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Quantification of Myocardial Mitochondrial Membrane Potential Using PET

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Abstract

Purpose of review: To present a method enabling *in-vivo* quantification of tissue membrane potential (Ψ_T), a proxy of mitochondrial membrane potential (Ψ_m), to review the origin and role of Ψ_m , and highlight potential applications of myocardial Ψ_T imaging.

Recent findings: Radiolabelled lipophilic cations have been used for decades to measure Ψ_m *in-vitro*. Using similar compounds labeled with positron emitters and appropriate compartment modelling, this technique now allows *in vivo* quantification of Ψ_T with positron emission tomography. Studies have confirmed the feasibility of measuring myocardial Ψ_T in both animals and humans. In addition, Ψ_T showed very low variability among healthy subjects, suggesting that this method could allow detection of relatively small pathological changes.

Summary: *In-vivo* assessment of myocardial Ψ_T provides a new tool to study the pathophysiology of cardiovascular diseases and has the potential to serve as a new biomarker to assess disease stage, prognosis, and response to therapy.

Keywords

mitochondrial membrane potential; tissue membrane potential; positron emission tomography; triphenylphosphonium; mitochondria; heart failure

Introduction

Mitochondria are double-membrane-bound cellular organelles found in almost all eukaryotic cells. They are involved in the production of the chemical energy required for cellular biomechanical reactions via the phosphorylation of adenosine diphosphate (ADP) into adenosine triphosphate (ATP). The mitochondrial membrane potential (Ψ_m), which

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Conflict of Interest

Dr. Normandin has a patent US20190125281A1 issued.

Dr. Alpert has a patent US020190125281A120190502 issued.

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Human and Animal Rights and Informed Consent

This article does not contain any studies with human or animal subjects performed by any of the authors.

represents the electric gradient across the mitochondria inner membrane, is at the very core of mitochondrial energy production, fueling the conversion of ADP to ATP. Alteration of myocardial Ψ_m is present in a wide range of cardiac pathologies and typically appear early in the disease process, often preceding symptoms [1–3]. Until recently, assessment of myocardial Ψ_m was only possible in isolated mitochondria and explanted hearts [4,5]. However, new methodological and technological advances enable *in vivo* quantification of membrane potential in absolute units of millivolts (mV). In this paper, the origin and role of myocardial Ψ_m will be overviewed. A method for membrane potential quantification with positron emission tomography (PET) will be presented. Finally, potential cardiovascular applications of this technique will be discussed.

Role of mitochondria and origin of Ψ_m

Mitochondria play several roles in cellular homeostasis and function, including hormonal, ion, and immune signaling, steroid synthesis, regulation of cellular metabolism, as well as apoptosis [6–8]. One of the principal function of mitochondria is the phosphorylation of ADP into ATP, providing the vast majority of the energy necessary for cell function, including cardiomyocytes' contraction [8]. The first steps of this process involve glycolysis and oxidation of fatty acids to produce Acetyl-CoA. Acetyl-CoA is the common intermediate driving the production of NADH and FADH₂ through the tricyclic (Krebs) cycle (Figure 1). NADH and FADH₂ then feed electrons to the electron transport chain (ETC). Redox reactions are taking place along the ETC and protons are translocated across the inner membrane, from the mitochondrial matrix into the intermembrane space, establishing a proton-motive force (p_m). This force is composed of the pH gradient (pH_m) and the mitochondrial membrane potential Ψ_m , with Ψ_m comprising the bulk of p_m . Ultimately, the ATPsynthase dissipates a small fraction of the energy stored in the p_m to convert ADP into ATP by letting protons cross back in the mitochondrial matrix. Critical to this process is the impermeability of the mitochondrial inner membrane to most ions, allowing accumulation of protons in the intermembrane space.

When the ETC function is optimized, there is maximal coupling between proton pumping and phosphorylation of ADP to ATP. This occurs when there is minimal proton leakage across the inner membrane. When levels of ADP increase, as seen in states of high energy demand, there is increased phosphorylation and utilization (and thus dissipation) of the proton gradient, stimulating the ETC function and oxygen consumption. Conversely, low ADP levels slow phosphorylation rate, leading to inhibition of the ETC [9]. It is thought that Ψ_m has direct inhibitory effects on the rate of electron transport [10], and when there is significant reduction of Ψ_m , the ETC is free to run at maximal rate and Ψ_m can be progressively restored. Collapse of Ψ_m via opening of the mitochondrial permeability transition pores (mPTP) and subsequent release of cytochrome C into the cytosol plays a central role in cellular apoptosis.

