Metabolomics

Citric Acid Cycle Intermediates in Cardioprotection

Gabor Czibik, MD, PhD;* Violetta Steeples, BSc;* Arash Yavari, BSc, MRCP, DPhil;*
Houman Ashrafian, MA, MRCP, DPhil*

Abstract—Over the last decade, there has been a concerted clinical effort to deliver on the laboratory promise that a variety of maneuvers can profoundly increase cardiac tolerance to ischemia and/or reduce additional damage consequent upon reperfusion. Here we will review the proximity of the metabolic approach to clinical practice. Specifically, we will focus on how the citric acid cycle is involved in cardioprotection. Inspired by cross-fertilization between fundamental cancer biology and cardiovascular medicine, a set of metabolic observations have identified novel metabolic pathways, easily manipulable in man, which can harness metabolism to robustly combat ischemia-reperfusion injury. (Circ Cardiovasc Genet. 2014;7:711-719.)

Despite a dramatic drop in the incidence of acute myocardial infarction over recent decades,1 coronary artery disease (CAD) is still the foremost cause of death globally, with >7 million people dying annually.2 In developed countries, reductions in exposure to major risk factors and the application of evidence-based interventions to patients with CAD have shifted the epidemiology of CAD toward older age.3,4 There are concerns that this positive trend may be obviated by the pandemic of obesity and type 2 diabetes mellitus. Advances in cardiac surgery have led to its application to an increasingly older subset of patients, with a correspondingly greater risk of myocardial injury. Sensitive means of detecting myocardial injury have broadened the at-risk population to include those undergoing major noncardiac surgical procedures (eg, vascular or major abdominal surgery), particularly in the elderly.5 Much of the effort to reduce CAD-related myocardial injury has focussed on reducing the burden of coronary atheroma and plaque stability. The delineation of robustly protective alleles reducing the burden of atherosclerosis (eg, PCSK96 and APOC37,8) has signposted potential drug targets where.13–15 Because patients with CAD are often elderly with numerous other comorbidities and confounding medical therapies,14 the protective potential of IPC may be compromised, motivating the exploration of new strategies to reduce ischemic cardiac injury and the associated healthcare and societal costs.

One such approach has been the application of metabolic strategies to modify the time course of myocardial injury. In the 1960s, preceding the pharmacological or mechanical reperfusion era, Sodi-Pallares proposed using a solution containing glucose, insulin, and potassium to prevent reperfusion arrhythmias.16 The rationale was to support myocardial energy generation through augmentation of ischemic glycolytic activity using pharmacological doses of glucose and insulin.17 This combination reduces circulating free fatty acids that potentially inhibit glycolysis18 (mitigating the Randle cycle via insulin’s inhibitory effect on lipolysis) and has been proposed to prevent arrhythmias through the restoration of intracellular potassium and stabilisation of membrane potential.16 The majority of experimental studies applying glucose, insulin, and potassium treatment have reported reduced infarct size19 with improved postischemic functional20 and energetic recovery,21 with subsequent successful translation to humans.22,23 Encouraged by his success and that of the first Diabetes Mellitus Insulin-Glucose Infusion in Acute Myocardial Infarction (DIGAMI) trial,24 the method gained popularity to reduce IR-injury after...
Traditional Functions of the CAC

The CAC, either in its complete or in its modified forms, is a metabolic hub in all aerobic and some anaerobic organisms (eg, green sulfur bacteria). The CAC has 2 principal purposes: (1) to harness intermediary metabolism to generate ATP (through oxidative phosphorylation and substrate-level phosphorylation not requiring a terminal electron acceptor, such as oxygen, eg, in hypoxia) and (2) to provide precursors for biosynthetic pathways. As traditionally envisaged (eg, in most biochemistry textbooks), the CAC begins with the condensation of oxaloacetate with acetyl-CoA, consisting of 8 sequential linear irreversible (eg, citrate synthase and the 2-oxoglutarate dehydrogenase complex) and reversible steps, ending with the regeneration of oxaloacetate (Figure 1A). These CAC steps are generally considered to take place predominantly in the mitochondrial matrix.

Acetyl-CoA derived from glycolysis (via pyruvate dehydrogenase), from β-oxidation of fatty acids, and from the breakdown of ketone bodies and amino acids, generate the reducing equivalents NADH and FADH₂ (reduced nicotinamide adenine dinucleotide and reduced flavin adenine dinucleotide, respectively), which, via the electron transport chain, synthesise ATP by oxidative phosphorylation. In addition there is a comparatively small but significant matrix substrate-level energy generation (guanosine triphosphate, GTP, convertible to ATP) not requiring oxygen.

