abscissa. Infusion rate: 5 ml/kg/h.

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# **Technical Considerations/Models**

# Chapter 11

# The Importance of the Determination of ATP and Catabolites

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Since 30 years there has been a great deal of interest in the regulatory properties of the high-energy phosphates (HEP) ATP and phosphocreatine (PCr), and their breakdown products. The determination and interpretation of these compounds can be full of pitfalls, mainly as a consequence of their high turnover rate and compartmentation. Three different methods to determine the myocardial HEP or breakdown products are compared:

1. Cardiac tissue can be rapidly frozen, and after appropriate extraction, the HEP can be determined either enzymatically or chromatographically. These methods are sensitive and selective, but evidently destructive.

2. ATP and PCr levels can be determined by <sup>31</sup>P-NMR in vivo and in situ.

Furthermore, as a result of the chemical shift of the inorganic phosphate, the intracellular pH can be estimated. This method is less selective and sensitive than the destructive measurements, but the only method available to directly correlate intracellular biochemical with physiological parameters.

3. The apolar products of ATP breakdown such as adenosine and its catabolites inosine, (hypo)xanthine and uric acid can be determined in the coronary effluent. This method is sensitive. However, it is an indirect assessment of ATP breakdown.

A complete understanding of myocardial HEP turnover and breakdown will require complementary measurements of these compounds with each of the three measured paradigms.

# Introduction

In the normoxic heart a delicate balance exists between ATP production in the mitochondria and ATP utilization by the contractile apparatus. In response to an increase in work, the heart increases its oxygen consumption and rate of ATP synthesis instantaneously. Throughout ischemia, when flow is inadequate and hence oxygen delivery is limited, the normal equilibrium is perturbed. Despite a decline in contractility, ATP utilization is greater than its production resulting in a decrease in ATP concentration. During reperfusion, the oxygen supply is restored, but neither ATP nor contractility are restored to their pre-ischemic values.

ATP and its catabolites are central to the regulation of energy metabolism. For this reason, it is essential to measure the concentration of these metabolites in the cell and be aware of the limitations of the techniques presently available.

Initially, we will give a brief overview of myocardial ATP metabolism, then discuss the techniques available to determine ATP and its catabolites and finally evaluate the results of these measurements.

#### Normoxia

The normoxic myocyte maintains remarkably constant ATP concentrations under different conditions of work. This observation suggests a tight coupling between the production of ATP by oxidative phosphorylation in the mitochondria, and its consumption by the contractile apparatus (Fig. 1, top panel). It is thought that ATP or its products of hydrolysis (ADP and inorganic phosphate) play a major role in mitochondrial regulation, either directly as ADP (17), or through cytosolic ratios such as the phosphate potential ([ATP]/[ADP][P<sub>i</sub>]; (see refs. 31,46,69), the [ATP]/[ADP] ratio (65) or the energy charge, ([ATP + 0.5 [ADP])/([ATP] + [ADP] + [AMP])) (see ref. 5). Other important candidates involved in mitochondrial regulation are the NAD/NADH ratio (23) and Ca<sup>2+</sup>ions (22,63).

# <u>Normoxia</u>



Fig. 1. Schematic view of ATP breakdown during normoxia, ischemia and reperfusion. PCr = phosphocreatine; E = energy.

Phosphocreatine (PCr) is in equilibrium with ATP and ADP via the creatine kinase reaction (64). PCr is thought to play a role either as an energy buffer, which keeps ATP levels constant during a sudden increase in energy demand (37), or as an energy shuttle (11,58). In the latter model, it is proposed that the energy produced by the mitochondrion is predominantly transported to the myofibrils as PCr. Recently these theories have been refined by the suggestion that PCr acts either as an ADP buffer or ADP transporter (39,40). However, some investigators report that, in a normoxic heart, ATP, PCr, and in turn, ADP, AMP and inorganic phosphate ( $P_i$ ) levels are remarkably constant, giving a constant ATP/ADP ratio, energy charge and phosphate potential at various workloads (8,25,63,73). Thus, the signal which enhances mitochondrial function as a result of increased cardiac work still remains elusive.

### Ischemia

During flow restriction and hence diminished oxygen delivery to the heart, the balance between ATP production and consumption is disturbed. Despite a rise in anaerobic glycolysis, and a decrease in contractility, ATP concentrations decline, with a concomitant rise in ADP and  $P_i$  levels. In turn, ADP is metabolized to AMP (by adenylate kinase), which subsequently is dephosphorylated by 5'-nucleotidase to adenosine. This apolar compound leaves the cell and enters the bloodstream, where it is further converted to inosine, hypoxanthine and uric acid (refs. 1,16,21,34,35; see Fig. 1, middle panel). The rapid decline in contractility has been attributed to a number of different process - inadequate mitochondrial ATP production, insufficient delivery of ATP (37), an increase of ADP (19), an increase of  $P_i$  (3,44), a decrease in the phosphate potential (18,31,41), developing acidosis (43), low oxygen tension (53), or a collapse of the coronary system.

It should be stressed that ischemia is only a relative term, indicating a restriction of coronary flow. However, the severity of ischemia depends primarily on the extent of imbalance between ATP production and ATP consumption. This is reflected by the rise in ADP concentration and the parallel fall in PCr concentration (35). The concentrations of ADP and AMP are determined in turn by the adenylate kinase reaction, an equilibrium reaction. As a result, a high concentration of AMP induces a high level of release of adenosine and its catabolites with a consequently steeper drop in ATP levels (see Fig. 2). According to this model, the fall in ATP, the decline in PCr levels and the purine release are a true reflection of the severity of ischemia (16,35).

# Reperfusion

During reperfusion (when both circulation of blood and adeqate supply of oxygen are re-established), the mono- and diphosphates are rapidly reconverted to ATP (Fig. 1, bottom panel). The ATP levels remain depressed, however, as a result of washout of adenosine and its catabolites (1,6,20,21). Unless the cell is irreversibly damaged, the PCr concentration recovers to greater than its pre-ischemic levels, reflecting low ADP levels (6,59). This could indicate an adequate restoration of mitochondrial activity accompanying depressed myocardial function. Evidence from the literature suggests that there is a correlation between the concentration of ATP at the end of ischemia or at the time of reperfusion and subsequent restoration of function (56,62,68). However, there are many other factors resulting from the severity of the ischemic insult such as accumulation of H<sup>+</sup> and lactate which may cause failure of functional recovery (55). Indeed, it has been demonstrated that restoration of ATP levels in itself does not necessarily increase function (57). Changes in the contractile apparatus or Ca<sup>2+</sup>-handling by the sarcoplasmic reticulum may also be involved. If the mitochondria are not able to generate sufficient ATP, as reflected by persistently elevated ADP and low PCr concentrations, then reperfusion may be incomplete (59). In this situation the mitochondria are irreversible damaged and the cells ultimately die.

# **Compartmentalization**

There are several distinct domains in the heart, in which one can expect to find different concentrations of high-energy phosphates. These domains can be divided into multi- and intracellular compartments (54,67). The multi-cellular compartment consists of the different cell types found within the heart, e.g., myocytes, smooth muscle cells, endothelial cells, fibroblasts, conductive cells and blood.

high-energy phosphates levels differ between the extracellular Although compartments, initially the bulk (>85%) is localized within the myocytes (67). This implies that changes in ATP and PCr concentrations of >15% should reflect differences within the myocytes alone. However, changes of metabolites normally present in low concentrations (uM) should be considered more carefully. For example, a small release of adenosine or one of its catabolites could be attributed to endothelial cells (60), and a fraction of the  $P_i$  measured by nuclear magnetic resonance (NMR) arises from the blood phosphate (14). Furthermore, different populations of myocytes are thought to exist within the heart (e.g., subepi- vs. subendocardial cells), which could give rise to non-uniform high-energy phosphate distribution. Finally, if redistribution of myocardial flow occurs (regional ischemia), large regional differences of high-energy phosphates can be expected within the heart (51).



Fig. 2. Correlation between intracellular phosphocreatine (PCr) and inosine release (left panel) and intracellular ATP and inosine release (right panel) from isolated perfused rat hearts during a 90% flow reduction. PCr and ATP were measured by <sup>31</sup>P-NMR, and inosine concentrations in the coronary effluent were assayed by HPLC.  $\Box$  = hearts made ischemic with pacing at 300 beats/min;  $\blacklozenge$  = hearts made ischemic with pacing at 300 beats/min; and  $\blacktriangle$  = hearts made hypoxic without pacing (35). n = 5-9.

The intracellular compartments can be subdivided in three separate entities. First, there are physically distinct subcellular structures with different high-energy phosphate levels, e.g., mitochondria (16,28,30,66). It has been shown that there are marked differences in high-energy phosphate concentrations in the cytosol and mitochondrion. ATP and PCr are considered to be primarily cytosolic, whilst free ADP levels are higher in the mitochondria (4,16,66). Secondly, there is a bound compartment. Nucleotides readily bind to proteins, and in myocytes about 80% of the ADP is bound to actin (61). Thirdly, enzymes can form multi-enzyme complexes, in which the product of one enzyme is channeled to another enzyme, which uses it as a substrate. In the myocyte, adenylate kinase and ATPases are all in close proximity of each other (39,70). Furthermore, ATP produced by membrane-bound glycolytic enzymes and/or glycogen is used predominantly by ionic pumps (13,71). For these reasons one should be careful in the interpretation of the results of gross high-energy phosphate and catabolite determinations.

#### Techniques

To determine high-energy phosphates and  $P_i$  concentrations in heart, two fundamentally different methods can be used. The first method is to freeze the heart rapidly in order to arrest metabolism, and then, after appropriate extraction, to assay the compound of interest. One consequence of this method is that the sample is destroyed. The second method is to monitor the ATP, PCr and  $P_i$  levels with <sup>31</sup>P-NMR. This can be done in-vivo, and is a non-destructive and non-invasive method.

# Determination of ATP in Extracted Tissue

Nucleotides are very unstable compounds in biological matrices. This means that during tissue sampling, ATP and PCr levels can fall rapidly with a concomitant rise in ADP and  $P_i$  levels. Wollenberger et al. (72) used aluminum tongs precooled in liquid nitrogen to "freeze-clamp" the tissue of interest within seconds. Hearse also repeatedly stressed the need for fast tissue enzyme deactivation (36). The frozen tissue can be extracted with perchloric acid (PCA) or trichloroacetic acid (TCA), with or without alcohol and EDTA. PCA is removed by precipitation of its potassium salt, or by extraction with an amine dissolved in freon. TCA is removed by ether extraction. Recoveries of 80 - 100% for both procedures have been obtained (34).

It should be stressed that slow freeze-clamping induces temporal tissue hypoxia with consequently lower PCr concentrations and hence increased ADP, AMP and  $P_i$  levels. The extent of this artefact depends on the rate of ATP turnover in the tissue (34,47). A high rate of ATP prior to freeze-clamping results in larger errors in ADP, AMP and  $P_i$  levels, than those from a tissue with a low ATP turnover rate. Freeze-clamping and subsequent acid-extraction result in the measurement of total nucleotides in cardiac tissue, independent of the form in which they exist and irrespective of their cellular location. To assess the nucleotide distribution within the cell, Sobel and Bünger isolated mitochondria from cells in a non-aqueous medium, to prevent ATP-hydrolysis, followed by an acid extraction (66). Mild extraction of freeze-clamped heart tissue with salt buffers can give an estimate of the free ADP available (61).

In neutralized samples, high-energy phosphates can be measured by enzymatic asays (10,50), bioluminescence (49,50), isotachophoresis (24) or high-performance liquid chromatography (HPLC, refs. 15,35). The differences between these techniques are not essential. The advantage of both HPLC and isotachophoresis is that, within a single run, several compounds can be determined. In principle, this leads to a more precise determination of metabolite ratios. Enzymatic determinations, however, can be more selective. Bioluminescence is specifically used for its sensitivity (50). In earlier days, the purity of the luciferine/luciferase enzymes caused problems.

Nowadays, with increased purification of these enzymes and inhibitors of enzymes such as adenylate kinase, the results of these assays are more reliable.

Adenosine and its catabolites in the coronary effluent are primarily measured by HPLC and enzymatic assays (10,33). Again, with these assays, the sample preparation is the most crucial step of the whole procedure. A small increase in ATP breakdown as a result of improper sample handling can result in a massive increase in adenosine and its catabolites (33).

# Determination of High-Energy Phosphates with <sup>31</sup>P-NMR

Nuclear magnetic resonance is a spectroscopic method which utilizes the magnet spin properties of an atomic nucleus (e.g., <sup>31</sup>P, spin = 1/2). By applying a varying magnetic field and detecting the energy transitions in the radio frequence range, the quantity of a particular nucleus, its chemical micro-enviroment (e.g., the phosphate component in either ATP, PCr or P<sub>i</sub>) and its macro-enviroment (whether a molecule is free or bound) can be monitored (26). This technique is completely non-invasive and can be applied to in-vivo systems (27,38). <sup>31</sup>P-NMR spectra have been obtained from isolated perfused hearts (6,29,35), from hearts in open-chested or intact animals by means of an internal or external surface coil (2,14,32,42,51) and from human myocardium using a spatial localization technique (12).

The sensitivity of this technique is, in part, dependent on the magnetogyric ratio and relaxation times of the nucleus under investigation and the strength of the magnetic field. The selectivity is dependent on the strength of the magnetic field. In a beating heart, the clearly observable peaks are  $P_i$ , PCr and the three phosphates of ATP with a very small contribution from ADP. The chemical shift of the  $P_i$  peak can determine the pH (26,29). In a blood-perfused heart, the  $P_i$  peak is contaminated with the 2,3-diphosphoglycerate from blood. It is possible to distinguish between the two by a spectral editing method (14).

Although the NMR technique does not have the sensivity and selectivity of the previously mentioned analytical techniques, its advantages are obvious. With this technique, it is possible to monitor biochemistry and physiology of the heart simultaneously. Furthermore, within one preparation, continuous measurements can be made. Also with NMR it is feasible to "magnetically label" specific phosphate groups and follow their transfer into another molecule as a result of an enzymic reaction (saturation transfer, refs. 26,64). This is a unique method for determining in vivo enzyme kinetics. Creatine kinase and ATPase activity have been studied extensively, and the results have helped clarify the role of these enzymes in the cell (7,9,25,45,48,52,64).

#### Conclusions

In the introduction, ideas concerning the importance of the regulation of ATP and its catabolites are described. Although there is still controversy about the exact mechanism of regulation of high-energy phosphate metabolism, there is a general consensus concerning the importance of measuring ATP and catabolite concentrations. One should, however, be aware of four problems:

1. In order to understand the regulatory properties of a compound or an enzyme (complex), knowledge of local subcellular concentrations is essential. At present, these concentrations cannot be determined.

2. Analytical assays in a tissue sample give an average measure of the compound in different compartments. In freeze-clamped and acid-extracted tissue, total (bound and soluble) nucleotides are determined, whilst with <sup>31</sup>P-NMR, only the soluble components are determined. Thus, determination of ADP and  $P_i$  by these two techniques

will generate very different values owing to their significant binding in vivo and breakdown of ATP and PCr on sample preparation.

3. Despite certain advantages in determining metabolite ratios (either in an extract or with an NMR spectrum), this method can be misleading. Energy charge calculations from extracts of freeze-clamped tissue, using total ADP levels, can lead to serious underestimation. Low PCr/ATP ratios can be used to indicate ischemia but, alternatively, a high PCr/ATP ratio can reflect either a normoxic tissue with high PCr levels, or a post-ischemic tissue with low ATP levels. Thus, absolute values of ATP are needed to distinguish between these two possibilities.

4. Relationships exist between biochemical parameters, the severity of ischemia and onset of reperfusion. Although the mechanisms linking the biochemical parameters and physiological functions are not always clear, some parameters (concentrations or ratios) can be used as indicators of ischemia and reperfusion damage. To truly evaluate the condition of a heart, some type of metabolic stress need to be used.

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# Chapter 12

# Cardiac Energy Metabolism Probed with Nuclear Magnetic Resonance

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Nuclear magnetic resonance (NMR) spectroscopy possesses great potential for studying myocardial energy metabolism. To ensure that the observed NMR signal predominantly originates from the heart, localization is required, which can be achieved by excision or exposure of the heart, or by means of sophisticated NMR localization techniques. A number of different atomic nuclei have been employed. <sup>1</sup>H NMR has been mainly used to follow lactate accumulation in ischemic or anoxic hearts. <sup>13</sup>C NMR has been applied to study the fate of different substrates in the citric acid cycle and amino acid pools, and the role of glycogen metabolism in ischemia or anoxia. <sup>19</sup>F, <sup>23</sup>Na and <sup>39</sup>K have been employed to investigate the consequences of altered energy metabolism for myocardial intracellular concentrations of  $Ca^{2+}$ ,  $Na^+$  and  $K^+$ . The most abundantly used nucleus for studying myocardial energy metabolism is <sup>31</sup>P. Numerous contributions have been made to the investigation of ischemia and reperfusion, protection of the heart against the consequences of ischemia and reperfusion, contractile failure, variation of highenergy phosphate levels over the cardiac cycle, regulation of oxidative phosphorylation and intracellular enzyme kinetics of both isolated perfused hearts and hearts in situ. Even human myocardial metabolism can be assessed by <sup>31</sup>P NMR, which is on the verge of becoming a clinical tool for investigating heart disease.

# Introduction

Nuclear magnetic resonance (NMR) spectroscopy offers unique possibilities to study non-destructively in vivo and in vitro myocardial energy metabolism. The technique relies on the response of atomic nuclei, which are polarized by a strong magnetic field, to a radiofrequency pulse. Many atomic nuclei behave, due to a property called "spin", as magnetic dipoles. In a magnetic field the tiny nuclear magnets will orient themselves either parallel or anti-parallel to the magnetic field (when the nucleus has a spin quantum number  $\pm 1/2$ ). Due to an unequal occupation of both orientations, a net magnetization results. With low-energy radiofrequency pulses, transitions between the two orientations can be induced, allowing for manipulation of the net magnetization. The magnitude of this net magnetization can be detected as an emitted radiofrequency (RF) signal when after the pulse the nuclei relax to their equilibrium orientation.

The so-called resonance frequency of the emitted radiowaves is characteristic of the observed nucleus and directly proportional to the magnetic field strength. Table 1 shows the resonance frequencies and some properties of nuclei, commonly observed in in vivo and in vitro NMR. When a particular nucleus is observed, the exact resonance frequency is determined by its chemical environment. This property, known as chemical shift, allows us to discriminate between identical nuclei in

different molecules or in molecules in different environments. The chemical shift  $\delta$  is normally expressed as  $\delta = (v_i - v_r) \cdot 10^6 / v_r$  ppm (parts per million) with  $v_i$  = resonance frequency and  $v_r$  = resonance frequency of a reference compound. The NMR signal from biological tissue is often a complex pattern of radiowaves, which is Fourier transformed into a frequency spectrum by computer.

Since the emitted signals are generally very weak, the measurement has to be repeated a number of times (signal averaging) to obtain a sufficient signal-to-noise ratio for reliable quantitative measurements. The signal intensity is in principal proportional to the number of observed nuclei, and higher at higher magnetic field strengths. However, upon signal averaging, signal intensity can be affected by other factors such as the spin-lattice relaxation time  $T_1$  and the measurement repetition time. The relaxation time  $T_1$  is a time constant which characterizes the return of the nuclei to their equilibrium distribution. When the next scan is performed before complete relaxation has occurred, the subsequent signal will be less than in the first measurement.  $T_1$ -relaxation times of nuclei in small molecules and ions in biological tissue normally range from a few hundreds of milliseconds to a few seconds.

# Localization

For a meaningful determination of myocardial energy metabolism with NMR, some form of localization is required to ensure that the observed signal predominantly originates from the heart. The most straightforward and until now most commonly used method is the excision of the heart. The Langendorff perfused or working heart preparation is placed in a glass tube and brought in the magnet center. The RF-transmitter and receiver coil surrounds the tube and detects signal from the entire heart. The possibilities of this approach have first been demonstrated by Gadian et al. (36) in 1976.

Hearts in situ have been studied by surgically introducing NMR coils to fit around the heart of mechanically ventilated animals (41), thereby unavoidably picking up signal from blood. Other invasive approaches to study myocardial energy metabolism in vivo with NMR include chronically implanted coils (61) and small surface coils introduced as a catheter coil (53). The surface coil normally is a flat, circular coil detecting signal, as a rule of thumb, from two halve spheres on either side of the coil with a radius equal to the coil radius. This approach is easily clinically applicable (103), although the signal from the heart will be contaminated with signal from blood, bone and skeletal muscle.

Nucleus	Natural abundance (%)	Relative sensitivity (%)*	Resonance frequency (MHz) at 1 Tesla magnetic field strength			
<sup>1</sup> H	99.98	100.0	42.58			
<sup>19</sup> F	100.0	83.0	40.05			
<sup>31</sup> P	100.0	6.6	17.24			
<sup>23</sup> Na	100.0	9.3	11,26			
<sup>13</sup> C**	1.1	1.6	10.71			
<sup>39</sup> K	93.1	0.051	1.99			

Table 1.	<b>Properties</b> (	of commonly	v observed	l nuclei in	in vivo	and in	vitro NMR	spectroscopy
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\* Relative sensitivity is the NMR sensitivity of nuclei relative to that of an equal number of protons.

<sup>12</sup> The most abundant carbon isotope <sup>12</sup>C (98.9%) is not magnetically active, whereas the nonradioactive isotope <sup>13</sup>C is. The surface coil method has been refined by combination with NMR-imaging derived slice selection (15,16). This method exploits the fundamental dependence of resonance frequency on magnetic field strength by applying a linear magnetic field gradient perpendicular to the plane of the surface coil, together with an RF pulse of limited bandwidth. In this way only a slice of tissue is excited instead of the entire half spherical volume, resulting in one dimensional localization. Other one dimensional localization methods make use of the inherent gradient of the RF field emitted by a surface coil (13).

Most elegant localization methods are the so-called volume selective excitation techniques (5) in which three orthogonal magnetic field gradients are used. Originally these methods were developed for coils that surround the subject and require high power RF-pulses. However, since safety considerations and technical limitations preclude their use for investigating human heart, alternative approaches have been developed. One of the most promising of these techniques is the so-called ISIS method (87), a three dimensional localization method which requires only low power RF-pulses and can be used with surface coils, and consequently is ideally suitable for studying human myocardial metabolism.

### Myocardial Metabolism

A number of different nuclei has been employed to study myocardial metabolism with NMR. Each nucleus has its own characteristics and is associated with particular molecules commonly probed with it. Generally, narrow resonance peaks are only obtained from nuclei in molecules that are fairly mobile. Nuclei in large molecules give rise to very broad resonance lines underlying the narrow peaks from mobile metabolites or ions.

# ${}^{1}H$

Despite its relatively high sensitivity, <sup>1</sup>H NMR has rarely been used to study myocardial energy metabolism. This is mainly due to the enormous signal arising from  $H_2O$ , which tends to conceal other resonances, and the ubiquity of protons in cellular components, which often results in very complicated spectra. Increasing the magnetic field strength not only increases sensitivity but also spectral dispersion which facilitates the interpretation of complicated spectra.

By saturating the  $H_2O$  resonance by selective irradiation with a very narrow frequency bandwidth prior to data acquisition, Ugurbil et al. (101) have been able to detect in isolated perfused rat hearts resonances corresponding with taurine, carnitine, creatine and phosphocreatine, glycerides, and especially in ischemic hearts, lactate. Lately, sophisticated methods for water suppression (14,47) and separation of overlapping peaks by editing techniques, which also frequently use selective irradiation (44,105), have been developed that allow a more reliable determination of tissue lactate in brain and skeletal muscle.

Only recently these methods have been applied to follow the time course of lactate accumulation in ischemic and anoxic isolated rat hearts (91). With a time resolution of 5 min, lactate was detected after 5 min in the ischemic hearts and after 10-20 min in anoxic hearts. Unfortunately, these measurements were performed at a relatively low field strength of 2 Tesla. At higher field strengths, a better time and spectral resolution can be obtained with a practical detection limit around 0.1 mM.

The chemical shift range of  ${}^{13}C$  is much wider than that of  ${}^{1}H$  (ca. 250 vs. ca. 15 ppm). Therefore,  ${}^{13}C$ -spectra can be easier interpreted than  ${}^{1}H$ -spectra. Unfortunately  ${}^{13}C$  is far less sensitive than  ${}^{1}H$  and has a natural abundance of only 1.1% (see Table 1). Therefore,  ${}^{13}C$  in vivo and in vitro NMR spectroscopy is limited to molecules that have been specifically enriched with  ${}^{13}C$  or are present at relatively high intracellular concentrations. In addition, nuclei which are close together by chemical bond or in space, can show homonuclear and heteronuclear interactions or couplings. As is the case with  ${}^{1}H{}^{-13}C$  interactions, this often results in splitting of the peaks in doublets, triplets etc. By simultaneous irradiation with  ${}^{1}H$ -resonance frequencies, this interaction can be "decoupled", which increases signal intensity and simplifies the spectra. However, decoupling requires the presence of a second coil for transmission of  ${}^{1}H$ -frequencies.

Nevertheless, <sup>13</sup>C NMR has been successfully applied to study metabolism of both isolated, perfused hearts and hearts in situ. The first natural-abundance <sup>13</sup>C spectrum of isolated rat hearts was reported by Bailey et al. (6). It showed only resonances attributable to lipids, which are present in sufficiently high concentrations. When [2-13C]acetate was offered as a substrate instead of glucose to investigate citric acid cycle activity, labeling of the C-2, C-3 and C-4 carbons of glutamate was observed. Since the C-3 carbon of alpha-ketoglutarate and hence glutamate (via transamination) can only be labeled in a second turn of the tricarboxylic acid cycle of previously labeled citric acid and since scrambling of the label occurs between the C-2 and C-3 of succinate, the steady state enrichment of glutamate at C-3 occurs at a slower rate than that at C-4. From this difference in incorporation rate, the relative rates of cycling in the tricarboxylic acid cycle and the transamination to glutamate can be estimated. When acetate in the perfusate was partly replaced by glucose, label also appeared in C-2 and C-3 of aspartate. This may reflect changes in transamination due to increased activity of the malateaspartate shuttle. From coupling patterns between <sup>13</sup>C at C-4 of glutamate with neighboring <sup>13</sup>C at C-3 (which could only be observed in HClO<sub>4</sub> extracts of the heart, due to better resolution), it was concluded that exogenous acetate provided all of the substrate oxidized in acetate-perfused hearts. Finally, Bailey et al. (6) failed to detect any incorporation of label in alanine via pyruvate formed by malic enzyme from malate, although considerable malic enzyme activity has been reported in rat heart (93). They suggested that the malic enzyme is either kinetically controlled to feed substrate effectively unidirectionally into the citric acid cycle, or is inhibited in vivo, or cannot convert the labeled malate pool due to compartmentation.

The observation that only glutamate carbons become detectably labeled in [2- $^{13}$ C]acetate-perfused rat hearts was essentially confirmed in a recent study (72), where also a minor additional resonance from acetylcarnitine was observed. About 90% of substrate oxidized was acetate. When propionate was added, which enters the tricarboxylic acid cycle via anaplerotic pathways, aspartate and malate appeared in the  $^{13}$ C spectrum, whereas glutamate resonances decreased. It appeared that under these conditions only 50% of the substrate oxidized was acetate. In guinea-pig hearts perfused with [3- $^{13}$ C]pyruvate as substrate, resonances were detected from glutamate, aspartate, alanine, citrate, malate, lactate and acetylcarnitine, whereas in [3- $^{13}$ C]lactate-perfused hearts only glutamate resonances were observed. This most likely reflects a difference in the malate-aspartate shuttle activity, since lactate can only be utilized effectively when sufficient NAD<sup>+</sup> is provided for oxidation to pyruvate (95):



During anoxia in guinea-pig hearts, which had been preperfused with [3-<sup>13</sup>C]pyruvate, substantial labeling of succinate occurred, whereas the intensities of labeled glutamate and aspartate simultaneously decreased. These results were interpreted as anaerobic metabolism of glutamate and aspartate to succinate yielding nucleoside triphosphates which may provide a supplemental mechanism to glycolysis for production of high-energy phosphates during anoxia or ischemia (18). Chance et al. (25) have presented a sophisticated mathematical analysis of <sup>13</sup>C labeling of tricarboxylic acid cycle intermediates, which allows calculation of total citric acid cycle flux.

Other authors have used D-[1- $^{13}$ C]glucose as substrate to study myocardial glycogen synthesis and subsequent glycogenolysis, and appearance of label in lactate during anoxia (83,84) or ischemia (66). Glycogen is one of those rare exceptions in that all of the  $^{13}$ C-carbons are NMR-visible, despite the very large mol. wt. as has been demonstrated for liver and heart glycogen (83,97). Apparently the molecule possesses high internal mobility.

It is clear that <sup>13</sup>C NMR offers unique possibilities and has distinct advantages over conventional <sup>14</sup>C-labeling techniques to study simultaneously the contribution of competing metabolic pathways and to elucidate specific pathways from preferential labeling. One may therefore expect that, despite the drawbacks mentioned before, <sup>13</sup>C NMR will become an important tool to study myocardial energy metabolism.

# <sup>19</sup>*F*, <sup>23</sup>*Na and* <sup>39</sup>*K*

<sup>19</sup>F, <sup>23</sup>Na and <sup>39</sup>K are being increasingly used to study myocardial intracellular ion concentrations. <sup>23</sup>Na and <sup>39</sup>K NMR (31,90) are employed to directly measure intracellular [Na<sup>+</sup>] and [K<sup>+</sup>] in conjunction with so-called shift reagents. These paramagnetic reagents weakly interact with the extracellular ions, thereby changing the resonance frequency of these ions and separating the signals from intra- and extracellular ions. Intracellular  $[Ca^{2+}]$  can be measured with <sup>19</sup>F NMR by loading the cell with a fluorinated  $Ca^{2+}$ -chelator (98). (Direct estimation of  $[Ca^{2+}]$  -and  $[Mg^{2+}]$  for that matter- using the magnetically active isotopes <sup>43</sup>Ca and <sup>25</sup>Mg is hardly feasible due to low sensitivity and low natural abundance.) Although <sup>19</sup>F, <sup>23</sup>Na and <sup>39</sup>K do not probe myocardial energy metabolism directly, they may become a powerful tool to study the consequences of altered energy metabolism for intracellular ion concentrations. Unless gated to the cardiac cycle, the time resolution of these measurements in isolated, perfused hearts, which varies from about 1 min for <sup>23</sup>Na to several min for <sup>39</sup>K and <sup>19</sup>F, is not sufficient to detect variations in intracellular concentrations during the cardiac cycle. However, it has been possible to observe substantial differences in intracellular [Na<sup>+</sup>] and [Ca<sup>2+</sup>] at different levels of cardiac performance (21,73,90,99) and after 10 min of global ischemia (73,99).

<sup>31</sup>P is the most abundantly used nucleus for studying myocardial energy metabolism. Figure 1 shows a <sup>31</sup>P NMR spectrum of an isolated Langendorff-perfused rabbit heart, illustrating the <sup>31</sup>P resonances one may find in heart. Since protonation of phosphate groups changes the chemical environment of the phosphorus nucleus, a number of the peaks in Fig. 1 show a pH-dependent chemical shift. Alternatively, the resonance frequency can be used to determine intracellular pH. Inorganic phosphate (P<sub>i</sub>) is a good candidate for intracellular pH-measurement (80), which is illustrated by the P<sub>i</sub>-peaks corresponding with an extracellular pH of 7.4 and an intracellular pH of 7.0 in Fig. 1.

Intracellular pH measurement of hearts in situ is often complicated by the (partially) overlapping resonances of 2,3-diphosphoglycerate (2,3-DPG) from blood and P<sub>i</sub>. Brindle et al. (19) have used an editing technique, which exploits the [<sup>1</sup>H-<sup>31</sup>P]-couplings in 2,3-DPG and the absence of these couplings in P<sub>i</sub>, to effectively suppress the resonances of 2,3-DPG. The P<sub>i</sub> resonance thus measured indicated a pH of approximately 7.3. Therefore they concluded that the P<sub>i</sub> signal obtained from normoxic hearts in vivo predominantly originates from blood, although spectra obtained from isolated heart at high workloads can show significant resonances from intracellular P<sub>i</sub> (35).

Another external factor which may influence chemical shifts is binding of metal ions. Especially the gamma- and beta-ATP resonances are sensitive to the binding of



Fig. 1. 81.0 MHz (4.7 Tesla) <sup>31</sup>P NMR spectrum of an isolated rabbit heart obtained from 128 scans with an interpulse time of 10 s. Methylene diphosphonate is an external reference, phosphomonoesters include sugar phosphates and AMP,  $P_i$  = inorganic phosphate, ext = extracellular, int = intracellular, GPE = glycerophosphoethanolamine, GPC = glycerophosphocholine, PCr = phosphocreatine, ATP = adenosine 5'-triphosphate, ADP = adenosine 5'-diphosphate, NAD(H) = nicotinamide adenine dinucleotide.

 $Mg^{2+}$  (42,48). From the spectrum in Fig. 1, it can be derived that >92% of ATP is complexed to  $Mg^{2+}$ -ions and, assuming the [ATP] to be 8 mM (37), the intracellular  $[Mg^{2+}]$  can be estimated (42) to be 0.5 mM. Only high resolution <sup>31</sup>P NMR spectra of heart extracts show ADP resonances resolved from ATP resonances. In in vivo spectra from intact hearts, the peaks of alpha- and beta-ADP unfortunately coincide with the peaks of alpha- and gamma-ATP, resp. The beta-ATP peak, however, is unique for ATP, so strictly speaking, the difference between beta- and gamma-ATP levels are normally two orders of magnitude higher than ADP levels. Nevertheless, free cytosolic [ADP] can be derived from the creatine kinase (CK) reaction. Combined measurement of O<sub>2</sub>-consumption and fluxes through the CK reaction (using magnetization transfer methods, which will be discussed below) have shown that CK fluxes in normoxic, perfused heart exceed the mitochondrial ATP synthesis rate by at least a factor of 3 (12,76,77) over a wide range of cardiac performance. This demonstrates that the CK reaction is sufficiently rapid to maintain the cytosolic reactants near their equilibrium concentration. It is therefore justified to derive [ADP]<sub>free</sub> from the CK equilibrium as [ADP] = ([ATP] [Cr])/([PCr] [H<sup>+</sup>] · K<sub>eq</sub>) in which  $K_{eq} = 1.66 \cdot 10^9 M^{-1}$  (at pH 7.2,  $[Mg^{2+}]_{rree} = 1 mM$  and an ionic strength of 0.25 M (68). [ATP], [PCr] and [H<sup>+</sup>] can be measured from <sup>31</sup>P NMR spectra and the creatine (Cr) concentration can be derived assuming a constant total pool of Cr + PCr of about 26 mM (37,69,106). Depending on the workload and the carbon substrate being oxidized, [ADP]<sub>free</sub> has been reported in the range from 10 um up to 132 um (35,77,106).

Ischemia and reperfusion. One of the first topics addressed with <sup>31</sup>P NMR was intracellular acidosis during ischemia and subsequent recovery upon reperfusion (39,45,51), which soon became combined with simultaneous measurement of left ventricular pressure in paced hearts (46). The metabolic consequences of global ischemia, which were investigated in these and many subsequent studies, comprise a fast decrease of the PCr level at the onset of ischemia, followed by a slower decrease of the ATP level and a concomitant decrease of intracellular pH and increase of the P<sub>i</sub> level. Garlick et al. (38) have suggested that glycogenolysis during ischemia is the major H<sup>+</sup>-producing reaction. They observed an attenuation of intracellular acidosis in glycogen-depleted hearts. However, as pointed out by Gevers (40), glycogenolysis and lactate-producing glycolysis is not associated with a net generation of  $H^+$ -ions, and acidification is rather the result of ATP hydrolysis. Bailey et al. (7) have pretreated glucose-perfused rat hearts with insulin resulting in increased glycogen content and increased accessibility of glycogen to phosphorylase during ischemia. They observed a decrease of intracellular pH to 5.7 as compared to 6.2 for untreated hearts, and a slower decline of ATP levels. The extremely low intracellular pH contradicts the commonly believed control of glycolysis by pH at the level of phosphofructokinase. Probably cessation of glycolysis in untreated hearts at pH 6.2 is rather a consequence of reduced accessibility (7). Seymour et al. (94) observed a better mechanical recovery upon reperfusion of insulin-pretreated ischemic heart, indicating that ATP maintenance is more important to recovery than maintenance of intracellular pH is. In protecting isolated, perfused rabbit hearts by hypothermia and cardioplegia against the consequences of global ischemia, Flaherty et al. (29) also found better functional recovery with higher ATP content at the end of the ischemic period, although intracellular acidosis and increased P<sub>i</sub> levels correlated inversely with recovery of postischemic ventricular structure and function.

Numerous other interventions have been applied to protect the isolated heart against the consequences of ischemia or hypoxia and reperfusion, including inosine infusion (49), adenosine deaminase inhibitors (27), perfluorocarbon cardioplegia (28), free radical scavenging (3,4) and treatment with prostaglandin (88), beta-blockers and calcium antagonists (58,65,67,71,81,85,89,92). The beneficial effect of treatment with calcium antagonists in most cases relied on an energy-sparing effect. However, we recently showed that pretreatment of rats with anipamil, a new calcium antagonist, did neither alter left ventricular developed pressure (LVDP) of the isolated hearts under normoxic conditions, nor the rate and extent of ATP and PCr depletion during ischemia, as can be seen in Fig. 2, whereas intracellular acidosis was attenuated. Upon reperfusion, hearts from anipamil pretreated animals recovered significantly better than untreated hearts, with respect to high-energy phosphate levels, return to low  $P_i$  levels (see Fig. 2), and left ventricular function and coronary flow. Intracellular pH also recovered rapidly to pre-ischemic levels, whereas in untreated hearts a complex intracellular  $P_i$ -peak indicated areas of different pH within the myocardium. Protection in this case cannot be attributed to an energy-sparing effect during ischemia, since a negative inotropic effect was absent during normoxia.

<u>Contractile failure</u>. The decline of contractile function of isolated rabbit hearts during the early phases of ischemia has been studied by Jacobus et al. (50). From comparison with respiratory acidosis, they concluded that intracellular acidification early in the ischemic period accounts for at the most a 50% reduction in LVDP and that other factors must also play a role. Since ATP levels did not change appreciably during their experiments, these cannot fulfil this role. Only when the [ATP] falls below about 35% of the normal level, contractility is affected (62,82).

Early contractile failure during hypoxia has recently been studied by several investigators using <sup>31</sup>P NMR. Allen et al. (1) have found in isolated, perfused ferret hearts that also during hypoxia with normal glycolysis rates, the decreased pH only partially accounted for the decreased LVDP. In this situation the amplitude of the calcium transients was unaffected (2). Therefore, they suggested intracellular  $P_i$  to be the major other factor that contributes to early contractile failure. When glycolysis was inhibited, no pH decrease occurred, but in this situation a decline in calcium transients has been previously reported (2). At the same time a decrease of 20% of the [ATP] during the first 5 min of hypoxia was observed. Using the CK



Fig. 2. Serial <sup>31</sup>P NMR spectra obtained from untreated (left) and anipamil-pretreated (right) rat hearts during control perfusion, 30 min of total ischemia, and 30 min of reperfusion. Numbered peaks include: 1) extracellular  $P_i$ ; 2) intracellular  $P_i$ ; 3) PCr; 4, 5 and 7) gamma-, alpha- and beta-ATP, resp.; 6) NAD(H).

equilibrium, the [ADP] could be derived and was used to calculate the free energy change of ATP hydrolysis:  $\Delta G_{ATP}$ . As originally suggested by Kammermeier et al. (52), Allen et al. (1) discussed the decline in  $\Delta G_{ATP}$  and consequent lack of energy to maintain sarcoplasmic reticulum calcium pump activity as a causal factor in the rapid and total contractile failure during hypoxia with inhibited glycolysis. However, a similar decrease in  $\Delta G_{ATP}$  was observed during hypoxia with normal glycolytic activity.

Using a method for eliciting and measuring maximal  $Ca^{2+}$ -activated tension in ferret hearts (74), Kusuoka et al. (64) concluded that during the early phase of hypoxia  $[P_i]$  is responsible for the depression of contractility and that pH plays only a minor role. Matthews et al. (78) suggested that the myocardial inotropic state may be directly responsive to the ambient  $pO_2$ , although their results are qualitatively similar to those of Allen et al. (1). Interestingly, they also reported that no significant decrease in the ATP-utilization rate occurred, despite an 80% reduction in mechanical activity during hypoxia.

Using a surface coil and an open-chest model (41), Bittl et al. (11) have investigated the effects of hypoxia in the heart of living rats. Unfortunately, due to the previously mentioned overlap of 2,3-DPG and  $P_i$  resonances, measurements of myocardial intracellular pH and  $[P_i]$  were not possible. However, based on the decrease of the steady state [high-energy phosphate] at different levels of hypoxia, they calculated with the data of Kusuoka et al. (64) at the most a 20% reduction in maximal Ca-activated pressure caused by  $P_i$  accumulation. On the other hand, they observed a 45% decrease in systolic pressure at 8%  $O_2$  ventilation, suggesting that other mechanisms than the  $[P_i]$  are involved too. They measured flux through the CK reaction and found a close match between this flux and cardiac performance at different degrees of hypoxia, whereas ATP levels fell only slightly during hypoxia. They suggested that the decreased turnover of ATP and PCr reflects the combination of impaired ATP synthesis and utilization and may be the energetic basis for hypoxic cardiac failure. However, their decreased ATP turnover is not consistent with the unchanged ATP utilization in isolated hypoxic hearts (78).

