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Endothelial mitochondrial ROS, un-coupled from ATP synthesis, determine both physiological endothelial activation for recruitment of patrolling cells, and pathological recruitment of inflammatory cells

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Abstract

Mitochondrial reactive oxygen species (mtROS) are signaling molecules, which drive inflammatory cytokine production and T cell activation. In addition, cardiovascular diseases, cancers, and autoimmune diseases all share common feature of increased mtROS level. Both mtROS and ATP are produced as a result of electron transport chain activity, but it remains enigmatic whether mtROS could be generated independently from ATP synthesis. A recent study shed light to this important question and found that during endothelial cell (EC) activation, mtROS could be upregulated in a proton leak-coupled, but ATP synthesis-uncoupled manner. As a result, EC could upregulate mtROS production for physiological EC activation without compromising mitochondrial membrane potential and ATP generation, and consequently without causing mitochondrial damage and EC death. Thus, a novel pathophysiological role of proton leak in driving mtROS production was uncovered for low grade physiological EC activation, patrolling immunosurveillance cell trans-endothelial migration and other signaling events without compromising cellular survival. This new working model explains how mtROS could be increasingly generated independently from ATP synthesis and endothelial damage/death. Mapping the connections between mitochondrial metabolism, physiological EC activation, patrolling cell migration and pathological inflammation is significant towards the development of novel therapies for inflammatory diseases and cancers.

Author's contribution

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Competing interests

The authors declare that they have no competing interests.

XL carried out the primary literature search and drafted the manuscript. PF, WY, KC, ML, KX, TG, and HW provided material input and helped critically reading the manuscript. XFY designed the study, provided field expertise and revised the manuscript. All authors read and approved the final manuscript.

Keywords

mitochondria; ROS; cardiovascular disease; cancer; vascular inflammation

Mitochondrial reactive oxygen species, induced by newly classified conditional danger signals, orchestrate inflammatory responses

Historically considered as merely cellular "powerhouses" that manufacture ATP and other metabolites, mitochondria are increasingly being recognized as "metabolic hub" that are involved in the crosstalk and signal integration of cell proliferation, death, differentiation, inflammation, and repair pathways. Remarkably, it is now increasingly appreciated that mitochondria serve as "sentinel" organelles that are not only capable of detecting cellular insults but also orchestrating inflammatory responses¹. One of the most well characterized mechanisms of such involves mitochondria-derived reactive oxygen species (ROS), which are often induced in response to cellular homeostasis disruptions including infections, sterile damage, and metabolic disturbances. Reciprocally, mitochondrial ROS (mtROS), as signaling molecules, convey these cellular insult signals to the rest of the cell by engaging in a variety of cellular pathways, such as inflammasome activation, proinflammatory transcription factor activation, apoptosis, autophagy, and DNA-based neutrophil extracellular trap formation^{2, 3}. The key cellular receptors that recognize the "threat" signals include Tolllike receptors (TLRs) and cytosolic sensing NLRs [NOD (nucleotide binding and oligomerization domain)-like receptors] that recognize a variety of pathogen-associated molecular patterns (PAMPs) and endogenous metabolites-related danger-associated molecular patterns (DAMPs)⁴. Together, ⁴ TLRs work synergistically with NLRs to recognize PAMPs and DAMPs, particularly in inflammation privileged tissues where inflammasome component genes for activating pyrotopsis (inflammatory cell death) are not constitutively expressed^{5–11}. Moreover, four additional DAMP receptors including transmembrane C-type lectin receptors, retinoid acid inducible gene I (RIG-I), AIM2 (absent in melanoma 2), receptor for advanced glycation end products (RAGE, also a receptor for high mobility group box 1(HMGB1)) have been characterized¹². Herein, these 6 categories of DAMP receptors were referred as classical DAMP receptors¹³. However, we are still far from uncovering the identities of DAMP receptors that are responsible for recognizing all the elevated endogenous metabolites-related DAMPs. In fact, there are as many as 41,993 different entries in current human Metabolome Database (http://www.hmdb.ca/). In order to have a sufficient binding affinities¹⁴, additional DAMP receptors must exist in order to transduce specific and effective signals from diverse human metabolite-related DAMPs¹³.