However, the CAC is more than a highly efficient sequence of spatially optimized reactions to degrade acetyl-CoA. Short sections of the CAC are close to thermodynamic equilibrium and are, therefore, bidirectional under different conditions. Krebs emphasized this reversibility by observing (albeit with relatively nonspecific inhibitors, such as malonic acid) that the enzymes interconverting succinate ↔ fumarate ↔ malate ↔

![Figure 1. The citric acid cycle (CAC). A. An overview of CAC reactions. Observe that apart from 3 irreversible steps (citrate synthase, isocitrate dehydrogenase, and 2-oxoglutarate dehydrogenase), the rest of the CAC reactions are reversible, allowing reverse flux under certain conditions (such as in anoxia). B. Anaplerosis and cataplerosis of the CAC. Anaplerosis replenishes CAC intermediates, whereas cataplerosis allows production of various biosynthetic precursors making the CAC as a central metabolic hub, connecting diverse metabolic pathways. CoA-SH indicates reduced coenzyme A; FADH₂, reduced flavin adenine dinucleotide; GTP, guanosine triphosphate; NAD⁺, oxidized nicotinamide adenine dinucleotide; NADH, reduced nicotinamide adenine dinucleotide; and QH₂, reduced coenzyme Q.](http://circgenetics.ahajournals.org/ by guest on November 2, 2017)
oxaloacetate are reversible. This reversibility facilitates the generation of precursors for glucose synthesis (from oxaloacetate via the pyruvate recycling pathway), fatty acid and cholesterol synthesis (from citrate), amino acid anabolism, nucleotides (via 2-oxoglutarate [2-OG]), and heme biosynthesis (via succinyl-CoA). In contrast to its perceived role as a perpetual rotor degrading acetyl-CoA, the CAC represents a highly versatile collection of reactions that can be concatenated in its entirety or in part to generate energy (oxidatively or without oxygen) and as a biosynthetic toolbox.

Exemplifying the versatility of the CAC as a whole and individual reaction steps per se, dominant mutations in 2 of the 3 isocitrate dehydrogenases (IDH1 and IDH2) have been found to promote glioma, acute myeloid leukemia, chondrosarcoma, and other tumours. In the traditional conception, acetyl-CoA is thought to progress unidirectionally through the CAC. Instead, IDH1 (primarily cytoplasmic and NADP+ dependant) and IDH2 (primarily mitochondrial and NADP+ dependant) along with IDH3 promote the conventional oxidative decarboxylation of isocitrate to 2-OG but may also promote the reverse (reductive carboxylation). Reductively metabolized glutamine has been proposed to be a cellular carbon source for fatty acid synthesis during hypoxia or impaired mitochondrial respiration (eg, FH deficiency). Although these observations, adduced from ex vivo cancer cell preparations, remain controversial and potentially confounded by technical aspects of metabolic tracer studies, there is little doubt that mutated IDH isoforms have an abnormal flux through CAC intermediates. The same flexibility provides potential therapeutic opportunities to mitigate hypoxia/ischemia.

Adequate anaplerosis is especially important because the steady state concentrations of CAC intermediates are typically low in relation to the fluxes through the CAC. Breakdown of amino acids replenish 2-OG, succinyl-CoA, and oxaloacetate; odd-chain fatty acid oxidation generates succinyl-CoA; carboxylation of pyruvate by pyruvate carboxylase and malic enzyme produces oxaloacetate and malate, respectively. (Figure 1B).

To maintain its activity and to compensate for metabolite leakage from the mitochondria, CAC flux can only be maintained if this cataplerotic activity constituting the removal of intermediates from the CAC is balanced by anaplerotic reactions replenishing CAC intermediates (Figure 1B). Breakdown of amino acids replenish 2-OG, succinyl-CoA, and oxaloacetate; odd-chain fatty acid oxidation generates succinyl-CoA; carboxylation of pyruvate by pyruvate carboxylase and malic enzyme produces oxaloacetate and malate, respectively. (Figure 1B). Adequate anaplerosis is especially important because the steady state concentrations of CAC intermediates are typically low in relation to the fluxes through the CAC. These uneven CAC intermediates are explained, in part, by the differential turnover rate of individual metabolic reactions, the differing thermodynamic/kinetic properties of each CAC step and differential mitochondrial inner membrane permeability. The measurement of both concentration and flux is technically challenging, even with dynamic tracer studies (for flux). For example, citrate, with the largest pool size (≈200 nmol/g), has a turnover time of minutes, whereas oxaloacetate with the smallest pool size (5–10 nmol/g) has a turnover of the order of 100x per minute. Consonant with metabolic control analysis, in the CAC, flux considerations dominate over steady-state thermodynamic principles per se.