Regional ischemia. The surface coil approach with both open and closed thorax has been used by other investigators to study the effects of ligation of the left anterior descending coronary artery (LAD). <sup>31</sup>P NMR spectra of ischemic regions in cat hearts showed similar changes as were observed in global ischemic rat or rabbit hearts (100). In rabbits ATP levels decreased together with PCr levels, although at a slower rate. After about 30 min of ligation, substantial amounts of high-energy phosphate were still present (71). In dog hearts the decrease of high-energy phosphates was relatively limited (17,43). The different results obtained for different species may well reflect differences in collateral flow. In dog hearts a correlation was found between regional contractile function and both pH and P<sub>i</sub>, whereas ATP levels correlated with recovery of regional contractile function (43). A closed-chest dog model allowed for long-term evaluation of the effects of permanent LAD occlusion (17). The results were quite diverse. In some dogs P<sub>i</sub> and pH returned to normal within 30 min, whereas in others PCr/P<sub>i</sub> ratios remained depressed for up to 5 days. In all dogs myocardial pH returned to normal within 15 h.

<u>Cardiac Cycle</u>. The variation of high-energy phosphate and  $P_i$  levels during the cardiac cycle has also been investigated with <sup>31</sup>P NMR. Using an NMR method, in which the data acquisition was gated to the heart beat, Fossel et al. (33) observed a considerable variation in high-energy phosphate and  $P_i$  levels during the heart cycle of an isolated working heart. They found both PCr and ATP levels to be maximal at minimal aortic pressure and minimal at maximal aortic pressure, whereas

 $P_i$  showed a direct phase relationship with the aortic pressure curve. However, using a saturation transfer method, Matthews et al. (76) demonstrated that in an isolated Langendorff-perfused rat heart only 2.5% of the total ATP is turned over during one heart cycle compared to about 35% in the other study (33). An explanation for the observed discrepancy may be a poor nutrient and/or oxygen supply in the isolated, working rat heart.

Gated <sup>31</sup>P NMR spectra of rat (61) and dog (54) hearts in situ did not reveal any variation over the heart cycle. Assuming that the total Cr pool remains constant, also during the cardiac cycle, variation of the [ADP] over the cardiac cycle cannot be concluded, although the  $[ADP]_{free}$  is believed to be a key regulatory factor in oxidative phosphorylation (24).

<u>Respiratory regulation</u>. The regulation of oxidative phosphorylation at different steady state levels of cardiac performance has been studied using <sup>31</sup>P NMR in isolated rat hearts perfused with different substrates (35) and in hearts in situ of different species. For isolated rat hearts perfused with pyruvate (+ glucose), ADP levels were generally very low and varied with myocardial O<sub>2</sub> consumption, which suggests a regulation of oxidative phosphorylation through ADP availability. However, with glucose as carbon substrate in the presence or absence of insulin, ADP levels were much higher and the phosphorylation potential, ADP/ATP ratio and [ADP] were relatively constant over the range of workloads examined. It has been previously reported that, even at low workloads, levels of citric acid cycle intermediates, pyruvate and mitochondrial NADH are at least an order of magnitude lower with glucose is used as substrate, carbon substrate delivery to mitochondria is the rate-limiting step in mitochondrial NADH generation and therefore oxidative phosphorylation. This suggestion has been substantiated by a combined <sup>31</sup>P NMR and NADH fluorescence study in which increased O<sub>2</sub> consumption was associated with increased mitochondrial NAD(P)H levels and not with P<sub>i</sub>, PCr, ATP or ADP (55).

When varying the rate pressure product over a wide range from  $5 \cdot 10^3$  to  $25 \cdot 10^3$  mmHg/min, Balaban et al. (8) also did not observe a change in PCr/ATP ratio in in vivo paced dog hearts using <sup>31</sup>P NMR surface coil spectroscopy with a catheter probe. From the constant PCr/ATP ratio a constant [ADP] is inferred. Chance et al. (23) studied transfer functions, which relate work and substrate concentrations of (possible) control chemicals, in various tissues in situ. They observed in dog hearts a very steep transfer function when relating work and [ADP], in line with other studies in this species (8,70). These results again suggest a respiratory regulation by mitochondrial NADH. However, this steep transfer function is not found in hearts in general since in neoate lamb (23) and cat (70) hearts a far less steep relationship between work and [ADP] is observed. The authors suggest that these species differences may be related to adaptation to different life-styles, i.e., an endurance-performance adaptation for dogs vs. a stalking and sprinting adaptation for cats.

When comparing the results in refs. 8 and 35, it is found that the [ADP] in dog hearts in vivo is comparable to the [ADP] in isolated rat hearts perfused with glucose/insulin. From et al. (35) presented preliminary data showing that fatty-acid perfused hearts have ADP levels that are lower than those in glucose-perfused hearts, but not sufficiently low to become rate limiting as is the case in pyruvateperfused hearts. Since fatty acids are considered to be the major carbon source for myocardial energy metabolism in vivo, a similar situation may be found for the heart in situ. The fact that different substrates may lead to different ratios of high-energy phosphates and  $P_i$  has been recognized before (79), but in addition, Zweier and Jacobus (106) observed an increase in contractility when switching from glucose to pyruvate as substrate. Energy Metabolism Probed with NMR

<u>Magnetization transfer</u>. Where [high-energy phosphates] do not change upon increased cardiac performance, there is still a well-established coupling between cardiac performance and myocardial  $O_2$  consumption and therefore net ATP synthesis (59). As mentioned before, magnetization transfer methods can be be used to measure fluxes through the enzymatic reactions which are involved in energy production, like the CK reaction:

PCr + ADP + H<sup>+</sup> 
$$\stackrel{\text{KI}}{\longleftarrow}$$
 ATP + Cr  
<sup>k-1</sup>  
and the ATP synthetase reaction: P<sub>i</sub> + ADP  $\stackrel{\text{k2}}{\longleftarrow}$  ATP.

. .

The magnetization transfer method (30) will be briefly outlined for the simplest case of a two-site exchange process, illustrated by the exchange of phosphate between PCr and gamma-ATP. In this case the CK reaction is treated as a pseudo first-order reaction by assuming [ADP],  $[H^+]$  and [Cr] to be constant, resulting in:

PCr 
$$\stackrel{k_{for}}{\longleftarrow}$$
 gamma-ATP

in which  $k_{for} = k_1 \cdot [ADP] \cdot [H^+]$  and  $k_{rev} = k_2 \cdot [Cr]$ . The flux in the forward direction is defined as  $F_f = k_{for} \cdot [PCr]$  and in the reverse direction as  $F_r = k_{rev} \cdot [gamma-ATP]$ . The equilibrium magnetizations of PCr and gamma-ATP are symbolized as  $M^o_{PCr}$  and  $M^o_{gamma-ATP}$ . When perturbed, the magnetization of, e.g., PCr  $(M_{PCr})$  returns to its equilibrium situation, when we assume the chemical exchange to be absent, as described by the following equation:

$$dM_{PCr}/dt = (M^{\circ}_{PCr} - M_{PCr})/(T_{1 PCr})$$

in which t = time and  $T_{1 PCr}$  is the intrinsic relaxation time  $T_1$  of PCr that would be observed in the absence of exchange. In the presence of exchange, this equation must be modified. PCr loses magnetization through transfer of phosphate groups to ADP to form ATP and gains magnetization from the reverse process which results in:

$$dM_{PCr}/dt = (M^{o}_{PCr} - M_{PCr})/(T_{1 PCr}) - k_{for} \cdot M_{PCr} + k_{rev} \cdot M_{gamma-ATP}.$$

By applying selective irradiation of a very narrow bandwidth to the position of the gamma-ATP resonance for a certain period of time, as illustrated in Fig. 3, it is possible to selectively saturate or nullify the gamma-ATP magnetization, resulting in:

$$dM_{PCr}/dt = (M^{o}_{PCr} - M_{PCr})/(T_{1 PCr}) - k_{for} \cdot M_{PCr}$$

In a steady-state situation, where  $dM_{PCr}/dt = 0$ , it therefore appears that

$$M_{PCr}^{s} = (M_{PCr}^{o})/(1 + k_{for} \cdot T_{1 PCr}),$$

which means that in the presence of saturation of the gamma-ATP resonance the magnetization or signal intensity of PCr ( $M_{PCr}^{s}$ ) is no longer equal to  $M_{PCr}^{o}$  but is reduced by a factor  $1/(1 + k_{for} \cdot T_{1 PCr})$  as can be seen in Fig. 3. After perturbation  $M_{PCr}$  returns to its equilibrium value according to an effective  $T_{1}$ , given by:

$$T_{1 \text{ eff PCr}} = (T_{1 \text{ PCr}}) \cdot M^{s}_{PCr}/M^{o}_{PCr}$$

The rate constant  $k_{for}$  can then be derived from:

$$k_{for} = (1 - M^{s}_{PCr}/M^{o}_{PCr})/(T_{1 \text{ eff } PCr}).$$

 $T_{1 \text{ eff PCr}}$  and  $M^{s}_{PCr}$ , the magnetization of PCr in the presence of long saturation of gamma-ATP, can by obtained from a so-called progressive saturation  $T_{1}$ measurement (34) in which  $T_{1}$  is derived from a series of measurements with repetitive pulses with varying interpulse delays, in the presence of irradiation of gamma-ATP.  $M^{o}_{PCr}$  can be obtained from a control spectrum in the absence of saturation.

 $k_{rev}$  can be obtained in an analogous way by selective irradiation of PCr and substitution of PCr by gamma-ATP and vice versa in the above equations and considerations. Since the complete measurement of both  $k_{for}$  and  $k_{rev}$  with  $T_1$ 's is a lengthy procedure, which requires at least 30 min, depending on magnetic field strength and heart size, intracellular enzyme kinetics can only be studied in hearts in a metabolic steady state.

With magnetization transfer methods, fluxes in the forward direction were found to be larger than in the reverse direction (77,86). Initially, these results were interpreted as an indication for compartmentation (86) and support for the "shuttle"



Fig. 3. <sup>31</sup>P NMR spectra of an isolated rabbit heart. Spectrum A represents 128 scans with a recycle time of 10 s. Spectrum B represents 128 scans with selective irradiation on gamma-ATP during 9.6 s in between scans. Spectrum C represents the difference between A and B.

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model of the CK reaction (9). However, as pointed out by several investigators (26,77), PCr is a substrate unique for CK, but ATP participates in many more reactions. This leads to an underestimation of the reverse flux. When pyruvate or glucose/insulin was used as substrate at high workloads, substantial differences in forward and reverse fluxes were present, whereas with glucose alone at lower workloads fluxes were equal (102). By using a multiple saturation transfer method in which both PCr and P<sub>i</sub> were saturated, including all reactions in which ATP hydrolysis is involved, Ugurbil et al. (102) could fully account for the observed discrepancies. However, Koretsky et al. (60) mentioned that this approach is not succesful in rat hearts in situ. In addition, they measured the influence of the adenylate kinase reaction by simultaneous irradiation of the beta-ATP resonance, but concluded that the activity of this reaction is too low to account for the observed discrepancy in forward and reverse fluxes in the heart in situ. Therefore, they suggested that for the heart in situ, ATP compartmentation is still involved.

Regulation of CK fluxes in relation to cardiac performance is still a matter of controversy. In isolated rat hearts, Matthews et al. (77) found little change in CK forward flux upon increasing cardiac performance, but observed a considerable change in PCr/ATP ratio, which was compensated by changes in  $k_{for}$ . In contrast, Bittl and Ingwall (12) reported a 3.5-fold increase in forward CK flux when changing from K<sup>+</sup>-arrest to a rate-pressure product (RPP) of 44.7 · 10<sup>3</sup> mmHg/min for isolated rat heart. Yet they observed very little change in PCr/ATP ratio, but increased flux was primarily caused by increased  $k_{for}$ . Also in the heart in situ, they observed a comparable relation between forward flux and RPP (10). Kupriyanov et al. (63) studied the same range of RPP (12), but found only a 1.5-fold increase in forward flux in isolated rat hearts, in agreement with earlier results (77). Using isolated rabbit hearts, we recently did not observe significant dependence of forward CK flux on RPP although necessarily a smaller RPP range was studied than in the rat heart studies. Since experimental conditions vary between the different studies cited, no firm conclusions can yet be reached concerning the relation of CK flux and cardiac performance.

Bittl and Ingwall (12) have interpreted their results as evidence for a direct coupling between energy production and energy transfer. However, Shoubridge et al. (96) reached a different conclusion. They depleted rat hearts of PCr by feeding rats a creatine analogue, which is not a substrate for CK. These hearts could perform normally despite a 90% depletion of PCr. Therefore, PCr does not seem an obligate intermediate of energy transduction in the heart. Fossel and Hoefeler (32) found with a similar approach (CK inhibition) that only at high workload isolated rat hearts were unable to maintain their peak-systolic and end-diastolic pressures.

ATP-synthesis rates have also been studied using magnetization transfer methods (76), despite the greater technical difficulty due to the relatively low  $P_i$  content of the heart. Kingsley-Hickman et al. (56) compared ATP synthesis rate and  $O_2$  consumption and found a surprisingly high P:O ratio of 6.3. They explained their results by invoking a unidirectional ATP synthesis which exceeded the net ATP synthesis by a factor of 2. However, Brindle and Radda (20) suggested that the glycolytic enzymes glyceraldehyde-3-phosphate dehydrogenase and phosphoglycerate kinase could also catalyze a significant  $P_i$ -ATP exchange in addition to the mitochondrial ATP-synthase. In a subsequent study in which these enzymes were inhibited, Kingsley-Hickman et al. (57) found a P:O ratio of 2.36 and concluded that the mitochondrial ATP-synthase works essentially unidirectionally.

Studies on human heart. Studies of human cardiac metabolism have been few and merely demonstrate the possibility to obtain localized <sup>31</sup>P NMR spectra of human heart (13,16). Only one study has been reported where <sup>31</sup>P NMR is used for diagnosis and therapeutic evaluation of a pediatric case of cardiomyopathy (103).
Since the technical difficulties in obtaining <sup>31</sup>P NMR spectra of human skeletal muscle are far less than for cardiac muscle, a number of investigators has studied skeletal muscle metabolism in patients with congestive heart failure (22,75,104). They found metabolic alterations which did not appear to be merely related to reduced blood flow, but also to impaired substrate availability and altered biochemistry. However, in the near future, <sup>31</sup>P NMR will also be used to study *myocardial* metabolism of patients with congestive heart failure and other heart diseases and will become an increasingly important tool for clinical investigation.

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# Chapter 13

# Positron Emission Tomography and Myocardial Biochemistry

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Positron emission tomography (PET) is a new method for the external quantification of regional myocardial blood flow, substrate fluxes and biochemical reaction rates. It takes advantage of physiologic tracers, tracer kinetic principles and the quantitative imaging capabilities of the tomograph. To date, the technique has been used primarily for the study of ischemic heart disease and cardiomyopathies. This report summarizes experimental and clinical investigations on substrate metabolism during the process of ischemic injury. Uptake of C-11 palmitate and F-18 2-fluoro-2deoxyglucose appear to be well-suited markers for the study of myocardial fatty acid and glucose metabolism, respectively. Observations with PET have shown noninvasively in vivo the known impairment of fatty acid oxidation and the enhanced glycolytic flux during ischemia. Some distinct findings unexpected from existing biochemical knowledge will be highlighted.

# Introduction

Positron emission tomography (PET) offers the opportunity for studying noninvasively regional myocardial tissue function. The technique owes its unique capabilities to several features which will be briefly outlined. For more detail on physics and instrumentation, the reader is referred to excellent recent reviews (16,29). This imaging device relies on the emission (in diametrically opposed directions) of two 511 keV photons during positron annihilation. Registration and localization of the annihilation event by a ring of external detectors occur through coincidence detection which allows electronic collimation and correction for photon attenuation. The cross-sectional images produced by the tomograph after appropriate reconstruction reflect a quantitative measurement of regional radioactivity.

The physiologic tracers used with PET are compounds labeled with positron-emitting isotopes of elements that are abundantly present in vivo, such as carbon-11, fluorine-18, nitrogen-13 or oxygen-15. Labeled substrates, substrate analogs and drugs can be synthesized in high specific activity without disturbing their biological properties. The local tissue concentrations of the tracers are in the pico- to nanomolar range and, therefore, they do not interfere with the process to be studied.

Combining these physiologic tracers with the quantitative capabilities of PET allows one to image the distribution, uptake and turnover of tracer in the myocardium. If a kinetic model correctly describing the tissue kinetics of the tracer is available, quantitative measurements of regional blood flow in ml/min/g tissue or myocardial substrates fluxes in mmol/min/g tissue can be obtained noninvasively. A successful example is the application to the myocardium of the operational equation initially developed to measure exogenous glucose utilization rates in the brain with the glucose analog F-18 2-fluoro-2-deoxyglucose or FDG (30,48). This agent traces the transmembraneous exchange of glucose and the hexokinase-mediated phosphorylation to glucose-6-phosphate. After phosphorylation to FDG-6-phosphate, the tracer becomes trapped in the cell. The kinetics of FDG in tissue are described by a three-compartment model, validated in isolated perfused rabbit myocardium (20), in canine hearts (31), and more recently in humans (42). As shown in Fig. 1, the estimates of exogenous glucose utilization determined externally with PET and the FDG model agreed well with those calculated by the Fick method (31).

Thus, PET appears potentially useful as an in vivo biochemical assay technique that will permit examination and validation of functional processes that have been described primarily with in vitro experiments. Although there is continuing interest in receptor binding studies (49,50) and quantification of regional myocardial blood flow (2,12,35,46), we will limit this review to the most intriguing application of PET which in our view relates to the study of myocardial substrate metabolism. The reader interested in the entire scope of PET studies of the heart is referred to recent detailed monographs (1,34,37).

#### Normal Alterations in Substrate Metabolism Demonstrated by PET

Unlike other organs, such as the brain, the heart meets its energy demands oxidizing several substrates (17,18). Selection of fatty acids (FA), glucose, lactic acid, ketone bodies and amino acids for oxidation depends largely upon their arterial concentrations which in turn are regulated by the dietary state, by physical activity and by hormones. For example, plasma FA levels are high in the fasting state (0.9 mM). Oxidation of FA therefore accounts for about half of the myocardial oxygen consumption. Arterial lactate levels often rise from 0.9 mM at rest up to 9.5 mM during exercise; lactate then becomes the major substrate for oxidation. Uptake and utilization rates of various substrates rapidly respond to changing demand in order to meet instantaneously changes in energy requirements of the cardiac muscle.

The heart's ability to increase substrate oxidation in response to greater demand and to resort to alternate substrates have been demonstrated noninvasively in vivo with PET. Fasted subjects were studied first at control and then again during states



Fig. 1. Utilization of exogenous glucose by canine myocardium measured by the Fick method and externally with the fluorodeoxyglucose (FDG) model and PET. LC = lumped constant. From Ratib et al. (31), with permission.

of increased cardiac work induced by rapid atrial pacing while plasma substrate concentrations remained unchanged (14). Compared to control, the expected increase in FA oxidation was observed using serial PET imaging and C-11 palmitic acid (CPA) as a tracer for FA metabolism. Although a tracer kinetic model for CPA is not available, its tissue kinetics have been extensively documented (21,22,38,39). Normal myocardium avidly extracts this long chain FA (16 carbon); the first transit retention fraction averages 65% (38,56). Maximal myocardial C-11 activity tissue concentrations are attained within 3 to 4 min. Thereafter, the activity typically clears from normal myocardium in a biexponential fashion, indicating fractional distribution of the tracer between two major pools of different sizes and with different turnover rates (Fig. 2). This is consistent with the known dual fate of FA (19,53): the late slow turnover pool is related to the endogenous lipid pool, consisting mainly of triglycerides. The early rapid phase of the time - activity curve relates to the release of C-11 carbon dioxide which is the end product of CPA oxidation (56). In the study by Henze et al. (14), during pacing at rates above 100 beats/min, the relative size of the early rapid phase increased while the relative size of the late slow phase declined. Similarly, the steepness of the slope of the early rapid phase increased while the relative oxidative pathways and was oxidized more rapidly.

The second entity demonstrated in vivo by PET studies is the heart's normal ability to alternate between substrates in response to changes in substrate concentrations in plasma. With combined use of tracers of glucose (FDG) and FA (CPA) metabolism, the UCLA group has documented the effect of substrate availability on substrate selection by normal myocardium. This was shown initially in open-chest dogs studied first after an overnight fast and then again after glucose/insulin infusion (32). Despite comparable blood flows and myocardial oxygen consumption, the fractional distribution of CPA in cardiac muscle changed in response to altered plasma substrate concentration. For example, the relative size of the early rapid clearance phase and its clearance rate declined or this curve component disappeared entirely when plasma FA levels were lowered and glucose and lactic acid concentrations increased by a factor two (14). This was accompanied by lesser amounts of C-11 carbon dioxide released from myocardium, indicating that less CPA oxidized. Observations in human myocardium, though still small in number, are similar (14) and consistent with the notion that as FA availability declines, the myocardium switches to alternate substrates (e.g., glucose and lactic acid).



Fig. 2. C-11 time activity curve derived from serial PET images after intravenous C-11 palmitic acid (CPA) injection. Note the peak activity (A) and the subsequent biexponential tissue clearance. The late phase is extrapolated back to the time (A) and from the intercept B, the fraction of CPA entering the early and late phase can be estimated. Biexponential fitting also provides the halftime (T 1/2) of each curve component.

#### Metabolic Studies in Myocardial Ischemia and Postischemic Dysfunction

The sequence of metabolic events initiated by ischemia has been extensively studied in invasive animal models or in vitro experimental systems (23,27,28). It appears that the amount of residual blood flow determines whether residual metabolic activity is maintained or whether ischemia will proceed to irreversible tissue injury. As oxygen availability decreases, the flux of FA through the beta-oxidative spiral falls. This provides substrate for esterification of FA with alpha-glycerophosphate resulting in increased triglyceride synthesis. Thus, the onset of ischemia is usually associated with enhanced glycolysis. The glucose is derived from endogenous (i.e., glycogen) and/or from exogenous sources. As ischemia progresses and tricarboxylic acid cycle activity decreases, pyruvate is converted to lactate and released from myocardium. Anaerobic glycolysis remains then the only source of ATP. The rate of lactic production and removal from the cell determines how long and at what rate glycolysis can be maintained.

Because beta-oxidation is most sensitive to oxygen deprivation, initial studies with PET used CPA as a tracer for identifying ischemia.

In animal experiments, both low-flow and high-demand ischemia affected the fractional distribution of CPA in tissue (22,33,39). The fraction immediately entering oxidative pathways decreased, its clearance rate from myocardium declined and a greater fraction was deposited in the endogenous lipid pool. These changes were seen as a consequence of an impaired transfer of palmitoyl-CoA into mitochondria and of greater "trapping" of CPA in the triglyceride pool. Also, in patients with high-demand ischemia induced by moderate atrial pacing, abnormal kinetics of CPA were observed in segments supplied by stenosed coronary arteries (13). The expected increase in tracer entering the rapid curve component and its clearance rate observed in normal myocardium, was attenuated in segments subtended by stenosed coronary arteries. Interestingly, these metabolic abnormalities occurred without electrocardiographic, or in some instances, functional (by two-dimensional echocardiography) correlates of acute myocardial ischemia.

These studies show the ability of the CPA approach to externally detect expected changes in FA metabolism as well as the sensitivity of the technique. Yet, some limitations should be recognized.

First of all, as discussed earlier, in the absence of an appropriate tracer kinetic model, studies with CPA remain largely qualitative.

Second, the initial uptake of CPA depends on the delivery, i.e., on blood flow. Thereafter, the activity clears rapidly from myocardium. This limits the usefulness of this tracer in severe ischemia because regional blood flow is then markedly reduced. Limited uptake of the tracer will preclude derivation of statistically adequate data, needed for construction of reliable tissue time - activity curves.

Third, Fox et al. (10) have shown that during severe ischemia and hypoxia, about 40 to 50% of the C-11 activity released from myocardium into the coronary sinus is represented by unaltered non-metabolized CPA. Back-diffusion competes with metabolic sequestration of the tracer, suggesting a block of the energy-requiring activation step of CPA to palmitoyl-CoA. This factor will confound the use of the tissue clearance-rate of C-11 activity during the early rapid phase of CPA kinetics.

These limitations prompted many investigators to resort to FDG for the detection of exogenous glucose utilization which is enhanced in ischemia and which presents itself on the cross-sectional image as a positive signal.

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Increases in FDG uptake were demonstrated in the isolated arterially perfused rabbit septum (24) as well as noninvasively in dogs with coronary stenoses and rapid atrial pacing (36). Findings in patients are similar. Camici et al. (6) studied patients with known coronary disease during and after exercise induced ischemia. FDG uptake was preserved in myocardial segments that were inadequately perfused during exercise. When injected after exercise, absolute FDG uptake was increased (compared to normal segments). The authors suggested that glucose utilization remained high in postischemic myocardium as a consequence of restoration of glycogen stores (7).

Following on this observation, several groups examined the recovery of metabolism from transient ischemic insult (11,45,55). Short periods of transient ischemia such as, in dogs, a complete coronary occlusion for 20 min or a critical stenosis for one hour resulted in prolonged recovery of FA metabolism despite return to baseline of myocardial blood flow (Figs. 3 and 4). Interestingly, the recovery of metabolism paralleled the slow recovery of regional function, a concept frequently referred to as "myocardial stunning" (3,15). Recent observations in postischemic myocardium using the tissue clearance of C-11 acetate as a marker for tricarboxylic cycle activity showed impaired FA oxidation in segments with nearly normal C-11 acetate kinetics (4,5). These findings in stunned myocardium suggest not only that repair of the biochemical machinery after ischemic damage takes time but also that different metabolic pathways may recover at different rates. Why glucose utilization remains predominant at the expense of FA oxidation during reflow is unclear but may reflect a prolonged "stunning" of beta-oxidation or the residual inhibitory influence of FA intermediates.



Fig. 3. PET imaging with C-11 palmitic acid (CPA) in a dog 24 h after a transient ischemic insult (1 h critical circumflex stenosis reducing transmural blood flow to 40% of control). Note the delayed clearance of activity from the lateral and posterior walls (arrows). The time of acquisition after intravenous CPA injection is indicated under each image.

#### Myocardial Infarction and Reperfusion: Metabolism as an Indicator of Tissue Viability

Estimates of size and mass of infarcted myocardium were obtained with PET and CPA in dog experiments (54) and in patients with chronic infarction (47,51). Tomographic estimates of infarct size by planimetry of the segmental defect in CPA uptake correlated well (r = 0.92) with enzymatic estimates by serial creatine kinase serum sampling.

More recently, the widespread use of thrombolytic agents for restoring blood flow in evolving myocardial infarction raised the interest in the metabolic fate of reperfused tissue. Early identification of reversible versus irreversible injury is critical for patient management.

In a canine model of reperfused infarction, ischemic but viable myocardium was characterized by a persistent glucose uptake which was absent in necrotic tissue, as shown in Fig. 5 (26). Schwaiger et al. (44) examined the recovery of metabolic abnormalities over a 4-wk period in dogs after a 3-h occlusion of the left anterior descending coronary artery followed by reperfusion. Enhanced glucose utilization and impaired fatty acid oxidation persisted for hours to days. Again, as these abnormalities disappeared, segmental function improved. In order to elucidate the mechanisms of increased glucose utilization in reperfused myocardium, the relative contribution of oxidative and non-oxidative glucose metabolism was studied by arterio-venous sampling for C-14 carbon dioxide and C-14 lactate after infusion of C-14 labeled glucose (43). The data show increased glucose utilization of which 75% entered glycolysis, most of it being released as lactate. This suggests that the energy produced from non-oxidative glucose metabolism is sufficient to maintain cellular viability but insufficient to allow a normal mechanical function.

Encouraged by these experimental findings, several groups started to submit patients with recent myocardial infaction to PET studies (8,9,25,40,41).



Fig. 4. Uptake and clearance of C-11 palmitic acid (CPA) in stunned myocardium, 24 h after critical stenosis of dog circumflex. The two curves were obtained from the study shown in Fig. 3. As compared with the normal septal and anterior walls, clearance of C-11 activity is delayed in the posterior and lateral walls. The halftime of the early phase is increased from 4.2 min in the normal zone to 7.2 min in the stunned zone. The tracer fraction entering the late phase is increased to 21% in the stunned as compared to 16% in the normal zone.

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Marshall et al. (25) compared regional glucose utilization and blood flow in 15 patients studied an average of two weeks after myocardial infarction. Different metabolic patterns were defined. In normal myocardium, blood flow and FDG uptake match closely. A marker of blood flow (i.e., N-13 ammonia) is used to identify the abnormal segment, and FDG to distinguish between presence and absence of metabolism in segments with reduced blood flow. In infarcted segments, flow and metabolism are concordingly decreased. By contrast, ischemic myocardium is characterized by a "mismatch", i.e., an increase in exogenous glucose utilization relative to blood flow. Segments with "mismatches" were present mainly in patients with post-infarction angina and more extensive disease. They corresponded in location to the sites of electrocardiographic changes and wall motion abnormalities occurring during anginal episodes. Consequently, 12 patients were studied <72 h after onset of symptoms (40,41). The same "mismatch" pattern was observed in about half of the segments indicating the presence of injured but viable myocardium in the infarction area for prolonged periods after the acute event.

These are important findings as PET could help in selecting patients for more aggressive therapeutic approaches. Reperfusion of an area of scar is not likely to be of much benefit; revascularization of ischemic tissue should improve segmental left ventricular function, as demonstrated recently in a study on the effects of coronary artery bypass grafting (52).

#### Conclusions

This review attempts to describe some of the features of PET as a tool for probing myocardial regional biochemistry. Observations by PET made in human heart are consistent with the knowledge on biochemistry derived from experimental in vitro systems. For the first time, PET offers the opportunity to test the validity of this knowledge directly in the living human heart. On the other hand, observations in patients showed some distinct differences to experimental findings. For example, augmented glycolytic flux persists in ischemically injured myocardium for longer periods than expected. Also, transient reductions in oxygen delivery to myocardium do promote prolonged impairments in FA utilization despite reoxygenation. The detection of tissue viability appears nowadays as a potentially clinically useful application of PET. However, these clinical investigations have remained largely qualitative and major aspects of myocardial metabolism such as protein synthesis and tricarboxylic acid cycle activity await further investigation.



Fig. 5. Myocardial glucose utilization rates (MR-Glc) are calculated by the fluorodeoxyglucose (FDG) model in tissue samples characterized as normal, ischemic but viable, or necrotic by histochemistry, electron microscopy and blood flow. Measurements are performed 4 h after reperfusion in dogs subjected to a 2-h period of coronary occlusion. Depressed glucose uptake indicates irreversible injury while normal FDG uptake is associated with viable tissue (\*\*p<0.01; \*\*\*p<0.001). From Melin et al. (26), with permission. TTC = triphenyl tetrazolium chloride.

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# Chapter 14

# Energy Metabolism and Transport in Neonatal Heart Cells in Culture

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Heart cells in culture are increasingly being used in studies at the cellular level in many fields of cardiac biology and pathology. This chapter reviews the metabolism of the energy-producing substrates - glucose and fatty acids (FA's) - pinpointing the findings in cultured heart cells and their contribution to understand uptake and subsequent distribution of these metabolites. The roles of glucose and FA's in energy production and balance have been determined mainly from studies of  ${}^{14}CO_2$  production from labelled substrates. Oxidation of glucose and FA's is incomplete if either of these substrates is present in excess, indicating that substrate utilization is controlled by a cellular mechanism. In addition, FA's in cellular reserves seem to be preferentially utilized over newly incorporated ones, implying that cellular lipids undergo continuous renewal. The recently developed technique for measuring oxygen consumption without disrupting the energy demands of the cardiomyocyte. The regulation of high-energy phosphate production and its transport from the mitochondria to the myofibrils via the creatine kinase system are also discussed in this chapter. Cultured heart cells have been used for studying oxygen deprivation. Under suitable conditions, such systems provide information on the sequence of events at the cellular level during anoxia. Kinetic studies of high-energy phosphate depletion and concomitant nucleoside and oxypurine formation during anoxic injury are presented. The repair mechanism via reoxygenation and reperfusion deserves detailed investigation.

# 1. Introduction

The relationship between cellular chemical energy and mechanical work in muscles is one of the most fascinating links between the classical disciplines of biochemistry and physiology. This chapter will be limited to studies on energy metabolism in cultures of neonatal heart cells, pinpointing findings that have furthered understanding of cardiac energy metabolism. The first part will deal with substrate utilization; the second, with certain aspects of high-energy phosphate metabolism and transport.

The heart-cell culture system is a stable preparation, in which both hormonal influences and diffusion barriers are absent. In addition, the composition of the extracellular environment may be controlled in this system, which allows investigations at the cellular level to be carried out. The cells in culture must, however, adapt to the new substratum and a continually changing environment owing to frequent changes of medium, with catabolic products that are never completely removed from the vicinity of the cells. In addition, cultured heart cells, which clearly resemble those in situ in their ability to contract spontaneously and rhythmically, are not required to act against the frictional resistance of blood vessels. Therefore, their energy requirements correspond to those for isotonic contractions. Consequently, there are less mitochondria in cultured cells than in cells in vivo. In spite of the reservations mentioned above, the basic metabolism, structure and electrophysiological properties of cardiomyocytes resemble those of the intact heart. Thus, studies with cultured heart cells, in which the fate of a substrate can easily be followed, have certainly extended our knowledge of cardiac physiology and biochemistry.

#### 2. Lipid Metabolism

There is a well-established dual role for lipids in the heart: a) they are the major functional-structural component of the sarcolemma, and b) they also serve as the major energy source of the myocardium, primarily an oxidative organ. For this reason, storage of lipids is much more important than that of glycogen, since they provide a much more efficient energy source (15,45).

Quantitative studies of heart metabolism in situ were greatly advanced when cardiac catheterization techniques were devised. Cardiac  $O_2$  consumption was shown to be too high to be accounted for by the oxidation of glucose, lactate and pyruvate (4,6). Non-esterified fatty acids (NEFA's) were shown to be the major energy-providing substrate in the heart (5,33).

#### Fatty Acid Uptake and Distribution

Fatty acids (FA's) are transported in the plasma in the form of complexes with albumin, which has several high-affinity FA binding sites. Less than 1% of the FA's circulates in the free state or in loosely-bound forms (110,132,135).

In cultured heart cells (28,76,77,101), two phases occur in the uptake of labelled FA's: a) Passive diffusion, predominating at FA levels of >200 uM; b) A saturable specific binding process ( $K_m$  about 10 uM). FA uptake in cardiomyocytes - as in liver cells - is dependent on the total [FA] (both free and albumin bound refs. 101,132). The various long-chain FA's compete for uptake. Since a specific binding protein could not be demonstrated, the transport system is probably located on the cell surface (101). Maximal uptake of palmitate occurs at palmitate: albumin ratios ranging from 7 to 10, differing markedly from the in vivo value of about 2. Albumin enhances both uptake and subsequent esterification of palmitate, but reduces that of oleate (76). This is due to binding of most palmitate molecules to low- and medium-affinity binding sites in albumin. With oleate (an unsaturated FA), albumin acts as an alternative uptake site to cardiac cells, thus lowering oleate uptake into cardiomyocytes (76). The uptake process is not energy dependent, but is probably facilitated (77).

Following uptake, FA's are immediately metabolized in the sarcoplasmic reticulum (20,113). A cytosolic protein of mol. wt. 12,000 that binds FA acyl-CoA has been demonstrated (63,64). It suppresses the detergent-like properties of acyl-CoA.

Detailed kinetic investigations of FA pathways have recently been carried out by our group (8). We found that most of the FA's were immediately esterified, NEFA only accounting for 5 to 10% of the the total FA incorporated. Diglycerides did not accumulate in this system, indicating that the FA-conversion rate exceeds uptake. On depletion of the cells of energy-providing substrates and uncoupling oxidative phosphorylation, FA esterification was inhibited by 80% but not accompanied by FA accumulation in the cells. Impairment of esterification led to export of both diglycerides and FA's. Under normal conditions, FA incorporation into triglycerides and phospholipids was about equal. Most of the triglycerides were located in the 100,000 g supernatant fraction, whereas phospholipids were generally associated with the particulate and membranous fractions.

#### Fatty Acid Oxidation

The mitochondrial route is usually considered the major pathway for the metabolism of exogenous FA, indicating that exogenous, and not endogenous FA's, are preferentially used for energy supply. However, some reports suggest that this is not the case - most FA's were found to be in the esterified form shortly after uptake (8,113). Also in skeletal muscle, FA's are not oxidized directly, but are first converted to the esterified form and generally stored in the vicinity of the mitochondria as lipid droplets (18,111,141).

Based on studies with perfused hearts, Shipp et al. (108) suggested that endogenous lipids provide energy, although it was not clear whether phospholipids (108) or triglycerides (72,73) are preferred. We demonstrated in cultured heart cells that endogenous lipids are preferentially used over FA's that have just entered the cell (82,83).

To undergo oxidation acyl-CoA thioesters are transported into the mitochondria after conversion to acylcarnitine at the outer mitochondrial membrane. On reaching the inner membrane, the carnitine is exchanged with the inner mitochondrial membrane carnitine by a transferase, the acyl group then being transferred to the mitochondrial matrix CoA and finally, beta-oxidation occurs (89,90,92).

In cardiomyocytes there are conflicting data regarding the importance of FA oxidation as compared to that of other substrates. According to Ross and McCarl (97), glucose, pyruvate and lactate contribute more to energy production than palmitate. Others (112) reported that more energy is derived from glutamine than from either glucose or FA's. The shift from lipid to carbohydrate metabolism in aging cultures (32) is thought to be due to low intracellular levels of the enzymes involved in FA metabolism (37,38) or to a postulated hypoxic glycolytic pathway in such cells (95). However, according to our data (30,82), FA's are the major energy source for cardiac myocytes; the decrease in FA oxidation upon aging could be eliminated by preincubating the cells with palmitate. The variation in cell populations in culture as a function of age may also be of significance (136). Clearly, this topic warrants reexamination in the light of these findings.

FA degradation occurs primarily via beta-oxidation involving four sequential reactions (47). FA acyl-CoA's of chain-lengths from  $C_4$  to  $C_{18}$  undergo oxidation via the "oxidase". Short FA intermediates of palmitate and stearate oxidation could only be found under special conditions, e.g.,  $O_2$  restriction (60,114). There were no differences in the oxidation rates (i.e., the amount of  ${}^{14}CO_2$  released) from  ${}^{14}C$ -palmitate labelled at either at C 1, 6 or 11, fed to intact rats (131) or given to perfused hearts (27,88).

Long-chain FA's are completely oxidized to  $CO_2$  in cultured heart cells (101). However, in agreement with earlier findings (88), we have reported a striking difference in the production of  ${}^{14}CO_2$  from either 1- ${}^{14}C$ - or 16- ${}^{14}C$ -palmitate in cultured heart cells (82,83), the release being several fold higher from the former. This implies that the two termini of the palmitate molecule have different catabolic fates (see ref. 39).

In hepatocytes total oxidation of  $1^{-14}$ C- or  $16^{-14}$ C-palmitate was almost identical, but considerably less  ${}^{14}$ CO<sub>2</sub> was produced from  $16^{-14}$ C-palmitate (16). These data imply the preferential incorporation of omega-C<sub>2</sub> units into ketone bodies (see ref.

9). If this also holds for heart cells, reutilization of the ketone bodies would be expected, since these are the energy source of choice in the myocardium (133).

Our studies revealed considerable release into the medium of water-soluble products from the <sup>14</sup>C-methyl terminus of palmitate (82,83,84). This extracellular radioactivity persisted even after long incubation periods, indicating that the released product did not undergo further metabolism. The metabolite was identified as 3-hydroxybutyrate (83,84), and by isomer separation (61), 80% of the radioactivity was shown to be associated with the L(+)-isomer.

The CoA derivative of L(+)-3-hydroxybutyrate is an intermediate in long-chain FA oxidation (128). It is thought that the L(+)-isomer of 3-hydroxybutyrate participates in beta-oxidation attached to the enzyme complex as the CoA derivative, while free 3-hydroxybutyrate released into the body fluids is the D(-)form. Although some studies indicated that mammalian tissues could metabolize L(+)-3-hydroxybutyrate, this route was thought to be unimportant (54,55). Furthermore, mitochondrial 3-hydroxybutyrate dehydrogenase, which oxidizes 3-hydroxybutyrate to acetoacetate, has an absolute stereospecificity for the D(-)-isomer (56). Based on the uneven distribution of radioactivity in both the  ${}^{14}CO_2$  and in the water-soluble 3-hydroxybutyrate, we suggested that L(+)-hydroxybutyryl-CoA is detached and deacylated to form the free acid, which is transported across the membrane, contributing to the water-soluble, radioactive 3-hydroxybutyrate in the extracellular medium. It may, however, be metabolized at a very slow rate, as indicated by the <sup>14</sup>C-DL-3-hydroxybutyrate disappearance from the medium (83). Furthermore, other radioactive metabolites were also released into the medium, particularly from 1-14Cpalmitate. However, under such conditions, the total excreted radioactivity was considerably lower and was distributed among several compounds, one of which was identified as D-3-hydroxybutyrate. This is consistent with earlier reports (27,13) of high [3-hydroxybutyrate] in the extracellular compartment in perfused hearts. It is also possible that some of the early reports in which 3-hydroxybutyrate was chemically determined, in fact unknowingly dealt with the L(+)-isomer.

#### Abnormal Fatty Acid Metabolism

Erucic acid (EA, delta-13 docosenoic acid;  $C_{22:1}$ ) is the major fatty acid contained in rapeseed oils. An EA-rich diet causes an accumulation of EA-rich triglycerides in heart-muscle cells, which leads to cardiac lipidosis if fed to rats and other animals. We have shown that cardiac cells have a decreased ability to oxidize EA, as studied by using EA labelled at both the C-14 (on the double bond) and the carboxyl carbons. <sup>14</sup>CO<sub>2</sub> originating from the carboxyl group appears rapidly, whereas that derived from C-14 lagged behind for about 12 h (82,85,86). Thus, apparently EA cannot be directly oxidized in the mitochondrion, but undergoes first chain shortening. This occurs in two phases: first to C<sub>20:1</sub> and then to C<sub>18:1</sub> (86). The C<sub>18:1</sub> produced may then be undergo beta-oxidation. Chain shortening to a lesser degree also occurs with other FA's, as lower labelled homologues, such as C<sub>14:0</sub>, C<sub>12:0</sub> and C<sub>10:0</sub>, derived from 16-<sup>14</sup>C-palmitate could be detected in heart cell cultures (Pinson et al., unpublished).