A series of research articles published from our laboratory recently addressed this important issue by using the endogenous lysophospholipid (LPL) family members as the prototype. LPLs are bioactive, lipid-derived metabolites that act through G-protein coupled receptors (GPCRs) and they are generated by regiospecific phospholipases on substrates such as membrane phospholipids and sphingolipids¹⁵. Some of the LPLs that have been identified include lysophosphatidic acid (LPA), lysophosphatidylcholine (LPC), lysophosphatidylethanolamine (LPE), lipophosphoglycan (LPG), lysophosphatidylinositol (LPI), and lysophosphatidylserine (LysoPS)¹⁶. The prominent roles of LPLs in chronic

inflammatory disorders such as coronary artery disease (CAD), hypertension, atherosclerosis and severe vascular diseases are well established ^{17-20. 18, 19}In our recent report, we proposed a conceptually innovative paradigm that G-protein-coupled receptors (GPCRs) of LPLs can be classified as novel conditional DAMP receptors for the following reasons¹³: (1) LPLs are elevated during cellular stress or under stimulation by cardiovascular risk factors²¹; (2) Basal level of LPLs mediate normal cellular functions while elevated LPLs initiate signaling cascade to either activate or dampen innate immune responses via LPL-GPCRs; (3) The expression of anti-inflammatory LPLs and their receptor are also elevated in response to inflammation and may play a role in resolution of inflammation; and we designate such LPL receptors as homeostasis-associated molecular patterns (HAMPs); and (4) LPL receptors and classically DAMP receptors reciprocally regulate the expression of each other, which is responsible for either progression or resolution of inflammation 1^{13} . Thus, LPL receptors serve as conditional DAMP receptors by modulating physiological activities and housekeeping functions under normal conditions, while contributing to DAMP-mediated signaling pathways during cellular stress. We consolidated this working model by demonstrating that LPLs "transform" resting endothelial cells (ECs) to conditional innate immune cells²² via caspase-1 activating-sirtuin 1 inhibiting pathway²³. In addition, we found that in addition to post-translationally activating inflammasome, long-term low dose cardiovascular disease risk factors such as LPLs are required for transcriptionally upregulating low levels of or missing inflammasome components in order to elicit inflammatory responses⁶, presumably via histone modifications²⁴ and noncoding RNAs regulation^{25, 26}.

Despite these significant advances, it was unclear how distinct GPCR in response to LPL stimulation could converge to activate the inflammasome and in turn activate ECs. Previous studies investigated the role of cytosolic ROS-generating systems such as nicotinamide adenine dinucleotide phosphate (NADPH) oxidase in this process^{27–29}. It has been shown that LPL could induce nicotinamide adenine dinucleotide oxidase (mitochondrial)/ NADPH oxidase (cytosolic)-related ROS in ECs³⁰. In addition, LPC-induced ROS production and inflammasome activation depends on sodium influx, which might contribute to the production of NADPH oxidase³¹. Furthermore, it is well proven that alteration NADH/NAD + ratio, by ischemia, mutation, dysfunction of cytochrome c or even low levels of ATP, can induce mtROS production (mainly superoxide and hydrogen peroxide) at the complex 1 of mitochondrial electron transport chain. Meanwhile, reverse electron transport due to alterations in redox potential also causes the generation of mtROS from complex 1. These alterations could be sensed by NADPH oxidase components gp91PHOX and p22PHOX to assemble inflammasome and regulate cell signaling^{32, 33}. Nevertheless, key NADPH oxidase components gp91^{PHOX} and p22^{PHOX} are found to be dispensable for inflammatory cytokine production, whereas blockade of mtROS reduces inflammatory cytokine secretion instead³. Moreover, mtROS downstream of T cell receptor signaling are also required for antigenspecific T cell activation³⁴, suggesting that membrane receptor signal-linked mtROS promote T cell cytokine generation³⁵, T cell proliferation³⁶ and T cell/regulatory T cell (Treg)^{37–40} metabolism⁴¹. In agreement with previous studies, we found that LPC could induce both mitochondrial ROS and cytosolic ROS in a similar manner in human aortic ECs²¹. Importantly, LPC-induced mtROS are found to be increased independent of NADPH