To maintain locally high metabolite concentrations, CAC reactions are compartmentalized within the mitochondria. There may be further spatial subcompartmentalization in the vicinity of individual CAC enzymes to maximize local effective substrate concentrations and reaction efficiency. Specialized carrier proteins catalyze the transport of nucleotides, amino acids, inorganic ions, fatty acids, ketone bodies, and cofactors across the impermeable mitochondrial inner membrane. These transport steps are important to maintain a separate inner mitochondrial pool and micromilieu for ATP production, for amino acid breakdown, for macromolecular and heme biosynthesis, and for heat generation. An example of such a mitochondrial carrier mechanism is the malate-aspartate shuttle (MAS). MAS is a key metabolic shuttle, active in the heart, which transfers electrons and protons from glycolysis-derived cytosolic NADH (to which the inner mitochondrial membrane is impenetrable) into the matrix for use by the electron transport chain using malate as an electron carrier. This involves a series of reactions involving the cytosolic reduction of oxaloacetate to malate, transport of malate into the mitochondrial matrix (in exchange for 2-OG), oxidation of malate to regenerate oxaloacetate with reduction of mitochondrial NAD+ to NADH. Oxaloacetate then undergoes transamination using glutamate as nitrogen donor to generate aspartate (and 2-OG), which exits the matrix via a glutamate–aspartate transporter (Figure 2). Although the influence of MAS in the heart in ischemia is complex and requires further clarification, its manipulation is likely to be important in ischemia-reperfusion injury.

The MAS is the most important although not the only means to transport net protons and electrons from the cytoplasm to the mitochondrial matrix for oxidation. Like the MAS, the glycerol-3-phosphate shuttle helps reoxidize cytosolic NADH to NAD+, making it available for glycolysis. Should this turnover fail, accumulating cytosolic NADH would act as a brake to glycolysis in the face of a diminishing reducible NAD+ pool. Reducing equivalents generated during glycolysis and more substantially by the CAC are converted to ATP through the transfer of electrons to molecular O2 by the electron transport chain. Any lack of oxygen increases the NADH/NAD+ ratio, substrate inhibits glycolysis/CAC, and substantially disrupts...
energy production. Accordingly, during hypoxia, anaerobic regeneration of NAD\(^+\) continues to occur, in part, through reoxidation of NADH by lactate dehydrogenase, which catalyzes electron transfer to pyruvate, rather than O\(_2\), to yield lactate. As oxidative phosphorylation and the closely coupled CAC flux are arrested, NADH accumulates. In the absence of molecular oxygen, even glycolytic acceleration may be inhibited by incomplete myocardial NADH reoxidation limiting the regeneration of NAD\(^+\) essential for continued glycolysis. As opposed to skeletal muscle, lactate dehydrogenase in the heart is predominantly the H-type, with increased susceptibility to form the abortive ternary complex lactate dehydrogenase-NAD-enol that competes with NADH.\(^42\) Thus, without alternative means to deplete NADH, the hypoxic metabolism would slow because of redox imbalance (increased NADH). As a corollary, modalities promoting NAD\(^+\) regeneration would potentially improve energetics.\(^43,44\)

**Noncanonical Functions of the CAC: Succinate Production in Hypoxia**

Ischemic cell death likely results from a rapid decline in intracellular ATP concentration, [ATP]\(_{ic}\), that is irreconcilable with viability.\(^45\) Maintenance of adequate cellular energy production is thus a fundamental and continual requirement for all tissues, with its disruption leading to significant clinical consequences. Metabolic studies in isolated hearts and muscle preparations and in natural states of extreme physiology (eg, diving mammals) have yielded important insights into cellular mechanisms, specifically emphasizing the CAC components, which maintain [ATP]\(_{ic}\) and have broadened our understanding of the CAC.

