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PERSPECTIVES

Rethinking Mitchell's Chemiosmotic Theory: Potassium Dominates Over Proton Flux to Drive Mitochondrial F₁F₀-ATP Synthase

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A Perspective on "ATP synthase K^+ - and H^+ -fluxes drive ATP synthesis and enable mitochondrial K^+ -"uniporter" function: I. Characterization of ion fluxes" & "ATP synthase K^+ - and H^+ -fluxes drive ATP synthesis and enable mitochondrial K^+ -"uniporter" function: II. Ion and synthase flux regulation"

Mitochondria are the dominant source of energy in the form of adenosine triphosphate (ATP) in most cells. In the mitochondrial matrix, the Krebs cycle is fueled by nutrients to reduce nicotinamide- (NADH) and flavin adenine dinucleotide (FADH₂)¹, which donate electrons to the respiratory chain (Figure 1). The ensuing electron transfer along complexes I-IV of the chain and onto oxygen (O_2) provides the energy to pump protons (H^+) from the matrix to the intermembrane space, generating a chemical (ΔpH) and an electrical potential ($\Delta \Psi_m$) across the inner mitochondrial membrane (IMM), which together constitute the protonmotive force ($\Delta \mu_{\rm H}$). According to the chemiosmotic theory developed by Peter D. Mitchell, $\Delta \mu_{\rm H}$ is the driving force for oxidative phosphorylation of adenosine diphosphate (ADP) to ATP at the F_1F_0 -ATP synthase (Figure 1)². This concept, for which Mitchell was awarded the Nobel Prize for Chemistry in 1978, has been accepted for more than 50 years and can be found in literally every textbook of biology.

In the current issue of Function, Juhaszova and colleagues³ substantially challenge—or rather expand, but do not tumble—this concept in revealing that in addition to H⁺, potassium ion (K⁺) flux through the F₁F₀-ATP synthase (working the same way as H⁺) provides the majority of energy to produce ATP (Figure 1). Why was this was overlooked for more than six decades? Presumably because the F₁F₀-ATP synthase has a > 10⁷-fold selectivity for H⁺ over other cations.⁴ But what was not sufficiently considered is that due to the 10⁶-fold higher cytosolic

concentration for K⁺ (~100 mM) than for H⁺ (~100 nM), such that K⁺ flux—driven mostly by the same high electrical driving force ($\Delta \Psi_m$) - could be comparable to H⁺ flux via the ATP synthase.

Employing a variety of experimental systems, including proteoliposomes containing purified mammalian F1F0-ATP synthase, planar lipid membranes, but also intact rat cardiac mitochondria, Juhaszova et al.³ elegantly demonstrate that for each H⁺, 2.7 K⁺ ions are transferred at the ATP synthase under physiological conditions. Since contraction of intramitochondrial volume hinders the activity of the respiratory chain, and K⁺ influx osmotically allows water to expand the matrix, such two-ion flux through the F₁F₀-ATP synthase not only increases ATP synthesis, but also improves its efficiency: Compared to H⁺ flux, K⁺ flux exhibited a 3.5-fold higher ATP synthesis, but only a 2.6-fold higher O₂ consumption rate.³ Although a "two-ion theory of energy coupling" was proposed previously by Nath,18 the models differ substantially: while Nath proposed a H⁺/K⁺ antiport within the F1F0-ATP synthase may maintain electroneutrality,¹⁸ the model presented here³ defines a H⁺/K⁺ symport via the ATP synthase, where K⁺ extrusion is accounted for by the distinct K⁺/H⁺ exchanger (KHE; Figure 1). Importantly, this novel concept, which allows an optimized matching of energy supply to demand, was corroborated by a minimal computational model comprising the "core" mechanism constituted by ATP synthase, driven by both H⁺- and K⁺-motive force, respiratory chain, adenine nucleotide translocator, phosphate carrier, and the K⁺/H⁺ exchanger in a parallel study published elsewhere.5

As if this discovery was not enough of a scientific earthquake, in a second manuscript, the same authors⁶ uncover that by this previously unrecognized K^+ flux, the F_1F_0 -ATP synthase

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Figure 1. The Janus-faced mitochondrial F1F0-ATP synthase as the master regulator of life and death. The electron transport chain (ETC) receives electrons from NADH and FADH₂ to translocate protons (H⁺) across the inner mitochondrial membran (IMM) to provide the driving force for the F1F0-ATP synthase to produce ATP. The current studies 3,6 suggest that in addition to the H+ motive force ($\Delta\mu_{\rm H})$, the even greater K⁺-motive force ($\Delta \mu_{\rm K}$) is harnessed to drive ATP production at the ATPase. Through its impact on mitochondrial volume, this optimizes ATP production during increased ATP demand. Pathological concentrations of Ca²⁺ and/or reactive oxygen species (ROS) trigger cyclophilin D (CyPD) binding to the ATPase and thereby the formation of a permeability transition pore (PTP), which can induce cell death⁹. K⁺ flux via the ATPase is under the control of survivalrelated protein Inhibitory Factor 1 (IF1), which in turn is regulated by the Bclfamily proteins Bcl-xL and Mcl-1, to constitute a mitochondrial ATP-dependent $K^{\scriptscriptstyle +}$ current (K_{ATP}) that protects against PTP opening during ischemia/reperfusion and other stress conditions. Organ protection during ischemia/reperfusion provided by the canonical mKATP activator diazoxide requires IF1. K+ influx via the ATPase is counterbalanced by K⁺ extrusion via the K⁺/H⁺-exchanger (KHE). In addition to the ATPase, also CCDC51 complexing with ABCB8 constitutes functional mK_{ATP}¹⁶, but without coupling to ATP production.