oxidase function, but rather as a result of Ca^{2+} entry from the cytosol to the mitochondria. MtROS production is determined by the rates of both mtROS production and disposal. We found that LPL do not significantly regulate the expressions of mitochondrial antioxidant superoxide dismutase 2 (SOD2) and uncoupling molecule uncoupling protein 3 (UCP3), although there is a slight expression increase of adenine nucleotide translocator (ANT), another mitochondrial uncoupling molecule, after LPL stimulation. Of note, there are multiple different species of mtROS, including superoxide and hydrogen peroxide, both of which seem to be upregulated by LPL in ECs⁴². Collectively, these results indicated that LPL triggers mtROS and subsequent transformation of EC to immune-like cells through means of stimulating cytosolic Ca²⁺ entry into the mitochondria. Given the fact that LPLinduced EC activation has been implicated as an initiation step of atherogenesis and tumor angiogenesis^{19, 20}, therapeutic agents that target mitochondrial calcium uniporter and mtROS might serve as novel therapeutic targets against atherosclerosis and cancer. However, two fundamental questions remain to be answered before these applications could be used are that, how mitochondria, in response to the stimulation of conditional DAMPs such as LPC, generate mtROS in non-damaged ECs without compromising cellular energy supply; and how non-damaged ECs recruit patrolling monocytes and other innate immune cells in physiological conditions to perform their immune surveillance $duty^{43}$. More specifically, it remains unclear how mitochondria could produce mtROS specifically for cellular signaling purpose independently from ATP synthesis, despite the fact that both ROS and ATP production are coupled to the mitochondrial electron transport chain activity⁴⁴ (Figure 1).

Reciprocal regulation of mtROS and mitochondrial proton leak drives EC activation

Our recent study shed light into the important question outlined above and uncovered a previously unrecognized role of mitochondrial proton leak in driving ATP-synthesisuncoupled mtROS for normal physiological immune surveillance purpose²¹. Jensen first observed the formation of ROS in the respiration chain⁴⁵, followed by the pioneer works by Chance and colleagues on the mitochondrial production of hydrogen peroxide^{46, 47}. As explained in our recent review¹, generation of mtROS mainly takes place at the electron transport chain located on the inner mitochondrial membrane during the process of oxidative phosphorylation⁴⁴. During mitochondrial respiration, the electrons originated from the Krebs cycle pass along the electron transport chain, which ultimately lead to the generation of H₂O or ROS due to complete or partial mitochondrial O₂ reduction, respectively. Meanwhile, protons (H⁺) are being pumped from the mitochondrial matrix to the intermembrane space, thus creating mitochondrial membrane potential across the inner mitochondrial membrane. This proton gradient was used to synthesize ATP when protons reenter the mitochondrial matrix through ATP synthase (also termed complex V). Thus, the production of mtROS via electron leak⁴⁸ and ATP is intimately linked. Nevertheless, mtROS production is incompletely coupled to ATP synthesis, since mtROS is also coupled to the proton leak process mediated by uncoupling proteins⁴⁹, through which protons in the intermembrane space leak across the inner membrane independently of ATP synthase (Figure 1). Thus, one would argue that, in addition to its physiological and pathological importance for energy generation and metabolic rate, adaptive non-shivering thermogenesis

(in brown adipose tissue)⁵⁰, body mass regulation^{48, 49} and increased heat for acute inflammation ^{49, 51}, proton leak might play a role in the mitochondria's decision to induce mtROS or produce ATP. In fact, a recent study suggested that mtROS is induced during proton leak *in vivo*⁵², and that mtROS are also required for the activation of proton leak. However, the role of proton leak in mtROS production was not addressed in this study. In our recent report²¹, we also demonstrated that during EC activation process induced by endogenous proatherogenic stimuli LPC, mitochondrial proton leak and mtROS are both induced. In addition, we found that LPC induce the expression of ANT, which mediates both basal and inducible proton conductance. Importantly, we were able to demonstrate that proton leak induction is used by ECs to induce ATP-synthesis-uncoupled mtROS production (Figure 1)²¹. Thus, it appears that a positive regulatory loop between proton leak and mtROS exists, which might be needed to induce high amount of ATP-synthesis-uncoupled mtROS during physiological EC activation and pathological inflammatory responses (Figure 1).