For decades, it has been recognized that hypoxia/ischemia confers substantial changes in CAC metabolites, most prominently succinate. For example, in 1970, while investigating the ability of CAC metabolites to improve cardiac performance in anoxia, Penney and Cascarano\(^46\) observed that perfusion of anoxic rat hearts with glucose and a combination of fumarate+malate+glutamate or oxaloacetate+2-OG (2-oxoglutarate) enhanced glycolysis and ATP levels over glucose perfusion alone, without increased glycolysis. In both CAC metabolite treatment groups, succinate levels were found consistently elevated. The authors concluded that mitochondrial energy production took place even during anoxia; however, the precise nature of this mitochondrial contribution to anoxic energy generation was undefined.\(^46\) Similarly, glutamate and aspartate were recognized to lead to increased succinate levels in skeletal muscle before the discovery of the CAC itself.\(^47\)

The importance of this mechanism was implied by its evolutionary conservation. Hochachka et al\(^48\) observed that in diving mammals, blood concentrations of aspartate and 2-OG (2-oxoglutarate) levels decreased, whereas those of lactate, alanine, and succinate increased after diving (a hypoxic manoeuvre). They proposed a model of 2 interlinked, oxygen-independent pathways (Figure 3),\(^48\) whereby NADH produced during glycolysis and through anterograde carbon flow through 2-OG dehydrogenase generating GTP and succinate is coupled to NADH depletion through reducing aspartate-derived oxaloacetate to malate. The malate is converted to fumarate and ultimately succinate contrasting with the traditional conception of carbon flow in the CAC (because in the context of lacking oxygen normal CAC flux ceases).\(^48\) Other mechanisms contributing to NADH depletion may include mitochondrial diaphorases.\(^49\) Increased succinate production has also been observed in hypoxic cardiac myocytes.\(^27\)

Although there are controversies about the pertinence of this model to IR injury,\(^43,50-53\) at least under some circumstances, loading of the CAC using glutamate, fumarate, and aspartate have generally been successful but may not be clinically tractable. This strategy is also not without jeopardy—the energetic benefit of channelling amino acids to succinate may be mitigated by the release of ROS at reperfusion resulting from a burst of FADH\(_2\) in the context of too sudden forward flow of succinate through succinate dehydrogenase (complex II).

**Therapies Modifying CAC Intermediates as Potential Therapies of IR**

Recognizing the potential value of modifying, specifically augmenting, CAC intermediates, several strategies have been investigated. One readily translatable is the indirect augmentation of CAC intermediates using glutamate/glutamine.\(^54\)
Glutamate/glutamine has exhibited variable promise in animal and human models of myocardial IR, especially as a constituent of cardioplegia in coronary surgery. Another approach is to augment CAC intermediates using anaplerotic precursors directly. Despite successfully increasing myocardial CAC intermediates, dipropionylcysteine ethyl ester and heptanoate failed to protect the myocardium, in part, perhaps because of the differing protocols of drug infusion and myocardial ischemia. In contrast, dippyruvyl-acetyl-glycerol decreases myocardial infarct size in the pig. This is consistent with the observation that fumarate-enriched cardioplegia results in complete functional recovery of immature myocardium.

**Signaling Roles of the CAC: Insights From Renal Tumours**

Notwithstanding the controversies surrounding the energetic merits of succinate generation through mitochondrial substrate-level phosphorylation, increased succinate production has been consistently identified in different tissues, in different models in hypoxia/IR. In addition to its metabolic role, elevated succinate seen ubiquitously in ischemia is likely to activate its cognate cell surface receptor—G-protein–coupled receptor 91 (Gpr91). Succinate acting through this receptor contributes to retinal neovascularization and acting through prostaglandin E receptor 2–prostaglandin E receptor 4 modifies post–hypoxia-ischemic injury in the brain, emphasizing the pleotropic contributions of CAC metabolites to IR injury.

The discovery of normoxic HIF-1α stabilization by elevated fumarate levels in hereditary HLRCC and elevated succinate in hereditary paraganglioma with pheochromocytomas resulting from deficient FH and SDH (succinate dehydrogenase) respectively, potentially indicates another signaling role for CAC metabolites in the response to IR. The mechanism underlying normoxic HIF-1α stabilization in these tumors is thought to be substrate competition of the accumulating CAC metabolites with 2-OG. The latter is a key cofactor for prolyl hydroxylase domain 1 to 3 proteins that regulate HIF-1α by hydroxylation under normoxic conditions, promoting its proteosomal destruction (Figure 4A).