#### 3. Carbohydrate Metabolism

# Glucose Uptake

Glucose-6-phoshate is at the crossroads of glucose metabolism, several different pathways being available from that point: glycolysis yielding pyruvate (and lactate) usually followed by complete oxidation via the citric acid cycle; glycogen synthesis via uridine diphosphoglucose; and the pentose-phosphate shunt. In the aerobic heart, glucose does not constitute an important energy source. Even under extremely high work load with glucose as the only substrate in the extracellular fluid, glucose oxidation only accounts for 40% of the energy required for contraction, the rest being derived from intracellular triglycerides (70,73).

In the only extensive study on glucose transport in cultured heart cells, Paris et al. (75) compared the metabolism of 3-O-methylglucose, which enters the cell but is not further metabolized, and 2-deoxyglucose, which undergoes phosphorylation by hexokinase but may not be subsequently used. They showed that the rate of transport is several-fold higher than the phosphorylation rate. The latter is limited by the rate of transport into the cell, reaching a steady state within a few min. It is dependent on the extracellular [carbohydrate]. The intracellular [glucose] is below the K<sub>m</sub> for hexokinase and, in turn, determines the phosphorylation rate, which is increased by countertransport and decreased by cytochalasin B. Thus, any changes in carbohydrate metabolism are reflected in the rate of 2-deoxyglucose phosphorylation. Glucose transport is not affected by FA's in the medium or by preloading the cells with FA's (75). These findings conflict with those reported by Neely et al. (68), who found that in the working perfused heart palmitate inhibited glucose uptake. This discrepancy is possibly due to the contribution of a regulation system in cultured heart cells, which responds to extracellular hexoses (75). The decreased transport in cultured cells following incubation with cycloheximide suggests that the synthesis of a carrier protein is involved and that its activity may be modulated.

Other studies (29,30) have pointed to the importance of the composition of the extracellular medium for glucose uptake. Supplementation of the medium with 20% serum resulted in glucose uptake at a constant rate of 3 umol/h/mg protein. The medium thus became depleted of glucose after 10 to 12 h, well before the next medium change. In the absence of serum, however, glucose was taken up at about half this rate (30). Serum in the medium stimulated both glucose transport and hexokinase activity (29). These rates of glucose uptake are an order of magnitude higher than those found in both perfused (30 umol/h/g of tissue, ref. 66) and in fetal heart (66 umol/h/g, ref. 17). Other investigators obtained data with cultured cells (about 1.5 umol/h/mg protein, refs. 74,95,140) which were very similar to the findings in perfused hearts. The differences may be attributed to variations in the distribution of various cell types since cardiomyocytes take up glucose at a much higher rate than non-muscle cells (97).

# Lactate Production

Glycolysis in the heart is regulated via the activities of phosphofructokinase and glyceraldehyde-3-phosphate dehydrogenase. The supply of glucose-6-phosphate, either following glucose uptake and phosphorylation or via glycogen degradation, is a crucial step in the pathway from glucose to pyruvate and lactate (69,90,92).

Under conditions of normal  $O_2$  supply, the heart does not produce lactate, glucose either being used for glycogen synthesis or undergoing complete oxidation. The lactate dehydrogenase (LDH)  $H_4$  isoenzyme pattern is present, and consequently lactate is oxidized to pyruvate. Lactate is only the end product of glucose degradation during  $O_2$  deprivation, when the energy is supplied to the ischemic area via glycolysis or in the partially anaerobic fetal heart.

Studies with cardiomyocytes have shown that most of the glucose incorporated is rapidly converted to lactate which is then released into the extracellular space (29-31,97). However, lactate production is not a result of  $O_2$  restriction in the culture medium (30). The released lactate is not derived from endogenous glycogen, but from newly incorporated glucose, and is not dependent on the [lactate] in the medium (30,31). However, serum in the medium promotes optimal glucose uptake and

lactate production (30). Non-muscle cells in culture only oxidize glucose to lactate, while cardiomyocytes also metabolize some glucose via the citric acid cycle (97). Thus, the amount of lactate produced also varies with the initial seeding density (129). Indeed, it is difficult to compare the amount of glucose converted to lactate in various experiments, since different methods employed for preparing cell cultures by various research groups affect the proportion of myocytes and non-muscle cells in the system. Our group (31) and Ross and McCarl (97) have reported similar values for lactate production, 40 umol/mg protein (more than 65% of the incorporated glucose) and 0.88 umol/h/mg protein, representing 75% of the glucose taken up from the medium, respectively - values which are 4-fold higher than Orloff and McCarl's data (74).

Lactate production may serve as a mechanism for removing excess glucose from cultured heart cells, since only a small amount is used in energy production and a mere 5% is stored as glycogen (30,31). The shift in the LDH isoenzyme pattern from the heart ( $H_4$ ) to the muscle type ( $M_4$ ) may reflect this (11). However, changes in the relative proportion of the different cell types in the culture cannot be ruled out (120). Randle et al. reported that in the non-working perfused heart significant amounts of glucose and pyruvate are converted to lactate (91). These findings are possibly related to the fact that cells in culture, in direct contrast to those in vivo, are not engaged in work against resistance. Only after the medium has been completely depleted of glucose and >65% has been transformed to lactate, does the [lactate] in the medium decrease in a linear fashion, by oxidation or by transamination (30,31,97).

#### Glucose Oxidation Pathways

The growth of heart cells in culture occurs in two distinct phases: an initial phase of cell division, the duration of which depends upon the plating density, until confluency is reached, and a non-dividing phase with continued cell growth (136). These two phases are reflected in the glucose oxidation via two pathways in cultured cells: a) the pentose-phosphate pathway, and b) glucose oxidation via the citric acid cycle and oxidative phosphorylation. The importance of both pathways has been studied in cultured cells - the oxidation of glucose has been estimated by measuring the release of  ${}^{14}CO_2$  from an appropriately labelled substrate, either directly in the culture flask (129), or by means of a simple multi-channel trapping device (82,87), or by the rather sophisticated "radiorespirometer" working along similar lines (98).

The *pentose-phosphate pathway* is very active in early embryonic heart and becomes less so as the heart approaches the adult form, when the division index decreases. The end product of this pathway, ribose-phosphate, is a precursor for nucleoside and subsequent DNA synthesis. Since ribose is formed from carbons 2 to 6 of the glucose molecule, the relative importance of each pathway has been studied by comparing the amount of  ${}^{14}CO_2$  released from  $1{}^{-14}C$ -glucose obtained via the pentose-phosphate shunt and the citric acid cycle to that produced from  $6{}^{-14}C$ -glucose via the citric acid cycle alone. Warshaw and Rosenthal (129) reported enhanced glucose oxidation in conditions of low-density plating. This presumably reflects increased flow through the pentose-phosphate pathway, which is inhibited when the cells reach confluency (96,129). We have reported (30,31) similar findings, as well as a link between this pathway and  ${}^{3}$ H-thymidine incorporation into DNA. Thus, DNA synthesis, in the wake of the pentose-phosphate pathway, increases in parallel with the flow through the pentose shunt, after a delay of several hours, which is consistent with the increased requirement for ribose-phosphate in proliferating cultures.

The level of *glucose oxidation* via the tricarboxylic acid cycle has been shown to remain constant during all the phases of heart-cell culture (31,129). However,

conflicting results have been reported (103): rates of glucose utilization varied in cultures of different ages. These were interpreted as a postnatally-induced decline of glucose as an energy source.

#### Pyruvate and Lactate Oxidation

We have reported that following glucose uptake from the medium, lactate is released back into the medium and reutilized within the cell either via oxidation or in further transformations. Rosenthal and Warshaw (96) found a lag phase in the time course of <sup>14</sup>CO<sub>2</sub> formation from 2-<sup>14</sup>C-pyruvate as compared to that produced from 1-<sup>14</sup>C-pyruvate, after which <sup>14</sup>CO<sub>2</sub> production from both substrates increased linearly, although <sup>14</sup>CO<sub>2</sub> formation from 1-<sup>14</sup>C-pyruvate occurred at a higher rate. This may be explained if CO<sub>2</sub> derived from carbon 1 is produced directly by the action of pyruvate dehydrogenase, while CO<sub>2</sub> from carbons 2 and 3 is formed via incorporation into acetyl-CoA followed by the citric acid cycle reactions.

Ross and McCarl (97) found that pyruvate was taken up at a rate of 4.8  $\rm umol/min/mg$  protein. About 50% of the incorporated pyruvate is reexported as lactate, the remainder being oxidized. They also confirmed an earlier finding (96) that  $\rm ^{14}CO_2$  release from 2- $\rm ^{14}C$ -pyruvate lagged behind that formed from 1- $\rm ^{14}C$ -pyruvate by about 10 min. The steady state for oxidation was reached within 20 min with the former substrate, and within 80 to 100 min with the latter. Lactate oxidation led to linear release of  $\rm ^{14}CO_2$  from 1- $\rm ^{14}C$ -lactate over a 2-h period. Subsequently the release rate fell (97). More  $\rm ^{14}CO_2$  was also produced from 1- $\rm ^{14}C$ -lactate than from 1- $\rm ^{14}C$ -pyruvate under similar conditions. As yet no satisfactory explanation has been given for this finding. Furthermore, 15% of the total lactate was found to be incorporated into cellular components - 2/3 of this into water-soluble molecules and 1/3 into lipids and proteins.

In conclusion, the above results show that in cultured heart cells, pyruvate is either oxidized or converted to lactate, but is not transformed into other cellular components, while lactate undergoes either oxidation or transformation to other compounds. Similar findings have been reported in the perfused heart (71,91), indicating that the LDH isoenzyme profile does not play a significant role in the regulation of pyruvate-lactate formation under low work load conditions.

#### 4. Oxygen Consumption

Oxygen consumption measurements are essential in studies of cellular metabolism. Although the current oxygraph may provide useful data, most of these experiments are carried out on cells that have been suspended by protease treatment and therefore consist of dying non-beating cells. Recently, Takehori et al. (115) have constructed a lucite attachment with a built-in  $O_2$  electrode, which allows  $O_2$  consumption to be measured in beating cultured heart cells attached to the substratum. With this device, they showed that  $O_2$  consumption was 47% higher in beating than in quies- cent cells, and that these beating rates were comparable to those observed in intact heart. In addition, the  $O_2$  consumption of quiescent cardiomyocytes was two-fold higher than in either cardiac non-muscle cells or in the  $L_6$  muscle cell line - a clear indication of the high level of energy metabolism in cardiomyocytes.

In adult cardiomyocytes,  $O_2$  consumption was two-fold higher than in neonatal cultured cells (10). It correlated with the higher mitochondrial creatine kinase (CK) in adult cells (100,118), implying that this CK plays a role in the coupling of energy production and utilization, presumably by preventing high-energy phosphate depletion during high cardiac work loads. However, the available data are scarce and much more research is required in this area.

#### 5. High-Energy Phosphate Transport

Over the last few decades, attention has been directed toward elucidation of myocyte energy metabolism, focusing on the roles played by CK and the creatine/ phosphocreatine ratio (Cr/PCr) in the maintenance of cellular energy levels and energy transport.

The finding that ADP undergoes phosphorylation by PCr (58) and the discovery of myosine ATPase (23) served to demonstrate the direct relationship between ATP utilization and mechanical work, and, consequently, the restricted role of PCr as a reservoir for high-energy phosphate for the coupling between energy utilization and production (12,44,58,65). However, in spite of the progress in this area, the mechanism of transformation of chemical energy into mechanical work is still not clear, although many of the biochemical events involved have been characterized (for review, see ref. 46).

Mitochondrial energy production is determined by the rate of ATP utilization, which is regulated via a cytosolic signal (53). Presumably, ADP formation and the Cr-PCr-CK system are involved in intracellular energy formation and in energy transport from the mitochondria to the myofibrils (100,104,106). CK bound to either myofibrils or membranes may be involved in phosphorylation of ATP, thus providing ATP at the sites where it is used (14,81,137,138,). According to this view, CK is one the key enzymes involved in cellular energetics.

Creatine kinase is a dimeric enzyme (19) consisting of two types of monomers - M and B. Of the three isozymes - MM, BB and MB - the MM form predominates in muscle (24,25). The existence of a fourth form of CK has been demonstrated in mitochondria (mCK, ref. 50). In rat heart, about 20% of the CK activity is located in the mitochondria, and about 75% in the cytosol (21). In human heart, these values are 10% and 85% (7). Electrophoresis (50), in situ histochemical staining (3,107), and immunological techniques showed that mCK is distinct from the cytoplasmic CK's (94). mCK is bound to the outer surface of the inner mitochondrial membrane and possibly via cardiolipin (36).

Mitochondrial CK plays a critical role in the regulation of respiration, enhancing both  $O_2$  consumption and PCr synthesis in isolated mitochondria, whereas PCr reduces  $O_2$  uptake owing to competition for ADP between mCK and oxidative phosphorylation (51,52,109,127). There are apparently several pools of ATP. The ATP pool which is preferentially used by mCK and the possible role played by the mitochondrial membrane in the regulation of ATP fluxes between the mitochondria and the cytoplasm remain to be elucidated (2,26,57,139).

Inhibition of oxidative phosphorylation with oligomycin in cultured heart cells leads to cessation of contraction within 30 to 60 min (106). This effect is correlated with the depletion in PCr levels, the ATP levels remaining virtually unchanged. In addition, when cells are cultured in Cr-containing medium, or are pulsed with Cr, PCr levels increase, while there is no change in ATP. Yet, there is no evidence for direct participation of PCr in contractile activity.

The inhibition of PCr synthesis by oligomycin suggests that it is dependent on oxidative phosphorylation, and that mCK is involved in the control of PCr levels in cultured myocardial cells. Indeed, this isozyme has been identified in rat myocardial cells. It is absent from cardiac non-muscle cells, and from the  $L_6$  muscle cell line (118).

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The relative proportion of the mCK isozyme increases upon aging, both in culture and in vivo. The BB form decreases under these conditions, and the MM isozyme predominates during development of myocardial cells. The relative isozyme distribution is similar in culture and in fresh tissue, although total CK activity is 4-to 5-fold higher in the latter: 1.5 U/mg protein in culture, 6.2 U/mg in fetal heart, and 7.3 U/mg in 1- to 6-d old heart (118).

Creatine appears to trigger mCK synthesis - if present in the culture medium, it leads to a 44% increase in mCK. It is reasonable to assume that myocardial energy demands increase postnatally. Energy production is regulated at the mitochondrial level by Cr concomitantly with the development of the Cr-PCr-mCK energy shuttle system (106,118). However, the cause-and-effect relationship of Cr in the coupling of energy production to energy utilization via resynthesis of PCr in the mitochondria remains to be demonstrated (62).

Analysis of the relative proportion of CK isoenzymes at various times before and after birth revealed striking changes - 14-d old fetal heart contains only BB isozymes, while in the 17-d old fetus, 80% of the enzyme activity is due to M forms (MB and MM isozymes). In mice, mCK is also undetectable in either fetuses or immediately after birth, and accounts for 1% and 9% of the total CK activity in 6-d old and adult animals, respectively. In the lamb, however, cardiac CK and Cr are detectable even prenatally (49), whereas the BB form is totally absent, and the level of the MB form is decreased from 24% to 5% during this period (25,48). In cultured striated muscle cells, transition from B to M forms is incomplete, with significant levels of MB and BB isozymes expressed throughout the period in culture (22,59,79,80). The M form is apparently characteristic of the post-fusion state (13,78). In addition, fusion and synthesis of contractile proteins can be stimulated by the supplementation of Cr to the culture medium (105,106).

# 6. Energy Metabolism during Oxygen Deprivation

Under conditions of  $O_2$  restriction or deprivation, the heart immediately switches to anaerobic glycolysis. By this means, it briefly maintains the high-energy phosphate levels, prior to a rapid decline in ATP and CP production with a concomitant increase in degradation products (35,93,102,116). Cultured heart cells have also been used as a model system for studying  $O_2$  deprivation at the cellular level (1,40,121). In some experiments, metabolic inhibitors were employed together with  $O_2$ deprivation (41).

Recently, an alternative approach has been taken to assess the effects of  $O_2$  deprivation in cultured cells. By combining anoxia with glucose deprivation, coupled with extreme reduction in the volume of extracellular medium, a true microenvironment with reduced flow is simulated, in which the products of cellular catabolism, such as lactate, remain in the vicinity of the cells and are not washed away (122,124,126). Indeed, anoxic injury, as reflected in higher rates of enzyme release and sarcolemmal transformations, is increased under these conditions (122-126).

Several studies relating to nucleotide and purine metabolism in anoxic heart cells in culture have shown that in chick fetal heart cells, anoxia induces a two-fold increase in production of adenosine and its metabolites. Surprisingly, the levels of ATP remained unchanged (67). In another investigation, it was demonstrated that hypoxanthine release precedes cellular enzyme depletion (119). Correlation between cellular ATP depletion and degradation of the cell membrane by phospholipase C has also been reported (42,43).

Recently, we studied the kinetics of high-energy phosphate depletion and the release of nucleotide degradation products in cultured heart cells under conditions of anoxia combined with volume restriction (122,123). Decrease in CP levels usually precedes that of ATP. There is marked acceleration of the decrease in the levels of both of these, when anoxia is combined with volume restriction of the extracellular medium. In the presence of glucose and without volume restriction, the glycolytic flux from glucose and the glycogen stores may provide sufficient ATP to delay its depletion for considerable periods. In perfused heart (69), under conditions of high flow, the glycolytic flux from cellular glycogen and supplementary glucose may even provide sufficient energy to maintain some contractile activity.

The irreversible stage in anoxic cells is presumably related to marked depletion in ATP content. Indeed, major cellular damage, as reflected in lysosomal enzyme release, only starts when 50% of the cellular ATP has been depleted (122). In addition, release of free arachidonate and palmitate was associated with ATP depletion - arachidonate accumulation started when the ATP levels reached <25% of the normal values (34). If the rate of ATP utilization exceeds the rate of ATP production in the absence of O<sub>2</sub> and glucose, ATP breakdown gives rise to intracellular adenosine, which may be recovered in the form of purine bases in the medium. The breakdown of ATP results in the formation of ADP and AMP, which is further deaminated to IMP or dephosphorylated to adenosine under anoxic conditions. Adenosine may be further degraded to inosine and then, in turn, to hypoxanthine, xanthine, and urate. The degradation of inosine presumably involves nucleoside phosphorylase which is located in endothelial cells (99). In our studies, adenosine was not detected in cultured cells during  $O_2$  deprivation, inosine levels increased slightly, while the [hypoxanthine] in the external medium was even decreased. However, anoxia combined with volume restriction, led to high levels of purine release, although inosine levels remained low, and hypoxanthine, xanthine, and urate release reflected the decrease in cellular ATP levels (122,123).

# 7. Conclusions and Perspectives

Glucose metabolism in cultured heart cells has been termed "impaired" or "hypoxic", since these cells produce lactate. Careful examination of the literature on both cultured cardiomyocytes and perfused hearts reveals that lactate production should not be considered as an abnormal response, but rather as an expression of the cellular energy requirements. In this respect, cultured heart cells are more akin to the heart at rest or under low work loads, in which lactate production also occurs. Much remains to be learned about the mechanisms of carbohydrate metabolism. Clearly, the shift from the  $H_4$  to the  $M_4$  forms of lactate dehydrogenase is an inadequate explanation for the lactate production. Other crossroads of enzymatic control - the active and inactive forms of pyruvate dehydrogenase, governing the flow of pyruvate into the citric acid cycle, and the enzymes activating glycogen synthesis and degradation - deserve further study.

In cultured cardiomyocytes, fatty acids constitute the main source of metabolic energy, as in the heart in vivo. The use of these cultures has furthered our understanding of FA transport across membranes. In addition, FA's derived by the cleavage of intracellular reserves were shown to be preferentially oxidized over newly incorporated FA's. Another highly significant achievement made with cultured heart cells was the discovery of the pathway for incomplete oxidation of FA's leading to the release of L(+)-3-hydroxybutyrate, an intermediate in beta-oxidation. It will be interesting to examine the physiological role of this pathway, bearing in mind that L(+)-3-hydroxybutyrate, when injected into suckling rats, may be used in the synthesis of cholesterol and other lipids, chiefly in the brain (130).

Little mention has been made in this review of interactions between carbohydrates, ketone bodies and lipid metabolism in cultured heart cells - surprisingly this field has suffered from almost complete neglect. Cells in culture provide an ideal system

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for carrying out such investigations owing to the ease with which they may be manipulated and controlled.

At present, only a tentative scheme can be drawn up for the metabolic pathways of lipids within the cell. The flow of substrates between organelles, the sites where they are stored and metabolized, and the (hormonal) control of these events remain to be elucidated.

The energy shuttle requires the presence of creatine kinase bound to mitochondria and to myofibrils. Phosphocreatine, synthesized in the mitochondria by mCK, diffuses to the myofibrils where the ATP level is maintained by the myofibrillar CK. One may speculate that PCr is not susceptible to non-specific enzymic reactions, and that this constitutes a distinct advantage in energy transport. Regulation of energy metabolism and the energy shuttle by the Cr-PCr-mCK system occurs concomitantly with the continuous energy demand of the myocardium imposed upon a high resting metabolism. This system is not ubiquitous. It seems that its development in myocardial cells usually occurs postnatally. Further studies in this field together with  $O_2$  consumption measurements in situ under various metabolic conditions should provide more information on the energy requirements of the cultured heart cell.

Finally, since the cultured cardiomyocyte is a rhythmically contracting cell, one may expect that ATP utilization and some enzymic activities fluctuate during cycles of cardiac contraction and relaxation. The cultured cell system is ideal for studies on this level, since the technical difficulties of freeze-clamping (117,134) should be easier to overcome than in the perfused heart.

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# Chapter 15

# **Energy Metabolism of Adult Cardiomyocytes**

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By now the basic features of the energy metabolism of isolated cardiomyocytes have been described. Unstimulated cardiomyocytes are quiescent and exhibit the characteristics of cardiac metabolism at basal energic demand. Their energetic demand, however, can be stimulated to the maximal rates of the physiological range. As oxidative substrates fatty acids are preferred to lactate and glucose. The uptake kinetics for these three substrates have been characterized. For glucose and lactate defined transport mechanisms are identified, for fatty acids a purely physical diffusion process is also discussed. Under anoxia, energy production falls short of energy supply. Already during the reversible phase of hypoxic injury, cardiomyocytes have some cytosolic enzyme release. It is hypothesized that both release of enzymes and development of rigor are causelly related to changes in the free energy change of cytosolic ATP hydrolysis rather than in absolute ATP concentrations. Loss of  $Ca^{2+}$  control follows energetic exhaustion. At cytosolic  $Ca^{2+}$  levels beyond 3 uM, it becomes irreversible. In detail the relation between energy metabolism and onset of irreversible hypoxic cell injury is not sufficiently understood.

# I. Introduction

Isolated adult cardiomyocytes have become an established tool in heart research. A number of ways has been described to isolate muscle cells from myocardial tissue, and primary cultures of these non-dividing cells have also been developed (34). This review is designed to summarize the current knowledge about the basic features of their substrate and energy metabolism.

The results demonstrate that today cardiomyocyte preparations can be obtained that exhibit the typical metabolic characteristics of the myocardium. Studying metabolic behavior of a purified myocyte preparation allows to identify the contribution of the cardiac muscle cell type to the complex metabolic reaction of whole tissue. Furthermore, the properties of the cardiomyocyte can be investigated individually, i.e., free of the immediate influence of adjacent muscle cells. This latter possibility renders the use of cardiomyocytes particulary helpful in studies on hypoxic cell injury since in this model inhomogeneities in the exogenous factors, acting on the individual cell, can be avoided.

# **II. Energy Metabolism under Aerobic Conditions**

# 1. Oxygen Consumption

In the absence of external stimulation, adult ventricular myocytes have a stable resting membrane potential and are mechanically quiescent. Therefore their oxygen demand is low. In newly isolated as well as in short-term cultured cardiomyocytes, this basal oxygen demand was found to be approximately 1 umol  $O_2/min$  per g wet wt. at 37°C (47,50,51,60). This value is close to the oxygen demand of a rat heart in cardioplegic arrest (3). With respiratory uncoupling, oxygen consumption of cardiomyocytes can be stimulated to a magnitude known for the heart under maximal catecholamine stimulation, i.e., 20 to 40 umol  $O_2/min$  per g wet wt. (60). This

demonstrates that isolated cardiomyocytes possess the full physiological range for variations of energy demand. The  $Q_{10}$  value for lowering the temperature from 37 to 27°C was determined as 1.8 (60), closely resembling the corresponding  $Q_{10}$  values of the arrested rat heart (3).

Isolated cardiomyocytes have also been stimulated to contract by electrical field stimulation. The metabolic cost per contraction was found to be similar to that in myocardial tissue under zero load conditions (25). The increment in energetic demand caused by mechanical activity under such unloaded conditions is proportional to the square of length changes, indicating that it accounts primarily for the elastic compression of the myofilaments (54).

# 2. Uptake of Substrates

The uptake of glucose into the myocardial cell is mediated by a specific transport system of which the affinity and the capacity can vary. In isolated cardiomyocytes insulin accelerates initial uptake rates of 3-O-methylglucose, a nonmetabolized glucose analogue, into isolated cardiomyocytes by a factor of 2 (14) to 9 or 10 (19,24). Interestingly, the  $V_{max}$  values obtained with insulin were identical in these studies, whereas the basal rates varied. In the unstimulated state, the glucose transporter has a low affinity for glucose.  $K_m$  values range from 7-9 mM (19,52). Insulin leaves the transporter as one of low affinity. Thus, insulin acts mainly through enlarging the capacity of the transport system. The onset of insulin action on stimulating glucose uptake is observed in the first 30 s, the effect is maximal in 1-3 min (8,14). The promptness of the reaction indicates that it is not dependent on protein synthesis. This is consistent with the hypothesis that additional transporter molecules are only recruited from internal stores (38).

Under hypoxic and other forms of energetic stress the uptake of glucose by the myocardium is greatly enhanced (37,42,52). In contrast to the action of insulin, the affinity for glucose is fivefold increased, possibly helping the ischemic cell to continue aerobic glycolysis when interstitial concentrations of glucose decrease. The specific effect of insulin on glucose uptake under hypoxic conditions is much reduced. It seems that the maximal number of glucose transporters mobilized under the short-term action of insulin or anoxia is the same. When cellular energy reserves are largely exhausted glucose transport slows down again (21).

Increased pressure work of myocardium was also reported to enhance glucose transport with an increased  $V_{max}$  rate and decreased  $K_m$  value (44). In contrast, when unloaded isolated cardiomyocytes were stimulated to contractions (and this caused a fivefold increase in oxygen demand)  $V_{max}$  rates remained unchanged whereas the  $K_m$  value decreased by 50% (55). Thus, increased energy turnover per se cannot be the cause for a stimulated glucose uptake under pressure work.

Lactate is a major oxidative substrate for the myocardium and isolated cardiomyocytes. The transport of lactate into the isolated myocardial cell has been analyzed by Kammermeier et al. (36). They demonstrated that lactate uptake is characterized by a sigmoidal dose dependence, indicating a carrier-mediated process. The half-maximal transport rate was found in the range between 5-10 mM, i.e., the affinity of the transport system is low. At physiological concentrations of lactate, 2.3 mM pyruvate inhibited the uptake of lactate, suggesting a competitive mechanism. The uptake of lactate was enhanced when the pH was lowered. But the increment in transport rate exceeded the increase in the concentration of the protonated acid by about tenfold. From this the authors concluded that, instead of favoring nonionic diffusion, protons are more likely involved in the mechanism of a carrier-mediated transport process, e.g., in form of an H<sup>+</sup>-lactate cotransport. Dennis et al. (13) have characterized the lactate transport system in the perfused rat heart, and came to similar conclusions.

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Long-chain nonesterified fatty acids are major substrates for myocardial energy production. In the extracellular space, fatty acids are non-covalently bound to albumin. It has first been demonstrated by Hütter et al. (32) that the oxidation of fatty acids in the working heart is a function of the concentrations of the fatty acid/albumin complex, and not of the steady state concentration of unbound fatty acids. For constant fatty acid/albumin ratios, saturable Michaelis-Menten type kinetics were observed. In isolated cardiomyocytes, it was shown that transsarcolemmal transport obeys the same kind of kinetics (53). The  $V_{max}$  values obtained for various fatty acid/albumin ratios were virtually identical with those obtained for fatty acid oxidation in the whole heart. This indicates that the rate-limiting process in both experimental systems is the same, namely transsarcolemmal transport. Recently similar kinetics have also been reported for the translocation of long-chain fatty acids through the endothelium (160 nmol/min per g tissue, equivalent to approximately 5 umol/min per g endothelial mass) is far greater than that for transsarcolemmal transport (35 nmol/min per g cardiomyocyte mass), at a palmitate/albumin molar ratio of about 1 (53,63).

Because of the saturable kinetics for fatty acid uptake by various animal cells, some mechanism of facilitated diffusion is generally assumed. But the validity of this conception has been questioned since the purely physical partitioning of fatty acids between the complex with albumin and a phospholipid bilayer obeys the same type of saturable Michaelis-Menten-like kinetics (45). Model calculations indicate that the transport rates in the liver cell could be fully accounted for by this purely physical mechanism. Such a calculation can also be made for the myocardial cell. Similarly, DeGrella and Light (11,12) have argued that cardiomyocytes in the absence of albumin take up fatty acids purely by diffusion. The rates observed would fully account for transport rates observed in vivo.

# 3. Substrate Oxidation

In short-term cultured cardiomyocytes, oxidation rates for physiological substrates (glucose, lactate, pyruvate, palmitate) were found constant for several hours (50,51). At physiological concentrations, fatty acids are preferred to lactate and glucose. At the excessive concentration of 5 mM, pyruvate is oxidized at a rate similar to that for exogenous fatty acids. Thus, in cardiomyocytes, physiological substrates are used in a pattern characteristic for the myocardium. For cultured cardiomyocytes, however, these exogenous substrates represent only a minor energy source. After 4 h in serum-supplemented medium 199, the cells contain enlarged stores of glycogen and triacylglycerols (50). When incubated in a simple saline medium with selected exogenous carbon sources, they preferentially oxidize fatty acids from endogenous triacylglycerol.

It was shown that in quiescent short-term cardiomyocytes endogenous lipolysis accounts for 94% of the calculated oxygen demand, when exogenous substrates are absent (50). The remaining share is covered by glycogen degradation. Consistent with the pronounced preference for lipids, only 7% of pyruvate dehydrogenase is in its active form. The supply of exogenous glucose together with insulin stops glycogenolysis and increases glycolysis, but pyruvate dehydrogenase activity increases only slightly. Superphysiological concentrations of pyruvate (5 mM) displace glucose and glycogen as oxidative substrates and induce an activation of pyruvate dehydrogenase. The contribution of endogenous lipolysis to oxidative energy metabolism is conversely reduced. At equimolar concentration, lactate is less effective than pyruvate (50).

In detail, the metabolic behavior of quiescent short-term cultured cardiomyocytes reflects the characteristics of cardiac energy metabolism at low demand (29,33). In

aerobic myocardium at low respiratory rates, concentrations of NADH and acetyl-CoA are elevated. This causes beta-oxidation rates to be low. Increased cytosolic long-chain acyl-CoA levels in the cytosol together with high concentrations of ATP and production of alpha-glycerophosphate favored by the reduced cytosolic state stimulate triacylglycerol synthesis. On the other hand, NADH and acetyl-CoA are known to promote conversion of pyruvate dehydrogenase into its inactive form (52). Another factor also reducing the activity of this enzyme consists in low mitochondrial  $Ca^{2+}$  levels in the quiescent cardiomyocytes (20). At low activity of this flux-regulating enzyme, fatty acids are by far the preferred substrate.

A reduced cytosolic state together with a low oxidation rate of pyruvate favors net lactate production in quiescent myocytes under fully oxygenated conditions (51). With glucose as sole exogenous substrate, they produce lactate in higher quantities than  $CO_2$  from exogenous glucose. As expected, this relation is inverted when the energy demand is increased. It is well documented that lactate is produced also by the normoxic heart if perfused with glucose as the sole substrate, especially at a reduced work load (61,67). In extrapolating these results, a relatively large lactate production from aerobic, quiescent cardiomyocytes is indeed predicted. In contrast to the behavior of quiescent cardiomyocytes not long after isolation, cardiomyocytes apparently have a truly enlarged glycolytic capacity after several weeks in culture. After 25 days in culture, the energy obtained by degradation of glucose was 30% aerobic and 70% glycolytic (R. Spahr, H.M. Piper & S.L. Jacobson, unpublished observations).

At a normal energetic state, the phosphofructokinase reaction is the major ratelimiting step in the Embden-Meyerhof pathway. Its activity is regulated predominantly by the cytosolic concentration of citrate and of fructose 2,6bisphosphate (31,52). The effect of both these regulators can be demonstrated also in quiescent cardiomyocytes. Citrate levels have been shown to correlate closely with glycolytic flux rates in experiments in which primarily the relative contribution of lipids to substrate oxidation is varied (51). Under insulin, the prevalent stimulus for phosphofructokinase activity is apparently fructose 2,6-bisphosphate (51).

# III. Energy Metabolism and Cell Injury under Hypoxic Conditions

# 1. Energy Production and Demand

It has been demonstrated that the individual heart-muscle cell will not stop mitochondrial respiration and increase glycolytic flux unless exogenous  $pO_2$  has dropped below 0.5 torr (68). This means that the cell as a whole is almost as sensitive to low oxygen tension as the isolated mitochondrion is. Conditions with an oxygen tension below this threshold level are usually referred to as "anoxia".

The hypoxic and the ischemic myocardium reduce their energy demand immediately after oxygen withdrawal (40). In the beating heart, the reduction in contractile activity, i.e., mechanical failure, accounts for most of this self-protective mechanism. This, however, is only part of the energy-saving mechanism, since myocardium, arrested prior to ischemia (5), and quiescent myocytes (48) also quickly reduce their energy demand under hypoxic conditions. In addition, the absolute saving in energy consumption depends on the prehypoxic energy expenditure (58). Except from the energy-conserving effect of early mechanical failure, little is known about the nature of these mechanisms.

In spite of this self-protective reduction of energy demand, anaerobic energy metabolism at the end always falls short of energy needs as indicated by the loss of high-energy phosphates (40). Cultured cardiomyocytes contain elevated amounts of glycogen, and they use glycogen for glycolytic energy production in substrate-free anoxia. But when ATP contents have fallen by about 70%, glycolytic energy

production almost stops even though still half of this substrate is left (48). It has been suggested by Kübler and Spieckermann (40) that such inability to use a glycolytic substrate is due to the lack of phosphorylation energy at the phosphofructokinase level. In the hypoxic perfused heart (26) and in anoxic isolated cardiomyocytes (48), irreversible injury can be postponed considerably if glycolytic flux is stimulated already at the onset of oxygen deprivation. Since enhanced glycolysis delivers more ATP to the cell, cellular ATP contents decrease at a slower rate. In these systems the waste products of glycolysis, lactate and protons, do not accumulate to the same extent as in ischemic tissue.

A contribution of mitochondria to the loss of energy in the hypoxic cell is well documented (23,56). In the non-respiring state, the mitochondrial ATP-synthetase acts in a backward direction, i.e., as an ATP-hydrolase. Thus, one major effect by which mitochondria contribute to the aggravation of hypoxic cellular injury consists in the inversion of a normal physiological function. The functional injury of mitochondria isolated from ischemic and hypoxic myocardium has been the subject of many studies, but it has not been demonstrated that these impairments are responsible for the functional inability of the heart after reperfusion. In isolated cardiomyocytes impairment of mitochondrial metabolism was not found very pronounced, when compared with other aspects of metabolic disturbance (9,41). And even in cells becoming hypercontracted in anoxia, the changes in mitochondrial ultrastructure are relatively moderate (4,57).

Other results also indicate that a loss of mitochondrial function determines not the "point of no-return" of hypoxic cell injury. When cytosolic  $Ca^{2+}$  just starts to rise in a hypoxic cardiomyocyte, resupply of ATP by resumption of oxidative phosphorylation prevents the impending loss of  $Ca^{2+}$  control (2). But when cytosolic free  $Ca^{2+}$  concentrations have risen to 3 uM, loss of  $Ca^{2+}$  control has become irreversible (2). It is probably the inability of mitochondria to resume sufficient ATP production at this  $Ca^{2+}$  level, defining it as a crucial threshold. Since this behavior is based on a normal functional property of mitochondria (7), it does not need specific mitochondrial injury to bring about cell death. In summary, according to current evidence mitochondria are not a "limiting structure" for cell survival.

#### 2. Enzyme Release

Release of cytosolic enzymes from the hypoxic perfused heart or from hypoxic isolated cardiomyocytes has been demonstrated to correlate with cellular ATP contents in certain experimental models (18,28,48). Already a small reduction in contents of energy-rich phosphates, as it may also occur in physiological stress, leads to a detectable release of enzymes (59). Studies on isolated cardiomyocytes in substrate-free anoxia have shown that this early enzyme release is due to protein loss from only the cytosolic compartment of cells which are still in the reversible phase of hypoxic injury (48,57). At this early stage, ultrastructural alterations are subtle (57); large holes in the sarcolemma can be ruled out as causing the protein leakage. The number of hypercontracted cells does not correlate with this early release of cytosolic enzymes.

There may be a causal relationship between protein release from hypoxic cells and a) increase in the number of subsarcolemmal vesicles (57); b) protrusion of cell surfaces into small sarcolemmal blebs (57); and c) rapid increase in fluidity of the hypoxic sarcolemma by an order of magnitude (16). Leakage of soluble proteins from reversibly energetically stressed cells has also been reported for other cells types (6,46,66,69). It has been hypothesized by Spieckermann et al. (59) that the heart cell releases cytosolic enzymes continuously, but at a low rate, and that the release through this unidentified channel is enhanced under conditions of energetic stress.
A comparison of energy depletion of isolated cardiomyocytes by anoxia and by glycolytic blockade, however, points to the free-energy change of cytosolic ATP hydrolysis (64) as a decisive factor for early enzyme release (49). This magnitude characterizes the thermodynamic availability of ATP and is determined by the ratio (ATP/ADP  $\cdot$  P<sub>i</sub>). Iodoacetate blocks glycolytic flux at the level of the glyceral-dehyde-3-phosphate dehydrogenase reaction. It traps free cytosolic P<sub>i</sub> by an enormous accumulation of phosphorylated intermediates. In comparison to anoxia, therefore, at a given ATP level in iodoacetate-treated myocytes, the (ATP/ADP  $\cdot$  P<sub>i</sub>) ratio will be higher, i.e., more phosphorylation energy will be available in the cytosol. After a loss of one third of initial ATP contents, release of lactate dehydrogenase from anoxic cells was indeed found three times higher then from iodoacetate-treated cells with the same ATP level (49).

In infarcted myocardium many cells are lysed. Therefore, a considerable part of the enzyme release from such tissue can be explained by mass loss from broken cells. Indeed, it has been shown that the size of the necrotic area correlates well with the amount of enzymes lost from infarcted myocardium (1). In the clinical setting, detection of cardiospecific enzymes in the circulation will generally indicate the occurrence of necrotic cells, because the moderate early release of cytosolic enzymes easily remains undetected due to the impeded washout from ischemic areas and the limited sensitivity of the methods in use.

# 3. Contracture Development

In the course of progressive energy breakdown, contracture development is observed in the globally-ischemic myocardium ("stone heart") and also in isolated, energydepleted cardiomyocytes. A number of studies indicate that the development of tension in the myocardial cell under hypoxia and ischemia (2,30), and also under other conditions of energy depletion (10), is caused by the formation of rigor complexes, i.e., of fixed crossbridges between actin and myosin filaments. Thus it is not initiated by a rise in cytosolic  $Ca^{2+}$ , which would lead through fast crossbridge cycling to sustained contraction. Recently, experiments deciding between these alternative mechanisms became feasible. By simultaneous monitoring of the cytosolic free  $Ca^{2+}$  concentration and cell shortening in single isolated cardiomyocytes, it was demonstrated that cell shortening of hypoxic cardiomyocytes precedes a rise in cytosolic free  $Ca^{2+}$  (2).

The factors causing rigor bond formation in the oxygen-deficient heart cell are still a matter of debate. Based on the finding by Weber and Murray (65) that, in vitro, rigor bonds between actin and myosin are formed in a  $Ca^{2+}$ -independent manner if the ATP concentration is below 100 uM, it has been hypothesized that in the hypoxic cell, too, rigor is caused by a lowering of the cytosolic ATP concentration below this level (23,30,39). However, the average ATP concentration in hypoxic heart tissue in contracture can be equal to or greater than 1 mM. It was therefore assumed that the cytosol contains a subcompartment next to the myofibrils in which the ATP concentration is critically reduced. So far, no direct evidence has been given for the thruth of this assumption. In the following an alternative explanation will be proposed.

The crossbridge cycle is driven by the energy liberated during ATP hydrolysis. Thermodynamically, the energy available from ATP hydrolysis is characterized by the free-energy change  $\Delta G$  (64), i.e.,  $\Delta G = \Delta G_0 + RT \ln [ATP]/[ADP] \cdot [P_i]$ . In this equation  $\Delta G_0$  signifies the standard free-energy change; R, the universal gas constant; T, the absolute temperature; and [ ] refer to the free metabolite concentrations. The rigor state is supposed to occur transiently in a normal crossbridge cycle and represents the state lowest in potential energy (15). It is therefore conceivable that single crossbridges remain trapped in the rigor state, if the average  $\Delta G$  value is too low to allow the cycling of all crossbridges to be

completed. In fact, tissue tension in the ischemic rat heart is found to increase after about 10 min (27), at which time the cytosolic  $\Delta G$  has been reduced from 60 to about 45 kJ/mol (17).