MtROS level dictates endothelial response and "activation scale"

Another important observation from our reports is that mtROS level dictates EC response²¹. By using mouse aortic microarray analysis, mouse plasma and aortic metabolomic analysis, the Seahorse Mitochondrial Stress test, flow cytometry mtROS detection, mtROS inhibitor mitoTEMPO, electron spin resonance, confocal microscopy, and human aortic endothelial cells, we found that lower dosage (10µM) of LPC induced ATP-synthesis-uncoupled mtROS, which does not lead to cell death but is required for low grade physiological EC activation. In contrast, higher dosage (>20µM) of LPC that presumably mimic pathological conditions, however, induces higher level of mtROS, which are associated with significant EC death²¹. 10µM LPC is effective in inducing mtROS and elicit physiological EC activation. It was found that this response is due to mitochondrial Ca²⁺ entry from the cytosol, as LPC-induced mtROS is completely blocked by Ruthenium Red, the general inhibitor for transient receptor potential vanilloid molecules (TRPVs) on the plasma membrane^{53, 54} and presumably also inhibitors of mitochondrial calcium uniporter on the mitochondrial membrane^{55, 56}. Our work have demonstrated for the first time that low dose endogenous LPC stimulated endothelial cells can achieve a physiological activation status without endothelial injury and damage, as judged by decreased ATP generation. The mtROS generation, coupled with increased proton leak but uncoupled from ATP synthesis, is the hallmark of this type of physiological endothelial cell activation, which is conceptually innovative²¹. Of note, high levels of mtROS drive pathological endothelial cell activation, which is in agreement with our previous publications. Our findings argue prominent roles of mtROS in driving both physiological EC activation and inflammation responses as well as initiating cell death in pathological conditions^{10, 23}. We speculate that during inflammation, the amount of conditional DAMPs or PAMPs in the cellular environment could be sensed by endothelial mitochondria⁵⁷, which reciprocally signal through various levels of mtROS to dictate the ultimate endothelial activation scales in both physiological and pathological conditions. Such mechanisms are not only important in recruiting and regulating normal physiological "patrolling" immune functions⁵⁸ and immune surveillance⁵⁹ for antimicrobiota and anti-tumor immunity⁶⁰, but also seems to be easily deranged during chronic inflammatory diseases, such as atherosclerosis (Figure 1).

Endothelial mtROS physiologically control patrolling cell immunosurveillance duty, which are "hijacked" during early atherogenesis to recruit inflammatory cells

It has been widely recognized that Ly6Chigh inflammatory monocyte recruitment into aorta in response to the stimulation of metabolic cardiovascular disease risk factors including hyperlipidemia⁶¹, hyperglycemia⁶², disturbed blood flow⁶³, hyperglycemia⁶⁴, and hyperhomocysteinemia^{64–67} plays significant roles in accelerating vascular inflammation and atherogenesis⁶⁸. Pathological endothelial activation and dysfunction⁶⁹ via generating a strong chemokine gradient are critical for recruiting inflammatory monocytes⁶². In our report, using monocyte adhesion assay, intravital microscopy, flow cytometry, and other biochemical assays as well as atherogenic apolipoprotein E deficient (Apo $E^{-/-}$) mouse model, we found that LPC-induced mtROS then contribute to aortic EC activation by regulating nuclear binding of activator protein-1 (AP-1) and inducing intercellular adhesion molecule-1 (ICAM-1) gene expression in vitro. Furthermore, mtROS inhibitor MitoTEMPO decreases aortic endothelial activation and aortic monocyte recruitment in atherogenic mice during early atherosclerosis *in vivo*²¹. In addition, along the same line, it has been shown that deficiency of superoxide dismutase 2 gene in $ApoE^{-/-}$ mice, which encodes a mitochondrial antioxidant protein, accelerates atherogenesis⁷⁰. Moreover, overexpression of thioredoxin 2 in ECs, which is another mitochondrial antioxidant enzyme, improves EC function and suppresses atherosclerosis development in $ApoE^{-/-}$ mice⁷¹. Furthermore, scavenging mtROS specifically in macrophages also decreases lesion formation in atherosclerotic mice⁷². Taken together, these studies suggested that in addition to promoting low scale physiological EC activation, mtROS promote atherosclerosis development by pathologically recruiting inflammatory cells to the lesions²¹.