This mechanism is supported by the finding that addition of excess 2-OG is able to out-compete fumarate, thereby reactivating prolyl hydroxylase domains and hydroxylating HIF-1α. The possibility that different metabolic states—signaled by varying levels of CAC intermediates—can regulate the central transcriptional coordinator of the cellular response to hypoxia, resonates with the observation that CAC metabolites are channelled toward succinate during ischemia and has important consequences beyond cancer, in other states of hypoxia/ischemia. Analogous HIF-1α augmentation by succinate may be germane to cardiac ischemia because mice overexpressing HIF-1α in the heart, when subjected to hypoxia, show signs of improved cardiac function.

---

**Figure 4.** Signaling functions of citric acid cycle (CAC) intermediates. **A**, The hypoxia-inducible factor (HIF)-1α-hypoxia pathway. The key in regulation of HIF-1α stability is an O2- and 2-oxoglutarate-dependent proline hydroxylation. When in excess, fumarate and succinate can out-compete 2-oxoglutarate and can stabilize HIF-1α protein in normoxia. **B**, The Nrf2-antioxidant pathway. The critical event in Nrf2 activation is when specific cysteine residues on Keap1 react with electrophiles, such as fumarate, and consequently Keap1 releases Nrf2. Keap1 indicates Kelch-like ECH-associated protein 1; Nrf2, nuclear factor (erythroid-derived 2)-like 2; PHD, prolyl hydroxylase domain; and VHL, von Hippel-Lindau tumor suppressor.
coronary artery ligation, exhibited a reduction in infarct size and improved postschismic recovery.69 Succinate-augmented HIF-1α levels may contribute to IPC because mice with reduced HIF-1α cannot be preconditioned.70,71

**Fumarate Is Cardioprotective via Activation of the Nrf2-Antioxidant Pathway**

As with succinate, the cardioprotective influence of fumarate has been recognized in both immature62 and adult hearts;72 however, it is likely that the contribution of fumarate to the coupled reductive pathway originally proposed by Hochachka et al68 would not suffice to account for the cardioprotective effect completely.73 Furthermore, the cardioprotective influence observed with exogenous nonesterified fumarate in adult rat hearts was relatively modest,72 raising the possibility of inadequate intracellular penetration of fumarate from the perfusate.72 Analogous to its role in normoxic HIF-1α activation, we wondered whether CAC metabolites, such as fumarate, may activate other cytoprotective transcriptional programmes.

To augment myocardial fumarate levels reliably, we generated a model of cardiac FH deficiency by crossing mice bearing LoxP sequences flanking exons 2 to 3 of the Fhl1 gene with mice carrying Cre recombinase under the ventricular cardiomyocyte-specific McI2v promoter. By interrupting the conversion of fumarate to malate, the resulting mice represented a genetic model of augmented cardiac fumarate concentrations. Mice with cardiac Fhl1 deficiency were viable and healthy (despite a disrupted CAC) until ≈3 to 4 months.72 The phosphocreatine to ATP ratio (a readout of energetics) was unaltered when compared with control hearts consistent with their viability. Successful Fhl1 knockout in the cardiac myocytes was confirmed at both mRNA and protein level. These hearts exhibited increased fumarate levels when compared with control (Fhl1/fl/fl, no Cre) hearts. The increases were significant and consistent though modest in magnitude (×1.6 fold of controls in total heart preparations).

When subject to ex vivo IR (40 minutes of global ischemia followed by 60 minutes of reperfusion), Fhl1 knockout hearts exhibited a marked reduction in mean infarct size from 37% in control hearts to 17% in Fhl1 knockout hearts. Assessment of markers of myocardial injury corroborated the robust cardioprotection in Fhl1 knockout hearts. A time-course of interstitial metabolite profiling detected reduced ischemic release of succinate, glutamate, and adenosine in Fhl1 knockout hearts, a pattern that has been associated with attenuated cardiac injury.73,74