In the globally ischemic and hypoxic heart, rigor is accompanied by muscle shortening. This is a gradual process, leading to shortened sarcomeres throughout the tissue. A gradual shortening of the myofibrils is also observed in single isolated myocytes under anoxia, if these are attached to a substratum (57). At an average ATP content of about 2 umol/g wet wt., these cells are in a square, but still polygonal form. The sarcomere length is reduced from the initial 1.9 um (slack length of sarcomeres) to 1.4 um (57). At 1.4-um sarcomere length, myosin filaments are already distorted. This sarcomere length, however, is also observed during forceful contractions in cardiomyocytes and is fully reversible. If the cells in this shortened square form are reoxygenated, they relengthen (57). This behavior demonstrates the reversible nature of this rigor state. In contrast to attached cells, free-floating single cells do not shorten in a gradual fashion, but rapidly shrink to a minimal sarcomere length of about 1.1 um or 60% their initial length (2,23). This difference of behavior can probably be explained by the complete absence of any force opposing rigor forces in non-attached cells. Such extremely contracted cells are usually unable to relengthen upon reoxygenation, although they can regain control of their cytosolic free Ca<sup>2+</sup> levels (2). Apparently, at 1.1-um sarcomere length, severe disruptions of the cytoskeleton have taken place, which normally disable the cells to reassume an elongated shape.

# 4. The "Point of No-Return"

From studies on short-term cultured surface-attached cardiomyocytes, which seem to form the most stable system currently available, it is now clear that, in the muscle cell itself, onset of enzyme release, irreversibility of structural cell injury, and cytolysis are separate events. In the heart, exposed to ischemia of hypoxia, these events occur in indistinguishably close temporal coincidence.

A sufficient criterion for impending cell death consists in the loss of cytosolic  $Ca^{2+}$  control. In isolated hypoxic cardiomyocytes, a rise in cytosolic  $Ca^{2+}$  is not the result of large membrane perforations. The  $Ca^{2+}$  accumulation in the cytosol enters the cell from outside, but the accumulation proceeds gradually and the disturbance of  $Ca^{2+}$  homeostasis can be reversed by early resumption of oxidative energy production. Therefore, it seems to be a shortage of energy which primes the heart cell for cell death. The sarcoplasmic reticulum plays a decisive role at the edge of irreversible loss of  $Ca^{2+}$  control: when its ability to sequester  $Ca^{2+}$  was blocked with caffeine, any recovery of  $Ca^{2+}$  control was abolished (2).

The exact role of energy metabolism in the process of progressive injury is still an open field for research. So far, in most studies average levels of high-energy phosphates have been determined and not the free-energy change of ATP hydrolysis. The latter is a parameter for the thermodynamically available energy. An apparent ATP threshold (1-2 umol/g wet wt.) for resuscitability has been described for ischemic dog myocardium (35,40), ischemic rat hearts (62), and anoxic cultured rat cardiomyocytes (48,57). But it is not definitely clear whether this threshold is a borderline under all circumstances. In energetically-poisoned cardiomyocytes, the ATP concentration may fall to much lower levels, before the cytosolic Ca<sup>2+</sup> concentrations rises (22). This, however, must not mean that a true recovery from these ATP concentrations is possible before the loss of Ca<sup>2+</sup> control. It has also been claimed that under certain conditions, ischemic hearts can recover from similary low ATP levels (43). But since in the latter study only some functional parameters for recovery were investigated, it does neither prove the possibility of true structural and metabolic recovery from very low energy levels.

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# Chapter 16

# Ischemia, High-Energy Phosphate Metabolism and Sarcomere Dynamics in Myocardium

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1. Effects of ischemia or anoxia on the dynamics of sarcomere contraction of myocardium are described on the basis of a model of excitation-contraction coupling in the cardiac cell.

2. A model of regional ATP supply to the components of the excitation-contraction coupling apparatus is proposed. This model emphasizes that ischemia/anoxia may affect various cell organelles to a different degree and with a different time-course.

3. The role of lowered ATP levels and accumulation of breakdown products of highenergy phosphates, and of glycogenolysis, in the development of abnormalities of force development and of sarcomere dynamics during ischemia/anoxia is discussed. It is proposed that early during ischemia an important effect of the rise of inorganic phosphate ions is to be anticipated; this effect is enhanced by intracellular acidosis and possibly magnesium accumulation. A striking series of recent findings suggests that the cell may be able to reduce calcium ion influx into the cytosol at low ATP concentrations.

4. The time-course of active force development and of force development between stimuli - the so-called unstimulated force during ischemia - and the mechanisms that may be responsible for the decrease of active force and for the development of unstimulated force during ischemia are discussed.

5. The role of accumulation of ions in the cell in abnormalities of excitationcontraction coupling is reviewed. It is proposed that the complex of ion-exchanging proteins at the surface membrane of the cell plays an important role in accumulation of sodium ions during ischemia which may underlie functional abnormalities that occur during reperfusion or re-oxygenation.

# Introduction

The effects of ischemia on contraction and relaxation of cardiac muscle function are still largely an enigma, understandably so because of the complexity of the effects of interruption of the flow through the coronary bed. The effects of stasis of coronary flow are evidently both to block the supply of nutrients and oxygen to the myocardium and to prevent washout of metabolites. Hence the ischemic insult is at least composed of the effects of impeded regulation of the extracellular milieu and of the derailment of the intracellular milieu as a consequence of the blockade of oxidative phosphorylation. Anoxia, which is often used as a model of ischemia, differs from ischemia itself because membrane-permeable metabolites are washed out of the extracellular space and hence also accumulate less inside the cell.

We wish to dissect the effects of components of this insult on the contractile process in this chapter. In the course of this "dissection", we will emphasize the effects of ischemia or anoxia or metabolic blockade on excitation-contraction coupling and on contraction. In particular the rapid initial changes of high-energy phosphate levels and of their breakdown products will be emphasized. We will then discuss the possible contribution of these concentration changes to the rapid loss of "contractility", to the failure of relaxation and to the mechanism that underlies the development of force in the unstimulated myocardium, i.e., the diastolic contracture.

The discussion will be based on the effect of anoxia on isolated trabeculae from the right ventricle of rat heart. These trabeculae were studied with the aid of laser diffraction techniques (61) to allow analysis of sarcomere behavior as a tool to evaluate both contractile processes and to investigate properties of the excitation-contraction coupling apparatus.

# **ATP-synthesis and Excitation-Contraction Coupling**

# Normoxia

ATP is predominantly generated by oxidative phosphorylation in the mitochondria of normoxic working myocardium. The average oxygen consumption of this process in a heart operating under normal conditions is 150 umol/s/L (57). Incorporation of phosphate ions (P<sub>i</sub>) in ADP yields 1 mmol ATP/s/L ( $\pm$  10%) depending on the main substrate used (57,70). Feedback mechanisms serve to maintain the levels of ATP, creatine phosphate (CP), ADP, AMP and protons as well as those of the breakdown products of AMP nearly constant in the working cell. Consequently, one may expect that the energy-requiring processes, i.e., excitation-contraction (EC) coupling, contraction and relaxation occur while the concentrations of ions and non-ionic metabolites fluctuate only modestly. Both the rate and magnitude of energy turnover by the EC-coupling apparatus and by contraction is thus expected to be independent of fluctuations of energy metabolism during normoxia.

Figure 1 depicts pathways of ATP supply to the most important components of excitation-contraction coupling and to the contractile filaments, that have emerged over the past decades. ATP is generated in the mitochondria by oxidative phosphorylation and exchanged for ADP by translocase at the inner mitochondrial membrane. Mitochondrial creatine kinase converts creatine in the intermembranous space into CP at the cost of ATP, and the released ADP returns from there to the mitochondrial core to be rephosphorylated. CP diffuses from the intermembranous space to the sites of ATP hydrolysis on the contractile filaments, the sarcoplasmic reticulum and the sarcolemma. Each of these structures is occupied with a creatine kinase that generates ATP close to the ATPases in the structure (5). The creatine formed in the process diffuses back toward the mitochondria.

It has been shown that this ATP supply to the ATPases at the cell's effector sites is complemented by localized glycolytic units. This is of importance because breakdown of glycogen is the main pathway for the generation of ATP during ischemia. Glycogen has been shown by ultrastructural techniques to be present in particles at specific strategic locations in the sarcoplasm, i.e., near the sarcolemmal Na<sup>+</sup>/K<sup>+</sup> ATPase, near the contractile filaments (the myosin ATPase and near the sarcoplasmic reticulum Ca<sup>++</sup> ATPase). Heilmeyer et al. (33,51) described in detail a particle isolated from skeletal muscle which contained glycogen, glycogenolytic enzymes, phosphorylase phosphatase and Ca<sup>++</sup> ATPase. Entmann et al. (17,18,23) described an SR-glycogen complex isolated from cardiac muscle, which also contains all the enzymes for glycogenolysis. These particles appear to be associated with the sarcoplasmic reticulum in situ as well as in the isolated microsomal fraction and seem to allow a specific functional coupling between ATP synthesis from glycogen and the SR. More recently, Weiss and Lamp have shown that local glycolysis may be a preferential source for ATP for membrane function (74).

The evidence that the protein-glycogen complexes are located near the ATPconverting enzymes could imply that during ischemia the rate of breakdown, and therefore the rate of depletion of glycogen, varies with location in the cell. Depletion of glycogen at a specific location could possibly have a large effect on the nearby ATPases while at other sites ATP supply and conversion will continue. This phenomenon may contribute to metabolic nonuniformity in the cell.

Another intriguing concept, emerging over recent years, is that ATP not only is a source of energy, but also regulates the function of membrane channels and ion exchangers (see Fig. 1). For instance the  $K^+$ -conducting channels mentioned above



Fig. 1. This figure shows a diagram of the excitation-contraction coupling system in the cardiac cell and a model of the structures involved in excitation-contraction coupling with ATP derived from oxidative phosphorylation and from regional glycogenolysis. Panel A illustrates that during the action potential, calcium enters the cell as a rapid flux followed by a maintained component of the slow inward current. The calcium inward current (indicated by the dashed line) does not lead directly to force development as the calcium ions that enter the cell are bound rapidly to the sarcoplasmic reticulum (SR) that envelops the myofibrils. The rapid influx of calcium is thought to trigger calcium-induced release of calcium by release sites of the sarcoplasmic reticulum. The released calcium ions activate the contractile filaments, and contraction ensues; relaxation follows because the cytosolic calcium is sequestered again by the sarcoplasmic reticulum and partly extruded through the cell membrane by the Na<sup>+</sup>/Ca<sup>++</sup> exchanger and by a low-capacity high-affinity Ca<sup>++</sup> pump. The force of a contraction is thus determined by the circulation of calcium from the SR to the myofilaments and back to the SR, and by the amount of calcium that entered the cell during the preceding action potential. The relaxation of the twitch depends on the rate of calcium dissociation from the contractile filaments and on the rates of calcium sequestration and extrusion. It is important to note that the Na<sup>+</sup>/Ca<sup>++</sup> exchanger is electrogenic such that calcium extrusion through the exchanger leads to depolarization of the cell. Panel B. Oxidative phosphorylation by the mitochondria yields ATP that is used to convert creatine (C) into creatine phosphate (CP) at the mitochondrial inner membrane. Both C and CP diffuse relatively freely through the outer membrane of the mitochondria and diffuse toward the myofilaments, the SR and to the sarcolemma. There, CP is converted into C while ADP is converted into ATP by creatine kinase (CK) thus supplying the enzymes involved in the excitation-concentration coupling process with energy. Enzymes involved in glycogenolysis are present in glycogen granules found at each of the structures that are relevant to excitationcontraction coupling. This warrants local supply of ATP irrespective of the ATP flux at other sites in the cell. Another important aspect of the role of ATP in the cell is that ATP (indicated by asterisks) controls the properties of the Ca<sup>++</sup> channel at the surface membrane (indicated by 1); the K<sup>+</sup> channels (indicated by 2); the Na<sup>+</sup>/Ca<sup>++</sup> exchanger (indicated by 3); and Ca<sup>++</sup> channels in the SR (indicated by 4); in such a way that, if the ATP level drops below 0.5 mM, K<sup>+</sup> channels open and "clamp" the membrane potential at the K<sup>+</sup> equilibrium potential. Then, Ca<sup>++</sup> fluxes through the sarcolemmal and SR channels are blocked while the Na<sup>+</sup>/Ca<sup>++</sup> exchanger is inhibited. Hence Ca<sup>++</sup> entry and efflux in a cell, in which the metabolism is blocked, is diminished and Ca<sup>++</sup>-induced cell damage is less likely to occur. For further explanation, see text.

are regulated by ATP (58,75). Similarly, ATP dependence of the  $Ca^{++}$  current in myocardium has been described (37). Another example with implications for ischemia is the observation that Na<sup>+</sup>/Ca<sup>++</sup> exchange in the squid axon is inhibited after cyanide blockade of oxidative metabolism; the exchange is restored by intracellular ATP (4). Regulation of SR calcium channels by ATP in skeletal muscle has been shown by Smith et al. (67). Recently Meissner's group also observed that the threshold at which Ca<sup>++</sup>-sensitive Ca<sup>++</sup> channels in cardiac SR open, increases if the [ATP] is lowered (62). The effect of reduction of the [ATP] on the K<sup>+-</sup> and Ca<sup>++-</sup> transport systems seems to be that the cell membrane potential is "clamped" near the K<sup>+</sup> equilibrium potential and that both calcium entry into the cell and Ca<sup>++</sup> release by the SR are suppressed.

### Ischemia

The situation described above that during normoxia EC-coupling and contraction are independent of fluctuations of the energy metabolism is suddenly lost when acute ischemia develops. An interruption of coronary flow exhausts the extracellular fluids acutely of oxygen. Oxygen flow into the oxidative phosphorylation stops. When oxidative metabolism stops, CP which normally is an intermediate of the high-energy phosphate (HEP) traffic, becomes the prime source of energy. The [CP] in the normoxic cells amounts to about 15 mM. It should therefore suffice to supply the cell following interruption of coronary flow for a period of 15 s, provided that the rate of consumption of ATP is initially unchanged.

The extremely high rate of conversion of ADP to ATP through hydrolysis of CP by the creatine kinases initially maintains the high ATP levels (5,38), but evidently at the expense of liberating  $P_i$  ions and creatine, and uptake of protons. The  $[P_i]$  and the pH rise. Because protons are strongly buffered in the cell (buffer capacity 70-80 mEq H<sup>+</sup>/pH unit; ref. 16), the pH rises only modestly.

The  $[P_i]$  rises rapidly following onset of ischemia as these ions are unbuffered. The initial rate of rise of the  $P_i$  level should logically equal the rate of ATP consumption in the cell with fully operative oxidative phosphorylation. As we have outlined above, the rate of ATP consumption and thus of  $P_i$  generation during normoxia, and hence immediately after onset of ischemia, equal about 1 mmol/s/L. Because  $[P_i]$  ions hardly permeate through the cell membrane, this implies that the intracellular  $[P_i]$  is expected to rise with 10 mM in 10 s.

As will be discussed below, the change of metabolite levels in itself has drastic effects on EC-coupling and on the contractile process. Consequently, the ATP expenditure by these processes will change rapidly after onset of ischemia as a result of changes in the metabolite levels. The decrease of the ATP consumption then evidently decreases in its turn the rate of generation of metabolites.

ATP levels fall when CP hydrolysis cannot keep up with ATP breakdown, and ATP metabolites are formed. The rise of the concentration of ADP, AMP and  $P_i$  accelerates anaerobic glycolysis rapidly after onset of ischemia (57), with a modest ATP yield, but at the expense both of a large increase in the [lactate] and of a large release of protons. The latter evidently causes a sharp decrease of the pH despite the ample capacity of the intracellular buffers (39). Moreover, the lactate production leads to a proportional and substantial increase in the osmolarity.

ATP levels decline rapidly during ischemia indicating that the maximal rate of ATP synthesis by anaerobic glycolysis (which amounts to about 0.3 mmol/s/L; ref. 63) fails to meet the ATP hydrolysis. ADP is partly converted to AMP, which allows a small amount of ATP to be regenerated through the adenylate kinase equilibrium (73). Most of the AMP, however, is degraded eventually into uric acid through adenosine, inosine, hypoxanthine and xanthine (28-30). The breakdown of the nucleotides in the cell is also accompanied by an increase in the cellular osmolarity.

# Effects of the HEP-Metabolite Levels on the Function of the Energy-Requiring Processes

The ATPases of the mechanisms that serve excitation-contraction coupling and contraction bind and hydrolyze ATP. They convert the available energy from these reactions into chemical or mechanical work and heat, while the products  $P_i$  and ADP are released. The rates of binding ATP and of release of the products are functions of the concentrations of ATP, ADP and  $P_i$ . The predictable effect of a decrease of the [ATP] and a rise of the [ADP] and  $[P_i]$  is thus to inhibit the energy-converting ATPases as is well-known for the contractile system (59,75). We will put relatively little emphasis on the concept that the phosphorylation potential (11,22,43) determines force development by ischemic cardiac muscle, as various authors have shown that the concentration of the breakdown products of hydrolysis is more important than the ATP/ADP  $P_i$  ratio (2,44).

The Km's of the ATP-hydrolyzing systems involved in EC-coupling are in the range of 1 mM (2). However, the depression of cardiac function occurs already at ATP levels that are substantially above the Km's of these enzymes (11). ATP shortage would, therefore, only influence these enzymes and the ATP-sensitive channels and ATP-sensitive exchangers, mentioned above, if spatial non-uniformity of the ATP distribution occurs in the ischemic cells, e.g., as a result of the high energy turnover in the contractile system. However, myocardium seems to be protected to some extent against such failure of its membrane functions by the localized ATP supply partially through glycolytic units (74).

The accumulation of ATP-breakdown products (e.g., adenosine), on the other hand, is known to affect sarcolemmal ion channels, and therefore to change the currents that flow through the cell membrane at rest and during the action potential. Hence, the initiating event of EC-coupling is depressed shortly after onset of ischemia. Because the action potential not only triggers contraction in the myocardium but also modulates the strength of contraction through variation of the influx of calcium during the plateau phase (53), a concomitant change in the strength of the contraction will ensue (see Fig. 1).

Many of the processes involved in EC-coupling and contraction are activated by the entry of  $Ca^{++}$  in the cell and subsequent release of  $Ca^{++}$  by the SR. Thus abnormalities of  $Ca^{++}$  handling by the cell are expected to affect cardiac function during ischemia and hypoxia (65). Activation of proteins by  $Ca^{++}$  invariably requires binding of  $Ca^{++}$  to anionic amino acid residues, such as carboxyl groups of tyrosine, as is the case with troponin C (48). It is well-known that protons compete with  $Ca^{++}$  for these binding sites but do not activate the protein to which they bind themselves (19). Similary Mg^{++} released during ATP breakdown competes for binding sites for  $Ca^{++}$  (20).

The rate and the efficiency at which the ATPases convert energy of ATP into other forms of energy, is further determined by the ionic conditions to which they are exposed. These effects are well-known for the contractile process, which, for example, is strongly depressed when the contractile filaments are exposed to high ionic strength (14). Similarly,  $Ca^{++}$  release by the SR is depressed in hypertonic solutions (24,36).

# The Effects of Ischemia or Hypoxia on Force Development by Trabeculae

Figure 2 shows that the typical response of a cardiac trabecula to hypoxia consists of a rapid decline of twitch-force development to low levels. The decline may be monophasic or may follow a more complex pattern with even an initial increase in force depending on the stimulus rate and on the loading conditions (Bucx et al., in preparation). The decline in active force development is accompanied by a slight increase in the unstimulated force. While unstimulated force develops occasionally, but not always, slight spontaneous activity of the sarcomeres may be observed microscopically or by laser diffraction methods (9). After 5-15 min of anoxia, unstimulated force rises steeply to levels that ultimately may exceed maximal twitch force by 50% (see Fig. 2).

This pattern is typical both for studies in which myocardium of isolated perfused hearts or of isolated superfused muscles is made ischemic or anoxic with maintained perfusion or in which oxidative phosphorylation has been stopped by  $CN^{-}$  ions, with or without suppression of anaerobic glycolysis (1). The rate at which the twitch force decays and unstimulated force develops, however, depends on the experimental paradigm that has been used.

The initial rise of twitch force has been attributed to a transient increase in the intracellular pH (1) due to breakdown of CP. The mechanism underlying the subsequent suppression of force development has not completely been elucidated yet. Both reduction of  $Ca^{++}$  release by the SR and depression of the contractile response to  $Ca^{++}$  are considered responsible. Decrease of  $Ca^{++}$  loading of the SR probably results from a diminished calcium influx into the cell (55). Two components of  $Ca^{++}$  entry into the cell have been studied particularly; both were depressed during ischemia. First, the amount of  $Ca^{++}$  entering during the action potential is reduced with the decrease of the duration of the plateau of the action potential. This occurs within a few min when hypoxia is combined with inhibition of glycolysis (49).



Fig. 2. This figure illustrates the time-course of active force (at 1 Hz stimulus rate, middle panel), unstimulated force (top panel) and sarcomere length (SL) in a cardiac trabecula of a rat during perfusion with a hypoxic Krebs-Henseleit solution without glucose at a temperature of  $30^{\circ}$ C. Force development during the twitches rapidly declined following onset of hypoxia followed by transient partial recovery and a further decline to zero. Unstimulated force initially rose slightly, which was accompanied by slight spontaneous activity of the sarcomeres; after 15 min, unstimulated force rose rapidly to a level that exceeded active force development in the control period by 50%. This force disappeared within 15 s after recoxygenation of the muscle (see inset); a few seconds later twitch force was apparent again. Spontaneous contractions immediately followed the electrically-induced twitches during the first 10 min after recoxygenation, and were accompanied by a Ca-dependent contracture that caused an increase of the unstimulated force. The signs of calcium overload subsequently disappeared and twitch-force recovery was complete after 30 min.

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One mechanism for the shortening of the action potential may be that adenosine, AMP or ADP activate a purinergic (P1) receptor and thereby increase the current through a K<sup>+</sup> channel (45). Weiss et al. (74) provided another mechanism for this phenomenon by evidence that glycolysis at the sarcolemma in a metabolically active cell is essential to keep the local ATP-sensitive K<sup>+</sup> channels (58) closed even during normoxia. Opening of these channels during hypoxia and interrupted glycolysis would cause a repolarizing current that shortens the action potential (49) and extracellular K<sup>+</sup> accumulation (66).

A reduction of the  $Ca^{++}$  current during the action potential as a result of the effect of lowered ATP levels such as described by Horie et al. (37) further decreases  $Ca^{++}$  influx during the action potential. This results in decreased loading of the SR and thus decreased  $Ca^{++}$  release. In a later stage of hypoxia, accumulation of Mg<sup>++</sup>, released as a result of ATP hydrolysis, further decreases SR loading (20). Eventually, at low ATP level, the  $Ca^{++}$  channels in the SR are less responsive (62), so that  $Ca^{++}$ -induced  $Ca^{++}$  release (53) is suppressed (see Fig. 1).

Secondly,  $Ca^{++}$  influx into the hypoxic cell in the period between action potentials is reduced. This has convincingly been shown in cardiac cells (31); it may result from a reduction of Na<sup>+</sup>/Ca<sup>++</sup> exchange at the lowered pH (60). Moreover, a reduction of the ATP concentration has been postulated to reduce Na<sup>+</sup>/Ca<sup>++</sup> exchange (4). The mechanism of this effect is unknown, but binding of ATP to the exchanger has been assumed to modify the affinity of the exchanger for Na<sup>+</sup> (54). Although these mechanisms are likely to play a role during ischemia, the channels and the exchanger are sensitive to ATP at levels around 0.2 mM. They therefore do not explain the rapid decrease of twitch force following onset of events mimicking ischemia.

The decrease of twitch force occurs too rapidly to be explained by a decrease of the free [ATP] (2,44). Both the rise in [P<sub>i</sub>] and the fall of intracellular pH have been shown to be important (44). The effects of P<sub>i</sub> and of protons occur so rapidly that they probably reflect a strong negative inotropic action on the contractile filaments (44). Protons and Mg<sup>++</sup> act by their competition with Ca<sup>++</sup> for binding to troponin C on the actin filaments (20), and P<sub>i</sub> probably through its inhibition of P<sub>i</sub> release from the crossbridge (44,75), which allows the development of the force-generating state of the bridge. Other possibilities that have been investigated are the accumulation of lactate and the concomitant rise in osmolarity; blockade of the lactate transport over the cell membrane does not mimic the effect of ischemia, however (15).

# The Effects of Ischemia or Hypoxia on Relaxation

The rate of relaxation of the twitch of cardiac muscle decreases shortly after onset of ischemia, as is illustrated in Fig. 3. The fact that this has not universally been found, though, may be related to the simultaneous decrease of the amplitude of the twitch which itself determines the rate of relaxation, as was found in rat cardiac trabeculae (Bucx et al., in preparation). The mechanism underlying the decrease of the rate of relaxation is located in a decreased rate of sequestration and exhaustion of Ca<sup>++</sup> from the cytosol. Ca<sup>++</sup> sequestration by the SR slows down (19), probably as a result of protons for the SR Ca<sup>++</sup>-ATPase. The effect of lowered pH alone is to decrease the rate of uptake of calcium ions into the sarcoplasmic reticulum (19).

Furthermore, acidosis causes a substantial Na<sup>+</sup> influx during ischemia and anoxia as a result of pH-dependent Na<sup>+</sup>/H<sup>+</sup> exchange over the cell membrane (6,41), while the resultant Na<sup>+</sup> accumulation is incompletely corrected as a result of inhibition of the Na<sup>+</sup>/K<sup>+</sup> pump (46). Intracellular Na<sup>+</sup> accumulation causes a diminished efflux of Ca<sup>++</sup> during the relaxation phase and hence is expected to further delay relaxation (64). The diminished efflux of  $Ca^{++}$  is not completely explained by the rise in Na<sub>i</sub> (26), suggesting dissociation of H<sup>+</sup>, Na<sup>+</sup> and Ca<sup>++</sup> transport over the cell membrane, as might be expected if the Na<sup>+</sup>/Ca<sup>++</sup> exchanger is inhibited at low ATP levels (4,54).

An increase of the heart rate causes an increased Na<sup>+</sup> and Ca<sup>++</sup> influx over time and therefore induces an increase in the intracellular Na<sup>+</sup> level. This again leads to slowing of relaxation and induces on the other hand more Ca<sup>++</sup> uptake by the sarcoplasmic reticulum. Ineffective extrusion from the cytosol at a high heart rate causes again the cytosolic calcium concentration to rise. It is also to be expected that, at a high heart rate, the amount of Ca<sup>++</sup> in the SR is increased. It is known that, under these conditions, spontaneous calcium release may occur (69). If such calcium release ensues during the relaxation phase, further prolongation of relaxation would occur, as has been observed in studies of sarcomere dynamics in hypoxic rat cardiac trabeculae (9).

An increase in the heart rate, e.g., by pacing, further delays relaxation and causes an increase in the diastolic pressure as has been shown by Apstein and Grossman (3). This phenomenon, which has been coined "demand ischemia", is consistent with the mechanism outlined above.

# Diastolic Compliance and Ischemia

Shortly after onset of anoxia in isolated heart-muscle preparations, a conspicuous rise in the unstimulated force follows, as was illustrated in Fig. 2. Many studies have been performed to clarify the source of this force, which also increases diastolic compliance. Two possible sources have been addressed in particular. First, it is conceivable that the diastolic force increases due to activation of the contractile filaments by a rise in the free intracellular [Ca<sup>++</sup>], as was discussed in the preceding paragraph. The second possibility is that the HEP levels and the levels of HEP metabolites have changed sufficiently to induce rigor.



Fig. 3. This figure illustrates the effect of 5 min of hypoxia and reoxygenation on the time-course of the twitch. The top panel of the figure shows the tracings of sarcomere length (S.L.) during control (N), during hypoxia (H) and during reoxygenation (R). The bottom panel shows the concurrent force transients during the twitches at N, H, and R. Note the occurrence of spontaneous late contractions manifest as sarcomere shortening transients during H and R. It is clear that the relaxation phase is protracted during H and R. For further explanation, see text. (From Bucx, Sethi, Ter Keurs; unpublished observations.)

The first possibility has been investigated in detail in isolated myocytes (12,31) and in trabeculae (2) with the use of calcium-sensitive luminescent or fluorescent substances. These studies have shown that the unstimulated force increased or the cells shortened clearly before an increase in the free Ca<sup>++</sup> level in the cell (2,31). It is therefore unlikely that the force is a result of the Ca<sup>++</sup>-mediated activation of the contractile filaments, especially because simultaneous acidosis of the cell reduces the sensitivity of the filaments to Ca<sup>++</sup>. The conclusion of these studies and of energetic studies was that the mechanical response to persistent anoxia is due to the development of rigor bonds between actin and myosin (35).

The hypothesis that rigor bonds exist between actin and myosin is supported by mechanical studies of the dynamic properties of the sarcomeres, as is illustrated in Fig. 4. The figure shows that the dynamic mechanical response of a rigor bond can be best compared to that of a simple stiff spring located in the S2 element of the crossbridge between actin and myosin. The spring should exhibit a stiffness which is independent of the rate (47,72) or frequency with which the length of the spring is altered. Such frequency-independent stiffness was indeed observed in our analysis of anoxic cardiac trabeculae. Figure 4 also shows that the stiffness of sarcomeres which develop a sustained calcium-dependent (10). Such studies are further required to establish the time-course of change of the elastic properties of myocardium during ischemia. This will allow us to unravel the contribution of calcium-dependent active crossbridge cycling and of development of rigor bridges during the rise in unstimulated force.

The question: "Why does rigor develop?" is still unsettled. Many studies have shown considerable rigor force development, e.g., 150% of maximal active force, while the ATP levels were still in excess of 0.5 mM. This contrasts the observations on skinned fibers both in skeletal muscle and in heart that rigor only develops, if the ATP concentration is lower than 50 uM (20). Three possible explanations are presently under investigation.

First, the rigor force may result from a complete rigor in a fraction (e.g., 90%) of the cells of the preparation, while the remainder of the cells is still relaxed. This would require severe non-uniformity of the cell properties throughout myocardium. Non-uniformity of the response of cardiac tissue during ischemia or anoxia (Bucx et al., in preparation) and of cardiac cells (68) to anoxia has been observed. The hypothesis requires that the cells in rigor are indeed devoid of ATP (<50 uM), while the remaining cells still contain a normal concentration of ATP. With a suitable ratio of cells with normal ATP levels to those without ATP, one would indeed expect that the residual ATP level in the bulk of the tissue could be 0.5 mM, while the myocardium behaves as in rigor.

Alternatively, rigor may have developed at much higher ATP levels than in isolated skinned muscle studies, because of the simultaneous elevation of HEP-metabolite levels. Prime candidates are  $P_i$  and ADP, as experiments (52) have shown that the ATP threshold for rigor force development is indeed elevated in the presence of increased concentrations of these metabolites. Cooperative interaction between Ca<sup>++</sup> binding at elevated Ca<sup>++</sup> levels and rigor-bond formation would further enhance the development of rigor bridges, as has been shown by Bremel and Weber (7). This question awaits further study of skinned cells at varied levels of ATP, ADP,  $P_i$  and Ca<sup>++</sup>.

Another hypothesis to explain the rigor under these conditions has been put forward by Gudbjarnason et al. (27) and has later been supported by the work of Bricknell et al. (8). The hypothesis states that during anoxia a block of transport of ATP may occur from the mitochondria to the cytosol. ATP would thus stagnate in the mitochondria, while ATP shortage would manifest itself at the sites of consumption, either as rigor or as dysfunction of the other ATP-consuming enzyme systems. Some evidence for the latter hypothesis has been obtained in studies of the ATP content of the mitochondrial and cytosolic compartments of cardiac (and other) cells. These studies suggested compartmentation of ATP in the mitochondria during hypoxia (21,34).



Fig. 4. The top panel illustrates a model of the structure of the crossbridge of myosin attached to actin prior to and during the power stroke (R and A). The deformation of the crossbridge that leads to force development is assumed to take place in the S1 head of the bridge; as a result of the power stroke the S2 link between myosin and the S1 head is stretched. Stretch of the elastic S2 link causes a force that pulls actin along myosin. The bottom panel shows the response of muscle force as a function of the frequency of sinusoidal length changes. Cyclic attachment and deformation of the bridge is assumed to occur during the calcium- and ATP-dependent contraction, while the bridge attaches and stays in state R during rigor. The muscle in the rigor state, therefore, behaves as a simple stretched spring, and is stiff at all velocities at which the spring is stretched. The cyclic behavior of the crossbridges manifests itself in a time-dependence of the force response upon stretch. The time-dependence is easily appreciated if the muscle is stretched with sinusoidal movements of small amplitude and the force response is expressed as a function of the frequency of the length changes as in the bottom panel. These properties are represented for the unstimulated muscle under control conditions, at maximal force development during hypoxia and during a caffeinemediated tetanus. It is clear from the illustration that the properties of the unstimulated muscle during control and hypoxia only indicate a difference in stiffness, while both are frequency independent, in contrast with the stiffness of the muscle during the caffeine tetanus which strongly depends on the frequency of the length perturbations. These observations confirm that the hypoxic muscle is in rigor, while crossbridge cycling takes place in the muscle during the tetanus. Tm = tropomyosin; Tn = troponin.

# Arrhythmias During Ischemia or Hypoxia

Accumulation of Na<sup>+</sup> causes, as we have seen, a diminished efflux of calcium through the Na<sup>+</sup>/Ca<sup>++</sup> exchanger. This effect is enhanced by the development of acidosis as a result of anaerobic glycolysis, which causes an increase in the Na<sup>+</sup> level in the cell (6,41,60). Hence, phenomena known to occur during calcium overload of the sarcoplasmic reticulum could develop (64). They include spontaneous contractions with concomitant transient depolarizations and consequently triggered arrhythmias. But they are depressed by other components of ischemia (13), e.g., inhibition by low ATP levels of Na<sup>+</sup>/Ca<sup>++</sup> exchange mediated currents that may underlie the transient depolarization (64).

Opie and his collaborators have shown (13) that under conditions of ischemia, several factors reduce the amplitude of the spontaneous after-depolarizations such as: lactate accumulation, acidosis and  $K^+$  accumulation in the extracellular space. These effects have been interpreted to mean that triggered arrhythmias which are presumably based upon the occurrence of spontaneous after-depolarizations are unlikely events in a severely ischemic region. However, in a region in which ischemia is less severe and where the inhibitory effects are less strong, this mechanism may induce arrhythmias.

Washout of the above metabolites during reperfusion of the ischemic tissue rapidly allows the spontaneous after-depolarizations to occur again. Moreover, the increased levels of intracellular Na<sup>+</sup> persist during reperfusion (25), which enhances the occurrence of spontaneous contractions and triggered arrhythmias (32), even if during ischemia no signs of Ca<sup>++</sup> overload were present. Hence, further Ca<sup>++</sup> entry may occur immediately following reperfusion or reoxygenation and this may lead to development of spontaneous contractions, transient depolarizations, arrhythmias, and ultimately cell death.

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# Chapter 17

# Myocardial Function and Metabolism during Ischemia and Reperfusion in Anesthetized Pigs

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Reduction of coronary blood flow in anesthetized pigs offers an attractive model of myocardial ischemia. In addition to data obtained in canine models, the study of myocardial ischemia in pigs yields important information for situations in which an extensive collateral coronary circulation is absent. In this chapter the effects of the most widely used anesthetic agents during normal and reduced porcine myocardial perfusion are reviewed. Recent data on myocardial function and metabolism during partial and complete coronary occlusion, and after reperfusion, are critically examined. Emphasis is given to the relationship between metabolic markers and recovery of myocardial function after reperfusion.

# Introduction

A large number of animal models is available to study myocardial ischemia and interventions which aim to protect jeopardized tissue or limit irreversible injury. The dog has usually been the animal of choice in in-vivo studies, but the use of domestic pigs has increased considerably during recent years (39,46). The pig is a suitable animal for cardiovascular research, in particular on myocardial ischemia and atherosclerosis, because its distribution of the coronary arteries (47) and its serum lipoprotein profile (21) are comparable to that of man. Moreover, at variance with dogs, in pigs atherosclerosis can not only be easily induced, but also develops spontaneously. Location and nature of these atherosclerotic plaques are comparable to those observed in man (35).

For the study of acute myocardial ischemia, it is important to know that hearts of domestic pigs, in contrast to those of dogs, lack an extensive collateral circulation. In pigs the residual flow is usually less than 5% after complete occlusion of a coronary artery (22). Thus the choice of the pig as an experimental animal in the study of acute myocardial ischemia may only be relevant for those clinical situations in which we are dealing with human hearts with low collateral flow as in previously undiseased hearts (32,47). Collateral growth can be induced in young growing pigs by placement of a plastic cylinder around a major coronary artery, which becomes flow-limiting as the animals grow older. With this technique collateral blood flow may become as high as 40% of the flow through the coronary artery before placement of the constrictor (23). The advantage of now having a preparation which has gone through episodes of myocardial ischemia for some length of time may be outweighed by the considerable loss of animals due to sudden cardiac death during this process of collateral induction. In our laboratory this amounted to 30% of the animals and occurred frequently in connection with excitement.

# Occurrence of Ventricular Fibrillation and the Effects on Myocardial Metabolism

In anesthetized pigs a complete occlusion of the left anterior descending coronary artery at its origin invariably leads to ventricular fibrillation in all animals within 30 min after the artery has been occluded (42). Ligation at a more distal site only slightly reduces the incidence of this arrhythmia (41). Using a flow reduction rather than a complete occlusion will decrease the incidence of ventricular fibrillation as a reduction of coronary blood flow to 25% of baseline causes ventricular fibrillation in only 30% of the animals. A still smaller decrease (to 50% of baseline) almost completely abolishes serious arrhythmias during the first 30 min (41-43).

The loss of animals can be avoided by defibrillation, but the metabolic changes that occur during the periods of ventricular fibrillation could invalidate the results obtained after such an event. However, when in pigs on cardiopulmonary bypass the myocardium was reperfused and sinus rhythm restored after ischemic ventricular fibrillation lasting 10 min, there was almost complete and immediate reversal of the changes observed in mitochondrial preparations isolated from the ischemic segments. These included partial uncoupling of oxidative phosphorylation, decreased oxygen uptake and loss of cytochrome oxidase activity (34). However, myocardial creatine phosphate (CP), which had decreased from 6 umol/g wet wt. to almost zero, and AMP and ADP, both of which doubled during ventricular fibrillation, recovered only partially (to 60%) after defibrillation. ATP (40% decrease during ventricular fibrillation) did not recover during the first 10 min after restoration of flow and sinus rhythm. Myocardial lactate levels (1 umol/g wet wt.) increased 20- and 30-fold during 5 and 10 min of ventricular fibrillation, respectively. They remained higher (5 and 7 umol/g wet wt., respectively) after 10 min of reperfusion and restoration of sinus rhythm. Because in most studies animals encountering a ventricular fibrillation during acute myocardial ischemia are only included for further study when sinus rhythm has been restored within 60 s, it is unlikely that, in particular in the ischemic tissue, significant damage may be caused during this short period. A critical evaluation of the effect of fibrillation on the metabolism of the adjacent non-ischemic myocardium has not been carried out.

# Anesthetics and the Effects on Cardiac Metabolism

The use of anesthetics is unavoidable in open-chest preparations. It is therefore important to realize that these agents may also affect cardiovascular responses to pharmacologic agents and modify the effects of myocardial ischemia and reperfusion. So far no evidence has been presented that anesthetics disturb the balance between oxygen supply and demand in normal perfused hearts in spite of effects on its most prominent variables. In pigs the most commonly used anesthetic regimens include pentobarbital, butyrophenones (azaperone, droperidol), alpha-chloralose, ketamine and inhalation anesthetics, such as halothane, enflurane and nitrous oxide. Neuromuscular blocking agents (pancuronium bromide and vecuronium bromide) are frequently combined with these drugs (37).

Pentobarbital reduces stroke volume by a direct cardiac depressant action, while heart rate and systemic vascular resistance increase. The drug, however, depresses the baroreceptor-induced reflex tachycardia as mean arterial pressure falls. Since the latter is a common feature during myocardial ischemia, the ischemia-induced responses may be modified accordingly. The butyrophenones, on the other hand, cause vasodilatation and thereby increase heart rate. A rather surprising finding has been that the combination of azaperone-metomidate anesthesia yields rather high arterial lactate concentrations (4 mM), possibly due to inhibition of cellular respiration which may account for the negative inotropic effects. The high arterial lactate levels make lactate the preferred substrate for the myocardium (60 - 70% extraction). Free fatty acid and glucose extraction are almost non-existent with this anesthetic regimen (5,43).

When two end-tidal concentrations of halothane (0.46 and 1.04 vol%) were tested in pigs, myocardial oxygen consumption decreased dose-dependently to 40% of the value obtained at 0 vol% due to the effects on myocardial contractility and mean arterial

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blood pressure. Myocardial lactate uptake was not affected (19). Arterial lactate levels increased up to 40%, possibly due to the inhibitory effect of halothane on hepatic lactate metabolism. Myocardial concentrations of ATP, CP and glycogen were not affected by halothane, suggesting that there was no interference with the energy balance. Inhibition of mitochondrial oxidative phosphorylation by halothane has been suggested to be responsible for its negative inotropic effects. However, on the basis of the data described above, it may very well be that the drug effects on cardiac work are predominantly caused by interference with myocardial cellular calcium fluxes. Halothane is one of the few anesthetic agents of which the effects on ischemic porcine myocardium have been studied. In one such study (20), it was demonstrated that halothane (0.32 vol% end-tidal) and fentanyl (50 ug kg<sup>-1</sup> as bolus followed by an infusion of 100 ug kg<sup>-1</sup>h<sup>-1</sup>) supplemented with nitrous oxide and pancuronium bromide had similar effects on arterial coronary venous differences of metabolic markers of myocardial ischemia (ATP catabolites and lactate production), cardiac output, myocardial contractility and left ventricular filling pressure after coronary artery blood flow was reduced by 60%. As halothane may induce malignant hyperthermia in a number of breeds, the use of enflurane, which has also less cardiodepressant actions, is preferable.