Oxidative stress and Ψ_m

Reactive oxygen species (ROS) are a normal by-products of the respiratory chain. Under physiological conditions, the ETC is efficient with minimal production of ROS [11]. At

small concentrations, ROS play an important physiological role as signal transduction molecules within the mitochondria and cells. However, when Ψ_m falls outside its optimal physiological range, either above or below, there is a significant rise in the production of ROS by the ETC; increasing Ψ_m by only 10 mV leads to a 70–90% increase in ROS production [12,13]. The respiratory chain complexes are very sensitive to oxidizing agents due to the vulnerability of their components to direct ROS attack. Their oxidative modifications manifest with a decreased enzymatic activity and dysfunction of the whole respiratory chain [14]. Oxidative damage to the respiratory chain complexes results not only in a decreased efficiency of ATP production but can also lead to further increased in ROS production, amplifying the ROS release [14]. Finally, oxidation of the inner membrane anion channels (IMACs) by ROS lead to their opening and partial dissipation of Ψ_m . Overall, in conditions associated with increased ROS, Ψ_m may be reduced, and because the ratio ATP:ADP decreases exponentially as a function of Ψ_m , relatively small mitochondrial depolarization elicits a large decreases in the ability to convert ADP to ATP [15].

Quantification of Ψ with PET imaging

In vitro quantification of Ψ_m has been performed for decades by measuring the distribution of various lipophilic cations in isolated mitochondria [4,5]. At equilibrium, the concentration of those compounds on each side of a membrane (\bar{C}) can be expressed by the Nernst equation (Eq.1).

$$\frac{\bar{C}_{in}}{\bar{C}_{out}} = e^{-\beta\Delta\Psi} \quad \text{Eq. 1}$$

Where Ψ is the membrane's electric potential and $\beta = \frac{zF}{RT}$ is the ratio of known physical parameters: z is the valence of the ionic probe, F denotes Faraday's constant, R is the universal gas constant, and T is the temperature in Kelvin. Assessment of Ψ_m with PET imaging rely on the same methodology, using lipophilic cations labelled with positron emitters such as ^{18}F -tetraphenylphosphonium (^{18}F -TPP⁺). Those tracers cross freely and with minimal interaction the cellular and mitochondrial phospholipid bilayer membranes, and their distribution at equilibrium abides the Nernst equation. Radiotracer uptake is therefore exponentially proportional to Ψ_m ; under physiological conditions, concentration of a monovalent lipophilic cations will be 3 to 10 times greater in the intracellular space compared to the extracellular space and 100 to 500 times greater in the mitochondrial matrix compared to the extracellular space [16].

An image voxel containing myocardium can be modeled into 4 compartments which include the plasma, interstitial space, cytosol, and mitochondria (Figure 2) [17]. The plasma and interstitial space compose the extracellular space. Using this model, the tracer concentration in a voxel at equilibrium can be expressed as:

$$\bar{C}_{PET} = (1 - f_{ECS})(f_{mito} \cdot \bar{C}_{mito} + (1 - f_{mito}) \cdot \bar{C}_{cyto}) + f_{ECS} \cdot \bar{C}_{ECS} \quad \text{Eq.2}$$

Where \bar{C}_{mito} , \bar{C}_{cyto} , and \bar{C}_{ECS} , represent the probe concentration at equilibrium in the mitochondria, cytosol, and extracellular space respectively, and f_{ECS} and f_{mito} represent

the extracellular space and mitochondrial volume fraction, respectively. Noting that plasma concentration (\bar{C}_p) is equal to \bar{C}_{ECS} at equilibrium, division of Eq. 2 by \bar{C}_p yields the following expression:

$$V_T = \frac{\bar{C}_{PET}}{\bar{C}_p} = (1 - f_{ECS}) \left(f_{mito} \cdot e^{-\beta(\Delta\Psi_m + \Delta\Psi_c)} + (1 - f_{mito}) \cdot e^{-\beta\Delta\Psi_c} \right) + f_{ECS} \quad \text{Eq. 3}$$

Where V_T , Ψ_m and Ψ_c represent the volume of distribution of the tracer, and the mitochondrial and cellular membrane potential, respectively. Because $V_T \gg 1$ and $e^{-\beta(\Delta\Psi_m + \Delta\Psi_c)} \gg e^{-\beta(\Delta\Psi_c)}$, Eq. 3 can be approximated by

$$V_T \approx (1 - f_{ECS}) \cdot f_{mito} \cdot e^{-\beta(\Delta\Psi_T)} \quad \text{Eq. 4}$$