Until recently, the important issue regarding how patrolling monocytes and other innate immune cells⁷³ cross endothelium and patrol the vessels and tissues under physiological homeostatic conditions remained much less understood⁷⁴. Although it has been reported that orphan nuclear receptor NR4A1 controls the differentiation of patrolling Ly6C-monocytes⁷⁵; and that lymphocyte function-associated antigen 1 (LFA1) mediates patrolling monocyte recruitment at homeostatic conditions⁷⁶, the molecular mechanisms underlying how ECs at non-dysfunctional/damaged conditions become activated and recruit patrolling monocytes are still unknown. Our study provided a framework which explains how mtROS could be increasingly generated independently from ATP synthesis and endothelial damage/ death²¹. We speculate that mtROS may be utilized by ECs to recruit patrolling monocyte for detecting and clearing harmful pathogens and malignant cells under normal physiological conditions, but this pathway is "hijacked" during pathological conditions such as atherosclerosis and autoimmune diseases, which mediates continuous inflammatory cell recruitment into arteries and tissues (Figure 1).

A new working model of physiological and pathological endothelial

activation

In summary (Figure 1), we proposed a new working model of EC activation: Under normal physiological conditions, low levels of DAMPs and our newly proposed HAMPs are important for mediating the patrolling immune cell adhesion and migration⁷⁷ by driving ATP-synthesis-uncoupled, but proton leak-coupled mtROS. As a result, mtROS are specifically induced for EC activation purpose without causing EC cell death. However, during pathological conditions such as atherosclerosis, higher levels of DAMPs lead to mtROS production that is coupled with decreased ATP synthesis and mitochondrial membrane potential (ψ m) collapse as a result of mitochondrial dysfunction. This leads to pathological EC activation with EC cell death, including apoptosis, pyroptosis²³ and pyropapoptosis (a hybrid form of pyroptosis and apoptosis) that we recently reported¹⁰. Consequently, this leads to pathological recruitment of Ly6C^{high} inflammatory monocytes^{64–66, 68}, dendritic cells^{77, 78}, B cells⁷⁹ and T cells⁸⁰. We propose that one of the major physiological functions of mitochondrial proton leak in ECs maybe fine-tuning mtROS generation for cell signaling purpose, but such mechanism is easily compromised in pathological conditions, which might lead to uncontrolled inflammation and cell death leading to the development of inflammatory diseases, such as atherosclerosis.

Concluding remarks and future directions

Further studies that map the connections between mitochondria and inflammation in ECs are of great interest, as endothelial mtROS seem to control patrolling cell immunosurveillance process under normal physiological conditions, which is important for controlling cancer cells and detecting infected pathogens. However, endothelial mtROS are "hijacked" during atherosclerosis development, which mediate harmful inflammatory cell migration to the lesions, contributing to early atherogenesis. Targeting mitochondrial metabolism thus may serve as novel therapies for cancers and inflammatory diseases including atherosclerosis. However, in order to translate this knowledge from benchtop to bedside, several imminent questions remain to be addressed: 1) Despite the importance of oxidative stress in the development of cardiovascular disease, antioxidant therapies such as Vitamin C and Vitamin E fail to demonstrate their therapeutic efficacy. This may be related to our incomplete understanding of the physiological and pathological roles of different ROS species originated from different subcellular organelles. In fact, only a small portion of known naturally occurring antioxidants *in vivo* are located in the mitochondria¹. Future studies that investigate the effects of mitochondria-targeting antioxidant in cardiovascular diseases are warranted. 2) It is still unknown how mitochondria are "hijacked" during pathological conditions and additional mitochondrial factors that contribute to diseases remain to be identified. 3) Potential side effects of mitochondria-targeting drugs need to be studied since mtROS also play significant housekeeping roles under normal physiological conditions. We hope that our review will encourage investigators to enter this important field and accelerate the pace of translational medicine.

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Figure 1. A new working model of physiological and pathological endothelial activation

Mitochondrial reactive oxygen species (mtROS), un-coupled from ATP synthesis, control endothelial cell activation in response to different levels of danger signals associated molecular pattern (DAMPs) stimulation. The levels of proton leak control the scales of mtROS generation in healthy mitochondria or dysfunctional mitochondria. *A*. MtROS, generated in modest levels of increased proton leak and uncompromised ATP synthesis and mitochondrial membrane potential (ψ m), control physiological EC activation and patrolling immune cell recruitment in response to low levels of DAMPs and our newly proposed anti-inflammatory homeostasis associated molecular patterns (HAMPs). *B*. MtROS, generated in high levels of increased proton leak, compromised ATP synthesis, and decreased ψ m, control pathological EC activation, EC pyroptosis (inflammatory cell death) and apoptosis, and increased proinflammatory cell recruitment in response to high levels of DAMPs.