Pancuronium bromide is the most widely used neuromuscular blocking agent, but its monoquaternary analog vecuronium bromide exerts less cardiovascular actions. Neither compound has any effect on myocardial substrate utilization (6).

# The Effect of Restricted Coronary Blood Supply on Myocardial Function and Metabolism

In open-chest anesthetized pigs with unimpeded coronary blood supply almost all oxygen is extracted by the heart; oxygen saturation values of less than 15% are not unusual in blood sampled from the great cardiac vein (in pigs sampling of the coronary sinus does not render representative values because this vein receives also blood from the hemiazygous vein). In view of this high myocardial oxygen extraction, it is not surprising that a relatively small (25 - 30%) reduction in coronary blood flow already results in some loss of regional myocardial function (44). A reduction in flow by 60 - 70% results in a complete loss of function.

A relation, although relatively weak, has been established between the loss of contractile function and the appearance of the ATP-catabolite inosine in the coronary venous effluent during the first minutes of ischemia (4). The importance of this relationship between function and metabolism is weakened by the observation that the coronary venous concentrations of not only inosine, but also of most other metabolic markers of myocardial ischemia, remain not constant during prolonged periods of flow impairment (5,43). Thus after a reduction in coronary artery flow by 75% of baseline, the inosine concentration in the great cardiac vein increases rapidly from 20 to 120 uM during the first 20 min of ischemia. It then decreases gradually to 60 uM during the next 40 min (5). A similar time course, although less pronounced, has also been described for other nucleosides and for glycolytic breakdown products (5,43). Extreme care must therefore be taken if one wants to correlate functional changes, which are usually persistent, with those in appearance of breakdown products of ATP and glycolysis. Not only the rate of oxidative phosphorylation is markedly reduced at the onset of anaerobic conditions, but also the rate of ATP utilization is profoundly affected. Sufficient ATP may be present in severely ischemic myocytes to maintain certain functions (such as active ion transport) for a significant period of time. A more reliable parameter is therefore the tissue's energy charge, (ATP + 1/2 ADP)/(ATP + ADP + AMP), which reflects the imbalance between ATP utilization and ADP phosphorylation.

As stated before, presumably due to its high arterial concentrations, lactate is the preferred substrate during azaperone/metomidate anesthesia and both glucose and

free fatty acids are not taken up (43). With this anesthetic regimen, myocardial lactate uptake ( $31 \pm 8 \text{ umol/min}$ ) changes into release with a maximum value of  $48 \pm 9 \text{ umol/min}$  after coronary blood flow in open-chest pigs was reduced by 60%. Furthermore, now a slight uptake of free fatty acids ( $1.8 \pm 0.5 \text{ umol/min}$ ) was found. Using an in-situ working swine heart, Liedtke et al. (16) found that under pentobarbital anesthesia a 50% flow reduction yielded a lower glucose consumption and a lower free fatty acid extraction. Tissue levels of CP, ATP and lactate were not affected after 30 min. A further reduction to 40% of baseline was required before CP (by 69%) and ATP (by 25%) started to decrease.

Janse et al. characterized the "border zone" during myocardial ischemia in isolated blood perfused pig hearts (9,10). Histochemistry in biopsies revealed that glycogendepleted cells interdigitated with cells rich in glycogen. Because of the small size of the samples (3 mm in diameter), one must conclude that the demarcation between normal and ischemic myocardium is sharp. A similar conclusion was drawn from electrophysiological mapping. Of interest is also the observation that no transmural electrical and metabolic gradients were present within the ischemic myocardium. This is in agreement with the spatial development of irreversible tissue damage following coronary artery occlusion (36). On the other hand, Fujiwara et al. found a spatial wave front phenomenon of irreversible damage spreading from the subendocardium to the subepicardium during the first 40 min of ischemia which had disappeared after 2 h (8). The wave front phenomenon was independent of collateral blood flow. It may be related to the higher wall stress for the subendocardial layers which poses a higher oxygen-demand on these myocardial segments. When occlusions, complete or partially, lasted longer than 24 h an excellent relation existed between regional residual blood flow and regional function (7,31).

In contrast to the metabolic effects on the jeopardized myocardium, those on the adjacent non-ischemic myocardium have not very well been studied. Savage et al. (31) showed that in conscious pigs function of the non-ischemic myocardium increased considerably during the first 15 min after a coronary artery occlusion and persisted during the first 48 h. Since these changes in regional function were accompanied by a 20% rise in heart rate, myocardial oxygen-demand must also have increased. On the other hand, myocardial perfusion (42%) was only significantly elevated after 24 h. Liedtke et al. demonstrated that a moderate level of ischemia (52% flow reduction) caused not only a decrease in oxygen consumption of the underperfused myocardium, but increased oxygen consumption and free fatty acid uptake of adjacent non-ischemic myocardium (17). There was no commensurate increase in  $CO_2$  production from labeled palmitate, suggesting that fatty acids were utilized for acylation of glycerol-3-phosphate (ultimately leading to the formation of triglycerides). Tissue levels of long-chain acyl-CoA and acylcarnitine appeared to be increased. We also observed elevated tissue levels of long-chain acylcarnitine in adjacent non-ischemic myocardium after 60 min of complete coronary artery occlusion in pigs (12). This may be related to increased utilization of fatty acids for triglyceride formation in that region (12). In agreement with this finding is the observation by Liedtke et al. that during the last part of the 40-min period of ischemia, the energy balance of the adjacent myocardium was maintained by a twofold increase in glucose uptake (17).

# Recovery of Function and Metabolism of Ischemic Myocardium after Reperfusion

Depending on the duration and severity of flow reduction, the size of the coronary collateral circulation and the oxygen demand at the onset of ischemia, myocardial tissue may be reversibly or irreversibly damaged (33). The ischemic injury is reversible if restoration of coronary blood flow allows myocytes to survive and resume their function. Recovery of contractile function upon reinstitution of flow is usually not prompt and may last from several days to weeks. Thus the study of the

recovery of function requires a preparation which will not deteriorate for prolonged periods of time. This excludes isolated heart preparations and the open-chest anesthetized animal for such studies unless early markers of recovery of function become available.

If in pigs a coronary artery is occluded for less than 2 min, recovery of function is complete within 5 min (30). In contrast, when the preceding period of ischemia exceeds 20 min, no sign of functional recovery is observed after 2 h of reperfusion (28,40). After 30 min of ischemic flow reduction to 40% of baseline, recovery of function is only 50% during the first hours (45). Lactate production occurs during early reperfusion after a partial occlusion, but does not persist with the postischemic reduction in function and myocardial oxygen consumption (18). With the longer periods of severe ischemia, there was not only no sign of any recovery of systolic wall function, but end-diastolic wall thickness was increased after two hours of reperfusion (24). The latter was accompanied by a large accumulation of calcium in the tissue and in isolated mitochondria, while the ability of the isolated mitochondria to produce ATP was impaired. The authors concluded from these data that 30 min of ischemia in pigs leads to irreversible damage. Observations in our own laboratory undermine this conclusion. We found that animals, subjected to the same period of coronary artery occlusion, also exhibited no sign or recovery of function after 2 h of reperfusion. However, they demonstrated almost complete recovery of regional systolic wall thickening assessed by 2-dimensional echocardiography two weeks later (15,28).

Murphy et al. also observed that after 2 h of reperfusion following 30 min of ischemia, oxygen consumption, the ADP/O ratio and respiratory control index of isolated mitochondria, and myocardial ATP content, were all approximately 50% of that of the control segment. These were all less than 20% of control parameters in myocardium, which had been reperfused for 2 h after 60 min of ischemia (24). In view of these data, it is of interest that the mitochondrial and total tissue calcium contents increased to a similar extent suggesting that myocardial  $Ca^{2+}$  accumulation is mainly due to active  $Ca^{2+}$  transport into the mitochondria energized by substrate oxidation.

It is questionable whether the ATP concentration in myocardium is a limiting factor in the recovery of function. Van der Giessen et al. showed that pretreatment with the stable prostacyclin analog Iloprost enhanced recovery of function following 20 min of ischemia independent of the ATP content of the hypoperfused tissue at the end of ischemia and after 2 h of reperfusion (40). Peng et al. also showed that the ATP content of myocardium, which had been ischemic for 30 min, was 45% of control after 10 min of reperfusion and 50% of control almost 2 h later. This suggests that these levels do not increase gradually during early reperfusion (24,27). The time-course of ATP levels during ischemia and reperfusion should be known before definite conclusions regarding the value of ATP concentration measurements as a critical parameter, determining cell survival, can be drawn.

Reperfusion of ischemic myocardium has been reported to exacerbate damage, (often) leading to an acceleration of necrosis of potentially reversibly injured myocardium. The mechanisms involved are, despite strenuous efforts, still not fully understood (29). The swelling of cells, which occurs when normotonic fluids are used for reperfusion, may be one of the factors involved. In isolated porcine hearts, swelling may be so severe that capillaries will be obstructed and results in the no-reflow phenomenon (11,38). With normotonic fluids, lactate levels remain high, and CP and ATP levels low during reperfusion, but with hypertonic fluids CP and ATP levels are partially restored, and lactate levels are lowered. In pigs the role of hydrolysis of membrane phospholipids and formation of oxygen-derived free radicals in development of reperfusion damage has also been extensively studied (2,3,26). In a porcine preparation on an extracorporeal circulation, tissue levels of CP and the energy

charge decreased during ischemia. They recovered only partially during reperfusion, but completely when the free radical scavengers superoxide dismutase or catalase were present in the perfusion medium (2). In this study the ATP values did again not reflect recovery of function. The energy charge, which was lowered by only 5% during ischemia, recovered promptly with reperfusion when the free radical scavengers were present. In this respect it is important to note that xanthine oxidase possesses almost no activity in porcine and human myocardium and is therefore an unlikely candidate for generation of oxygen radicals during ischemia and reperfusion in these species (25).

Recently we have reported that the loss of integrity and function of the sarcolemmal and sarcoplasmic reticulum membrane, detected bv protein phosphorylation assays and freeze-fracture electron microscopy, may represent one of the earliest derangements causing irreversible myocardial damage (1,12,14,15,28). Several processes may be involved in the morphological changes: stretching of the bilayer by severe contractions and osmotic pressure changes, breakdown of membrane protein and lipids by proteases, phospholipases and free radical attack, rearrangement of phospholipid and protein molecules in the bilayer by lipid blebbing and fusion processes (12-15,28). The lipid blebbing and membrane protein aggregation, induced by solidification of negatively charged phospholipids, hexagonal II phase separation of phospholipids and neutralization of the charge of membrane proteins, may all be initiated by the increases in cytosolic  $H^+$  and  $Ca^{2+}$  concentrations. The membrane disruption that follows may cause the leakage of cytosolic proteins and bring the cellular  $Ca^{2+}$  movement further out of control. Spoiling of ATP by Ca<sup>2+</sup>-activated myofibrillar ATPase, excessive mitochondrial Ca<sup>2+</sup> accumulation and  $Ca^{2+}$  activated pumping ATPases in sarcolemma and sarcoplasmic reticulum are induced by the  $Ca^{2+}$  overload in the reperfusion phase. These data lend further support to the hypothesis that parameters other than the disturbance of energy metabolism should be taken into account in determining the extent of recovery after prolonged reperfusion.

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# Chapter 18

# The Energetics of "Stunned" Myocardium

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We review and report on experiments that address the mechanisms underlying "stunning" and "conditioning" of postischemic myocardium. Short coronary occlusions cause a long-lasting reduction of regional contractile function in the absence of cell necrosis. Regional oxygen-demand is lower as a consequence of reduced myofibrillar ATP turnover which leads to higher-than-normal phosphocreatine-levels. This sequence of events, coined "stunning", conditions the myocardium in such a way that a subsequent coronary occlusion is tolerated much better. Several arguments are provided that "stunning" is probably not caused by the only moderate reduction of the tissue ATP concentration. The hypothesis is put forward that the molecular mechanism of "stunning" is associated with a change in the sarcolemmal calcium ion homeostasis which reduces the calcium availability of the myofibrils. We hypothesize further that "stunning" may not be a special form of reversible ischemic damage but that it may be the unmasking of a pathophysiologic mechanism to protect against cardiac overload.

# A New Concept

In 1975 Heyndrickx et al. (7) described an observation, the significance of which became more important with time after the discovery: a short lasting coronary occlusion that did not lead to cell death produced a regional contractile dysfunction that did not recover for a surprisingly long time (days), whereas the electrocardiogram normalized within minutes. It was soon noted that the tissue concentration of ATP, that was moderately depressed by the short coronary occlusion, stayed depressed for a similarly long time (28). The intimate association between contraction and ATP was so suggestive of a causal relationship that a false sense of understanding prevailed, which inhibited awhile a true insight into the nature of "stunning". Recently, however, the buzz-word has entered the clinical laboratory because clinicians become aware of the fact that regional contractile function recovers slowly after successful thrombolysis and may therefore not be a good early end point to judge the success of the procedure. Clinicians have also reported cases of coronary occlusion with greatly impaired regional function, where expected to be dead (26). These cases were classified as "hibernating" myocardium by which is meant that the afflicted myocardium ceases to function and uses the little available energy to maintain its structure. These observations lead to the concept *that the heart may possess and utilize mechanisms that enable it to down-regulate its regional energy expenditure in the presence of reduced supply (37).* 

This is a relatively new concept which may soften the hard dogma of the supplydemand concept that dominated the pathophysiology of coronary artery disease for a long time. It would also be prudent not to overemphasize this new concept because the hard reality is that many patient's myocardium cannot be salvaged by reperfusion because it failed to enter the "stunning" or "hibernating" state within the allotted time window of reversible damage. However, *it appears to be of great*  scientific and practical interest to study the processes that enable the heart to cut its contractile engine in times of reduced supply in order to maintain structural integrity. Evidence that such mechanisms exist came from experiments where coronary occlusions of short duration were repeated. These experiments showed that the first occlusion left a trace, induced a memory and set the stage for the following occlusions whose impact on the myocardium was then much less injurious. One of the most convincing experiments was reported by Jennings' and Reimer's group (19,29), where a coronary occlusion of 40-min duration that always produced large infarcts, did not produce sizeable infarcts when the 40-min occlusion was preceded by a 10-min occlusion. A longer (total) ischemia time had in essence greatly reduced the ischemic damage, because the first occlusion had "conditioned" the myocardium in a way that is not mechanistically clear. The experiment does not say that infarcts are avoidable, if myocardium is "conditioned". However, the ultimate cell death is significantly delayed. Delayed-but-not-prevented is the important message (35,36) that we have learned from 10 years of hard, but not always successful, labor to reduce infarct size (36). The observation that a few drugs can delay, but not prevent, infarcts hints at these mechanisms that the heart can utilize to defend itself against undue demand.

All experiments described in this review point toward a change that occurs in the perception that we have of the role of the heart in general circulatory physiology: classical dogma has it that the heart under physiological conditions is always the obedient slave to the needs of other organs, ready to accept short-term energetic deficits, unable to refuse a demand even if selfinterest (in pathologic states) would dictate a slow-down. The classical observation for the physiologic energy deficit is the reaction of the coronary vasculature to a postextrasystolic beat: the potentiated beat produces a larger power output, but coronary flow increases only with the next beat, i.e., the deficit is first, payment later. This dogma holds also for pathological states: if a coronary occlusion occurs in the "unconditioned" state the afflicted myocardium does not down-regulate its needs. It continues to spend more (in futile attempts of contraction) than it receives. It beats itself, literally, to death.

The research of the last 10 years has modified this dogma in that we became more aware of situations where the heart shows the limited ability to defend itself against injury, or to protect itself against excessive loads. Besides stunning, other possible mechanisms of defense are the down-regulation of beta-receptors in congestive heart failure (17), the myocyte atrophy in chronic ischemia (33), the release of natriuretic factor at volume overload (3), and the development of a collateral circulation in chronic ischemia (34,35).

In the following paragraphs, we will describe the experiments that have led to this concept that cedes the myocardium a limited escape from the rigors of the control system.

# The Role of Adenine Nucleotides in "Stunned" Myocardium

As said above: the relationships between contractile function and ATP are so close that a situation where both are decreased is almost selfexplanatory. However, it has been pointed out that there is no close relationship between the tissue [ATP] and contraction (6,20,23,27) other than a very general one in the sense that a very low ATP content does not enable a high ATP-turnover. It is the turnover of ATP rather than its tissue concentration that is closely related to contraction. There is, however, also good evidence that the tissue [ATP] may control its turnover, as shown by Kammermeier et al. (12), because ATP-utilizing enzymes require a high energy level. We became interested in the problem some years ago, when we studied the myocardial synthesis of ATP from salvaged precursors. The question was: can tissue-ATP levels that had been decreased by reversible ischemia be restored by selective administration of nuclosides? And: does the restoration of ATP improve the The Energetics of "Stunned" Myocardium

function of "stunned" myocardium? The background of this problem was the observation that the de-novo synthesis of ATP is very slow (about 3 nmol/h per g heart muscle; ref. 40), requiring about 4 days for nearly complete replenishment. We infused adenosine, 5-amino-4-imidazolecarboxamide riboside (AICA-riboside) and ribose into the left coronary artery of in situ beating dog hearts (anesthetized, open chest; ref. 18) after a period of 45 min of ischemia, and obtained needle biopsies at regular intervals after the beginning of infusion. We measured the tissue concentration of adenine nucleotides and nucleosides with HPLC and we measured the specific radioactivity of the nucleotide pool after infusion of radioactive adenosine,  $^{3}$ H-AICA-riboside and of  $^{14}$ C-glycine (for de-novo synthesis).

Stimulation of ATP-synthesis by ribose was described by Zimmer in the rat heart (20). Phosphoribosylpyrophosphate is in short supply in the postischemic heart and the exogenous supply of ribose increases the ATP-synthesis. AICA-riboside is the immediate precursor of IMP. It was also shown to increase the ATP pool size and to accelerate the postischemic repletion of ATP (18,38). Adenosine is phosphorylated to AMP, which is converted to ATP by the adenylate-kinase reaction. As Table 1 shows, adenosine infusion is the most rapid means to replenish postischemic ATP. It increases the rate of ATP-synthesis over that of the de-novo rate by a factor of 200. It was the only substrate that increased the tissue [ATP] significantly during the first 3 h of reperfusion. Ribose and AICA-riboside increased the rate of synthesis, but not to an extent that would have resulted in increased steady-state levels.

We then tested the hypothesis whether repletion of ATP resulted in improved regional postischemic contractile function (8). We tested this by implanting two pairs of sonomicrometer crystals into the subendocardium, one into a potentially ischemic, the other into a control region. We then studied the return of function after a period of 45 min of coronary occlusion in the presence or absence of exogenous (intracoronary infusion) adenosine and AICA-riboside. Although adenosine (but not AICA-riboside) caused significant increases in the tissue [ATP], postischemic regional function was not improved. AICA-riboside significantly depressed regional function via the synthesis of AICA-triphosphate (41). We concluded from this experiment that the dysfunction of "stunned" myocardium may have little to do with the tissue [ATP].

As we know now, other variables may enter the equation. Our own experiments (5) have shown that the coronary endothelium rapidly deaminates adenosine. Others have shown that endothelium predominantly phosphorylates adenosine (21). The true [adenosine] at the myocyte level following exogenous infusion is therefore not known. Although it is feasible that a part of the synthesized ATP resides also in the endothelium, it is not very probable that all of it is concentrated in endothelial cells. Given a volume fraction of endothelium of 3%, the final "endothelial" [ATP] after adenosine infusion would reach over 60 mM, i.e., clearly not compatible with solubility and osmolality. Another complicating factor is the inhibition of adenosine

Table 1. Purine nucleotide de novo synthesis and accelerated synthesis rates after 3 hours of intracoronary substrate infusion in previously ischemic and nonischemic myocardium

	De novo	Ribose	AICA-riboside	Adenosine	
Postischemic	$3.1 \pm 0.3$	$15.1 \pm 0.4$	27 ± 3	268 ± 45	
Nonischemic	$1.5 \pm 0.3$	9.8 ± 0.5	28 ± 4	$219 \pm 36$	

Rate of incorporation of precursors into the canine myocardial ATP-pool during precursor-infusion at reperfusion following a 45-min occlusion period. Adenosine is the most potent precursor. The rate of ATP synthesis from adenosine is 180-times faster than the de-novo synthesis without exogenous infusion of precursors. Data are given in nmol/h per g wet wt. AICA-riboside = 5-amino-4-imidazolecarboxamide riboside.

kinase by one of its co-substrates, i.e., adenosine itself (22). The infusion of relatively large amounts of adenosine into a coronary artery, which resulted after 3 h in a significant increase of ATP levels in the dog heart, may have been the fortuitous outcome of activation of adenosine deaminase in the endothelium that reduced the actual [adenosine] at the myocytes to levels that favor phosphorylation. Own experiments in isolated buffer-perfused rat hearts (Görge and Schaper, unpublished) that may not be entirely comparable with in situ beating blood-perfused dog hearts, showed that adenosine was phosphorylated only at effective concentrations below  $5 \cdot 10^{-5}$  M.



Fig. 1. Reduced regional O<sub>2</sub>-consumption leads to down-regulation of blood flow. Subendocardial blood flow (upper panel, radioactive tracer microspheres) was measured in a control region (Con) and in an ischemic region (Occ 1, Occ 3) that was reperfused after three occlusions lasting 10 min each. Subendocardial blood flow was also measured at 0.5, 1, 2, and 3 h after reperfusion. Subendocardial blood flow falls to about 60% of that in a control region. Regional transmural oxygen consumption was measured from the transmural flows multiplied with the regional (arterio-)venous oxygen saturations. The figure shows that "stunning" was accompanied by reduced regional MVO<sub>2</sub> that led to a down-regulation of blood flow. \* P < 0.05 vs. Control; \*\* P < 0.001 vs. Control.

In conclusion, we have provided evidence that intracoronary infusion of adenosine into a region of "stunning" results in an ATP increase in that region, which is much more than can be accounted for by an increase only of the endothelial pool of ATP. On the other hand, the increase in myocyte ATP may have been too small to definitively refute the hypothesis that changes in the ATP level are causative in "stunning".

# Defective Energy Utilization vs. Energy Supply as a Cause of "Stunning"

One of the symptoms of "stunned" myocardium is the reduction of subendocardial blood flow after the hyperemic phase of reperfusion (see Fig. 1). We investigated the question of a microcirculatory disorder as a cause of dysfunction. We studied the distribution of left ventricular blood flow with radioactive tracer microspheres and the saturation of coronary venous blood with a differential oximeter. The oxygen saturation in two epicardial veins was studied, i.e., one draining from a previously ischemic region, the other from a normal region. About 1 h after onset of reflow following 10 min of occlusion, no differences in oxygen saturation between the two sampling sites were observed and it was assumed that these values were representative for the regions drained. The subendocardial blood flow determined at this moment had fallen by 25% below the value of normal neighbor-territories of the same heart. This leads to the conclusion that the "stunned" subendocardium consumed less oxygen, i.e., the general relationship between work output and oxygen consumption was maintained and blood flow was down-regulated to meet the lower demand. In order to exclude a primary microcirculatory disorder, we infused adenosine and we were able to show that dilatory reserve was normal, which makes "plugging" of microvessels as a cause of subendocardial flow reduction very unlikely. A microcirculatory disorder should also maintain the ischemic state well into the reperfusion period and it should further deplete ATP, which in fact does not occur.

Our experiments lead us to believe that the reduced  $O_2$ -uptake of "stunned" myocardium is not the expression of an energetic deficit, but rather a disturbance of energy utilization. In another series of experiments, we intracoronarily infused CaCl<sub>2</sub> into a "stunned" region. We could demonstrate that the region contracted normally again, even regained its maximal level of systolic shortening (10,37). Function returned to normal levels after stopping calcium infusion and the responses did not fade after several cycles. In yet another experiment, postextrasystolic potentiation showed almost complete restoration of function in a segment of "stunned" myocardium (see Fig. 2). These experiments demonstrate clearly that energy supply is not yet limited and that the contractile machinery is not damaged: vascular adaptation to increased demand is intact, mitochondria convert enough energy, myofibrils contract to the same maximal level if stimulated. We conclude therefore that "stunning" may be an expression of altered electromechanical coupling and that the most likely organelle, that underwent the change, is the sarcolemmal membrane. We tend to believe that "stunning" is not a defect or a damage, but rather a functional response to stress, perhaps related to down-regulation of voltage-dependent calcium-channels. We look upon "stunning" as an unmasking of a physiological response that is better developed in other organs (i.e., skeletal muscle), because the defense against myocardial ischemia was never put under the pressure of natural selection, because coronary heart disease develops relatively late in life.

# **Repeated Coronary Occlusions do not Produce Additive Effects**

Several groups (9,16,19,29,38) have studied repeated short coronary occlusions for a variety of reasons: to study the influence of several drugs on ischemia in the same heart (2), to study the behavior of the nucleotide pool, and to study the

development of infarcts (19). We have tried repeated ischemia to test a hypothesis that eventually proved to be wrong. We and others (9,38) have shown that only one short occlusion, lasting between 3 and 5 min, is sufficient to produce a measurable fall of myocardial ATP. Given the known long time requirements for ATP de-novo synthesis, we argued that repeated occlusions of relatively short duration must therefore lead to cumulative ATP depletions that may cause irreversible tissue damage. This appeared to be an attractive clinical hypothesis, because repetitive anginal attacks (such as in unstable angina) may lead to infarction even without complete coronary artery occlusion. Cases were reported (30) where patent coronary arteries existed shortly after a myocardial infarction. Alternative explanations are of course possible in these cases. Our experiments soon showed that repeated occlusions did not produce cumulative losses of ATP. The relatively largest loss occurred after the first occlusion, when that lasted for 10 min, or after the first 5 occlusions, when these lasted for 3 min. More occlusions had an increasingly diminishing effect, even 40 occlusions of 3 min interspersed by 3 min of reperfusion did not produce irreversible damage. From previous experiments it was quite clear that the first occlusion(s) had reduced the myocardial O<sub>2</sub>-consumption, that "stunning" had occurred which changed the energetic situation at the onset of the next occlusion. The myocardium at risk will hence become less ischemic with the next occlusion. The analysis of nucleotide breakdown products confirmed this: less ADP, AMP, and relatively much less adenosine were produced compared with the first occlusion. We hypothesized that in repeated occlusion the myocardium "learns" how to stop the contractile process quicker in order to conserve energy and we expected an earlier and more complete "bulging" with repeated occlusions. This was difficult to show, but in these experiments systolic shortening was close to zero during the first occlusion whereas bulging was -20% in repeated occlusions.



Fig. 2. Effect of postextrasystolic potentiation (PESP) on segment function of dog heart. Systolic segment shortening was measured with subendocardial sonomicrometer crystals. Following occlusion, systolic shortening falls quickly below zero within 1 min. A properly spaced extrasystole leads to PESP of contraction that elicits a systolic shortening up to 2 min of coronary occlusion. At 1 min into the occlusion, an almost normal contraction can be elicited (open circles). Solid circles: effect of PESP on non-ischemic region contraction. LAD = left anterior descending coronary artery.

# The Energy Balance of Ischemic Myocardium

When a coronary artery is suddenly occluded, the contractile function becomes weaker after 6 to 8 beats, thereafter contraction ceases and the affected ventricular wall shows systolic outward movement known as "bulging". Although it is understandable that the risk region ceases to contract when the substrate supply is abruptly cut, the still unanswered question is why the tissue [ATP] stays constant for a few minutes. Several hypotheses have been forwarded:

1. The decline in contraction following occlusion follows roughly the time course of fall of phosphocreatine (PC). This would support the hypothesis that only ATP synthesized from PC in close proximity to the myofibrils is the substrate for contraction. It is the "phosphocreatine shuttle" (11,32) that is responsible for the energy transfer between mitochondria and myofibrils. The fall in  $PO_2$  in the mitochondria produces fewer molecules of PC, on-site phosphorylation of ADP at the myofibrils falls and contraction stops. The high cytosolic [ATP] remains untouched for several minutes, because its accessability for the myofibrils is low (1).

2. The inorganic phosphate that is produced by the cleavage of high-energy bonds may combine with cytosolic calcium, which reduces the availability of calcium ions at the myofibrils, thereby reducing the strength of contraction (15).

Whatever the precise mechanism for the cutting of the contractile engine is, two processes occur at the same time but in different directions: one that continues to respond with contraction to the stimulus of depolarization; and the other tries to reduce the electromechanical coupling.

Figure 2 shows an unpublished experiment where extrasystolic depolarizations caused potentiated beats during the course of declining contractility following coronary occlusions. The record shows that potentiated beats can be elicited up to 3 min following coronary occlusion, i.e., absolute lack of energy cannot have been the only cause of contractile failure. It is feasible that the amount of calcium ions available to the myofibrils had also decreased. The extrasystolic beats had provided an extra amount of calcium ions, which in turn had produced the potentiated contraction. This experiment hints at the existence of regulation of contraction in ischemia that is not exclusively dictated by the energy needs.

On the other hand, ischemic myocardium continues to respond to the stimulus of depolarization. However weak and inefficient, it accumulates a debt and proceeds toward irreversible injury.

That ischemic myocardium continues to beat during reversible ischemia is not obvious from sonomicrometer records. The nature of this method is, however, that small differences in regional contraction are amplified. It is intuitively clear that the two segments studied (an ischemic and a control segment) may differ only a little in the strength of contraction in order to obtain a large effect in the ischemic segment, because both regions are connected to each other in a tug-of-war situation.

We approached this problem by comparing the systolic bulging with the bulging caused by intracoronary infusion of procaine amide (see Fig. 3). It is apparent that the procaine hearts bulged more, indicating that hidden in the systemic bulge weak contractions are "buried".

The importance of this discussion about the mechanisms of "stunning" and "conditioning" lies in the fact that, if the heart would better utilize the mechanisms of ischemia-related-electromechanical uncoupling, infarcts can almost be completely avoided.

Our experiments in the canine heart showed (36) with the exception of the subendocardium of the posterior papillary muscle, that collateral flow is perfectly capable to maintain the structure of the myocardium, if all attempts at mechanical activity were reduced to zero. Our experiments also showed that the heart has the ability to adjust to the lower level of oxygen supply as provided by collateral flow. Jennings' and Reimer's group (19,29) has shown that "conditioning" significantly delays infarction. Our experiments with intracoronary verapamil, that did not produce infarctions following 3 h of left anterior descending coronary artery occlusion, support our hypothesis that a partial electromechanical uncoupling with reduced  $O_2$ -uptake is the immediate mechanism of protection. The following energetic balance shows that ischemic myocardium can reduce, however imperfect, its metabolic-electromechanical coupling, but that adverse stimuli can override this protective effect.



Fig. 3. Systolic shortening following coronary occlusion and reperfusion (upper panel), and following intracoronary procaine infusion in the same protocol (lower panel). "Bulging" in ischemia is much less pronounced than with procaine infusion, indicating some active force production in the ischemic region. Function completely returns to normal after stopping of procaine and does not produce "stunning" which indicates that "stunning" is not caused by mechanical factors. Seg. short. = segment shortening.

The Energetics of "Stunned" Myocardium

Under normal basal conditions (anesthesia with physiologically low heart rates), coronary flow in a canine heart is about 50 ml/min/100 g. This represents 8 ml of extractable  $O_2$ /min which is converted into a rate of 22 umol ATP/min per g wet wt. Shortly after coronary occlusion, tissue perfusion by collateral flow is maintained at a rate of 25 ml/min/100 g for the subepicardium and of 8 ml/min/100 g for the subendocardium. These represent an aerobic ATP production of 9 vs. 3.5 umol/min per g wet wt., respectively. These rates of aerobic ATP production are supplemented with glycolytic ATP production in the order of 0.2 umol/min per g of high-energy phosphate bonds. The rate of metabolic spending in ischemia is, however, faster. When we contrast the rate of fall of the tissue [ATP] with the rate of ATP production, we can define the degree of "overspending". This rate of overspending is surprisingly small in the subepicardium (3.5% more energy is spent than received *per minute*).

Repetitive occlusions (which unmask the existence of energy-saving mechanisms) reduce the amount of overspending by mechanisms that are as yet not well understood. The indicator for reduced overspending is the tissue [ATP] which is the result of ATP production and ATP utilization. The influence of repeated occlusions on tissue ATP is greatly blunted, which is most probably due to the O<sub>2</sub>-sparing effect of "stunning" that results from changes in Ca<sup>2+</sup> homeostasis. Another possible mechanism would be a decline in Ca<sup>2+</sup> sensitivity of the sarcomeres (31).

# Controversies

Not all workers in this field believe that "stunned" myocardium is a desirable, useful and probably physiological mechanism that one should amplify and utilize in order to delay infarction or to reduce infarct size. Several reports (4,25) show evidence of a cellular damage by oxygen free radicals because radical scavengers can prevent "stunning". However, beta-adrenoreceptor blockade was also shown to prevent "stunning" (14) as well as treatment with adenosine and with calcium channel antagonists (13,24). The unifying theory would be that calcium homeostasis is involved in the phenomenon, because all mechanisms mentioned utilize calcium as a final common pathway including adenosine which is a calcium chelator.

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# Chapter 19

# **Energy Depletion due to the Calcium Paradox**

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Reperfusion of an isolated mammalian heart with a calcium-containing solution after a brief calcium-free perfusion results in an excessive influx of calcium into the cells and irreversible cell damage: the calcium paradox. A rapid depletion of myocardial high-energy phosphate stores is one of the characteristics of the calcium paradox. During the first 30 seconds of calcium repletion in isolated rat heart, intracellular creatine phosphate and ATP levels decreased by 65% and 45%, respectively, and creatine, ADP and AMP levels increased by 15%, 85% and 2800%, respectively. The effect of different energy states of the heart on the capacity to develop the calcium paradox was studied in the anoxically perfused (with and without substrate) and reoxygenated rat heart. The presence of either ATP or electron transport appeared to be a prerequisite for the occurrence of the calcium paradox. It has been suggested that acidification of the cytosol, as a result of hydrolysis of ATP and accumulation of calcium by mitochondria, is an important factor in the development of the calcium paradox. By using phosphorus nuclear magnetic resonance spectroscopy, the time course of intracellular pH, measured from the chemical shift of the intracellular inorganic phosphate peak, was investigated in isolated rabbit heart. Intracellular pH during the calcium paradox varied from 7.1 to 7.0, indicating that acidification of the cytosol does not play a causal role in the development of the calcium paradox.

# Introduction

Ringer demonstrated that contraction of the heart rapidly ceases when calcium is removed from the extracellular fluid (22). Zimmerman and Hülsmann reported that reintroduction of calcium into the extracellular fluid does not result in recovery of contraction of the heart, but in irreversible cell damage (28,29). They introduced the term "calcium paradox" to describe this phenomenon. The calcium paradox is characterized by an excessive influx of calcium into the cells (1), depletion of endogenous high-energy phosphate stores (2), severe contracture of the myofibrils (8), and rapid loss of intracellular constituents (29).

The influx of calcium into the myocardial cells may be divided into an early, relatively small gain in cytosolic calcium, and a subsequent, massive influx (18). Evidence has been provided that entry through the slow channels (17-19) and via the sodium-calcium exchange mechanism (3,10,18,23,26,27) are involved in the primary gain in calcium. Additional factors that may be responsible for the early gain in cytosolic calcium are the calcium pumps of the sarcolemma and the sarcoplasmic reticulum, whose activities are decreased by calcium-free perfusion (15,21). The raised cytosolic calcium may then trigger a number of events, including energydependent mitochondrial calcium accumulation (24) and activation of various ATPases (11), contracture-mediated disruption of intercalated discs (8), disruption of the sarcolemma (20), loss of intracellular constituents (29), and a secondary uncontrolled entry of calcium (18). Neither the sequence of these events nor their relative importance is known at present.

# **Energy Depletion during the Calcium Paradox**

A rapid decline of myocardial high-energy phosphate stores is one of the characteristics of the calcium paradox. This has been demonstrated by the freeze-clamp method (2) and by using phosphorus nuclear magnetic resonance ( $^{31}P$  NMR) spectroscopy (5).

Table 1 shows data, obtained with the freeze-clamp method, on creatine phosphate (CP), creatine and adenine nucleotide (ATP, ADP and AMP) levels in isolated rat heart during calcium-free perfusion and the calcium paradox (2). During calcium-free perfusion, when no energy is required for mechanical activity, CP and ATP levels increased by 20% and 10%, respectively. During the first 30 seconds of calcium repletion, myocardial CP and ATP levels decreased by 65% and 45%, respectively. In the same period there was an increase in creatine (15%), ADP (85%), and AMP (2800%). During continued reperfusion with calcium, the concentration of all compounds gradually decreased. The effluent fluid contained large amounts of creatine and AMP, and relatively small amounts of CP, ATP and ADP.

The finding that myocardial CP and ATP levels were decreased after 30 seconds of reperfusion with calcium, whereas creatine, ADP and AMP levels increased during this interval, points to a sudden consumption of high-energy phosphate stores. When the sum totals of ATP + ADP + AMP (Ad Nu) and of CP + creatine are considered, it is clear that there was hardly any loss of these intracellular compounds up to the first 30 seconds of reperfusion. After this period a decrease of these sum totals can be explained by loss into the extracellular space and by further degradation of AMP.

#### **Energy Dependence of the Calcium Paradox**

By perfusing isolated rat hearts at 37°C under anoxic conditions with and without substrate, and by reoxygenating hearts after an anoxic period, the effect of different energy states of the heart on the capacity to develop the calcium paradox has been studied (24).

After 35 min of anoxic perfusion (30 min with calcium; 5 min without calcium) in the *presence* of glucose (Fig. 1a), the hearts still contained 15  $\pm$  3 umol ATP per g dry heart tissue (mean  $\pm$  S.D.; n = 4), which is 80% of the amount of ATP present

Perfusion sequence		C	CP	Creatine	ATP	ADP	AMP	Ad Nu
+Ca	15 min	30	± 2	32 ± 5	18.8 ± 0.8	$3.2 \pm 0.4$	0.21 ± 0.04	22.2
-Ca	4 min	36	± 3	32 ± 3	20.8 ± 1.4	3.3 ± 0.3	0.18 ± 0.06	24.3
+Ca	0.5 min	12	± 5	37 ± 8	11 ± 2	6.1 ± 0.6	5 ± 3	22.1
	1 min	5	± 3	33 ± 5	6 ± 3	4.7 ± 0.5	5 ± 2	15.7
	2 min	4	± 2	23 ± 5	5 ± 2	$3.2 \pm 0.4$	$1.8 \pm 0.5$	10.0
	4 min	1.6	± 1.2	$10 \pm 5$	$2.9 \pm 1.0$	$2.4 \pm 0.8$	$1.1 \pm 0.5$	6.4

Table 1. Effect of calcium-free perfusion (-Ca) and reperfusion with calcium-containing solution (+Ca) at 37°C on creatine phosphate (CP), creatine and adenine nucleotide levels in isolated rat heart

Results are expressed as umol/g dry heart tissue (mean  $\pm$  S.D.; n = 4). Ad Nu = Sum of adenine nucleotides.

at the end of the control perfusion period (see Table 1). No measurable amount of creatine kinase (CK) was released from the hearts during this period of anoxic perfusion. Reintroduction of calcium to the anoxic glucose-containing perfusate resulted in a massive release of CK from the hearts.

After 35 min of anoxic perfusion (30 min with calcium; 5 min without calcium) in the *absence* of glucose (Fig. 1b), the myocardial ATP content was only 2% (0.39  $\pm$ 0.13 umol/g dry heart tissue) of that at the end of the control perfusion period. The absence of glucose and calcium in the perfusate caused a slow release of CK. After reintroducing calcium to the anoxic glucose-free perfusate, enzyme release increased slightly, indicating that the calcium paradox did not occur in hearts subjected to anoxic substrate-free perfusion. However, reoxygenation resulted in a prompt and massive release of CK. This oxygen-induced calcium paradox was completely inhibited when KCN was present during the last 15 min of anoxic perfusion and during reoxygenation. This indicates that reactivation of the electron transport



Fig. 1. Effect of reperfusion with calcium after a calcium-free period (shaded bar), in the anoxic rat heart (a: in the presence of glucose; b: in the absence of glucose), on the release of CK. a) Reperfusion with calcium resulted in an immediate and massive release of CK. b) Massive release of CK did not occur on reperfusion with calcium, but only on reoxygenation of the heart. In the presence of KCN oxygen-induced CK release was completely inhibited. c) Effect of reoxygenation after anoxic, glucose-free perfusion on the release of CK from isolated rat heart: Reoxygenation resulted in a relatively slight release of CK. Values are given as mean  $\pm$  S.E.M. (n = 8). Reproduced with permission from ref. 24.

system is responsible for the sudden cell damage. CK release on reoxygenation after a merely anoxic substrate-free perfusion is much less pronounced than with a preceding calcium-free period (Fig. 1c).

These data show that the calcium paradox fails to occur in the absence of energy, but occurs as soon as the electron transport system is reactivated (Fig. 1b). Electron transport is not a prerequisite for the calcium paradox; in the absence of electron transport the paradox occurs, provided that ATP is present (Fig. 1a).