Where Ψ_T is the total tissue membrane potential, defined as $\Psi_m + \Psi_c$. Eq. 4 can be rearranged as:

$$\Delta\Psi_T = \frac{1}{\beta} \ln \left[\frac{(1 - f_{ECS}) \cdot f_{mito}}{V_T} \right] \quad \text{Eq. 5}$$

From Eq. 5, we note that 3 parameters are required to quantify Ψ_T : 1) the extracellular space fraction, 2) the mitochondrial fraction, and 3) the tracer volume of distribution. The f_{mito} parameter is currently not measurable non-invasively. It has to be assumed constant and a value of 0.25 can be used for human studies [18]. It is important to note that mitochondrial fraction may decrease in disease state, ultimately leading to an underestimation of membrane potential (less negative). The volume of distribution V_T can be determined kinetically [17] or by taking the ratio between the voxel concentration measured by PET imaging and the plasma concentration at secular equilibrium [19,20].

Quantification of extracellular space

As stated above, measurement of extracellular volume (ECV) is critical to accurately quantify membrane potential. This is because there is a significant extracellular space variability among healthy individuals and several diseases are associated with extracellular space expansion. Failure to account for ECV leads to significant underestimation of Ψ as well as increased variability between subjects [21]. ECV can be assessed with computed tomography (CT) or magnetic resonance imaging (MR). In both cases, images are acquired before and after administration of a contrast agent (iodine-based for CT and gadolinium-based for MR) which distributes in the extracellular space. At the present time, MR allows more robust and precise quantification of ECV without additional ionizing radiation but may be less accessible. In that context, Ψ_T imaging represents an attractive application of hybrid PET/MR scanners. CT, on the other hand, is readily available and acquisition can be performed on a conventional PET/CT system. Detailed methodology of ECV quantification is beyond the scope of this review and can be found elsewhere (see [22]).

In vivo quantification of myocardial membrane potential

Fukuda et al were the first to attempt *in vivo* quantification of Ψ_T in the 1980s [23]. Following a bolus injection of the lipophilic cation ^{11}C -triphenylmethylphosphonium (^{11}C -TPMP), they measured myocardial tracer concentration in dogs with PET imaging as well as in mice and rats using autoradiography. Accounting for the extracellular volume, they calculated the Ψ_T using the Nernst equation with plasma and myocardial tracer concentrations. Over 25 years later, Gurm et al reported an attempt to non-invasively quantify Ψ_m using PET imaging in a swine model of ischemic heart disease [24]. They used the monovalent lipophilic cation ^{18}F -TPP⁺, which passively crosses membranes without significant interaction. Based on the Nernst equation, they reported an average Ψ_m of -91 mV (Table 1), which appears significantly discordant with previous measurements. This difference can be explained in part by the fact that extracellular volume was not taken into account, leading to an underestimation of Ψ_m . Furthermore, both Gurm et al and Fukuda et al used bolus radiotracer injection but assumed that imaging was performed at secular equilibrium (i.e. constant tracer concentration in the different compartments), which was not the case. Such assumption may lead to erroneous quantification of membrane potential [21,25]. More recently, Alpert et al reported a successful method to quantify Ψ_T in swine [17]. Using a bolus injection of ^{18}F -TPP⁺ and a dynamic PET acquisition, they estimated V_T in the myocardium using Logan analysis [26] and calculated Ψ_T using the Nernst equation, accounting for ECV. Subsequently, Alpert et al reported that intracoronary infusion of BAM15, a proton gradient uncoupler affecting specifically the mitochondrial membrane without depolarizing the cellular membrane, leads to partial and reversible decreased in ^{18}F -TPP⁺ concentrations (Figure 3) [19]. These results confirmed that myocardial ^{18}F -TPP⁺ concentrations measured by PET are sensitive to temporal change in Ψ_m .

Pelletier-Galarneau et al were the first to demonstrate the feasibility of Ψ_T *in vivo* quantification in humans [20]. Using a bolus plus infusion of ^{18}F -TPP⁺ and hybrid PET/MR imaging, they reported an average Ψ_T of -160.7 ± 3.7 mV among 13 healthy volunteers [20]. The corresponding average Ψ_m was -123 mV, in excellent agreement with explanted heart measurements (-118 mV) [27]. Importantly, Ψ_T showed very low variability between subjects, suggesting that this method could prove useful to detect relatively small pathological changes and that small sample size could be sufficient to test hypothesis. In addition, high-quality parametric images can be generated with this method, allowing assessment of regional variability in Ψ_T (Figure 4). Other radiotracers have been used in pre-clinical studies to assess mitochondrial membrane potential, including ^{18}F -fluoropentyl-triphenylphosphonium (^{18}F -FPTP), ^{18}F -fluorobenzyl-triphenylphosphonium (^{18}F -FBnTP), and ^{18}F -MitoPhos, all of which share the property of being monovalent lipophilic cations [28–31]. So far, only ^{18}F -TPP⁺ has been used in humans.