is also a major candidate for the long-sought mitochondrial ATPdependent K⁺ channel (mK_{ATP}), which is a central downstream effector of a phenomenon termed ischemic preconditioning, where repetitive brief episodes of ischemia and reperfusion of an organ reduce necrosis after a subsequent longer phase of ischemia with reperfusion by delaying the opening of the mitochondrial permeability transition pore (mPTP),^{7.8} an event that dissipates the mitochondrial membrane potential and—if nonreversible induces cell death.⁹

K_{ATP} channels, located on the sarcolemma and the IMM, are controlled by the metabolic state of a cell: when the cellular ATP/ADP ratio drops, activation of sarcolemmal KATP channels hyperpolarizes the cell membrane, reducing its excitability to reduce ATP demand,¹⁰ whereas activation of mitochondrial K_{ATP} channels (mKATP) optimizes ATP production through mitochondrial volume regulation,¹¹ as described above. Although after its first description in the early 1990s¹², the electrophysiological and pharmacological properties of the mKATP were extensively characterized,¹¹ its molecular identity has long remained elusive. It was initially proposed that, akin to its sarcolemmal counterpart, mKATP comprised K+-selective pore-forming subunits from the Kir6.x family; however, this model was discarded as genetic ablation of Kir6.x channels did not suppress mKATP responses.¹³ Subsequently, the renal outer medullary K⁺ channel (ROMK) evolved as a potential pore-forming subunit of mKATP based on a proteomic screen and in vitro evidence,¹⁴ but cardiacspecific knock-out of ROMK later revealed that it is dispensable for cardioprotection and mK_{ATP} responses.¹⁵ Recently, a protein with previously unknown function (CCDC51) was identified to form a channel with mK_{ATP} -like properties when associating with the ATP Binding Cassette protein 8 (ABCB8),¹⁶ which had already been shown to modulate mK_{ATP} activity.¹⁷ Since knock-out of CCDC51 in vivo confirmed its essential role to regulate mitochondrial function in unstressed conditions and protect from necrosis during ischemia/reperfusion,¹⁶ CCDC51 and ABCB8 are currently the most accepted candidates in the field to constitute the mK_{ATP} (Figure 1).

The second study by Juhaszova et al.⁶ in this issue of Function delineates the endogenous and exogenous regulation of the F₁F_o-ATP synthase in its function as a K_{ATP} channel. The survival-related protein Inhibitory Factor 1 (IF₁) is regulated by Bcl-family proteins, in particular Bcl-xL and Mcl-1, but not Bcl-2, through interaction at a BH3-like domain, which increases chemo-mechanical efficiency of the F₁F_o-ATP synthase to function as mK_{ATP} (Figure 1).⁶ Furthermore, the cardioprotective effect of diazoxide, the canonical mK_{ATP} activator, is shown to be mediated by IF₁. By applying Bayesian phylogenetic analysis, the authors conclude that IF₁ is likely an ancient Bcl family member that evolved from bacteria resident in eukaryotes and prevents excessive ATP consumption through the reversal of the ATP synthase to maintain the protonmotive force.⁶

The authors need to be applauded for providing groundbreaking results with fundamental implications for cellular bioenergetics and survival. First, these observations identify K^+ import via the F_1F_0 -ATP synthase as one central mechanism by which the rate of ATP turnover in the cytosol is matched by ADP phosphorylation in mitochondria. Second, they assign the F_1F_0 -ATP synthase a central role in cardioprotection, where mitochondrial $K^{\scriptscriptstyle +}$ influx via the $F_1F_0/K^{\scriptscriptstyle +}$ uniporter elevates the threshold to elicit ROS-induced permeability transition. Of note, by coupling mitochondrial K⁺ influx to ATP production, the subsequent K⁺ extrusion via the KHE at the expense of protonmotive force is energetically counterbalanced, which is not the case when K⁺ enters mitochondria via CCDC51/ABCB8 (Figure 1), thereby avoiding the production of futile heat through "uncoupled" K⁺ leak. This led the authors to suggest that CCDC51/ABCB8-related K⁺ flux may play a rather "fine-tuning" role compared with ATP synthase-dependent mKATP. However, since CCDC51 knock-out prevented most (but not all) of the cardioprotection provided by diazoxide,¹⁶ the herein suggested role of the ATP synthase as mKATP still needs to stand the in vivo test (of time), for instance in mice deficient of IF_1 .

In ancient Roman myth and religion, Janus is the god of beginnings, transitions, duality and endings, deciding over war and peace, or translated to biology—over life and death. Since in the past decade, the F_1F_0 -ATP synthase has already evolved as a central component of the mPTP under the control of cyclophilin D,⁹ the novel data presented in this issue of *Function*^{3,6} deservedly assign the mitochondrial F_1F_0 -ATP synthase a title as a *Janus-faced* enzyme, since in addition to its canonical role to produce ATP—utilizing both H⁺ and K⁺ flux—and its increasingly recognized role as a component of the mPTP,⁹ it also accounts for the protection from permeability transition through its novel function as an mK_{ATP} channel (Figure 1).

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Conflicts of interest

None declared in the context of this publication.

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