# Intracellular pH during the Calcium Paradox

Hydrolysis of ATP and accumulation of calcium by mitochondria (supported by electron transport or ATP) with deposition of insoluble calcium phosphate, are accompanied by a release of protons (4,9). Hence, it has been suggested that acidification of the cytosol, with a consequent stimulation of cytosolic and lysosomal (phospho)-lipases and proteases, plays an important role in the development of the calcium paradox damage (11,24). <sup>31</sup>P NMR spectroscopy was used to investigate the time course of intracellular pH during the calcium paradox in isolated rabbit heart at 37°C (25). Intracellular pH was measured from the chemical shift of the intracellular inorganic phosphate ( $P_i$ ) peak.

Until the end of the calcium-free period (10 min), intracellular pH amounted to 7.1 in all hearts (n = 6). Figure 2 shows <sup>31</sup>P NMR spectra of one of the hearts, taken

5



Fig. 2. <sup>31</sup>P NMR spectra of a rabbit heart obtained from 128 scans and taken between 5 and 10 min of control perfusion (+Ca), and between 5 and 10 min of the subsequent calcium-free perfusion (-Ca). Resonances are assigned as follows: 1, methylene diphosphonate (external reference compound); 2, sugar phosphates; 3, extracellular P<sub>i</sub>; 4, intracellular P<sub>i</sub>; 5, creatine phosphate; 6, gamma phosphate of ATP; 7, alpha phosphate of ATP; 8, NAD(H); 9, beta phosphate of ATP.

during control perfusion (+Ca), and during the subsequent calcium-free perfusion (-Ca). From the position of the intracellular  $P_i$  peak, intracellular pH during control perfusion was calculated to be 7.1. In this experiment the intracellular  $P_i$  peak was absent in the spectrum that was taken during calcium-free perfusion, most likely as a result of incorporation of  $P_i$  into high-energy phosphates, since myocardial CP and ATP levels increase during calcium-free perfusion (2).

Reperfusion with the calcium-containing solution resulted in a rapid decline of intracellular CP and ATP levels. During the first 100 seconds of reperfusion, intracellular pH in the heart used for Fig. 3 varied from 7.1 to 6.9. In the subsequent spectra the intracellular P<sub>i</sub> peak, and also the CP and ATP peaks, were no longer perceptible. Intracellular pH values (mean  $\pm$  S.D.) of all six hearts amounted to 7.1  $\pm$ 0.1 (0 to 20 s); 7.1  $\pm$  0.1 (40 to 60 s); 7.0  $\pm$  0.1 (80 to 100 s).

These results demonstrate that there was no appreciable fall of intracellular pH during reperfusion with calcium-containing solution. It may by argued that pH data were obtained only during the first 100 s of reperfusion, i.e., the period that the intracellular  $P_i$  peak was perceptible. Figure 4, however, shows that the calcium paradox damage develops so rapidly that the release of enzymes, which is one of



Fig. 3. <sup>31</sup>P NMR spectra of a rabbit heart obtained from 6 scans and taken between 0 and 20 s (a); 20 to 40 s (b); 40 to 60 s (c); 60 to 80 s (d) and 80 to 100 s (e) of reperfusion with calcium containing solution. Resonances are assigned as follows: 1, intracellular  $P_i$ ; 2, creatine phosphate; 3, 4 and 5, the gamma, alpha and beta phosphate groups of ATP, respectively. Intracellular pH wasmeasured from the chemical shift of the intracellular  $P_i$  peak, and amounted to 7.1 (a); 7.2 (b); 7.0 (c); 6.9 (d) and 6.9 (e). The vertical line indicates the position of the  $P_i$  peak at pH 7.0. Reproduced with permission from ref. 25.

the characteristics of the calcium paradox, is maximal between 40 and 80 s of reperfusion. The rapid disappearence of  $P_i$  from the cytosol during the calcium paradox may be the result of leakage into the extracellular space and uptake by mitochondria together with calcium.

As mentioned before, both hydrolysis of ATP and accumulation of calcium by mitochondria are accompanied by a release of protons. A third process, however, that has to be taken into account, is the breakdown of CP, which is a proton-consuming reaction (9,11). These opposite effects, in combination with the buffering capacity of the cell, may explain why the calcium paradox is accompanied by little or no net change in intracellular pH.

# **Concluding Remarks**

At present we do not know which energy-dependent processes are responsible for the occurrence of the calcium paradox. The recent observation by Haworth et al. (12), however, that ATP depletion inhibits calcium influx in isolated myocardial cells, may explain why the calcium paradox does not occur in hearts subjected to anoxic substrate-free perfusion (see Fig. 1b). Reoxygenation and reactivation of the electron transport system could then promote calcium entry through the slow channels and in exchange for intracellular sodium, which increases during calcium-free perfusion as a result of a decrease in sodium/potassium-ATPase activity of the sar-



Fig. 4. CK release from isolated rabbit heart during the first 4 min of reperfusion with calciumcontaining solution, after 10 min of calcium-free perfusion (37°C). CK activity was analyzed as described before (25). Results are expressed as mean  $\pm$  S.D. (n = 5).

colemma (16). There is strong evidence that the slow channels (14) and the sodiumcalcium exchange mechanism (6,7) are modulated by ATP. This hypothesis does not explain, however, why the calcium paradox is much more pronounced than the damage that occurs upon reoxygenation after a merely anoxic substrate-free perfusion (ref. 13, cf. Fig. 1b and c).

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# **Clinical Implications**

# Chapter 20

# Human Purine Metabolism

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The purpose of purine metabolism in man is to maintain an optimal level of the nucleotides in the tissues. The nucleotides play an important role in nearly all biochemical processes, including energy metabolism, DNA and RNA structure, and regulation of many metabolic pathways through allosteric effects on enzymes, or through the adenylate energy charge. The nucleotides regulate their de novo and salvage synthesis and the interconversions between AMP and GMP. Uric acid is the waste, degradation endproduct of purine nucleotides in man. Uric acid is hardly soluble in physiological fluids. Therefore, hyperuricemia and hyperuricosuria are associated with precipitation of tophi in joints and other tissues and calculi in the urinary tract. Inborn errors in purine metabolism include the x-linked superactivity of 5-phosphoribosyl-1-pyrophosphate synthetase (gout and uric acid lithiasis), the complete deficiency of hypoxanthine-guanine phosphoribosyltransferase (Lesch-Nyhan syndrome) and the partial deficiency of this enzyme (gout and uric acid lithiasis), adenosine deaminase (combined immunodeficiency), purine nucleoside phosphorylase (T-cell immunodeficiency), xanthine oxidase (xanthinuria) and myoadenylate deaminase (muscle disease). A transport defect for urate in the renal tubules, manifested in hypouricemia, has also been reported.

#### Importance

Purine nucleotides play an important role in nearly all biochemical processes (14,24,31,49). Adenine nucleotides are active in the storage and transfer of metabolically available energy, ATP being the universal currency of energy in biological systems. Another important function of the nucleotides is the carrying of a wide variety of groups, and their transfer to appropriate acceptors (glycosyl group, alcohol phosphate, sulfate, hydrogen, hydride ion, electrons, and alkyl group). In addition, the purine nucleotides form structural units. They are the activated precursors of DNA and RNA, the bulk of cellular nucleotides being in the form of these polymers. Nucleotides also participate as structural units in low molecular weight compounds, such as histidine and certain vitamins (folic acid, thiamine and riboflavin). Adenine nucleotides are components of three major coenzymes: NAD<sup>+</sup>, FAD, and CoA, and of the important methyl donor, 5'-adenosylmethionine. Other nucleotides, such as cAMP and cGMP, have physiological functions as mediators of hormone action (regulatory signals). The nucleoside adenosine is considered to have an hormonal-like "retaliatory metabolite" role in matching energy demands to the synthesis of ATP in the heart muscle (32) and probably in some other tissues. Last but not least is the role of the nucleotides in the regulation of many catabolic pathways through allosteric effect on the enzymes, and their effect on many pathways of intermediary metabolism through the energy charge, ([ATP] + 0.5][ADP])/([ATP] + [ADP] + [AMP]), in the cells (2).

#### Metabolism

The pathways of purine nucleotide metabolism operate to maintain an optimal level of the various nucleotides in the tissues. The pathways of purine metabolism include (14,24,31,38,49,58,59): synthesis of nucleotides, either from small nonpurine molecules, by the energetically costly de novo pathway, or from preformed purines by the economical salvage pathways: interconversions between the nucleotides; and degradation of excess nucleotides. The degradation of nucleotides may furnish purine bases or nucleosides for salvage nucleotide synthesis in tissues in greater need for the nucleotides than the source tissue. Not all pathways operate in all tissues, and certainly not at the same intensity (see tissue characterization, below).

#### De Novo Biosynthesis of Purine Nucleotides

The purine ring is assembled from glycine (C-4, C-5, and N-7), from the amino nitrogen of aspartate (N-1), from the amide nitrogen of glutamine (N-3 and N-9), from activated derivatives of tetrahydrofolate (C-2 and C-8) and from  $CO_2$  (C-6) (Fig.1).

The purine ring structure is being assembled on a ribosyl moiety and when completed it is the nucleotide IMP. This is in contrast to the pyrimidine ring, which structure is completed before its attachment to the 5'-phosphoribosyl moiety to form OMP. The first-committed step in the de novo pathway is the formation of 5'-phosphoribosylamine from 5-phosphoribosyl-1-pyrophosphate (PRPP) and glutamine, catalyzed by glutamine-PRPP amidotransferase. The amide group of glutamine displaces the pyrophosphate group attached to the C-1 of PRPP. The other components of the purine skeleton are introduced in nine additional reactions, yielding finally IMP (Fig. 2; for full details, see refs. 14,19). IMP is the parent purine nucleotide molecule, but as a nucleotide it has no role in metabolism, except for being the precursor of AMP and GMP. AMP is synthesized from IMP in two steps. The first step is amination, accomplished through the attachment of aspartic acid to form the intermediate adenylosuccinate. This reaction is catalyzed by adenylosuccinate synthetase (EC 6.3.4.4). GTP is required in this reaction, providing a potential regulatory mechanism. Adenylosuccinate lyase (EC 4.3.2.2) cleaves fumaric acid from adenylosuccinate to form AMP. GMP is formed from IMP also in two steps. In the first step, IMP is oxidized to XMP, catalyzed by IMP dehydrogenase (EC 1.2.1.14) with NAD<sup>+</sup> as hydrogen acceptor. XMP is aminated by the amide group of glutamine to form GMP. ATP is consumed in this reaction, being cleaved to AMP + PPi. The conversion of GMP and AMP to their respective di- and triphosphates



Fig. 1. The sources of the carbon and nitrogen atoms of the purine ring.

occurs in two successive steps, catalyzed by respective kinases, requiring ATP. AMP, ADP and ATP are interconvertible. Adenylate kinase (EC 2.7.4.3) catalyzes the reaction ATP + AMP  $\implies$  2ADP. The deoxyribonucleotides are synthesized by direct reduction at the 2'-carbon in the ribose moiety of ADP and GDP, rather than by synthesis of the molecule using the 2'-deoxy analog of PRPP. This reduction is a complex reaction, catalyzed by ribonucleotide reductase and NADPH as a cofactor (ref. 26; Fig. 3).

#### Interconversions between AMP, IMP and GMP

AMP and GMP can be reconverted to IMP, in one step reactions, catalyzed by other enzymes than those involved in their synthesis from IMP. GMP is converted to IMP by GMP reductase (EC 1.6.6.8), whereas AMP is converted to IMP by adenylate deaminase (EC 3.5.4.6). The last reaction plays a major role in ATP degradation, as well as in the activity of the purine nucleotide cycle.

#### The Purine Nucleotide Cycle

This cycle (AMP - IMP - adenylosuccinate - AMP; ref. 29) operates probably in many tissues. However, the activity and role of this cycle were investigated mainly in skeletal muscle. The roles attributed to the activity of the cycle in the muscle include (29): conservation of nucleotides in the tissues (by prevention of the formation of diffusible nucleoside and base degradation products from AMP or IMP); conservation of the adenylate energy charge (by conversion of excess AMP to IMP); activation of ATP generation by glycolysis (by the formation of ammonia) and by the tricarboxylic acid cycle (by generation of intermediates for the cycle); and neutralization of acid production due to ATP degradation (by production of ammonia). Indeed, the activity of adenylate deaminase, the rate-limiting enzyme of this cycle, is the highest in the skeletal muscle from all tissues (29). The absence



Fig. 2. Pathways of purine metabolism in the human. Not all tissues contain all pathways (see text). Interrupted lines represent regulation by feedback inhibition.

of this cycle is well manifested in the condition of hereditary deficiency of muscle adenylate deaminase (refs. 10,50; see below). The activity and role of this cycle in the heart muscle is not yet fully clarified (52,62).

The rate of activity and possible roles of the guanine nucleotide cycle (GMP - IMP - XMP - GMP) have not been evaluated.

# Salvage Nucleotide Synthesis

The de novo synthesis of IMP consumes the equivalent of six high-energy phosphodiester bonds (by ATP hydrolysis). Additionally two high-energy bonds are consumed for IMP conversion to GMP, and one bond for the conversion of IMP to AMP. Thus, de novo synthesis of AMP or GMP is energetically very costly. Therefore, not surprisingly, once formed, the purine ring is recycled many times between the base (or nucleoside) and the nucleotide forms, before being further degraded to uric acid, the waste-endproduct to be excreted in the urine. The recycling of the purine bases or adenosine to the corresponding nucleotidic form is called "salvage nucleotide synthesis", since these pathways salvage the purines from the unreasonable (from the energy point of view) further degradation. Two general mechanisms perform salvage of preformed purine compounds: the phosphoribosylation of bases and the phosphorylation of nucleosides.

The phosphoribosyltransferases. Two enzymes in human tissue can phosphoribosylate purine bases. Hypoxanthine-guanine phosphoribosyltransferase (HGPRT; EC 2.4.2.8) phosphoribosylates hypoxanthine and guanine with PRPP to yield IMP and GMP, respectively. Another enzyme, adenine phosphorybosyltransferase (APRT; EC 2.4.2.7), phosphoribosylates similarly adenine to AMP. Physiologically, the salvage of hypoxanthine and guanine is much more important than that of adenine. This is indicated by the finding of PRPP accumulation in HGPRT-deficient tissues (36), but not in APRT deficiency (22,40,41,54). It is accepted that the consumption of PRPP by the salvage enzymes is greater than that by the de novo pathway, and accordingly that the proportion of nucleotides produced by salvage is greater than



Fig. 3. Enzyme system for the reduction of ribonucleoside diphosphates to deoxyribonucleoside diphosphates.

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that produced de novo. In contrast to the above, recent studies raised questions concerning the physiological efficiency of hypoxanthine salvage in various tissues, such as erythrocytes (5,34), liver (56), and skeletal (64) and heart muscle (62). The inefficient activity of HGPRT demonstrated in these tissues was attributed to low availability of substrate PRPP (34), to inhibition of HGPRT by the physiological content of cellular nucleotides (56), or to inefficient activity of adenylosuccinate synthetase (7). Evidently, the finding that a considerable quantity of hypoxanthine and guanine is degraded daily to xanthine and uric acid (see section on degradation, below), indicates a certain inefficiency of HGPRT. The physiological significance of adenine salvage is more seriously disputed, mainly in view of the questionable production of adenine in human tissues and the lack of accumulation of PRPP and absence of increase in purine production in APRT deficiency. However, the finding of 2,8-dihydroxyadenine urolithiasis in APRT deficiency (8,40) furnishes evidence that adenine is formed in the human, probably as a by-product of the synthesis of polyamines from 5'-adenosylmethionine (40), and that it is rapidly salvaged to AMP.

Adenosine kinase. The salvage of nucleoside occurs mainly for adenosine and 2'deoxyadenosine and is catalyzed by adenosine kinase (EC 2.7.1.20). Another enzyme, deoxycytidine kinase can phosphorylate 2'-deoxyadenosine and 2'-deoxyguanosine to the respective deoxynucleotides. The salvage of adenosine is probably the main salvage operating in erythrocytes (34), heart (62), and skeletal muscle (64). The main source of adenosine is probably the transmethylation reaction (from 5'adenosylmethionine). In some tissues, such as heart muscle, adenosine may be generated also from AMP, when this substrate accumulates during severe ATP degradation (62).

<u>Salvage cycles.</u> The formation of hypoxanthine, guanine and adenosine from nucleotides, and their recycling to nucleotides constitute cycles, which functions are not yet fully clarified. One possibility is that formation of hypoxanthine, guanine and adenosine in one tissue (e.g., liver and skeletal muscle) and the salvage of these purines to the respective nucleotide in another tissue (e.g., erythrocytes and brain), provide a system for transport of purines from "purine-rich" to "purine-poor" tissues (59). The adenosine futile cycle (AMP - adenosine - AMP could function in certain tissues, such as the heart during ischemia, as a mechanism for fast generation of the vasodilator adenosine. However, this role of the adenosine cycle has not been verified yet (62).

# Regulation of Purine Biosynthesis

Substrate 5-phosphoribosyl-1-pyrophosphate is the most important regulator of purine synthesis. PRPP synthetase (EC 2.7.6.1) catalyzes its formation from ribose-5-phosphate and ATP. Ribose-5-phosphate is generated by the pentose phosphate pathway, by either the oxidative or non-oxidative directions, as well as from ribose-1-phosphate, produced by degradation of the purine nucleosides inosine and guanosine. There is evidence that in many tissues, the availability of ribose-5-phosphate is saturating for PRPP synthesis under physiological Pi and nucleotide concentrations (42). However, this is not the case in skeletal and heart muscle. In these tissues, probably due to low activity of the oxidative pentose phosphate pathway, ribose-5-phosphate availability is limiting for PRPP synthesis and therefore also for purine synthesis (62,63).

In accord with the main purpose of the various pathways of purine nucleotide synthesis, which is the maintenance of optimal nucleotide concentrations, the nucleotide endproducts regulate purine metabolism at several levels. The rate of the de novo pathway is controlled by feedback inhibition at several sites. The firstcommitted step in the pathway, catalyzed by glutamine-PRPP amidotransferase, is sensitive to inhibition by various purine nucleotides, especially by AMP and GMP, in a synergistic (cooperative) manner. These inhibitors are competitive with substrate PRPP, which concentration is a major regulator of the amidotransferase activity (18). PRPP promotes conversion of the enzyme from the large inactive form (MW 270,000) to the small (MW 133,000) active form, whereas the purine nucleotides, alone or in combination, convert the small form of the enzyme to the large form. The activity of PRPP synthetase, which is therefore regulating the activity of the amidotransferase, is also subjected to feedback inhibition by the nucleotides, particulary AMP, ADP, GMP and GDP (16,51). At the physiological cellular content of nucleotides and Pi, the activity of PRPP synthetase is low, but there are several indications that the regulation of this enzyme, which is not committed for purine nucleotide synthesis than that of the first-committed amidotransferase. The most conclusive indication for this is the flamboyant purine synthesis in patients with the hereditary (x-linked) superactivity of PRPP synthetase, due to resistance to feedback inhibition by nucleotides (45,47,48,60), or other molecular alterations (ref. 4; see below).

The branching point in the synthesis of AMP and GMP from IMP is also regulated by the respective nucleotides. Adenylosuccinate synthetase is subjected to feedback inhibition by AMP, whereas IMP dehydrogenase is subjected to feedback inhibition by GMP. Moreover, GMP is a substrate for the synthesis of AMP (IMP conversion to adenylosuccinate), and ATP is a substrate for the synthesis of GMP (XMP conversion to GMP). This reciprocal substrate relation balances the synthesis of adenine and guanine ribonucleotides from IMP.

Another level at which the nucleotides control their biosynthesis is the product inhibition of the salvage pathways. IMP and GMP inhibit HGPRT, whereas AMP inhibits adenosine kinase.

The activity of ribonucleotide reductase, which is essential for the generation of substrates for DNA synthesis, is subjected to a complex regulation, aimed at providing an optimal balance between the deoxyribonucleotides for DNA synthesis (refs. 26,49; Fig. 3).

#### Degradation of Purine Nucleotides

ATP constitutes the largest nucleotide pool. When its degradation (energy consumption) exceeds the capacity of the cell to regenerate ATP from ADP, AMP is formed. The main pathway for AMP degradation is probably through deamination to IMP. This has been demonstrated in liver (55), in erythrocytes (5), and in skeletal (64) and heart muscle (62). IMP is the main nucleotide substrate for degradation to non-nucleotide purines. Surplus IMP, formed from AMP, or from excessive de novo production (see section on inborn errors, below) is degraded by 5'-nucleotidase (EC 3.1.3.5) to inosine. The possibility that non-specific nucleotidases, such as alkaline phosphatase, have a role in nucleotide dephosphorylation has been raised, but there are several indications that the activity of the specific nucleotidases is more important in this respect than the non-specific enzymes (3). GMP can be similary dephosphorylated to guanosine. The extent of this reaction is not known, but xanthine production in xanthinuria indicates that it may have quantitative significance (see xanthinuria, below). Inosine and guanosine are degraded by purine nucleoside phosphorylase (EC 2.4.2.1) to hypoxanthine and guanine, respectively. Guanine is deaminated by guanase (EC 3.5.4.3) to xanthine, and hypoxanthine and xanthine are degraded by xanthine oxidase (EC 1.3.2.3) to uric acid, which is the waste-endproduct of purine metabolism in man (Fig. 2). The experiments of nature in humans, the hereditary deficiency of purine nucleoside phosphorylase (11,25), of adenosine deaminase (12,25), and of xanthine oxidase (17), furnish evidence supporting the above pathway of degradation. In purine nucleoside phosphorylase deficiency, the main purine metabolites in urine are inosine, deoxyinosine and guanosine (25), indicating the flow of degradation in normal subjects through these

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metabolites. In xanthinuria, about 79% of total oxypurines excreted are in the form of xanthine (17), deriving from hypoxanthine by the residual xanthine oxidase activity, but mainly from guanine. This last finding indicates a significant flow of degradation from GMP through guanosine to guanine. That AMP degradation does not proceed to a significant extent through adenosine to inosine is indicated by the absence of decrease in urinary uric acid in hereditary adenosine deaminase deficiency (25). This finding also indicates that adenosine, even if formed from 5'adenosylmethionine (transmethylation reactions), does not contribute to uric acid formation, either due to insignificant formation, or more probably, due to efficient salvage.

Uric acid, the waste-endproduct of purine metabolism in the human, is not an ideal endproduct, since it is hardly soluble in body fluids. Excess uric acid in blood or urine tends to precipitate, as sodium hydrogen urate in joints or other tissues (tophi formation), or as uric acid, sodium urate or ammonium urate in calculi in the urinary tract (9,44). Interestingly, the renal handling of urate is also unlike that of classical endproducts. The renal handling of urate includes four components (43): a) urate is almost totally filtrable; b) the filtrable urate is almost totally reabsorbed in the proximal tubule; c) there is urate secretion; and d) there is subsequently postsecretory urate reabsorption. The urinary uric acid derives from the urate secretion which escapes reabsorption. In recent years, an increasing number of subjects with hereditary renal hypouricemia, due to defects in renal uric acid transport, have been reported (43). Most of these defects appear to be in the presecretory reabsorption site.

#### Tissue Characterization

The various tissues differ in the presence and intensity of operation of the above pathways of purine metabolism (59). The glutamine-PRPP amidotransferase, the firstcommitted step in the de novo pathway, is not active in erythrocytes and polymorphonuclear leukocytes, and its activity in the brain is low (1,19). In contrast, the amidotransferase is very active in liver and in skeletal muscle (6,53,63). These two tissues are probably the main source of preformed purines for salvage synthesis in other tissues. The circulating erythrocyte is devoid of many parts of purine metabolism (30). In addition to absence of almost all of the de novo pathway, it lacks adenylosuccinate synthetase. Thus, adenine nucleotides can be synthetized in this tissue only by salvage from adenosine (and adenine?). The erythrocyte lacks also xanthine oxidase. This enzyme which catalyzes the last two steps in purine degradation, is found only in liver, jejunal and rectal mucosa, and colostrum, and is absent from most other tissues (17,33). Some pathways may be markedly more active in specific tissues, due to a specific role, e.g., the purine nucleotide cycle in skeletal muscle. This cycle has an important role in energy metabolism in the skeletal muscle tissue (see below). HGPRT, the important salvage enzyme, is active in human brain markedly more (10 fold) than in other tissues (37). Indeed, the brain is the main tissue affected in HGPRT deficiency. The physiological activity of enzymes in a tissue may be deduced from the maximal activity of the enzyme in the tissue extract  $(V_{max})$ , the affinity of the enzyme to its substrates  $(K_m)$ , the availability of substrates, and presence of inhibitors and modifiers. The physiological activity of pathways in the intact tissue may be gauged by fluxes of labelled precursors. Although much of the above information is known for purine metabolism in many tissues, an accurate knowledge of the actual (physiological) fluxes of the pathways of purine metabolism, especially that of de novo versus salvage synthesis in the various human tissues, is not yet available.

#### Inborn Errors

#### **PRPP** Synthetase Superactivity

Several families have been reported with excessive purine production due to hereditary superactivity of PRPP synthetase. In the first case reported in 1972 (45,47,48,60), the enzyme superactivity was characterized to reflect resistance to feedback inhibition by nucleotides. In the other cases, the enzyme alterations were found to be increased specific activity (increased  $V_{max}$ ), increased affinity (low  $K_m$ ) for substrate ribose-5'-phosphate, or combination of such abnormalities (4). The pattern of inheritance is x-linked recessive (61). The affected males exhibit uric acid lithiasis and juvenile gout. Treatment by allopurinol is very effective. The excessive purine production characteristic to all mutants of superactive PRPP synthetase, emphasizes the role of both the enzyme PRPP synthetase and product PRPP in the regulation of purine synthesis.

#### HGPRT Deficiency

Complete hypoxanthine-guanine phosphoribosyltransferase deficiency is causing the x-linked (recessive) Lesch-Nyhan syndrome. The clinical syndrome was first reported in 1964 (27). It is characterized by cerebral palsy with choreoathetosis and spasticity, self mutilation, mental and physical retardation, and severe purine overproduction. Purine overproduction is attributed to accumulation of PRPP, due to the absence of salvage nucleotide synthesis from hypoxanthine and guanine, and to reduced feedback inhibition, due to lower cellular content of IMP and GMP. The last mechanism has not been conclusively assessed. HGPRT deficiency, underlying the Lesch-Nyhan syndrome, was revealed in 1967 (39). Allopurinol effectively reduces the uric acid level in blood and urine, and by this prolongs the life of the patients, who otherwise are prone to develop severe renal insufficiency (gouty nephropathy). But allopurinol treatment in HGPRT deficiency is associated with increase in blood and urine oxypurines, associated in some of the patients in xanthine urolithiasis (46). No specific treatment is available for the neurological manifestations, the etiology of which is not yet clarified.

Partial HGPRT deficiency, first reported in 1967 (23), causes only excessive purine production, resulting in uric acid lithiasis and juvenile gout, but apparently does not cause significant neurological abnormalities (21).

#### APRT Deficiency

Complete deficiency of APRT is associated with 2,8-dihydroxyadenine urolithiasis (8,22,40,54). Adenine, which is probably produced as a by-product of the synthesis of polyamines from 5'-adenosylmethionine, is normally salvaged by APRT to AMP. In the absence of the enzyme, adenine is oxidized by xanthine oxidase to the insoluble 2,8-dihydroxyadenine. The pattern of inheritance is autosomal recessive. There is no accumulation of PRPP in tissues and purine production is normal. These findings suggest that adenine production and salvage are probably of low quantitative significance.

#### Muscle Myoadenylate Deaminase Deficiency

Myoadenylate deaminase deficiency is a relatively benign disorder, first described in 1978 (10), characterized clinically by muscle fatigue, following excercise. Purine production is normal. The pattern of inheritance is autosomal recessive. This disorder furnishes evidence for the importance of the nucleotide cycle in skeletal muscle. It was suggested (50) that during intensive muscle work, the AMP formed

from ATP degradation, is converted to IMP, which accumulates stoichiometrically. Consequently, at rest, when ATP regeneration exceeds ATP degradation, the IMP is reconverted to AMP and to ATP. As discussed above, the activity of this cycle is of several advantages to the muscle tissue. The absence of this cycle is the cause for the manifested abnormalities.

# Xanthine Oxidase Deficiency (Xanthinuria)

The enzyme deficiency is either complete or partial (17). In normal subjects the enzyme is found in liver, small intestinal and rectal mucosa, and colostrum or breast milk (17,33). Xanthine and hypoxanthine replace uric acid as the endproduct of purine metabolism. Accordingly, there is hypouricemia and hypouricosuria. Total purine production is decreased due to enhanced salvage of accumulating hypoxanthine. Xanthine urolithiasis is common. Formation of xanthine from guanine indicates the presence of GMP dephosphorylation to guanosine and subsequently to guanine (see above). Xanthine myopathy has also been reported.

# Adenosine Deaminase Deficiency

The deficiency of adenosine deaminase (12,25) causes abnormalities in nucleoside metabolism which are selectively toxic to lymphocytes, resulting in combined immunodeficiency (cell-mediated T-cell- and humoral B-cell-immunodeficiency). Adenosine and deoxyadenosine levels are elevated in plasma and urine. dATP accumulates in peripheral mononuclear cells, especially in T-lymphocytes, and inhibits ribonucleotide reductase (DNA synthesis). The accumulation of adenosine results in accumulation of S-adenosylhomocysteine, inhibiting transmethylation (15,25), which is essential to lymphocyte multiplication and function. Purine production is normal, indicating absence of AMP degradation through adenosine in normal subjects. Inheritance is by an autosomal recessive pattern.

# Purine Nucleoside Phosphorylase Deficiency

Children, affected by this disease, have severe cell-mediated immunodeficiency (11,25). dGTP accumulates in T-lymphocytes and inhibits ribonucleotide reductase. The disorder is inherited by an autosomal recessive trait. The degradation of purines is blocked at the nucleoside level, the main degradation products in urine being inosine, guanosine and deoxyinosine. Accordingly, there is hypouricemia and hypouricosuria. Total purine production is increased due to the absence of salvage of hypoxanthine and guanine (which formation is blocked). This disorder furnishes evidence concerning the quantitative significance of HGPRT activity and concerning the main pathway of nucleotide degradation.

# Inborn Errors in other Pathways of Metabolism affecting Purine Metabolism

<u>Glucose-6-phosphatase deficiency</u>. Purine overproduction in Von Gierke's disease was attributed first to increased production of PRPP. The accumulation of glucose-6-phosphate results in acceleration of the oxidative pentose phosphate shunt, increasing ribose-5-phosphate generation, leading to increased production of PRPP (20). Increased nucleotide degradation, brought about by activation of adenylate deaminase, was suggested as the underlying mechanism (13,35). The AMP deaminase is activated due to the decrease in Pi (consumed by the enhanced glycogenolysis). The clarification of the mechanism(s) underlying purine overproduction in glucose-6-phosphatase deficiency is important for the understanding of the regulation of purine synthesis.

<u>Glutathione reductase superactivity.</u> This enzyme alteration was proposed to cause purine overproduction in some gouty black Americans by activation of the pentose shunt (28). However, the same condition was found in glucose-6-phosphate dehydrogenase deficiency without purine overproduction (57).

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# Chapter 21

# Diagnosis of Ischemic Heart Disease with AMP-Catabolites

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Lack of oxygen induces rapid breakdown of myocardial energy-rich phosphates, such as ATP. The purines adenosine, inosine, (hypo)xanthine and urate appear in the myocardial effluent during ischemia or reperfusion. In this paper I summarize experiments on the isolated rat heart, made temporarily ischemic; on the anesthetized pig with a stenosis of a coronary artery; in patients undergoing an atrial pacing stress test or coronary angioplasty. In the animal experiments, ischemic myocardium releases mainly inosine; the human heart, on the other hand, produces predominantly hypoxanthine during periods of ischemia. When an ischemic event follows relatively quickly an earlier one, purine production during the second period is substantially lower. Experimental animals or patients thus cannot serve as their own controls in studies on drug efficacy under these conditions. In certain clinical settings, the assay of hypoxanthine seems superior to that of lactate to demonstrate myocardial lack of oxygen. I conclude that myocardial purine production is a promising biochemical indicator for ischemic heart disease.

# Purines as an Indicator of Ischemia?

Among the coenzymes, nucleotides play an important role in, e.g., bioenergetic processes (17). The nucleotide ATP is essential in heart muscle, because it is the fuel for the pump. The product, ADP, will be rephosphorylated. Heart muscle needs a continuous supply of large amounts of oxygen for this process - oxidative phosphorylation. ATP turnover is very high; there is a delicate equilibrium between ATP supply and demand (19). Figure 1 shows how energy-rich phosphates form purines. These can leave the heart. Adenosine is a physiological vasodilator (ref. 3; see Chapter 8).

The ischemic heart produces adenosine, inosine and hypoxanthine, a fact known for decades. We showed that the ischemic pig heart releases inosine and hypoxanthine (8). Relatively late the quantitative importance of xanthine and urate production became clear (33,34). The enzyme xanthine oxidoreductase, of which the oxidase form could play a role in atherogenesis (see ref. 34) and reperfusion damage (25), is responsible for this conversion. In this chapter I will demonstrate - using examples from animal and clinical studies - that AMP catabolites are promising markers of myocardial ischemia.

# Purine Production by Isolated, Ischemic Rat Heart

Figure 2 shows the time-course of coronary flow and purine release in the isolated, rat heart (Langendorff preparation), with flow reduced temporarily by 75%. After 10 min of ischemia, the heart releases more than six times as much purine as during control perfusion. This increases to 16 times early during reperfusion. Reactive hyperemia occurs, possibly caused by adenosine release.

We concluded from balance studies that purine production is a good measure for ATP breakdown in ischemic heart (1,37). A caveat! During drastic reduction in coronary flow - complete cessation of flow is the extreme example - catabolite release during ischemia goes down (Fig. 3). If we reduce the cardiac output of the isolated, working rat heart by 94%, purine release does not increase, but decreases. Only during reperfusion release of adenosine plus catabolites increases in that case.

#### **Inosine Production during Repeated Coronary Occlusion in Pig Heart**

Figure 4 shows an experiment in which we reduced the left anterior descending coronary artery flow of the pig repeatedly by 60%. Arterial inosine did not show significant changes. However, during the first stenosis the heart released substantial amounts of inosine (39). During the second flow reduction, inosine release was much



Fig. 1. Myocardial ATP breakdown. Because of their polarity, the adenine nucleotides ATP, ADP and AMP, do not pass the cell membrane. The heart releases the purine nucleosides adenosine and inosine, and the oxypurines (hypo)xanthine and urate into the blood stream. P = phosphate.



Fig. 2. Purine release from isolated non-working rat heart, paced at 300 beats/min. A roller pump restricted flow. We measured AMP-catabolites in the coronary effluent by high-pressure liquid chromatography (7,16). The release of purines (adenosine + inosine + hypoxanthine + xanthine + urate) increased during ischemia and reperfusion. Means  $\pm$  SEM, n = 10 (control), n = 6 (ischemia), n = 3 (reperfusion).

less, suggesting a depleted myocardial adenine nucleotide pool. (We did a similar observation on myocardial lactate release.) We concluded that this model is not useful for interventions in which the animal serves as its own control.

#### Hypoxanthine Production by Human Heart during Atrial Pacing

The assay of purines in human blood is difficult, because the concentrations are so low. From Fig. 5 it is clear that the inosine concentration in human blood is lower



Fig. 3. Purine release from isolated working rat heart, paced at 300 beats/min. Perfusion took place with a preload of 10 cmH<sub>2</sub>O and an afterload of 75 cmH<sub>2</sub>O ( $37^{\circ}$ C). Ischemia was induced with a ball valve in the aortic outflow tract (29). - - Coronary flow. We measured AMP-catabolites in the coronary effluent by high-pressure liquid chromatography (7). Means ± SEM (n = 7). Note that reduction in flow in this preparation leads to increased purine efflux during reperfusion.



Fig. 4. Arterial concentration and myocardial balance of inosine in the anesthetized open-chested pig (39). We induced myocardial ischemia repeatedly by reduction of the flow in the left anterior descending coronary artery to 40% of control. Arterial inosine (measured enzymatically in deproteinized blood) did not change. Myocardial ATP breakdown, reflected by inosine release, during the second stenosis was much less than during the first one. Therefore, this model cannot be used for the study of interventions during myocardial ischemia in which the animal serves as its own control. Means  $\pm$  SEM (n = 12). \* P<0.05 vs. control. °P<0.05 vs. first flow reduction.

than 1 uM. It would be nice to prepare plasma or serum before deproteinization, because it circumvents the disturbance during chromatography by the relative high amounts of erythrocyte nucleotides (ATP etc.). However, Fig. 5 shows that during the preparation of plasma or serum the purine concentration measured increases. Therefore we deproteinize the blood immediately. We remove the disturbing nucleotides with an Al<sub>2</sub>O<sub>3</sub>-column, before performing high-pressure liquid chromatography. (Recently, we mixed some inhibitors of adenosine metabolism [dipyridamole and erythro-9(2-hydroxy-3-nonyl)adenine] immediately with the blood samples. This prevents the formation of AMP-catabolites during plasma preparation, and makes the tedious Al<sub>2</sub>O<sub>3</sub>-step unnecessary.)

During an atrial pacing stress test of patients undergoing a heart catheterization for diagnostic purposes, the arterial hypoxanthine concentration remains constant. In patients with significant narrowing of one or more coronary arteries, the coronary sinus hypoxanthine blood concentration exceeds the arterial one by a factor of two during pacing (Fig. 6). Patients with little or no angioscopically demonstrable atherosclerosis do not show such purine production. In earlier work (31) we described that patients developing angina pectoris during such a test, also show hypoxanthine production. Table 1 summarizes the effect of a repeated atrial pacing stress-test on hemodynamic and biochemical variables. Twenty-five minutes after the first test we repeated the pacing until heart frequency was maximal. The release of lactate and hypoxanthine was lower during the second pacing procedure. We concluded from that experiment, like we did from the pig experiment described in Fig. 4, that patients are not useful as their own control in this type of set-up. Kugler studied both the effect of pindolol (22) and nitroglycerine (21) on lactate and purine metabolism during two pacing stress tests. Because he failed to submit a control group (without drugs) to the same procedure, his conclusions from the lower production of lactate, inosine and hypoxanthine are dubious. It is noteworthy that Ihlen et al. (18) found reproducible lactate production, if they repeated the pacing procedure after a recovery period of 20 to 40 min. Whether a different patient population can explain the discrepancy, remains to be answered.



Fig. 5. Concentrations of AMP-catabolites in human blood, plasma and serum. We measured the purines by high-pressure liquid chromatography (16). In plasma and serum the total amount of these compounds was considerably higher than in blood, presumably due to nucleotide breakdown during blood processing. We omitted the urate concentration, because in human blood, plasma or serum it is much higher (200-300 uM) than that of the other purines. Data represent values of 5-6 non-resting volunteers (means  $\pm$  SEM). \*P<0.01 vs. blood; °P<0.05 vs. plasma.

# Hypoxanthine Production during Coronary Angioplasty

Also during percutaneous transluminal coronary angioplasty marked hypoxanthine (and lactate) production is demonstrable. Immediately after deflation of the occluding balloon, the levels of these compounds increase in the coronary sinus or the great cardiac vein (Fig. 7). The amount released after subsequent balloon deflations is still large. The metabolic derangements, however, normalize rather quickly (36).

#### Purine Production, a Biochemical Marker for Ischemia

Table 2 summarizes studies in with AMP-catabolites used to diagnose ischemic heart disease. Various investigators did their determinations in plasma or serum, without adequate precautions against catabolism (cf. Fig. 5). Buhl et al. (6) demonstrated increased excretion of hypoxanthine and xanthine in urine of cardiac patients. However, they were unable to detect raised oxypurine levels in peripheral venous blood, as described in Sovjet studies (12,28).



Fig. 6. Hypoxanthine production by the human heart during ischemia. An arterial (art) and coronary sinus (cs) blood sample was taken from patients undergoing heart catheterization, both at rest and during a rapid atrial pacing stress test. Pacing was stopped when the patient developed progressive chest pain or an A-V block. A stenosis of at least one coronary artery, exceeding 50%, was considered a significant obstruction. This study shows that pacing-induced myocardial hypoxanthine release is a marker for ischemic heart disease (ref. 16, see also ref. 31). Mean values  $\pm$  SEM and statistically significant differences (P<0.05) are presented.

Table 1.	Effect of a	repeated a	atrial pacin	g stress	test on hemod	vnamic and	hiochemical	variables
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Variable	P <sub>1</sub>	P <sub>2</sub>	
Heart frequency (beats/min)	144 ± 5	144 ± 5	
Left ventricular pressure (mmHg)	$137 \pm 4$	$137 \pm 4$	
Peak + dp/dt (mmHg/s <sup>-1</sup> )	$2470 \pm 160$	$2440 \pm 140$	
Vmax (s <sup>-1</sup> )	70 ± 4	$70 \pm 4$	
Coronary blood flow (ml/min)	$189 \pm 15$	$186 \pm 17$	
Lactate efflux (umol/min)	$23 \pm 3$	$14 \pm 3^*$	
Hypoxanthine efflux (nmol/min)	74 ± 10	$26 \pm 5$	

 $P_1$  = first pacing test;  $P_2$  = second pacing test. Means ± SEM (n=10). \*P<0.05 vs.  $P_1$  (see ref. 35).

Hopefully it will be clear from this review that an increasing number of cardiological clinics show interest for these biochemical indicators. The same holds for disciplines such as obstetrics (2,4,14,15,24,27,30,38) and surgery (refs. 9,11,23,28; Chapter 22). High-pressure liquid chromatography is probably the method of choice to measure purine concentrations. Sample preparation needs improvement to make the analysis routinely applicable.



Fig. 7. Hypoxanthine and lactate release from human heart after percutaneous transluminal coronary angioplasty (36). Means  $\pm$  SEM (n = 15). GCV = great cardiac vein.