Potential clinical applications

Alterations in mitochondrial function, ROS production, and Ψ_m are seen in a broad range of pathologies, including diabetes, cancer, various degenerative diseases, and myopathy [29,32,33]. In cardiovascular diseases, mitochondrial dysfunction is thought to play a

critical role in the development and progression of ventricular arrhythmias, heart failure, cardiotoxicity, and reperfusion injury just to name a few [34–37]. Non-invasive mapping of membrane potential could provide a new biomarker in mitochondrial-related diseases. In addition, it has the potential to play a role in the assessment of response to therapy, especially with the advent of new mitochondrial therapies.

Chemotherapy-induced cardiotoxicity

Chemotherapy-induced cardiotoxicity is a multifactorial complication of chemotherapy with interactions between the nature and dosage of the therapy, environmental factors, as well as genetic factors [38]. It is a relatively frequent side effects of many anti-cancer agents, including anthracyclines such as doxorubicin [39]. As the damages observed in chemotherapy-induced cardiotoxicity are often irreversible, it is imperative to detect cardiotoxicity early in order to minimize long-term morbidity and mortality, especially given the trend of improved cancer survival. The mechanisms underlying doxorubicin-induced cardiotoxicity (DIC) are incompletely understood. A key role has been attributed to mitochondrial dysfunction with mitochondria being the most injured intracellular organelle from exposure to doxorubicin [40]. While early research suggested that DIC was mediated through redox recycling of doxorubicin and subsequent mitochondrial and cellular damage, recent work has identified multiple alternative pathways of mitochondria mediated DIC [41]. Doxorubicin has a high binding affinity with cardiolipin, a phospholipid present in the mitochondrial membrane, which is involved in maintaining mitochondrial structure and function as well as cell survival [42,43]. Cardiolipin is particularly important for normal ETC activity, which is disrupted by doxorubicin binding to cardiolipin. The resulting increased electron transfer to oxygen molecules and generation of free radicals damages the mitochondrial membrane, leading to reduced ATP levels [40,41,43]. While the pathway of DIC through inhibition of the ETC can be observed as a result of acute mitochondrial exposure to doxorubicin [41], chronic effects of doxorubicin have been related to binding of doxorubicin to topoisomerase 2 β , which is largely present in mitochondria, resulting in DNA double-strand breaks and transcriptome changes leading to defective mitochondrial biogenesis and ROS formation [40,44–46]. Further suggested DIC mechanisms include calcium dysregulation following increased mitochondrial ROS generation and failed energy production as well as activation of apoptotic pathways [40,47]. The mitochondrial dysfunction underpinning DIC, particularly inhibition of ETC, can be associated with depolarization of Ψ_m . In fact, depolarization of Ψ_m following exposure to doxorubicin has been observed in *in vitro* studies in postnatal rat cardiac myocytes and human colon and breast cancer cell lines [47,48]. Importantly, these molecular changes occur early in the disease stage before irreversible functional impairment can be observed.

At present, screening for cardiotoxicity is done mostly by serial assessment of left ventricular ejection fraction (LVEF) and echocardiographic strain imaging [49], both of which reflecting changes that are occurring late in the disease process, after irreversible damages have occurred. In that context, non-invasive measurement of membrane potential could represent an interesting biomarker of cardiotoxicity, with preliminary data supporting this hypothesis. Indeed, McCluskey et al showed that myocardial concentration of ^{18}F -MitoPhos, a lipophilic cation, decreased significantly 48h after a single infusion of

doxorubicin, while other biomarkers (e.g. troponins) remained unchanged [31]. Further studies are required to confirm whether change in membrane potential can predict cardiotoxicity and alter patient management.