In other models of fumarate augmentation, either using FH-deficient tumors or increased cellular fumarate through the application of cell permeable fumarate analogues, activation of the Keap1-Nrf2-antioxidant pathway has been observed (Figure 4B).31,75 Nrf2 is a transcription factor that coordinates the defense against oxidative stress by activating antioxidant response element genes.76 In unstimulated cells, Nrf2 is sequestered in the cytoplasm by Keap1, promoting Nrf2’s proteosomal destruction.77 On cellular overload with ROS or electrophiles, reactive cysteine residues on Keap1 are modified, resulting in release of Nrf2, its translocation to the nucleus, and heterodimerization with small Maf proteins to transactivate antioxidant response element genes.77 By acting as an electrophile via its unsaturated bond, fumarate (but not succinate which is saturated) binds to the reactive cysteine residues of Keap1, forming an irreversible chemical modification of S-(2-succinyl)-cysteine (2SC), termed succination.73 Proteomic assessment of FH-deficient murine embryonic fibroblasts found a variety of proteins altered by this novel post-translational modification, many of which had metabolic functions and were mitochondrial in location.79 Although the functionality of succination remains generally undefined, the effect of succinated cysteine residues binding an obligatory \([\text{Fe}_2\text{S}_2]\)^2+ cluster in the CAC enzyme aconitase 2 has been shown to be a dose-dependent inhibition of aconitase activity.79 The iron–sulfur cluster of aconitase 2 is also highly sensitive to oxidation by superoxide, acting as a mitochondrial sensor of oxidative stress.80

Systematic expression profiling of our FH-deficient hearts, consistent with the cancer literature,31,75 supported the observation that fumarate augmentation upregulated several canonical Nrf2 target and coregulated genes, including glutathione S-transferases, methylcytenetetrahydrofolate dehydrogenase, NAD(P)H:quinone oxidoreductase 1 and, perhaps importantly, heme oxygenase 1 (Hmox1). Examination of succination in Fhl1 knockout hearts with an anti-2SC antibody revealed substantially increased succination. This supports the hypothesis that succination of Keap1 may be the likely mechanism for the observed reductions in Keap1 protein and the enhanced stabilization/translocation of Nrf2.

Substantial evidence supports the hypothesis that Hmox1 activation is robustly cardioprotective. This inducible cytoprotective enzyme catabolizes the pro-oxidant heme to free iron, carbon monoxide (CO) and biliverdin, with the latter then converted to bilirubin.81 Bilirubin itself is a potent antioxidant, whereas carbon monoxide is a vasodilator with additional anti-inflammatory and antiapoptotic properties.81,82 Both CO and bilirubin protect against cardiac IR injury; the former via activation of p38 MAPK and Akt kinases,24 the latter likely via an antioxidant mechanism.84 Although mice heterozygous for Hmox1 were found to be susceptible to IR injury,65 cardiac-restricted overexpression of Hmox1 has been shown to attenuate infarct size both ex vivo and in vivo, with reduced oxidative stress, inflammatory cell infiltration,85 left ventricular dysfunction, and apoptosis.86

Upregulation of Hmox1, coupled with heme resulting from the cataplerosis of succinyl-CoA predicted by our own computational modeling53 and that of others,67 raised the possibility of Hmox1 contributing to the cardioprotection. To address this, we injected control and Fhl1 knockout animals with vehicle or zinc deuteropopyrin 2-4-bis ethylenglycol (ZnBG), a specific heme oxygenase inhibitor86,87 before ex vivo IR. Pretreatment with ZnBG had no effect on infarct size in control hearts, but negated the cardioprotection observed in Fhl1 knockout hearts.

To assess whether our findings in Fhl1 knockout mice were translatable, we used dimethyl fumarate (DMF), an orally available and cell-permeable version of fumarate67 already in clinical use to treat patients with psoriasis and multiple sclerosis.76 Treatment of atrial cardiomyocyte-derived HL-1 cells with DMF induced nuclear translocation of Nrf2. When
administered orally to wild-type mice, dimethylfumarate recapitulated much of the Nrf2 target profiling observed in untreated Fh1 knockout hearts. When subject to IR, hearts from DMF-pretreated wild-type mice displayed a ∼75% reduction in infarct size to 9.3% when compared with 36.9% in vehicle pretreated wild-type hearts, accompanied by an improvement in postischemic coronary flow. To determine whether DMF’s infarct-sparing effect translated to an in vivo model, wild-type mice pretreated with oral DMF or vehicle underwent left anterior descending coronary artery ligation with subsequent evaluation of infarct size. A significant reduction in infarct size was observed in the DMF-treated group. This protective effect of DMF was absent in Nrf2 knockout mice, indicating the critical role played by Nrf2 in mediating DMF-conferred cardioprotection.32