Purine	Body fluid	Clinical situation	Ref.
Ado	A-CS plasma	APST	10
UA	Serum	AMI, IHD	13,26
Hx	A-CS blood	APST	16,31,32
Hx	Peripheral venous plasma	AMI, ischemia, AP	12,28
Ado, Ino, Hx	A-CS plasma	ACBG	11
Ino, Hx	A-CS plasma	APST	20-22
Hx, Xan (UA)	Urine	Cardiac arrest, AP, AMI	6
Ado, Ino, Hx, Xan	A blood	ACBG	5
Hx	A-CS blood	2 x APST	35

Table 2. AMP-catabolites in blood and urine as marker for myocardial ischemia in patients

A = arterial; ACBG = aorta-coronary bypass grafting; Ado = adenosine; AMI = acute myocardial infarction; AP = angina pectoris; APST = atrial pacing stress test; CS = coronary sinus; Hx = hypoxanthine; IHD = ischemic heart disease; Ino = inosine; UA = urate; Xan = xanthine.

In summary, I conclude that myocardial purine production is a promising biochemical indicator for ischemic conditions, encountered frequently in patients with heart disease.

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# Chapter 22

# Myocardial High-Energy Phosphates during Open-Heart Surgery

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To obtain a quiet and bloodless operation field during open-heart surgery, periods of myocardial ischemia are deliberately introduced by aortic cross-clamping. As a consequence, an unbalance between consumption and production of cellular high-energy phosphates is created. This results in a rapid fall of creatine phosphate levels, followed by a decrease in the tissue content of ATP. Low levels of ATP are related to the loss of cellular function and the onset of cell death. The precise mechanism of action is, however, still unclear. Although tissue ATP content as a prognostic index of ischemia-induced damage recently has been challenged, most protective measures taken during open-heart surgery aim at the conservation of cardiac high-energy phosphate pools in the ischemic tissue. To this end, commonly electro-mechanical activity is abolished rapidly by intracoronary infusion of an ice-cold crystalloid or sanguineous cardioplegic solution immediately after aortic cross-clamping. All kinds of variations with respect to the application of specific pharmaca and mode of administration, have been reported. Assessment of the tissue content of ATP and creatine phosphate in left ventricular biopsies, obtained during cardiac operations, provides a useful parameter to estimate the adequacy of the protective measures applied.

# Introduction

This chapter will review selected aspects of high-energy phosphate homeostasis in the human heart with particular emphasis on the situation during open-heart surgery. To date many cardiac defects are repaired operatively. During most of the surgical procedures global cardiac ischemia is deliberately imposed to provide the surgeon a quiet and bloodless operation field. The consequent lack of oxygen will impair adequate replenishment of cardiac high-energy phosphate stores. It has been generally accepted that failure to keep intracellular ATP within safe limits will result in loss of cellular function and ultimately in cell death. Therefore, measures have to be taken to prevent cardiac high-energy phosphate depletion during the period of global ischemia during open-heart surgery. In the present overview attention will be paid to:

- production and consumption of high-energy phosphates in the normoxic heart;
- disturbed high-energy phosphate homeostasis during ischemia;
- high-energy phosphates as index for myocardial damage during open-heart surgery;
- measures to maintain cardiac high-energy phosphates at adequate intracellular levels.

#### Cardiac High-Energy Phosphate Homeostasis during Normoxia and Ischemia

Energy production and energy consumption are tightly coupled in the working myocardial cell. The need for ATP dictates the rate of conversion of ADP and inorganic phosphate into ATP, while substrates and molecular oxygen are utilized in amounts exactly matching that required to produce ATP. At normal workload the heart consumes approximately 5 umol  $O_2$ /min per g wet wt. If 6 mol ATP are produced per mol  $O_2$ , than about 30 umol ATP are aerobically produced per min per gram. The intracellular content of ATP is in the order of 5 umol/g wet wt., implicating that each 10 s the equivalent of the total intracellular ATP pool will turn over. This figure stresses the significance of an unimpeded supply of molecular oxygen to the myocardial cells.

As indicated in Fig. 1, exogenous carbohydrates, such as glucose and lactate, and lipids like fatty acids, serve the myocardium as readily oxidizable substrates. Under normal physiological circumstances fatty acid oxidation provides more than 50% of the energy required for cardiac function (43). Lactate appears to be the preferential substrate when the extracellular concentration of this substrate exceeds 5 mM, a situation easily encountered during heavy physical exercise. The contribution of glucose to cardiac energy production depends on both the circulating level of insulin and the workload of the heart. The supply of exogenous substrates to the heart is not rate limiting under normal circumstances. Endogenous substrates are present in the form of glycogen, highly-branched glucose polymers, and triacylglycerols, molecules in which fatty acids are complexed to glycerol. In the absence of extracellular substrates, the amount of glycogen and triacylglycerol will be sufficiently high to meet the myocardial energy demands for about 60 and 40 min, respectively (42).



Fig. 1. Schematic representation of myocardial energy homeostasis. KC refers to Krebs-cycle activity, CP to creatine phosphate, and C to creatine.

Energy-consuming reactions take place in localized areas of the myocyte. Over 60% of the amount of ATP produced is utilized for myofibrillar contraction and relaxation; the remainder of ATP is consumed by ion pumps at the sarcolemma and sarcoplasmic reticulum and a variety of anabolic and repair processes. Under aerobic conditions approximately 98% of the cellular production of ATP takes place in the mitochondrial matrix. Quantitatively, glycolytic ATP production in the cytoplasmic compartment is of minor importance if oxygen supply to the heart is sufficient.

Traditionally the role of creatine phosphate, the intracellular concentration of which is 2-3 times that of ATP, has been considered to serve as buffer for cellular ATP. More recent findings suggest that creatine phosphate is involved in transporting energy from the site of production to the myofibrillar utilization reaction (Fig. 1; refs. 3,26).

A completely different picture emerges when supply of molecular oxygen is insufficient to maintain the production of ATP. Mechanical activity ceases rapidly. Since repetitive contraction and relaxation of the myofibrils are quantitatively the most important energy-consuming processes in the myocyte, cessation of mechanical activity can be considered as an appreciable intrinsic energy-sparing measure. The tissue content of creatine phosphate falls within one minute to very low levels. The decrease in the cardiac content of ATP is relatively slow; e.g., the overall content is halved after 20 to 30 min following the onset of ischemia (14). Mitochondrial ATP production is very low or negligible, depending on the residual supply, if any, of oxygen to the cell. Initially, the degradation of glycogen is accelerated and glycolytic activity is increased, while the majority of ATP is produced by converting glucose into lactate. Due to subsequent depletion of the endogenous glycogen pool and inhibition of the glycolytic flux by acidification of the intracellular compartment, the significance of the glycolytic production of ATP will become less during extended ischemia. In addition to reduced production of ATP with become ress oxygen-deprived cells, the depletion of the intracellular content of ATP is most likely further accelerated by induction of futile energy-consuming processes; e.g., accelerated turnover of the triacylglycerol pool (37) and stimulation of  $Ca^{2+}$ dependent ATP-ases due to intracellular Ca2+-overload (27). Reduction of the intracellular ATP pool in ischemic tissue is accompanied by increased levels of ADP and AMP. Concomitantly, the content of purine bases and oxypurines such as adenosine, inosine, hypoxanthine and xanthine increases as a consequence of further degradation of AMP (Fig. 1). Finally, ATP levels in the oxygen-deprived myocyte will fall beyond a critical limit. With ATP levels below this point, functional recovery of the heart is not possible despite readmission of oxygen by reperfusion. It has been suggested that the critical limit of ATP is in the order of 10 umol/g dry wt. or about 1-2 umol/g wet wt. of tissue (5,20).

It is noteworthy that the degradation products of the adenine nucleotides are released from the heart upon reperfusion. This loss hampers rapid resynthesis of the (partially) depleted adenine nucleotide pool. Commonly, creatine phosphate levels return to or even exceed pre-ischemic values upon readmission of flow. It should be kept in mind that low post-ischemic ATP levels do not necessarily imply that the turnover of ATP is impaired. Stable post-ischemic ATP values most likely reflect adequate mitochondrial activity to keep pace with the rate of ATP consumption.

Although considerable evidence has been provided that ATP is a major factor in, and possibly a determinant of, tissue injury, the precise relationship between depletion of ATP and loss of cellular function has not been elucidated. Theoretically, a variety of causal relationships can be proposed. First, ATP is required for contraction and relaxation of myofilaments. Therefore, low intracellular levels of ATP will in a direct manner hamper adequate mechanical activity. Findings of Ellis et al. (12), who observed a close relation between post-ischemic recovery of

contraction and repletion of tissue ATP stores in the dog heart are, in favor of this notion. The time required for complete recovery of both indices was found to be at least 2 weeks after the start of reperfusion following 2 h of ischemia. Second, Jennings et al. (24) have emphasized the important role of ATP in maintaining fluid and ion homeostasis in the heart. Their findings suggest that the cell is incapable to control properly the cellular water and ion balance when ATP levels dropped below a certain critical level. Swelling of cells will occur, especially after reinstallation of blood flow to the previously ischemic tissue, resulting in irreversible damage of cellular structures and impediment of post-ischemic blood flow. In particular, calcium ions are thought to accumulate in ATP-depleted cells most likely resulting in hypercontraction of the myofilaments, calcium loading of mitochondria and activation of Ca2+-dependent proteases and phospholipases (27). Third, ATP is required for the turnover and/or resynthesis of phospholipids, being important constituents of myocardial cellular membranes. Impaired phospholipid homeostasis in the cardiac cells might be closely related to the occurrence of irreversible damage (41). Further weight is given to the role of cellular ATP in preserving cardiac phospholipid homeostasis by the finding of Chien et al. (7) that depletion of ATP in isolated neonatal cardiac cells by metabolic blockers accelerates degradation of phospholipids.

Although the notion that the tissue content of ATP can be used as a prognostic index to estimate the extent of ischemia-induced damage and the adequacy of myocardial protection has been generally accepted, this concept was recently challenged by Buckberg and associates (6,32). They reported that the ability of cardiac tissue subjected to extended ischemia to recover contractile function, was not dependent on tissue steady-state ATP levels. In addition, they claimed that the extent of functional recovery was mainly determined by the conditions of reperfusion and the composition of the reperfusate. The duration of the preceding ischemic episode (32) and measures taken during ischemia (45) seem to be less decisive. Their experimental results, which suggest that the post-ischemic capacity to regenerate cellular ATP pools rather than the absolute level of ATP is the main determinant of cell viability, underline the complexity of high-energy phosphate homeostasis under pathophysiological conditions and the complex relationship between ATP and loss of cellular function.

# The State of the Heart during Open-Heart Surgery

Routinely, the heart of the patient subdued to open-heart surgery is in an empty-beating state during the initial period of cardiopulmonary bypass. The energy demand is low. To obtain a quiet and bloodless operation field the aorta is cross-clamped just distal to the coronary ostia resulting in global cardiac ischemia. Related to the type of operation, various surgical procedures can be followed. In case of correction of congenital anomalies and valve operations, the heart is subjected to one or more prolonged periods of global ischemia. Basically two different techniques are at hand for aorto-coronary bypass surgery, i.e., intermittent aortic cross-clamping or continuous aortic cross-clamping. The former technique is characterized by repetitive short periods of ischemia with intervals for reperfusion. When the latter technique is applied, all distal anastomoses are performed while the heart is in an extended global ischemic state.

Removal of the aortic clamp allows blood, molecular oxygen and substrates to return to the previously ischemic heart. The heart fibrillates spontaneously or is empty-beating. When body temperature has returned to the physiological range and the heart has resumed normal electro-mechanical activity, the patient is weaned from cardiopulmonary bypass. To prevent loss of cardiac function and irreversible cell damage due to the preceding ischemic period a variety of protective measures are commonly taken. In general, these measures are aiming at the conservation of cardiac high-energy phosphate pools in the ischemic tissue.

#### Assessment of High-Energy Phosphates in Human Heart

To assess the content of high-energy phosphates in cardiac tissue of patients subjected to open-heart surgery, full thickness biopsies of the free wall of the left ventricle can routinely be obtained with the use of an air-driven drill (20) or the Tru-Cut<sup>R</sup> needle (Travenol Laboratoria; ref. 35). In biopsies, weighing in the range of 5-10 mg wet wt., sufficient tissue is present to assess the high-energy phosphate content. A variety of very sensitive chemical assays are available, e.g., luciferase technique, fluorometry and high-performance liquid chromatography. The advantage of the latter technique is that the whole spectrum of high-energy phosphates and related substances can be analyzed in one single run.

To avoid artificial changes in the tissue content of ATP and creatine phosphate the tissue specimen has to be immediately frozen in liquid nitrogen or comparable coolant, and stored at a temperature below -80°C until biochemical analysis.

From the two biopsies, one obtained prior the aortic cross-clamping and the second after restoration of blood flow, the effect of the intermediate period of ischemia and the efficacy of the protective measures applied can be estimated. It should be kept in mind that parts of the hearts under investigation are in a diseased state prior to the induction of global ischemia. Since the amount of tissue taken from the heart represents approximately 0.05% of total cardiac mass, there is a reasonable chance that the value measured in the biopsy is not representative for the whole left ventricle. The authors have explored this possible drawback of the biopsy technique in more detail (39,40). Figure 2 shows histograms of ATP and creatine phosphate values measured in pre-ischemic cardiac biopsies taken during cardiopulmonary bypass in 70 patients. The interindividual variation was found to be considerable, i.e., 35% for ATP and 37% for creatine phosphate. Figure 3, showing ATP and creatine phosphate levels in pairs of pre-ischemic biopsies taken from adjacent areas within 2 min prior to aortic cross-clamping in 14 patients, indicates that in some patients marked spatial differences were encountered. The intra-individual coefficient of variation for both ATP and creatine phosphate was found to



Fig. 2. Histograms representing the individual values of ATP and creatine phosphate in left ventricular biopsies obtained during the pre-aortic cross-clamp period in patients on cardiopulmonary bypass (n = 70). Tissue values are grouped in cohorts of 2 and 5 umol g dry wt. of tissue for ATP and creatine phosphate, respectively. n refers to the number of values belonging to one specific cohort (after ref. 40, with permission).
be 11%. Since the median values were not significantly different, ATP and creatine phosphate can be used as index to assess the effect of ischemia in groups of patients. Conclusions for individual casus are obviously marred by the appreciable intra-individual variation.

A disadvantage of the biopsy technique is the invasive character of this procedure. An alternative non-invasive technique to study high-energy phosphate homeostasis in the heart is the nuclear magnetic resonance (NMR) technique (23,46). Despite promising results obtained in animals (28), to our knowledge this technique has not been used to monitor cardiac high-energy phosphates during open-heart surgery.

#### Preservation of High-Energy Phosphates during Open-Heart Surgery

In general, reduction of the aortic cross-clamp time and, hence, duration of ischemia, and lowering the temperature of the heart, which will result in slowing the rate of energy-consuming metabolic processes, can be considered as an effective measure to preserve myocardial high-energy phosphate levels. Improved surgical skills have reduced considerably the time duration needed to perform the repair procedure. Application of cardioplegic solutions have shown to delay appreciably the intracellular depletion of high-energy phosphates (20), due to the rapid cessation of electro-mechanical activity and the low temperature of the heart.

As indicated earlier, aorto-coronary bypass surgery has been performed with the use of two different techniques, i.e., intermittent aortic cross-clamping or continuous aortic cross-clamping in combination with administration of cardioplegic solutions. An extensive study has been performed by the authors and associates (18,44) in patients operated upon with intermittent aortic cross-clamping either at 34 or 25°C body temperature, and with continuous aortic cross-clamping in combination with repetitive administration of ice-cold St. Thomas' Hospital cardioplegic solution. No significant differences could be observed regarding clinical outcome, post-operative



Myocardial High-Energy Phosphates during Open-Heart Surgery

hemodynamic recovery and release of cardiac enzymes between the various techniques employed. However, analysis of cardiac tissue specimen obtained during the operation revealed that the cardioplegia procedure prevented completely the reduction of ATP during the ischemic period. In contrast, the level of ATP decreased to 72 and 71% of control in the intermittent cross-clamping groups with a body temperature of 34 and 25°C during bypass, respectively. The content of creatine phosphate returned to the pre-ischemic level after a reperfusion period of about 20 min following the release of the last aortic cross-clamp. In an earlier study it was shown that the administration of the St. Thomas' Hospital cardioplegic solution during continuous aortic cross-clamping in patients who underwent isolated aortic valve replacement (15) prevented completely the reduction of creatine phosphate in the ischemic heart. In this study the second biopsy was taken within 3 min after release of the aortic clamp. A drop of 50% in cardiac creatine phosphate content occurred, when the heart was only topically cooled. The tissue level of ATP remained stable irrespective of the protective measure applied. These observations are in line with the findings of Balderman et al. (2). They observed a rapid restoration of creatine phosphate immediately upon release of the aortic clamp in previously global ischemic dog heart, when treated with a cold cardioplegic solution during the period of oxygen-deprivation. In contrast, post-ischemic creatine phosphate restoration was delayed when the temperature of the heart was not kept within safety limits during the preceding ischemic period.

Other studies have also indicated that measurements of cardiac contents of high-energy phosphates provided a sensitive metabolic indicator of ischemic damage of the human myocardium. Findings of Balderman et al. (1) have shown that ATP levels are excellently preserved in human hearts treated with a cold, crystalloid cardioplegic solution when the total ischemia did not exceed 90 min. Creatine phosphate values significantly fell by about 60%, but returned to normal levels 30 min after reperfusion. However, preservation of high-energy phosphates was incomplete when the aortic cross-clamp time exceeded 90 min. Cunningham et al. (8) explored the efficacy of a blood cardioplegic solution to preserve human cardiac ATP levels during aortic cross-clamping lasting up to 120 min. In operations without technical failures, the ATP content was found to increase by about 50% during the ischemic period and returned to pre-ischemic values after 30 min of reperfusion. Although no satisfactory explanation could be offered for the unexpected rise of ATP during the period of aortic cross-clamping, their results clearly indicate the usefulness of sanguineous cardioplegia. Kaijser et al. (25) have reported that the temperature of the heart should not decrease beyond  $15^{\circ}$ C when blood-cardioplegia is applied. Silverman et al. (36) have shown that the efficacy of crystalloid cardioplegic solutions in patients undergoing myocardial revascularization to maintain cardiac ATP at its pre-ischemic level was greatly enhanced when the cardioplegic solution was given through the completed distal left anterior descending anastomosis in addition to infusion via the aortic root.

Recently, explorations have been made whether addition of pharmaca to cardioplegic solutions stabilize cardiac adenine nucleotide homeostasis during the ischemic episode. Nifedipine used as additive to the St. Thomas' Hospital cardioplegic solution in patients undergoing valve replacements, was found to be effective to prevent accumulation of degradation products of adenine nucleotides in the oxygen-deprived heart (17). Earlier studies of the same research team (16) have shown that lidoflazine in combination with intermittent aortic cross-clamping in patients subjected to aorto-coronary bypass surgery was as effective as cardioplegia to preserve myocardial high-energy phosphate levels.

Despite the marked cardioprotective properties of most crystalloid and sanguineous cardioplegic solutions routinely used in the clinical setting, attempts have been made to interfere more directly with cardiac high-energy phosphate homeostasis. Basically, three different approaches to manipulate directly adenine nucleotide and creatine

phosphate levels can be discriminated. First, addition to the cardioplegic solution of substances potentially active to promote resynthesis of high-energy phosphates. Second, the use of inhibitors of pathways involved in degradation of adenine nucleotides and related substances. Third, addition of high-energy phosphates, such as ATP and creatine phosphate, themselves to cardioplegic solutions.

Resynthesis of ATP can occur via 4 distinct pathways, one being the salvage pathway requiring inosine and/or hypoxanthine and ribose for resynthesis of AMP via adenylosuccinate, the second is phosphorylation of adenosine, the third ribophosphorylation of adenine and the fourth pathway is de novo synthesis of purines which procedes very slowly and requires substantial amounts of energy. Although promising experimentally obtained results have been described with respect to stimulation of the salvage pathway (10,13,47), data on this subject in the clinical setting are lacking.

In considering the use of inhibitors of pathways of adenine nucleotide and purine degradation experimental studies have indicated that application of these compounds may be a useful measure to preserve cardiac high-energy phosphate levels (see, for instance, ref. 9,19,22,38). There is, however, a paucity of information obtained in clinical studies to advocate the use of blockers of adenine nucleotide degradation in the operation theatre.

The effect of addition of creatine phosphate and ATP to a crystalloid cardioplegic solution has been extensively studied by Robinson et al. (29,30) in an isolated rat heart model. High-energy phosphates added to St. Thomas' Hospital cardioplegic solution significantly improved post-ischemic myocardial performance and reduced the loss of enzymes from the intracellular compartment after both 40 min of ischemia at 37°C and 270 min of oxygen deprivation at 20°C. ATP showed a relatively narrow dose-response curve. Optimal effects were obtained at 0.1 mmol ATP per liter cardioplegic solution. Higher concentrations of ATP affected negatively post-ischemic cardiac performance. Creatine phosphate exerted its maximal protective effect at relatively high concentrations, i.e., at least 10 mM (29). The beneficial effects of ATP and of creatine phosphate were found to be additive, since these substances together at their individual optimal concentrations provided significantly better myocardial protection than either substance alone. In addition to improved mechanical function and better maintenance of cellular integrity, high-energy phosphates as additives to cardioplegic solutions showed marked anti-arrhythmic effects (29).

The precise mechanisms underlying the beneficial effects of high-energy phosphates added to cardioplegic solutions remain to be established. With respect to ATP, a number of studies have indicated that extracellular ATP acts as a cardioplegic agent, depressing contractility of the heart (20). The cardioplegic action might be due to the calcium and magnesium chelating properties of ATP. Besides, it has been proposed that ATP also exerts other protective effects probably unrelated to the depression of cardiac contractility during the ischemic period. A direct intracellular site of action of ATP is not very likely. Myocardial cellular membranes are believed to be impermeable to extracellular ATP, ruling out the possibility that exogenous ATP will replenish or enhance the intracellular high-energy phosphate pool in a direct manner. Since degradation of exogenous ATP readily occurs in the coronary vascular and interstitial space (21), the beneficial effect of ATP might be mediated by its degradation products. However, Robinson et al. (30) concluded that the intact ATP molecule is required for maximal myocardial protection, since perfusion with ATP degradation products, such as adenosine and inorganic phosphate, failed to exert similar beneficial effects. The observation that the protective effects of ATP and creatine phosphate are additive (30) makes a common site of action less likely. Since creatine and inorganic phosphate, present at similar concentration in the cardioplegic solution as creatine phosphate, failed to protect the heart against loss of cellular integrity and post-ischemic depressed cardiac performance, intact creatine phosphate molecules are obviously required for the established beneficial action (29).

Extracellular creatine phosphate may exert its protective effect via an intracellular and/or extracellular site of action. It is generally believed that creatine phosphate cannot directly enter the myocyte, although findings of Breccia et al. (4) suggest that this polar compound can cross the sarcolemmal barrier under certain circumstances. In addition, Down et al. (11) have reported that intravenous injection of creatine phosphate (50 mg/kg body wt.) resulted in increased intracellular levels of creatine phosphate in the normoxic rat heart. Saks et al. (34) were unable to substantiate these findings in dogs. However, addition of creatine phosphate to a cardioplegic solution resulted in significantly enhanced post-ischemic cardiac ATP and creatine phosphate levels both in rats and human (34).

Robinson et al. (29) have pointed out that creatine phosphate may also exert a beneficial action by donating its high-energy phosphate group to plasmalemma-bound ADP with subsequent formation of extracellular ATP. The role of extracellular ATP is hypothetical, but maintenance of the content of ATP localized at the interstitial site of the sarcolemma might be qualitatively of great importance. Recently, Saks et al. (33) have hypothesized that the protective action of extracellular creatine phosphate may include 1) inhibition of sarcolemmal 5'-nucleotidase resulting in a less marked release of adenine nucleotide degradation products from the (post-)ischemic heart; 2) inhibition of accumulation of ADP-induced aggregation of platelets. Ronca et al. (31) have further extended the list of putative beneficial modes of action. They suggest that the high-energy phosphate protects ischemic and reperfused myocardial cells by reducing free oxygen radical formation and by decreasing the peroxidative damage of cell membranes. Unfortunately human studies dealing with ATP and creatine phosphate administration during open-heart surgery are scarce. Saks et al. (34) have reported that creatine phosphate (in the range of 8-10 mM) added to a sanguineous cardioplegic solution significantly reduced the incidence of post-ischemic arrhythmias in patients who underwent valve operation. Obviously, more clinical studies are needed to rate the experimentally obtained findings at their true value.

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# Chapter 23

# Apparent Inosine Incorporation and Concomitant Haemodynamic Improvement in Human Heart

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Haemodynamic effects of inosine were studied in five patients suffering from prolonged, otherwise intractible cardiogenic shock, and in six patients catheterized for coronary angiography. Heart performance was improved by inosine as evidenced by enhanced cardiac index (by  $63 \pm 29\%$ , p < 0.002; mean  $\pm$  SEM) in patients with extremely low cardiac output, and increased LV  $dp/dt_{max}$  (by  $22 \pm 7\%$ ) in patients with normal perfusion during angiography. Inosine exerts a positive inotropic effect in man. No data, so far, are available concerning the fate of exogenous inosine in human myocardium. Therefore, during inosine infusion in patients undergoing diagnostic cardiac catheterization, arterio-coronary sinus (a-v) differences of purine nucleosides were determined simultaneously with haemodynamic measurements. Before infusion the a-v difference of inosine, hypoxanthine and xanthine across the heart was nil, whereas urate production was evident. During infusion arterial inosine increased, exceeding the coronary sinus concentration by a maximum of  $200 \pm 52$  uM (p < 0.05) at the fourth minute. At that time the a-v difference in hypoxanthine, xanthine and urate amounted to  $16 \pm 11$ ,  $10 \pm 3$  (p < 0.05), and  $15 \pm 14$  uM, respectively. Thus there seems to be a substantial uptake of inosine by the human heart, concomitant with haemodynamic improvement.

#### Introduction

Severe left ventricular (LV) failure and shock following myocardial infarction (MI) lead to high mortality. This creates a continuous need for an effective drug, increasing cardiac output and improving peripheral perfusion. The efficacy of a variety of inotropic drugs currently used in the therapy of cardiogenic shock is not satisfactory. Sympathomimetics and their derivatives produce marked improvement in LV function. But they exert many known untoward effects, including increased myocardial oxygen consumption and infarct size. Digitalis, the traditional drug used in congestive heart failure, is virtually ineffective in patients with advanced pump failure. All these facts encourage an active search for other positive inotropic drugs. In view of this need, inosine, an inotropic substance naturally occurring in the body, is a promising agent. The myocardial cell produces this nucleoside by both deamination of adenosine and dephosphorylation of IMP (5). The heart releases it in large quantities after exposure to hypoxia, anoxia or ischaemia. Then ATP breakdown is stimulated (4,26,28,29). Inosine augments myocardial blood flow (3,18,19) and increases contractility of both normal (3,18,19,22) and ischaemic heart muscle (32,33) in experimental animals. Nevertheless, there are conflicting results concerning its inotropic action in different species. Its positive effect is evident in the dog (11,22,

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30,32), pig (10,33), rabbit (23), and frog (6). In contrast, the effect of inosine is absent in the guinea pig (17) and even negative in the rat (17,34).

#### Inosine for Cardiogenic Shock in Man

Inosine limits the size of experimental MI (10,12) and counteracts experimental LV failure (30,31). This encouraged us to apply the endogenous nucleoside in the management of low cardiac output states in man. Two female and 3 male patients 25-82 years, suffering from prolonged, otherwise intractable cardiogenic shock due to MI (4 patients) and congestive cardiomyopathy (1 patient), were studied (9). All suffered from circulatory failure characterized by clammy skin, mental obtundation and anuria. Cardiac index (CI) ranged from 0.58 to 1.63 l/min/m<sup>2</sup>. Before the inosine study was performed, all the patients had received conventional pharmacotherapy (dopamine, dobutamine, noradrenaline, glucagon, digoxin, hydrocortisone, etc.) with no signs of improvement, however. The haemodynamic data were obtained from a Swan-Ganz catheter. Cardiac output was measured in triplicate by the thermodilution technique and computed using a Hewlett-Packard cardiac output computer. The pressure readings were obtained by means of external transducers and recorded simultaneously with the heart rate. Inosine (Trophicardyl i.v., Laboratoire Innothéra, France) was infused in a dose of 5 mg/kg/min (4% solution). Haemodynamic variables were recorded prior to and at the 6th minute of infusion.

Inosine administered to patients who had not responded previously to conventional pharmacotherapy resulted in an increase in the CI (by  $63 \pm 29\%$ , p<0.002; mean  $\pm$  SEM) and the stroke volume index (by  $55 \pm 15\%$ , p<0.001). Pulmonary capillary wedge pressure and heart rate were not significantly affected (Table 1). Despite a high increase in CI in each case, in 2 patients blood pressure was unobtainable throughout the whole study. In another one it was unchanged and in the remaining two slightly increased by inosine. None of the patients survived; we decided to infuse inosine at the end-stage of shock after conventional therapy had failed, to improve performance of the heart. Furthermore, the delay between the first symptoms of shock and nucleoside administration was considerable.

Variable	Incre	eased	by	Р
HR (beats/min)	5	±	6	NS
PCWP (mmHg)	0.8	±	0.8	NS
CI (ml/min/m <sup>2</sup> )	680	±	14	< 0.01
SVI (ml/m <sup>2</sup> )	6.1	±	1.9	< 0.001

Table 1. Influence of inosine (5 mg/kg/min) on heart rate (HR), pulmonary capillary wedge pressure (PCWP), cardiac index (CI), and stroke volume index (SVI) in 5 patients suffering from prolonged, otherwise intractable, cardiogenic shock

#### Studies on Inosine Incorporation by Human Myocardium

Inosine exerts a positive inotropic effect in man. It improves significantly myocardial performance in patients with extremely low cardiac output. No data, so far, have been available concerning the fate of this agent in man. Therefore, we designed a study (21) in which inosine was infused to 6 patients undergoing diagnostic coronary catheterization in the course of coronary heart disease. We attempted to answer the following questions:

- 1. Is exogenous inosine taken up by the human heart?
- 2. Is the inotropic effect of inosine related to its myocardial uptake?

The protocol of inosine administration was similar to the one used in the previous study. The nucleoside (Trophicardyl) was infused at a rate of 5 mg/kg/min into the peripheral vein. The haemodynamic variables were recorded before and at the 6th minute of infusion. The first derivative of LV pressure (LV dp/dt<sub>max</sub>) was obtained from a resistance-capacitance differentiating circuit (Siemens-Elema). Arterial and coronary sinus blood was sampled for a purine assay before and after 2, 4 and 6 min of the procedure. Inosine, hypoxanthine, xanthine and uric acid were determined by the high-performance liquid chromatography (HPLC) method described in detail by De Jong et al. (21) and Harmsen et al. (14).

Inosine administration had no significant effect upon heart rate, LV end-diastolic pressure (Fig. 1) as well as on LV systolic and diastolic pressures. The contractility of the heart, as measured by LV dp/dt<sub>max</sub>, increased significantly by  $22 \pm 7\%$  (Fig. 1), varying from 6% to 44% in individual patients. During inosine infusion, a substantial increase in the arterial nucleoside concentration as well as that of its catabolites hypoxanthine, xanthine and urate was observed (Fig. 2). Figure 3 presents the arterio-venous differences of the nucleosides under study. There was no a-v difference in inosine, hypoxanthine and xanthine in control measurements except for urate. Urate was produced by each heart, indicating the presence of xanthine oxidase in the myocardium of the patients studied. During inosine infusion part of the arterio-coronary sinus differences for hypoxanthine, xanthine and urate were 16 ±11



Fig. 1. Influence of inosine (5 mg/kg/min i.v.) on heart rate, left ventricular end-diastolic pressure (LVEDP) and LV dp/dt<sub>max</sub>. Results obtained in five patients, catheterized for coronary angiography (means  $\pm$  SEM). C = control; Ino = inosine.

uM, 10  $\pm$  3 uM and 15  $\pm$  14 uM, respectively (Fig. 3). The large a-v difference of inosine at the same time (200  $\pm$  53 uM) suggests that this nucleoside is partially converted to catabolites and partially taken up by the myocardium.

#### Discussion

The present findings demonstrate that inosine exerts a positive inotropic effect in man. It improves significantly myocardial performance in patients with extremely low cardiac output. The same is true for patients with normal peripheral perfusion. The positive inotropic action was proved by an increase in cardiac index and stroke volume index in patients suffering from shock and by an increase in LV dp/dt<sub>max</sub> in patients undergoing diagnostic cardiac catheterization. The LV end-diastolic pressure measurements (as obtained from the pulmonary capillary pressure in patients suffering from shock and from the LV cavity in patients during catheterization) suggest that inosine doesn't affect the preload. Afterload also seems unaffected by the nucleoside, since there were no significant changes in blood pressure in both studies.



Fig. 2. Inosine-induced increase in arterial purine concentrations. The concentrations of inosine, hypoxanthine, xanthine and urate were measured prior to and at the second, fourth and sixth minute of i.v. inosine administration at a dose of 5 mg/kg/min.

Inosine Incorporation and Haemodynamic Improvement

The metabolic cost of inosine action is expected to be relatively low. The increase in stroke volume in patients suffering from shock, and the increase in LV  $dp/dt_{max}$ in the catheterized patients, were not accompanied by a parallel, significant increment in the heart rate. This is in agreement with our previous studies on open-chest dogs with an acutely implanted coronary flow probe (8) and on closedchest dogs with acute LV failure (31). In these inosine enhanced contractility and myocardial blood flow without raising myocardial oxygen consumption. These results were subsequently published by Smiseth (30). Thomas and Jones (32) found in the dog model that inosine enhanced contractility of both ischaemic and nonischaemic myocardium. However, the content of high-energy phosphates remained unchanged.

The mechanism of inosine action is not completely understood. We reported (11) that the nucleoside-induced increase in contractility is mediated in part by the  $\beta$ -adrenergic system and in part by an independent mechanism. To elucidate further the fate of inosine in the heart in that study, the nucleoside was administered in a dose of 5 mg/kg/min i.v. This dosage was shown to increase the inosine content in the myocardium up to the amounts observed during experimental coronary occlusion or cardiac arrest (19,20,27).



Fig. 3. Apparent uptake of inosine and its catabolites by the human heart during i.v. inosine administration. Note that before infusion the arterio-coronary sinus difference of inosine, hypoxanthine and xanthine across the heart was nil, whereas urate production was evident (lower panel). Means  $\pm$  SEM (n = 5-6), \*p<0.05, \*\*p<0.01 vs. baseline value.

The present results show that under baseline conditions the heart produces substantial amounts of urate. The a-v difference across the heart is about 50 uM (Fig. 3), which is in agreement with the preliminary data of Nelson (25). The finding suggest that the human heart contains a relatively active xanthine oxidoreductase. Although this enzyme has been demonstrated in the heart of several species, there are not many data available concerning the human myocardium. The oxidase form of xanthine oxidase can generate oxygen free radicals, playing an important, causative role in the development of coronary heart disease. During inosine infusion the arterial and coronary sinus urate levels rose, apparently due to an increased supply of its precursors hypoxanthine and xanthine (Fig. 2). The a-v difference of urateacross the heart disappeared (Fig. 3), which could indicate that xanthine oxidoreductase is inhibited by inosine or its catabolites.

Simultaneously with the high a-v inosine difference at the second and fourth minute of nucleoside administration, the increase in contractility was observed. At the sixth minute, this apparent myocardial inosine uptake was absent, but the positive inotropic effect was still the same. Whether redistribution of inosine among various intra- and extracellular compartments takes place during this phase of the infusion, remains to be studied. Inosine promotes enhanced nucleotide resynthesis in the myocardium (2,15) via breakdown to hypoxanthine, subsequent ribophosphorylation, and conversion to AMP. We therefore speculate that inosine incorporation into adenine nucleotides is responsible for the increased contractility of human heart.

It is noteworthy that no arrhythmia, ischaemia or other side effects were observed. Inosine, in contrast to catecholamines and digitalis, proved to be a cardiotonic agent without arrhythmogenic properties (1,13). Furthermore, this nucleoside could protect the myocardium against toxic effects of catecholamines (7,16,24).

In view of the need for effective therapeutic intervention to improve myocardial performance in cardiogenic shock, inosine is worth considering. It is an endogenous substance, free of side effects. Its helpful action, both in experimental myocardial infarction (10,12) and experimental acute LV failure (30,31), has been well proved. The initial clinical results obtained with inosine are promising and encourage further trials focused on its mechanism of action. Prolonged, early instituted administration of inosine seems to be the next logical step to assess the value of this therapy in the management of cardiogenic shock.

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# Chapter 24

# Cardioplegia and Calcium Entry Blockade

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Cooling and induction of electromechanical arrest (cardioplegia) limit myocardial damage during open-heart operations. Many studies provide evidence for the use of calcium entry blockers, as adjuncts to cardioplegia to optimize the protection. This paper reviews their possible mechanism of action, application time, and efficacy during hypothermia. A major conclusion is that negative effects are virtually absent, apart from short-term negative inotropic responses. There is an increasing body of positive evidence for their efficacy. A new development is the use of these drugs for regional cardioplegia during dilation of coronary arteries (coronary angioplasty).

## Necessity for Cardioprotection

In his presidential address, entitled "Intellectual creativity in thoracic surgeons", Spencer (69) analyzed 8 breakthroughs in cardiovascular surgery over the last 30 years. One of the subjects concerned myocardial preservation. Additional protection of the heart by calcium entry blockers is clear in many animal studies. The time has come for more extensive clinical work.

Buckberg et al. (9) pointed at the diminished heart function after extracorporeal circulation and the importance to protect the myocardium adequately. They asked: "Why should a heart which did not need pharmacologic or mechanical support before operation require these interventions after the cardiac lesion has been corrected?" Their methods, including avoidance of prolonged hypothermic cardiac arrest, sharply diminished the postoperative use of inotropic drugs and reduced mortality from 12 to 3% (9). Mortality within one month after aorta-coronary bypass surgery at my institution is around 1% (49). Although these figures are relatively low, one has to try to reduce mortality further. This holds a fortiori for certain other cardiovascular operations (68).

Morbidity should and can improve as well. Full restoration of myocardial function can take months after bypass grafting (66). Nowadays, induction of electromechanical arrest (cardioplegia) combined with hypothermia limits the initial insult. It ameliorates thereby recovery after cardiac operations. However, improvements are still possible; certain types of cardioplegia fail to protect the heart adequately from ischemic damage. In the last few years, substantial research has been carried out on interventions involving calcium metabolism. This chapter updates a review on calcium antagonists and cardioplegia (21).

## Calcium

The myocardial  $Ca^{2+}$  content in mammals is about 2 mmol/kg. The largest part resides in internal stores near the endoplastic reticulum. During diastole, the cytosolic  $Ca^{2+}$  concentration is presumably lower than 0.0001 mM. However, for active tension development, a  $Ca^{2+}$  concentration of about 0.01 mM is necessary. Therefore, calcium from subcellular organelles or from outside the cell moves to the

cytosol. This accomplishes the change from resting to active status. In the blood, the ionized  $Ca^{2+}$  level is about 2.5 mM.

Calcium ions can enter the cell by passive diffusion, by exchange with sodium, by voltage-activated transport, and possibly also by exchange with potassium. Voltage-activated transport takes place via the slow calcium channels. Per beat, only little calcium enters (reversibly) the cell. If there is a lack of oxygen, however, the cardiac cell membrane becomes more permeable for  $Ca^{2+}$  ions from the extracellular space. An increased influx into the ischemic tissue during reperfusion will ensue (57). Excessive accumulation leads to ischemic contracture, the "stone heart" syndrome, a life-threatening event. It is therefore imperative to prevent calcium accumulation, whatever the route. The phenomenon of the calcium paradox illustrates the importance of calcium homeostasis for the myocyte.

#### **Calcium Paradox**

The calcium paradox occurs when hearts are perfused with a solution containing less than 0.05 mM Ca<sup>2+</sup>, followed by one with a normal calcium concentration (see Chapter 19). Then tissue disruption follows, with enzyme release and contracture. Abrupt massive calcium influx to intracellular compartments seems responsible for the process (29). The clinical relevance of this discovery did not become clear until later: in a number of surgical procedures, calcium supply to the heart stops. This possibly initiates the calcium paradox during reperfusion. It is unlikely that the slow channels provide a major route of Ca<sup>2+</sup> entry during the calcium paradox (57). Nevertheless, calcium entry blockers can prevent some of the damage, which occurs because of the calcium influx (65).

#### **Calcium Entry Blockers**

The term *calcium antagonist* is in use since 1969 for drugs with negative inotropic properties that can be counteracted by calcium. In the definition of Fleckenstein (28), a calcium antagonist specifically blocks excitation-contraction coupling by mitigating the effects of  $Ca^{2+}$ . Drugs like verapamil, nifedipine and diltiazem are calcium antagonists that are valuable to treat angina pectoris and hypertension. They inhibit the slow, inward calcium flux, responsible for the plateau phase of the action potential. Alternative names include "slow-channel inhibitors", "calcium channel inhibitors" or "calcium entry blockers". The negative inotropic effect of these compounds can be ascribed to this decreased, slow calcium influx. Figure 1 shows in schematic form a classification of various calcium modulators. From Table 1, it is clear that cardioprotective calcium entry blockers form a heterogenous group of drugs. Nifedipine, a 1,4-dihydropyridine, is the "mother" compound of a series of drugs such as nicardipine, niludipine, nimodipine, nisoldipine, and nitrendipine. These belong to the most potent vasodilators on the market, with varying tissue specificity. Verapamil, a compound with a structural resemblance to papaverine, has as its "congeners" gallopamil, tiapamil, and anipamil.

The heart catabolizes quickly its energy-rich phosphates during lack of oxygen. Then ATP, which is essential for contraction, breaks down to the level of the purines adenosine, inosine, hypoxanthine, xanthine, and urate. These metabolites leave the cell. Catabolism of the adenine nucleotide pool by more than 75% prohibits resuscitation of an ischemic heart (42) As shown by many authors, verapamil can inhibit ATP breakdown in ischemic and reperfused heart (28,57). In addition, energy conservation is possible with diltiazem, nifedipine, nisoldipine and bepridil (22-24,41,74).