Heart Failure

Mitochondrial dysfunction is a hallmark of heart failure (HF), both with preserved and reduced ejection fraction [50]. Several alterations in mitochondrial function have been reported in heart failure, including ATP production, glucose metabolism, increased oxidative stress, dysregulation of calcium metabolism, protein modifications, and apoptosis [51,52]. In addition, therapies improving mitochondrial function have been shown to improve outcomes of patients with HF. For instance, the Q-SYMBIO trial showed that coenzyme Q10 (CoQ10), a molecule targeting oxidative stress and mitochondrial dysfunction, improves symptoms and reduces major adverse cardiovascular events in patients with HF [53]. Other molecules such as ACE inhibitors have also been shown to improve mitochondrial function [54], reinforcing the importance of mitochondrial dysfunction in HF. In the failing heart, there is increased production of ROS by the mitochondria [55,56]. Excess ROS is associated to cellular dysfunction, DNA damage, and can lead to cell death. In the myocardium, ROS can directly impair contractile function and activate pathways leading to hypertrophy, necrosis, and apoptosis [57]. Excess ROS was also shown to dissipate the Ψ_m via different mechanisms [58]. Ψ_m is directly related to the capacity of the ETC to pump protons across the inner mitochondrial membrane, and is therefore necessary for ATP generation [58,59]. The decrease of Ψ_m in HF had been confirmed in animal model of HF with decreased activity of the ETC complexes lead to decreased Ψ_m , which in turn resulted in reduced mitochondrial Ca^{2+} concentration and impaired myocyte contractility [35]. Imaging of membrane potential could provide insight in the pathophysiological processes of the disease, help with development of new therapeutic interventions, and allow for therapy optimization.

Ischemia-Reperfusion Injury

Acute myocardial infarction is one of the leading causes of death and disability worldwide. Treatment of acute ischemia relies on timely reperfusion. However, myocardial reperfusion itself may further damage the affected cardiomyocytes, a phenomenon known as ischemia-reperfusion injury (IRI), and it has been estimated that approximately 50% of the final infarcted area is related to IRI [60]. The complex mechanisms of IRI involve different biological pathways and are not yet fully understood. However, a key role has been attributed to opening of the mPTP during reperfusion [61]. This results in immediate depolarization of Ψ_m , disruption of ATP production, production of ROS, and the release or activation of several pro-apoptotic proteins like cytochrome C [62,63]. Furthermore, uncoupling of the ETC results in reversal of ATPase, which switches from synthesis of ATP to active hydrolysis. The resulting rapid decline of intracellular ATP concentration leads to disruption of ionic and metabolic homeostasis and the activation of degradative enzymes, in turn resulting in irreversible damage to the cell [64]. Triggers for mPTP opening include Ca^{2+} overload, rapid normalization of pH, oxidative stress, and mitochondrial depolarization, all conditions that occur during post-ischemic reperfusion [64,65]. Non invasive assessment of membrane potential could provide a new tool to study the pathophysiology of IRI and assess the effect of new therapies.

Conclusion

In vivo quantification of Ψ_m with PET imaging relies on the extension of well-established chemistry principles and benchtop *in vitro* methods. With radiolabelled lipophilic cations, such as ^{18}F -TPP⁺, and appropriate compartment modelling, *in vivo* quantification of the tissue membrane potential (Ψ_T), a proxy of Ψ_m , in absolute units of millivolts can be achieved. Studies have demonstrated the feasibility of measuring myocardial Ψ_T in both animals and humans and that myocardial ^{18}F -TPP⁺ concentration measured by PET are sensitive to temporal changes in Ψ_m . In addition, Ψ_T showed very low variability between subjects, suggesting that this method could be used to detect relatively small pathological changes. As alteration in mitochondrial function and membrane potential is ubiquitous in cardiovascular diseases, this imaging technique could provide a new tool to better understand the pathophysiology of those diseases and has the potential to serve as a new biomarker to assess disease stage, prognosis, and response to therapy. Further studies are needed to establish these potential roles.