These findings do not preclude the possibility that the cardioprotective potential of fumarate is also dependent on other pathways. Motivated by findings in the cancer literature, we examined whether increased fumarate was sufficient to activate the HIF hypoxia pathway but found no evidence to support this.89 There are several possibilities for the apparent discrepancy: (1) Fumarate levels of Fh1 knockout murine embryonic fibroblasts are 100-fold higher than in wild-type murine embryonic fibroblasts,79 suggesting that an estimated 2.2-fold increase in fumarate observed in Fh1 knockout cardiac myocytes (1.6-fold in whole hearts) might be insufficient to out-compete 2-OG; (2) prolyl hydroxylase domain 2, the major HIF-1α hydroxylase,36 has a much higher Kd for fumarate than for 2-OG; accordingly, fumarate levels have to exceed 2-OG greatly for binding to prolyl hydroxylase domain 2; (3) as an alternative mechanism to substrate competition, Sullivan et al91 proposed that elevated levels of fumarate resulted in increased succination and inactivation of the antioxidant glutathione to form GSF, which—by acting as an inhibitory substrate to glutathione reductase—enhances mitochondrial ROS levels, serving to stabilize HIF-1α. In Fh1 knockout cardiomyocytes, the modestly elevated fumarate is unlikely to have altered superoxide levels directly. Notwithstanding these considerations, endogenous antioxidant protection via activation of the Nrf2-dependent arsenal of cytoprotective genes is desirable in the setting of reperfusion injury, where oxidative stress is recognized as a major pathogenic contributor. The preclinical evidence for cardioprotection using the orally available, cell-permeable, and clinically safe DMF provides a tangible agent for clinical testing in the setting of acute coronary syndromes or predictable myocardial injury (eg, cardiac surgery).5,32

Conclusions and Future Directions

As outlined above, there is substantial evidence, accumulating over almost a century, that CAC intermediates change dramatically during the course of IR injury. The recognition that these changes, most prominently succinate production, may be beneficial has been substantially enhanced by the recognition that succinate and related metabolites have additional and perhaps more potent effects than ATP generation. These include cell surface receptor activation (eg, Gpr91), HIF-1α activation, and Nrf2 liberation from Keap1. However, relatively simple strategies to exploit the metabolic benefits of CAC intermediate (eg, modification through the provision of feedstock amino acids) have proved clinically challenging with variable efficacy. Alternative metabolic strategies using similar CAC-related products that will augment Hmox1 activity (either through increasing its substrate haem or increasing enzymatic activity) may be germane and are the subject of investigation.

Sources of Funding

This work was supported by the British Heart Foundation. A.Yavari is supported by the UK Department of Health’s National Institute for Health Research.

Disclosures

H. Ashrafian has received an unrestricted research grant from SBI-ALA, Japan. The other authors report no conflicts.

References


Circulation


Berra E, Benizri E, Ginouvès A, Volmat V, Roux D, Pouysségur J. HIF prolyl-hydroxylase 2 is the key oxygen sensor setting low steady-state levels of HIF-1alpha in normoxia. EMBO J. 2003;22:4082–4090.


KEY WORDS: cardioprotection ■ citric acid cycle ■ fumaric acid ■ oxidation-reduction ■ nuclear factor (erythroid-derived 2)-like 2 ■ reperfusion injury
Citric Acid Cycle Intermediates in Cardioprotection
Gabor Czibik, Violetta Steeples, Arash Yavari and Houman Ashrafian

doi: 10.1161/CIRCGENETICS.114.000220
Circulation: Cardiovascular Genetics is published by the American Heart Association, 7272 Greenville Avenue, Dallas, TX 75231
Copyright © 2014 American Heart Association, Inc. All rights reserved.
Print ISSN: 1942-325X. Online ISSN: 1942-3268

The online version of this article, along with updated information and services, is located on the World Wide Web at:
http://circgenetics.ahajournals.org/content/7/5/711

Permissions: Requests for permissions to reproduce figures, tables, or portions of articles originally published in Circulation: Cardiovascular Genetics can be obtained via RightsLink, a service of the Copyright Clearance Center, not the Editorial Office. Once the online version of the published article for which permission is being requested is located, click Request Permissions in the middle column of the Web page under Services. Further information about this process is available in the Permissions and Rights Question and Answer document.

Reprints: Information about reprints can be found online at:
http://www.lww.com/reprints

Subscriptions: Information about subscribing to Circulation: Cardiovascular Genetics is online at:
http://circgenetics.ahajournals.org/subscriptions/