Many studies demonstrate the salutary effect of calcium antagonists on myocardial energy metabolism during transient ischemia. Nevertheless, their mechanism of action is not fully clear. Calcium overload leads to excessive ATP consumption by ATPases Cardioplegia and Calcium Entry Blockade

and reduced mitochondrial ATP production. The resulting exhaustion of energy-rich phosphates will ultimately cause myocardial necrosis. Blockade of slow  $Ca^{2+}$  channels will restore calcium homeostasis with normal ATP consumption and adequate ATP synthesis. Protection due to reduction of work, to vasodilatation, or to interference with organelles, rather than to direct action on  $Ca^{2+}$  fluxes across the cell membrane, cannot be excluded. Indeed, the degree of negative inotropy observed before ischemia, seems to determine how much ATP is catabolized during ischemia (24). And also much the heart recovers (74).

#### **Calcium Entry Blockade during Heart Operations**

With cold chemical cardioplegia during heart operations, the tolerable duration of ischemia increases from less than 1 to 3 h. In experimental studies, reversible ischemia of up to 24 h appears possible (34). Effective protection has 3 facets: 1) energy sparing by induction of instaneous electromechanical arrest with compounds such as potassium; 2) slowing down of energy-consuming and degradative reactions by hypothermia; and 3) combating harmful changes caused by ischemia and reperfusion with protective agents (34). The control of calcium movement seems to be critical to protect the heart effectively (see Chapter 22). In this setting, calcium entry blockers could be helpful. From the available literature, it is clear that many experiments with animals and some clinical trials demonstrate calcium antagonists to have a protective effect on the heart.

#### **Animal Studies**

Table 2 shows the effect of diltiazem, lidoflazine, nifedipine, and verapamil as adjuncts to potassium cardioplegia. In all studies carried out in myocardium at 37°C,



Fig. 1. Inhibition and stimulation of calcium uptake by various agents. This scheme is based on a recent classification by Godfraind (31).

calcium entry blockers protected against damage caused by ischemia. Somewhat less clear is the effect of these drugs on the hypothermic heart. Many investigators reported additional protection by calcium antagonists. In a number of studies, however, the drugs did not significantly ameliorate ischemia-induced damage during hypothermia (see Table 2). At lower temperatures, calcium antagonists also didn't prevent damage caused by the calcium paradox (3). Nayler (58), on the other hand, showed that the protective effects of hypothermia and nifedipine are additive. However, in hearts arrested with potassium, this additive effect was absent at temperatures below 28°C (57). Calcium entry blockers introduce a discontinuity around this value into the Arrhenius plot of the maximal contraction frequency vs. temperature (62).

If the ischemic damage is virtually nil or the protection by hypothermia alone already adequate, calcium antagonist administration may be redundant. However, even if the specific cardioplegic effect is diminished, the calcium antagonist may be useful by lowering the pre-ischemic energy demand (22). Also, reperfusion conditions may be improved. It is noteworthy that calcium antagonists used as adjuncts to potassium cardioplegia don't give negative effects. Moreover, hearts arrested by means other than potassium cardioplegia also showed protection by verapamil, diltiazem, nifedipine, and lidoflazine. One exception should be noted: diltiazem combined with magnesium cardioplegia was detrimental to postischemic function, despite a rapid restoration of energy stores (73). High magnesium cardioplegia could induce heart block (10). Using calcium entry blockers, one has to keep in mind that a (reversible) negative inotropic effect can persist for some time. This could cause problems in the hospital setting.

#### **Clinical Trials**

Details on studies of calcium entry blockers used during cardiac operations are scarce (Table 3). Clark et al. (19) reported the effectiveness of nifedipine in a multicenter trial. They selected 205 highest-risk patients from a large open-heart population. The investigators employed cold potassium cardioplegia. Nifedipine influenced hemodynamics in a positive sense, but the authors provided no statistical details. Nifedipine seemed to preserve myocardial levels of ATP also in this study. However, no significant reduction in myocardial creatine kinase release occurred.

Calcium entry blocker					
	Selective			Nonselective	
Benzothiazepine	Dihydropyridine	Phenyl- alkylamine	Benzylphenylamine- pyrrolidine	Diphenylalkyl- piperazine	Phenyl- alkylamine
Diltiazem	Felodipine Nicardipine Nifedipine Niludipine Nisoldipine Nitrendipine	Anipamil Gallopamil Ronipamil Verapamil	Bepridil	Lidoflazine	Tiapamil

Table 1. Some cardioprotective calcium antagonists, acting at the plasma membrane

Calcium entry blockers, selective for slow calcium channels in myocardium, are also called slow channel blockers. The nonselective agents, mentioned in this table, act at similar concentrations on calcium and fast sodium channels. See also Figure 1.

#### Cardioplegia and Calcium Entry Blockade

Flameng et al. (27) found that the addition of nifedipine to the cardioplegic solution could prevent the ischemia-induced degradation of nucleotides, as it occurred when myocardial cooling was inadequate. However, clinical outcome didn't improve. In an earlier trial, Flameng et al. (26) showed that lidoflazine prevents in a dose-dependent way high-energy phosphates and glycogen breakdown. In addition, lidoflazine maintained ultrastructure and left ventricular stroke work index better. In this investigation, patients were pretreated with the drug. Operation conditions included intermittent aortic clamping and hypothermic conditions.

		Temperature			
Drug	Preparation	of heart (°C)	Variables tested	Result	Reference
<b>D</b> <sup>1</sup>	<b>T 1</b> , <b>1</b> , <b>1</b> ,	20 27	<b>TT</b> 1 1 / 1 1		
Diltiazem	Isolated rat heart	20 vs. 37	Hemodynamics/mitoch.	NS vs. +	76
	Dog, CPB	1	P/hemodynamics	±	70
	Isolated rat heart	10	P/arrhythmias	+	71
	Dog, CPB	20 vs. 28	Hemodynamics	NS vs. +	48
	Dog, CPB	37	P/hemodynamics	±	1
Lidoflazine	Dog, CX ligation	30	Hemodynamics	NS	33
	Dog, CPB, CX ligation	20	Hemodynamics	+	45
	Dog, CPB	37?	Cerebral flow	+	75
Nifedipine	Dog. CPB	10	Hemodynamics/edema	+	52
1	Dog. CPB	10	Hemodynamics	+	16
	Dog. CPB	10.37	Hemodynamics	+	18
	Dog. CPB	24	$Ca^{2+}$ homeostasis	+	7
	Dog. CPB	12. 21	P/hemodynamics	NS	43
	Isolated rabbit heart	15-37	Hemodynamics	+ >28°C	57
	Isolated rat heart	20 vs. 37	Hemodynamics/enzymes	+ vs +	77
	Isolated rat heart	15	P/hemodynamics	+	54
	Dog. CX ligation	30	Hemodynamics	+	33
	Isolated rabbit heart	14	Hemodynamics	+	46
	Isolated rat heart	20	P/hemodynamics	+ /-	6
	Isolated rat heart	37	P/hemodynamics	NS/+	78
	Isolated rat heart	20-37	Hemodynamics/enzymes	+. >31°C	30
	Pig, CPB	28	Hemodynamics/lactate	±,	55
	Guinea pig, CPB*	9 vs. 34	Hemodynamics	NS vs. +	14
	Dog, CPB	21	Arrhythmias	+	25
Verapamil	Dog. CPB	37?	Hemodynamics/mitoch	+	61
•	Isolated dog heart	4	Hemodynamics	+	56
	Isolated rabbit heart	14	Hemodynamics	+	46
	Dog, CPB	37?	Cerebral flow	+	75
	Isolated rat heart	20-37	Hemodynamics/enzymes	+. >34°C	37
	Dog, CPB	37	P/hemodynamics	+	51
	Dog, CX ligation	18	Mitoch.	+	79
	Isolated rat heart	37	P/hemodynamics	NS/+	78
	Dog, CPB	9	P/hemodyn./structure	+	50

Table 2. Efficacy of calcium entry blockers given as adjuncts to potassium cardioplegia to prevent damage due to myocardial ischemia

<sup>+</sup> indicates protection additive to potassium cardioplegia, and NS means protection not additive to cardioplegia. <sup>\*</sup> Heart-lung model. CPB = cardiopulmonary bypass; CX = circumflex coronary artery; mitoch. = mitochondrial function; P = myocardial high-energy phosphates. Hicks et al. (40) studied the effect of verapamil added to the cardioplegic solution and nifedipine instituted postoperatively. This regimen, for which the rationale remained unexplained, resulted in a reduction in postoperative levels of serum creatine kinase. From a more recent study, conducted without suitable controls, Hicks and DeWeese (39) concluded that verapamil potassium cardioplegia protected well against postoperative abnormalities in cardiac conduction. Kaplan et al. (44), however, did not advocate the addition of verapamil to potassium cardioplegia. In their small group of patients, it increased the need for pacing and inotropic drugs.

Vouhé et al. (72) studied the effects of diltiazem in conjunction with profound hypothermia during aortic cross-clamping. They found it safe; the drug reduced the release of myocardial enzyme and probably improved hemodynamic function immediately after bypass. Christakis et al. (15) also observed an improvement in biochemical variables and cardiac index with diltiazem cardioplegia. However, they reported potent negative inotropy and prolonged electromechanical arrest. Similarly, Barner et al. (5) noted that the addition of diltiazem to cold blood potassium cardioplegia prolonged the atrioventricular node recovery time.

From these trials and from preliminary data (35), the overall picture emerges that calcium entry blockers can contribute to protection during open-heart operations.

#### **Calcium Entry Blockers for Regional Cardioplegia**

In the last decade, percutaneous transluminal coronary angioplasty (PTCA) has established itself in the armamentarium of the cardiologist. He introduces a catheter through a systemic artery to dilate a stenotic artery by controlled inflation of a balloon. For some time only a proximal stenosis in a single vessel could be treated, provided the patients had stable angina and normal ventricular function. Today treatment is less conservative. Serruys et al. (67) reported that they use PTCA also for patients with unstable angina and diminished ventricular function. Various obstructed vessels can be dilated. The total period of occlusion can exceed 10 min.

Drug	Variables tested	Result	Ref.	
Diltiazem	Hemodynamics/enzyme release	+	72	
Dimuzoni	P/hemodynamics/enzyme release	+	15	
	Hemodynamics/enzyme release	±	5	
Lidoflazine	Hemodynamics/P/glycogen/EM	+	26	
Nifedipine	Hemodynamics/enzyme release	+/±	19	
ľ	Hemodynamics/P	NS/+	27	
Verapamil	Enzyme release	+	40	
	Hemodynamics/enzyme & lactate release	NS	44	
	Hemodynamics	*	32	

Table 3. Prevention by calcium entry blockers of damage during/after aorto-coronary bypass grafting (clinical trials)

+ indicates added protection during bypass grafting or valve replacement. \* A negative effect was reported, but see criticism by Clark (17). Abbreviations: EM = electron microscopy; NS = no significant improvement; P = myocardial high-energy phosphates.

After PTCA, substantial amounts of lactate and ATP catabolites are demonstrable in the myocardial efflux (63). It is therefore obvious that protection of the heart during such an intervention is useful. Local cardioplegia with calcium antagonists is now being tested. Calcium entry blockers can prevent arterial spasm during coronary angioplasty. It seems as if the time necessary for dilation can be prolonged safely by intracoronary application of nifedipine (67). In fact, Pop et al. (63) reported that this drug reduced lactate production during the procedure.

Diltiazem, administered intracoronarily, reduced severity and time of onset of ischemic ECG changes as well as anginal pain during PTCA (47). However, this antiischemic effect remained unexplained. Protection of the heart with calcium entry blockers during angioplasty is only in the initial research phase.

#### **Calcium Antagonists as Insurance Policy**

Hearse (35) stressed that proper hypothermia is relatively easy under in the heart laboratory conditions. That is certainly not always possible in the clinical setting, where one observes warm spots. Also, problems exist because of a varying degree of cooling. Calcium antagonists could then act as an insurance policy against regions of poor perfusion or unexpected warming (35), when administered adjunct to hypothermic cardioplegia. Furthermore, some of these drugs protect against vasoconstriction by microscopically small particles in unfiltered cardioplegic solutions (36). This is the reason Henry (38) called them "particle antagonists". He made it clear that the action of calcium entry blockers goes beyond just another cardioplegic maneuver. These drugs seem to play an important role in protecting the heart during the periods before and after cardiopulmonary bypass. During reperfusion, the cardioplegic protection by hypothermia and potassium is no longer effective. It becomes important to interfere pharmacologically with minimal disturbance of pump activity. Calcium antagonists could exert their salutary action partly by protecting the vessel wall. Nifedipine attenuates the strong contracture of isolated coronary arteries, produced by a cardioplegic solution (14). Finally, some of these drugs could be beneficial to inhibit rhythm disturbances during reperfusion (37,71).

#### Administration: What, and How Much?

One has to keep in mind that certain calcium antagonists (verapamil, diltiazem) affect the electrical properties of the heart. Nifedipine and verapamil show a similar ability to protect the rat myocardium against global ischemia. However, they exhibit significant differences in mode and characteristics of recovery during early reperfusion (78).

The potency to inhibit calcium influx and to dilate coronary vessels is high for the dihydropyridines. The relative potency is nifedipine > diltiazem > verapamil. Another point of consideration is the composition of the cardioplegic solution to which a calcium antagonist is adjunctive. St. Thomas' Hospital solution contains 16 mM  $Mg^{2+}$ , a physiologic calcium blocker (2). Some cardioplegic solutions, on the other hand, are devoid of this ion.

It is sensible to give calcium antagonists in a moderate dose together with the cardioplegic solution before open-heart operation. For nifedipine, this is probably 200 to 300 ug/L, using about 1.5 L of cardioplegic solution (19,27). A high concentration of calcium antagonist strongly depresses the heart. Problems during weaning from extracorporeal circulation or in the postoperative period could occur. Difficulties could arise during the reperfusion phase if calcium antagonists wash out very slowly. The biological half-life for i.v. administered verapamil and diltiazem is 4 to 5 h, that of nifedipine 1 to 2 h (53). Verapamil and bepridil accumulate in

muscle cells (60). The human heart extracts the former rapidly (32). The latter binds very strongly to membranes. It will take about 1 h to remove bepridil from myocarcardial tissue. In contrast, nifedipine and diltiazem permeate more slowly (60) and their displacement is rapid (see ref. 34). Bepridil and nisoldipine wash very slowly from the tubing system used in a perfusion apparatus (24). The long-lasting cardiac depression by nifedipine, observed (6) in an NMR-machine with long tubings, could be due to drug release. Therefore, it is important to check the affinity of calcium antagonists for the extracorporeal circuit to which they have access.

#### When, How, and to Whom?

When should calcium entry blockers be given to minimize ischemic damage effectively? Prophylactic use, i.e., administration before flow reduction, is strongly advisable (20,22,59). Calcium antagonists, given i.v. to animals before ischemia or as a component of hypothermic hyperkalemic cardioplegia, hypothermic normokalemic cardioplegia, or normothermic normokalemic cardioplegia, show salutary effects on the heart in most cases. Preusse et al. (64) claimed that pretreatment with verapamil followed by calcium-free cardioplegic solution has a "membrane-labilizing" effect. However, because of the risk of creating the calcium paradox, one shouldn't use calcium-free cardioplegia.

The calcium antagonist should be present before reperfusion starts. It is debatable whether it makes sense to treat patients per os for several weeks preoperatively. Some groups treat only high-risk surgical patients with these drugs. On the basis of their study with nifedipine, Casson et al. (12) suggested to continue the drug up to the time of operation. Verapamil pretreatment before aortic crossclamping may potentiate the myocardial preservation achieved by cardioplegia (4).

#### Conclusion

Many studies, mainly conducted in animals, show that calcium entry blockers are useful in open-heart surgery as adjunct of potassium cardioplegia and hypothermia. Optimal conditions have to be determined in clinical trials before their routine use can be advocated. To create local cardioplegia, calcium antagonists given intracoronary before transluminal angioplasty, are promising. Let's consider Buckberg's words: "The future does not lie in providing the value of cardioplegia, but in understanding better the deleterious effects of ischemia and reperfusion injury so that current cardioplegic techniques can be modified to avoid these effects" (8).

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# Chapter 25

# The Effect of Anesthetics on Myocardial Adenine Nucleotides

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The major interest in the effect of anesthetics on high-energy phosphates (adenine nucleotides) in the heart stems from the search for the mechanism of the negative inotropic effect of anesthetics and the interest in preserving myocardial energy during cardiac ischemia. Although the number of studies published is rather small, the results when the preparation has been stable are consistent. Anesthetics do not appear to change the concentration of high-energy phosphates in normal hearts. There is some evidence that barbiturates in high concentrations may interfere with myocardial energy synthesis and decrease high-energy phosphates in hypoxic or ischemic hearts and additional evidence that perhaps cardiac depressant anesthetics such as halothane may preserve high-energy phosphates in ischemic hearts. However, much more investigation is necessary in the ischemic heart in order to come to definite conclusions.

## Introduction

Why should there be interest in the effect of anesthetics on high-energy phosphates (adenine nucleotides) in the heart? In my opinion, there are basically two reasons. First, all potent inhalation anesthetics are dose-related myocardial depressants. However, the mechanism of this depression remains elusive (12,18). It seems likely that one or more sites lie in the supply of energy for myocardial muscle contraction or in the use of that energy as depicted in Fig. 1. Note that the myocardial high-energy phosphates, adenosine triphosphate (ATP) and creatine phosphate (CP), lie in the middle of these two opposing processes. Consequently, investigators interested in the mechanism of the negative inotropic effect of anesthetics have sought to measure the concentration of the storage form of energy for the heart, these high-energy phosphates, in an attempt to determine whether interference in the energy supply for the work of the heart by anesthetics might result in a decrement in the major energy source for heart muscle contraction, the ATP and CP.

The prevalence of patients needing anesthesia and surgery with ischemic heart disease in the western world dictates the second reason for the interest in the effect of anesthetics on myocardial high-energy phosphates. Inasmuch as the continuing high energy demand of the heart dictates that oxidative metabolism be maintained for adequate synthesis of the energy supply of the heart (ATP and CP), the ultimate pathophysiologic defect produced by myocardial ischemia must be a deficit of these energy sources (6). In order to decrease the mortality and morbidity of surgery and anesthesia in these high-risk patients, anesthetics which preserve high-energy phosphates, particularly in the face of ischemia, would be desirable. Consequently, all published reports on the effects of anesthetics on myocardial high-energy phosphates have been related to either the negative inotropic effect of anesthetics or the relationship between anesthetics, ischemia and myocardial high-energy phosphates.

#### **Anesthetic Effects**

The paucity of published investigations of the effect of anesthetics on myocardial high-energy phosphates is certainly due to the difficulty in assay. Until recently (see Chapters 12-14), accurate measurement necessitated tissue excision and rapid freezing to avoid the breakdown of ATP and CP. In addition, the biochemical analysis was tedious and demanding. More recently, the availability of nuclear magnetic resonance (NMR) and positron emission tomography has enabled studies to be done noninvasively. However, again, the techniques are expensive and cumbersome.

The earliest published study of the effect of anesthetics on myocardial high-energy phosphates investigated the effect of halothane in cultured neonatal rat heart cells. Stong et al. (19) found no change in ATP pool size in their cultured heart cells under the influence of as high as a 5% concentration of halothane. However, they did document a decrease in the incorporation of inorganic phosphate ( $P_i$ ) into the ATP and suggested that halothane was decreasing the rate of synthesis of ATP as a possible mechanism for the negative inotropic effect. However, they admitted that the tight feedback control between energy supply and energy demand could also account for the decreased incorporation of  $P_i$  into ATP under the influence of halothane (13,20,21). The author, working together with the editor of this book, studied the effect of halothane on myocardial function and metabolism in intact open-chest swine (14). A marked depression in left ventricular function as indicated by more than 50% decrease in peak contractile element velocity and maximum velocity of shortening, and a 40% decrease in cardiac output and stroke volume, was accompanied by absolutely no change in tissue concentrations of ATP or CP. In a study using one of the new techniques, magnetic resonance spectroscopy, Murray et



Fig. 1. Cardiac energetics: TCA = tricarboxylic acid; A' $\Gamma P$  = adenosine triphosphate; CP = creatine phosphate; ADP = adenosine diphosphate; P<sub>i</sub> = inorganic phosphate; CPK = creatine kinase. Used with permission from ref. 13.

al. examined the effect of halothane on high-energy phosphates in a rabbit Langendorff preparation (15). In addition, they also examined intracellular pH and the creatine kinase reaction. Clinically relevant concentrations of halothane (0.016 mM) produced a significant decrease in left ventricular developed pressure, beta ATP and CP. However, 60 min of perfusion in a separate group of rat hearts without halothane also produced a significant decrease in high-energy phosphates. Although the authors reported a decrease in the forward rate constant for the transfer of phosphate from CP to ATP by halothane, no mention was made of whether that effect was also time dependent.

In a far superior study, McAuliffe and Hickey also examined the effect of halothane on high-energy phosphates in the heart (11). Inasmuch as they are pediatric anesthetists, they used neonatal rabbit hearts, prepared as a Langendorff preparation with an intraventricular balloon for measurement of performance and varying of preload. 1.5% halothane produced a 50% decrease in left ventricular pressure and dP/dt, but there was no change in ATP, CP,  $P_i$  or intracellular pH. In addition, in a separate series of hearts, they correlated measured oxygen consumption with varying oxygen demand induced by changing the volume of the left ventricular balloon. Both with and without halothane, there was a good correlation between oxygen demand and oxygen consumption. Consequently, they concluded that the decreased oxygen demand produced by halothane was matched by decreasing high-energy phosphate supplies and that there was no evidence of a primary effect of halothane on energy supply in the neonatal rabbit heart.

Preliminary studies from our laboratory at the University of Texas Medical School in Houston have also suggested that anesthetics do not produce an effect on myocardial high-energy phosphates. Using an isolated perfused working rat heart, we have previously shown that halothane produced a dose-related decrease in glucose utilization along with the negative inotropic effect (2). In further ongoing investigations, we have also measured the concentrations of ATP and CP under the influence of negative inotropic doses of both halothane (1.6 volumes %) and isoflurane (2.4 volumes %). These equipotent anesthetic concentrations depressed aortic pressure, cardiac output, coronary flow, and myocardial oxygen consumption (Table 1). Note that the effect of halothane was greater than that of isoflurane. However, even with the marked depression of cardiac function, there was no change in tissue ceoncentrations of ATP, CP or the calculated energy charge ([ATP] + 0.5 [ADP])/([ATP] + [ADP] + [AMP]) (Table 2).

It would appear, therefore, that the potent inhalation anesthetics, halothane and isoflurane, do not interfere in the maintenance of normal myocardial high-energy phosphate levels as a mechanism for their negative inotropic effect.

Variable	Control	1.6% H	2.4% I	
MAP (mmHg)	82 ± 17	$49 \pm 10^{*+}$	$61 \pm 13^{*+}$	
Cardiac output (ml/min)	$54 \pm 8$	$39 \pm 10^{*+}$	$50 \pm 10$	
Heart rate (beats/min)	$259 \pm 40$	$246 \pm 48$	$238 \pm 33$	
Coronary flow (ml/min)	$16 \pm 4$	$11 \pm 5^{*+}$	$14 \pm 5$	
$O_2$ consumption (min/g)	$8.2 \pm 0.6$	$3.6 \pm 0.3^*$	$6 \pm 10$	

Table 1. Effects of halothane and isoflurane on working rat heart

Means  $\pm$  S.D. MAP = mean arterial pressure; \*p<0.05 vs. Control; \*p<0.05 vs. 0.8% halothane (H) or 1.2% isoflurane (I).

#### Anesthetic Effects on the Ischemic Heart

The cardiac anesthesia/surgery laboratories from Duke University Medical Center investigated the interaction of potent inhalation anesthetics and myocardial ischemia. Halothane produced no change in high-energy phosphate levels either before or after ischemia in a working rat-heart preparation (16). Time to ischemic contracture was also unchanged in this preparation by halothane. On the other hand, in a Langendorff preparation, isoflurane increased tissue content of ATP before ischemia and at the time of ischemic contracture, although the time to ischemic contracture was not different (4). Following reperfusion with the Krebs-Henseleit buffer. functional recovery and metabolic restitution of ATP levels were greater in the hearts treated with isoflurance. In their last publication (and only full paper), the same group investigated the effects of enflurane in the Langendorff perfused rat heart (5). The effect was intermediate between isoflurane and halothane. There was no effect of enflurane on pre-ischemic CP or ATP concentrations nor on highenergy phosphates at the time of ischemic contracture. In addition, there was no prolongation of the time to ischemic contracture. However, following reperfusion, both recovery of peak pressure and tissue ATP and CP were improved in the animals treated with enflurane.

Another author in this volume investigated the effect of thiopental on myocardial tissue levels of ATP and CP during total ischemia (17). Ruigrok and co-workers showed that 100 mgL<sup>-1</sup> of thiopental significantly decreased myocardial concentrations of CP before ischemia and at various times during ischemia, and concentrations of ATP only during ischemia. They also examined creatine kinase release and the myocardial ultrastructure of hearts with and without thiopental treatment during hypoxia and low flow ischemia. Thiopental decreased creatine kinase release during hypoxia and during low-flow ischemia at a normal pH, but had no effect in total ischemia or low-flow ischemia at a low pH. The authors hypothesized that increased hydrogen ions produced by a low pH perfusate or total ischemia favored the entry of thiopental into the cell where the well-known barbiturate inhibition of mitochondrial function reduced ATP production. At normal pH, the cardiodepressant effect of the anesthetic decreased myocardial oxygen demand so that there was no net effect on high-energy phosphates.

From the laboratories of Kumazawa in Yamanashi Medical College Japan, came a series of investigations looking at the interaction between various anesthetics and ischemia in a rat-heart/lung preparation. Neither morphine (2 mgkg<sup>-1</sup>) nor fentanyl (100 ugkg<sup>-1</sup>) had an effect on myocardial ATP concentration (8). In a companion study, the same group examined the effect of fentanyl (1.2 ugml<sup>-1</sup>) and the same dose of fentanyl plus diazepam (3 ugml<sup>-1</sup>) in both an ischemic and non-ischemic heart/lung preparation (10). ATP concentrations were measured after 10 min of global ischemia and 20 min of reperfusion. There was no difference in the ATP concentrations between the control and the two anesthetic groups. Likewise, there were minimal differences in cardiac function although in the fentanyl group, recov-

Condition	Adenine triphosphate (umol/g protein)	Creatine phosphate (umol/g protein)	Energy charge
Control 1.6% Halothane 2.4% Isoflurane	23.8 ±0.2 19.3 ±2.0 25.6 ±1.2	$\begin{array}{rrrr} 21.1 & \pm 1.2 \\ 20 & \pm 2 \\ 21.9 & \pm 1.2 \end{array}$	$\begin{array}{rrr} 0.817 & \pm 0.005 \\ 0.76 & \pm 0.03 \\ 0.818 & \pm 0.009 \end{array}$

Table 2. No effect of halothane and isoflurane on high-energy phosphates in working rat heart

The energy charge is ([ATP] + 0.5[ADP])/([ATP] + [ADP] + [AMP]).

Anesthetics on Myocardial Adenine Nucleotides

ery from ischemia to a cardiac output of 30 ml·min<sup>-1</sup> and an aortic pressure of 80 mmHg was faster. In a similar preparation, the effects of two concentrations of thiopental (10 and 100 ug·ml<sup>-1</sup>) were investigated (7). Although there were no significant changes in ATP and CP concentrations produced by either dose of thiopental, the 100 ug·ml<sup>-1</sup> dose did decrease the concentrations of both high-energy phosphates and significantly increased concentrations of ADP and AMP with a resultant decrease in the myocardial energy charge (see above). In addition, recovery was delayed by this concentration of thiopental. Thus, their results agreed with those reported by Ruigrok et al. (17). Finally, the effects of halothane and enflurane in the same model were investigated (20). The animals were given 1% halothane and 2% enflurane from calibrated vaporizers to the excised heart/lung preparation. Neither anesthetic produced a change in the concentration of ATP, ADP or AMP compared to the control.

Another Japanese worker investigated the effect of thoracic epidural anesthesia in a basally anesthetized open-chest dog preparation before and after complete occlusion of the left anterior descending coronary artery (22). Although thoracic epidural anesthesia decreased the ischemia-produced elevation of the ST segment and increased myocardial pH, there were no changes in ATP or CP either before or after the coronary artery occlusion. Occlusion did markedly decrease the concentration of both ATP and CP in the control basally anesthetized animal and after thoracic epidural anesthesia.

#### **Summary and Conclusions**

Although the number of investigations has been rather limited, and predominantly has examined the effect of halothane, the results are consistent concerning the effect of the potent inhalation anesthetics on myocardial high-energy phosphates. In spite of marked depression of cardiac function produced by these compounds, no change in tissue level of ATP or CP has been ascribed to the pharmacologic effects of the anesthetics themselves. However, it is becoming apparent that measuring tissue homogenate concentrations of the high-energy phosphates may not accurately reflect pharmacologic effects. Several groups of investigators have recently demonstrated that glycolytically produced ATP and the ATP synthesized as a result of oxidative phosphorylation may subserve different functions (1,3,23,24). Glycolytic ATP appears to be necessary for the maintenance of electrical integrity of the cardiac sarcolemma and particularly governs the state of the potassium channel (24), while oxidatively based ATP may be more important for the major demands of the contractile proteins (23). Thus, analysis of total ATP and CP may not reveal a specific effect mediated through glycolysis. This may be relevant for anesthetic effects inasmuch as our laboratory has demonstrated a specific interference in glycolysis produced by halothane (2). Obviously, further investigation along these lines with anesthetics would appear to be fruitful.

The development of ischemic contracture has been used to quantitate the degree of ischemia. Bricknell et al. demonstrated that maintenance of glycolytically produced ATP was more important in delaying or preventing ischemic contracture than oxidative ATP (1). They speculated that the energy necessary for sarcoplasmic reticular  $Ca^{++}$  uptake might be preferentially derived from glycolysis, especially during ischemia. There is considerable ongoing controversy about the differential effects of the volatile anesthetics halothane, enflurane and isoflurane on the cardiac sarcoplasmic reticulum (18). These differences might explain the different effects of the three anesthetics on ischemia and ischemic contracture (4,5,16). However, considerably more studies are necessary to document and explain these effects.

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# Chapter 26

# **Regeneration of Adenine Nucleotides in the Heart**

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The fast adenine nucleotide degradation, which takes place in the ischemic heart, contrasts with the slow regeneration after ischemia. There are four pathways for adenine nucleotide regeneration:

1. De novo synthesis. This is a slow process. IMP is built from small molecules in a number of steps. The process can be accelerated to some extent by administering ribose. Ribose is incorporated in 5-phosphoribosyl-1-pyrophosphate (PRPP) and the amount of PRPP is rate-limiting. Administration of 5-amino-4-imidazolecarboxamide riboside (AICAriboside), which can be converted to an intermediate of this pathway, has also been tried for a faster regeneration, but this compound produced a deterioration of heart function. The conversion of IMP, end product of both de novo synthesis and salvage pathway, to AMP is very slow.

2. Salvage of hypoxanthine or inosine. Inosine has to be catabolized to hypoxanthine before it is incorporated. IMP is formed from hypoxanthine in one step, with subsequent conversion to AMP. PRPP is rate-limiting, so adding ribose enhances the incorporation rate. Inosine has hemodynamic effects.

3. Adenine ribophosphorylation. Adenine is incorporated in AMP in one step. PRPP is again rate-limiting. No nucleotide conversion is necessary. Toxic effects of long-term adenine administration have been described.

4. Adenosine phosphorylation. This is the fastest pathway for adenine nucleotide regeneration. Adenosine causes renal vasoconstriction together with systemic vasodilation (hypotension). These adverse reactions limit the practical value of adenosine.

Even though there is limited clinical experience with some of these compounds that enhance postischemic ATP pool repletion, none has been examined for this purpose in humans.

## Introduction

Ischemia leads to a rapid fall in the adenine nucleotide content of the heart. The rate of ATP decrease depends on factors like tissue perfusion (collaterals), heart function, temperature, species. The dephosphorylated breakdown products of ATP, the purine nucleosides and oxypurines, leak out of the myocardial cell. A simple scheme of ATP breakdown is shown in Fig. 1. The amount of adenine nucleotides will be roughly halved within the first hour of ischemia. A return of flow will cause a restoration of the adenine nucleotide levels (if the cells are not irreversibly damaged), but this takes days to accomplish.

The effects of ischemia on a cardiac myocyte are complex and their mechanisms have not fully been unravelled. There is a good correlation between the effect of myocardial protection during ischemia and the postischemic content of adenine nucleotides. Thus it is plausible that a fast restoration of ATP levels will promote the recovery of the postischemic heart. To regenerate ATP in the heart, four pathways are available, as was pointed out by Goldthwait (14) 30 years ago:

- 1. De novo synthesis;
- 2. Salvage of hypoxanthine or inosine;
- 3. Adenine ribophosphorylation;
- 4. Adenosine phosphorylation.

A scheme of these pathways is shown in Fig. 2.

#### **1. De Novo Synthesis**

This pathway involves the production of the purine skeleton from small molecules in 10 steps (see Fig. 3). The first step is the formation of 5-phosphoribosyl-1-amine (PRA) from 5-phosphoribosyl-1-pyrophosphate (PRPP) and an amino group donor like glutamine. The last step is the formation of IMP. There are no branching points between PRA and IMP. The conversion of IMP, end product of both de novo synthesis and salvage pathway, to AMP or GMP is slow.

Little is known about the value of enhanced guanine nucleotide regeneration. Adenylosuccinate synthetase catalyzes the reaction: IMP + aspartate + GTP  $\rightarrow$  adenylosuccinate + GDP + P<sub>i</sub>. The activity of this enzyme is very low (5). Adenylosuccinate



Fig. 1. Adenine nucleotide catabolism: ATP loses its phosphate groups. AMP can be converted to either adenosine or IMP. They are both broken down to inosine. This is catabolized to hypoxanthine, xanthine and urate.



Fig. 2. Pathways for adenine nucleotide regeneration: 1. De novo synthesis; 2. Salvage of hypoxanthine and inosine; 3. Adenine ribophosphorylation; 4. Adenosine phosphorylation. Pathways 1 to 3 use 5-phosphoribosyl-1-pyrophosphate (PRPP). Pi = inorganic phosphate, PPi = inorganic pyrophosphate, R-1-P = ribose-1-phosphate, S-AMP = adenylosuccinate.

lyase catalyzes the reaction: Adenylosuccinate  $\rightarrow$  AMP + fumarate; the same enzyme is also involved in one of the last steps of the de novo synthesis. Zimmer et al. (39) estimated that 0.04% of the nucleotide pool can be regenerated per hour this way. The rate of ATP synthesis is enhanced after ischemia (26), and limited by the concentration of PRPP (11). Ribose induces a considerable enhancement of the PRPP-concentration and of the rate of adenine nucleotide synthesis in heart (38), although the ribokinase activity in most tissues is low (2). There is some clinical experience with ribose. Zöllner et al. (40) described the successful symptomatic treatment of a patient suffering from an AMP-deaminase deficiency with high oral doses of ribose (maximum 60 g per day). Segal et al. (29) described a hypoglycemic effect of ribose. Human lymphocytes, exposed to extremely high ribose concentrations (25-50 mM), showed inhibition of their DNA repair synthesis (41).

Another possibility is to give 5-amino-4-imidazolecarboxamide riboside (AICAriboside), which can be phosphorylated to its ribotide (AICAR). This intermediate in the pathway becomes IMP in two steps. Swain et al. (31) showed that a 24-hour infusion of AICAriboside caused an accelerated repletion of ATP and GTP pools in postischemic canine myocardium. In an isolated cat heart, Mitsos et al. (27) described an



Fig. 3. De novo synthesis: Formation of IMP from small molecules in 10 steps. IMP is converted to AMP. AICAR = 5-amino-4-imidazolecarboxamide riboside, FH4 = tetrahydrofolate, PRA = 5-phosphoribosyl-1-amine, PRPP = 5-phosphoribosyl-1-pyrophosphate, S-AMP = adenylosuccinate.

increase in postischemic left ventricular developed pressure, but this was accompanied by a considerable increase in left ventricular compliance. In an open-chest dog model, Hoffmeister et al. (18) observed a marked impairment of regional function by postischemic AICAriboside. This deterioration progressed in time. We conclude that the value of accelerating de novo synthesis is limited.

#### 2. Salvage of Hypoxanthine or Inosine

Hypoxanthine can be utilized to restore the ATP content of the heart. The enzyme hypoxanthine guanine phosphoribosyltransferase catalyzes the reaction: Hypoxanthine + PRPP  $\rightarrow$  IMP + PP<sub>i</sub>. PRPP is rate-limiting (38). Harmsen et al. (16) found that adding ribose stimulates the hypoxanthine incorporation rate. (For a discussion about the conversion of IMP to the other nucleotides, we refer to the part on de novo synthesis.) Although the rate of adenine nucleotide production from this pathway is faster than the de novo synthesis, it is limited by the availability of PRPP and the activity of adenylosuccinate synthetase. Hypoxanthine enters the cell by simple diffusion. As a substrate for xanthine oxidase (XO), hypoxanthine may generate free oxygen radicals in species showing XO activity in the heart. Its presence in the human heart has to be clearly demonstrated. Addition of a xanthine oxidase inhibitor could be important.

Inosine has been described as a cardioprotective agent (1,13,32). Mammalian cells lack inosine kinase (35). Therefore inosine has to be broken down by nucleoside phosphorylase: Inosine +  $P_i \rightarrow$  hypoxanthine + ribose-1-phosphate, before it is incorporated in the adenine nucleotides. As nucleoside phosphorylase is not located in the cardiomyocyte (28), the ribose-1-phosphate can be used for PRPP production, but presumably not in the myocyte (See Fig. 4). It is possible that inosine incorporation in the myocyte can be enhanced by ribose addition, but as far as we know this has never been tried.



Fig. 4. Inosine has to be broken down to hypoxanthine within the endothelium before incorporation in adenine nucleotides of the myocytes. cl = capillary lumen, ec = endothelial cell, hx = hypoxanthine, ino = inosine, is = interstitial space, mc = myocyte, Pi = inorganic phosphate, PRPP = 5-phosphoribosyl-1-pyrophosphate.
In contrast to hypoxanthine, inosine does effect contractility, heart rate and blood flow. Juhász-Nagy and Aviado (22) found an increase in myocardial contractility in dog heart, while Hoffmeister et al. (17) observed a decrease in rat heart. There is conflicting evidence about the effect on heart rate (21,22). Coronary vasodilation has been reported by many authors (21,32). Inosine inhibits the uptake of adenosine in the heart (25), both nucleosides use the same carrier to enter the cell (36). Although inosine is used in Mediterranean countries as a cardiotonic, the effects of this nucleoside on humans have not extensively been investigated. Recently, De Jong et al. (7) described the apparent inosine uptake by the human heart together with an increase in dP/dt. We conclude that inosine and hypoxanthine are possibly valuable to the postischemic heart, but more research on this subject is necessary.

#### 3. Adenine Ribophosphorylation

Little research has been devoted to this pathway. The enzyme adenine phosphoribosyltransferase catalyzes the reaction: Adenine + PRPP -> AMP + PP<sub>i</sub>. The adenine concentration in blood is very low (24). If adenine is administered, the amount of PRPP becomes rate-limiting. Addition of ribose can enhance the incorporation rate. Ward et al. (33) gave adenine + ribose to a dog heart, in which the ATP levels had been halved. They repleted ATP in 24 hours, while it took 5 days without treatment. In isolated ventricular cells, Dow et al. (8) compared the incorporation rate of adenosine, of adenine + ribose, and of hypoxanthine + ribose. They showed that adenosine incorporation is twice as fast as adenine + ribose, and twenty times as fast as hypoxanthine + ribose is. Adenine lacks hemodynamic effects; it has no effect on adenosine transport (25). Clifford and Story (6) described growth disturbances in young animals fed a high adenine diet. Recently Yokozawa et al. (37) showed that metabolic abnormalities resembling chronic renal failure could be produced by long-term feeding of adenine to rats. Human experience with adenine is available from its use as an erythrocyte preservative in blood banks. We conclude that adenine could be a valuable compound, provided its safety for postischemic ATP regeneration can be proved.

### 4. Adenosine Phosphorylation

This pathway provides the fastest possibility to regenerate adenine nucleotides. The enzyme adenosine kinase catalyzes the reaction: Adenosine  $+ ATP \rightarrow AMP + ADP$ . Adenosine has been used for cardioprotection with varying success (9,15,18,19,20,23, 30). Gerlach et al. showed that adenosine incorporation is very fast in endothelial cells (12). Breakdown of adenosine in blood is very fast: a half life of 10 seconds has been estimated. It is possible that most of the infused adenosine will be either catabolized or incorporated in the endothelial cells, with little adenosine incorporated in the myocardial cells. Infused adenosine is also taken up very rapidly by other cells (especially in the lungs). Adenosine transport is mediated by a carrier. Dow et al. (8) reported that in isolated cardiac myocytes adenosine transport is rate-limiting for adenosine incorporation.

Adenosine exerts other effects mediated by purinergic receptors. It is a potent vasodilator and plays presumably an important role in coronary vasoregulation (4). Some groups described the slowing of atrioventricular impulse conductance and a decrease in heart rate (3,34). Foker et al. (10) reported that adenosine caused a vasoconstriction of the renal vessels. In humans, adenosine is used against arrhythmias (superventricular tachycardias). We conclude that adenosine is potentially useful although it has many side effects.

#### Conclusion

There is no clinical experience with enhanced ATP regeneration in the postischemic heart. For this purpose inosine, or adenine in combination with ribose, seem the most promising compounds.

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