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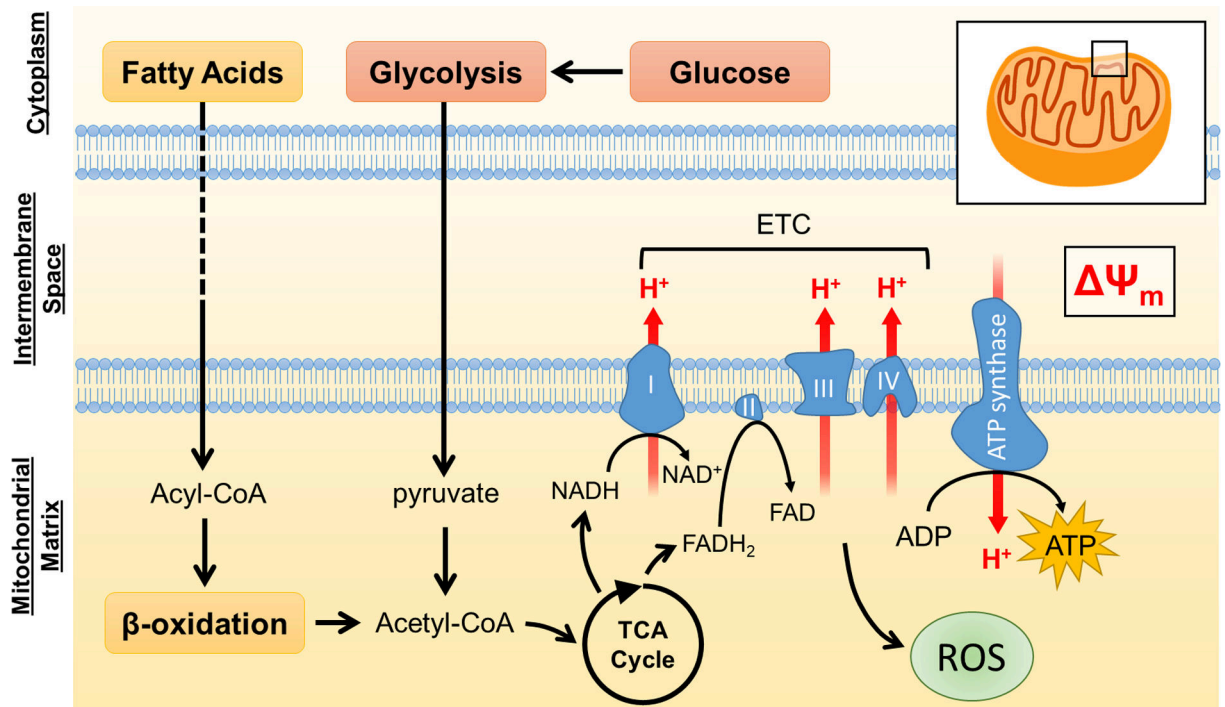


Figure 1:
Schematic representation of the electron transport chain and generation of the mitochondrial membrane potential (Ψ_m); ADP, adenosine diphosphate; ATP, adenosine triphosphate; ROS, reactive oxygen species; TCA, tricyclic cycle.

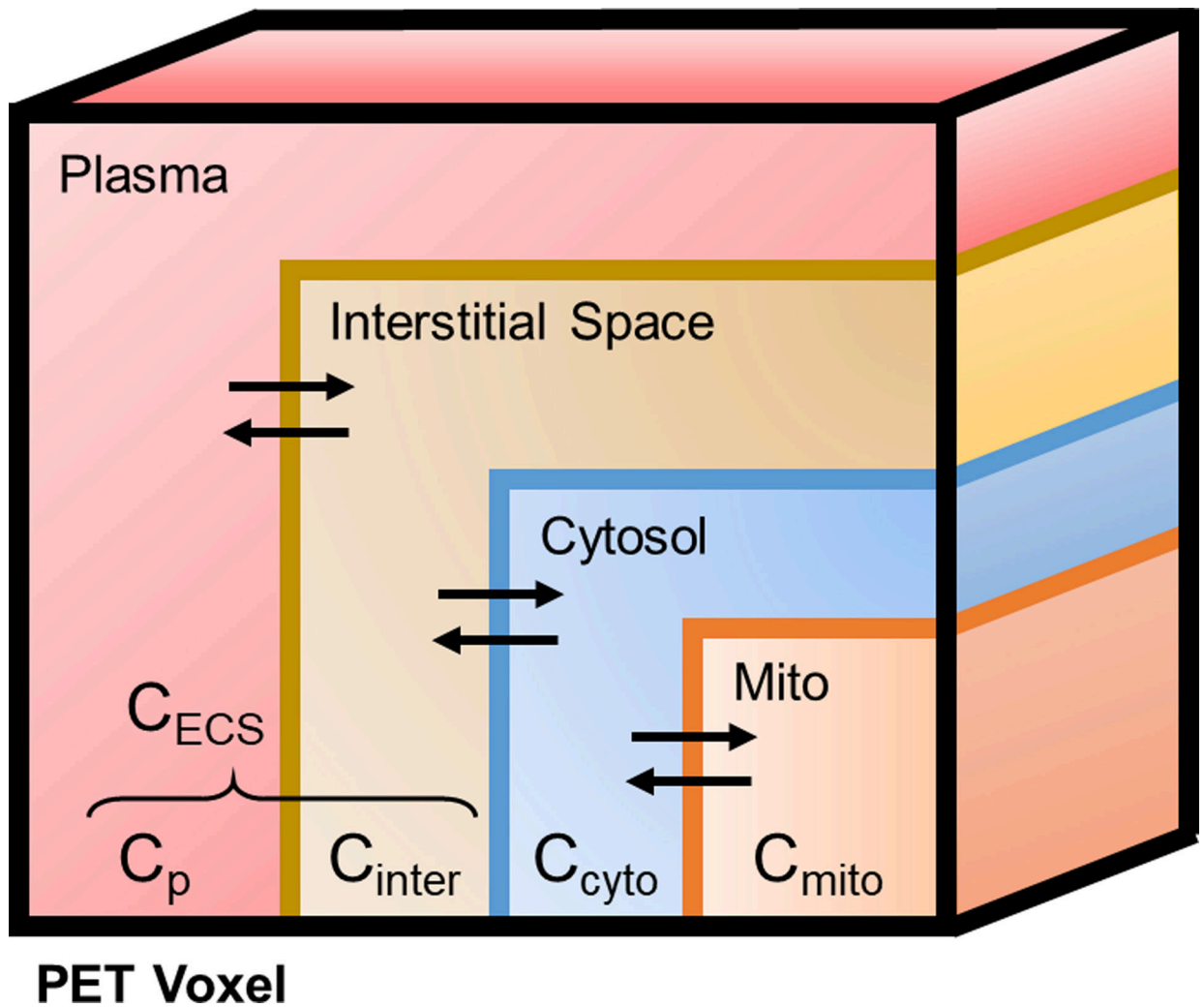


Figure 2:

Compartmental model for a lipophilic cation in a positron emission tomography (PET) image voxel. C_p , C_{inter} , C_{cyto} , C_{mito} , and C_{ECS} represent the tracer concentration in the plasma, interstitial space, cytosol, mitochondria, and extracellular space respectively. (Adapted from [17•].)

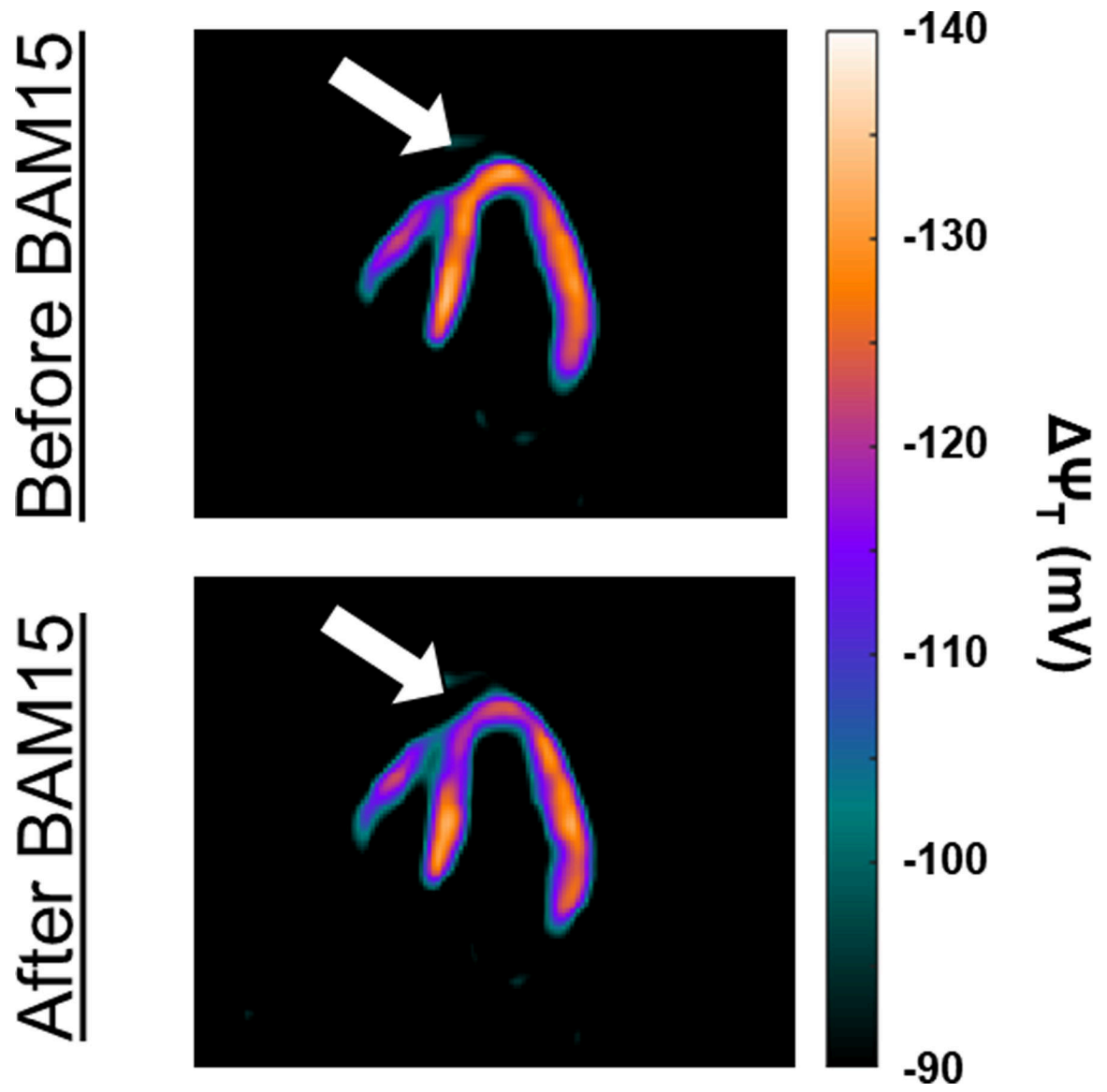


Figure 3: Effect of BAM15, a selective mitochondrial proton uncoupler, on tissue membrane potential (Ψ_T). Vertical long axis of a pig heart before (**top**) and after (**bottom**) intracoronary infusion of BAM15 in the left anterior descending artery demonstrating partial depolarization in the septal and apical segments. (Data from [19].)

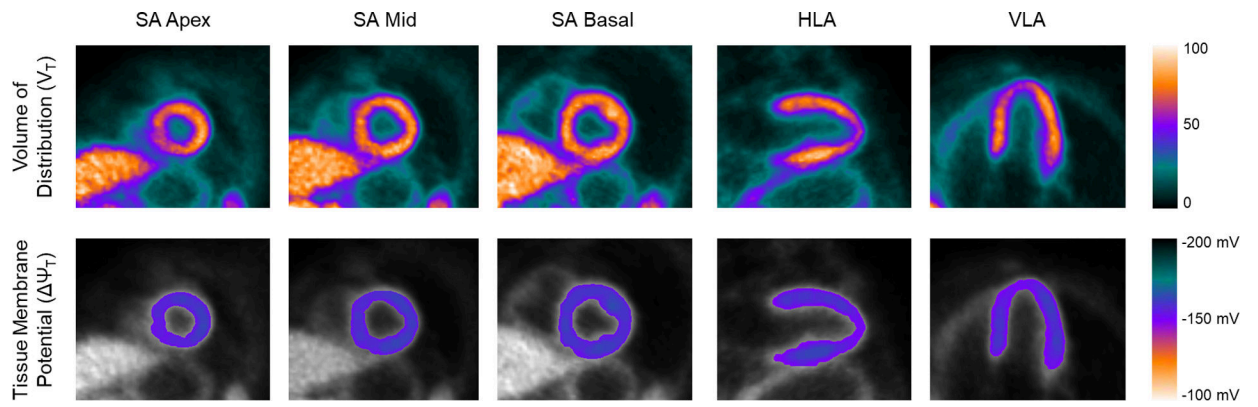


Figure 4: Parametric images of the volume of distribution (V_T) and tissue membrane potential ($\Delta\Psi_T$). HLA, horizontal long axis; SA, short axis; VLA, vertical long axis. (Reproduced by permission from Springer Nature from: Pelletier-Galarneau et al. *Eur J Nucl Med Mol Imaging*. 2020) [20••].

Table 1:*In vivo* assessment of membrane potential*

Authors	Year	Species	Ψ_m (mV)	Ψ_T (mV)
Fukuda et al es	1986	Dogs	-	-148.1 ± 6.0
		Rats	-	-146.7 ± 3.8
		Mice	-	-139.3 ± 5.8
Gurm et al [24]	2012	Swine	-91 ± 11	
Alpert et al [17]	2017	Swine	-	-129.4 ± 1.4
Pelletier-Galarneau et al [20]	2020	Human	-	-160.7 ± 3.7

* Disparities in observed values can be attributed to methodological and technical differences, species variability, and partial depolarization induced by anesthesia.