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Mitochondrial Regulation of Airway Smooth Muscle Functions in Health and Pulmonary Diseases

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Abstract

Mitochondria are important for airway smooth muscle physiology due to their diverse yet interconnected roles in calcium handling, redox regulation, and cellular bioenergetics. Increasing evidence indicates that mitochondria dysfunction is intimately associated with airway diseases such as asthma, IPF and COPD. In these pathological conditions, increased mitochondrial ROS, altered bioenergetics profiles, and calcium mishandling contribute collectively to changes in cellular signaling, gene expression, and ultimately changes in airway smooth muscle contractile/proliferative properties. Therefore, understanding the basic features of airway smooth muscle mitochondria and their functional contribution to airway biology and pathology are key to developing novel therapeutics for airway diseases. This review summarizes the recent findings of airway smooth muscle mitochondria focusing on calcium homeostasis and redox regulation, two key determinants of physiological and pathological functions of airway smooth muscle.

Keywords

Airway smooth muscle; mitochondria; calcium; contraction; proliferation

Airway smooth muscle (ASM) resides in the trachea and bronchial tree up to the terminal bronchioles. It is the primary determinant of the airway pathophysiology via its regulation of diameter of the bronchi and bronchomotor tone. In addition to contractile function, ASM cells secrete extracellular matrix, cytokines/chemokines and undergo excessive proliferation, most notably under airway pathologies. To perform diverse functions, ASM transitions between a “mature” contractile phenotype and a proliferative/ synthetic phenotype characterized by a tendency to grow and/or synthesize extracellular matrix and other biologically active proteins. ASM plasticity plays a central role in both lung development and in disease conditions such as asthma, chronic bronchitis and emphysema [1, 2]. The modulation of this phenotypic switch is accomplished by a complex, coordinated interaction among external stimuli such as growth factors, extracellular matrix and ASM cells with

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external stimuli, eliciting a network of intracellular signaling cascades in ASM cells leading to transcriptional activation and protein translation of phenotype-specific genes. Each of the functions of ASM involves smooth muscle contractile proteins, cytoskeleton, intermediate filaments and organelles such as the sarcoplasmic (endoplasmic) reticulum and mitochondria. In this review, we discuss recent findings on ASM mitochondria focusing on their role in the most important aspects of ASM physiology and pathology, such as ASM contractility and proliferation/remodeling.

In recent years, mitochondria have been increasingly recognized as important organelles in ASM pathophysiology and homeostasis due to their indispensable roles in cellular energy production, redox regulation and calcium homeostasis. The dominant role of mitochondria is to produce the cellular energy in the form of ATP that is required for contraction, relaxation and synthetic functions of ASM. This is achieved via oxidative phosphorylation (OXPHOS) of glucose, pyruvate, and NAD and is known as aerobic respiration (Figure 1). Calcium-dependent dehydrogenases in the inner mitochondrial matrix reduce NAD^+ and FADH to NADH and FADH₂ in response to increases in mitochondrial calcium ($[\text{Ca}^{2+}]_m$) [3, 4]. The redox energy from NADH and FADH₂ is transferred to oxygen (O_2) via the electron transport chain (ETC). Following the acceptance of an electron, each complex in the ETC pumps protons (H^+) from the matrix into the intermembrane space. At the end of the ETC in complex IV, the two donated electrons are used to convert O_2 into H_2O [5–7]. The increase in the H^+ gradient in the intermembrane space, also known as the proton motive force, drives the influx of H^+ back into the matrix through the F_1F_0 ATP synthase proton pump. F_1F_0 ATP synthase uses the proton motive force to generate ATP from ADP and inorganic phosphate [5–7]. Newly synthesized ATP is transported into the intermembrane space in exchange for ADP through the adenine nucleotide translocator, where it is then transported across the outer mitochondrial membrane through the voltage-dependent anion channel (VDAC) to the cytosol [8]. Therefore, calcium concentration in the mitochondria is an important regulator of the ATP generating function of mitochondria. During the electron transfer, a small percentage of electrons leak from the ETC and reduce oxygen leading to the production of superoxide, a major form of mitochondrial reactive oxygen species (ROS). Although physiological levels of ROS can function as a second messenger for cell signaling, increased ROS production is detrimental causing DNA and protein damage, and ultimately cell death.

In addition to serving as a powerhouse of ASM cells and source of ROS, mitochondria also contribute to cellular calcium homeostasis by their ability to temporally store calcium. This ability of mitochondria makes them the temporal storage unit of cellular calcium besides the sarcoplasmic reticulum (SR) or endoplasmic reticulum (ER), which is a major calcium store inside of ASM cells. There is an interplay between the SR/ER and mitochondria with regard to calcium regulation in part due to anatomic juxtaposition. Increase in cytosolic calcium mitochondrial calcium uniporter (MCU) that resides in the inner mitochondrial membrane is responsible for the uptake of calcium into mitochondria matrix, while the release of calcium from mitochondria back to the cytosol occurs via a mitochondrial sodium-calcium exchanger (NCX) [9, 10]. Therefore, mitochondria function as the calcium buffering system of the cells by rapidly taking up and releasing calcium.

Taken together, the functional role of mitochondria in ATP synthesis, ROS generation and regulation of calcium homeostasis support the hypothesis that healthy mitochondria are indispensable for ASM physiology. Mitochondrial dysfunctions associated with decreased ATP, increased ROS and calcium mishandling contribute directly to pulmonary aging and lung diseases including pulmonary fibrosis (IPF), asthma and chronic obstructive pulmonary disease (COPD) [11–13].

Mitochondria in the regulation of Ca^{2+} dynamics and contractility in ASM

Calcium is a master regulator of contractility of ASM. As a secondary messenger, calcium transduces signals from the cell surface to enzymatic components inside of the cell. Therefore, Ca^{2+} is intimately involved in “excitation-contraction coupling” of ASM.

In a resting condition, ASM cells maintain relatively low intracellular Ca^{2+} concentration. Upon exposure of ASM cells to contractile agonists, the activation of membrane receptors such as G-protein-coupled receptors (GPCRs) triggers a signaling cascade followed by a rapid rise in intracellular calcium ($[\text{Ca}^{2+}]_i$). In ASM cells, a rise in intracellular Ca^{2+} concentration activates downstream contractile signaling. Calcium binds to calmodulin and the Ca-CaM complex activates calmodulin kinase, which in turn activates myosin light chain kinase (MLCK). Phosphorylation of myosin light chain by MLCK is a critical first step in the initiation of actin-myosin cross-bridge cycling and contraction. Among GPCRs, Gq-coupled receptors in ASM are of particular interest due to their prominent role in promoting ASM contraction (Figure 2).

Activation of contractile agonist receptors coupled to a Gq-type of G protein activates the β isoform of phospholipase C (PLC- β), which produces two second messengers, inositol 1,4,5-triphosphate (IP_3) and 1,2-diacylglycerol (DAG), via hydrolysis of phosphatidylinositol 4,5-bisphosphate (PIP_2). IP_3 diffuses through the cytosol and binds to its specific receptor (IP_3R) located on the SR, thereby mobilizing the IP_3 -sensitive pool of intracellularly stored Ca^{2+} , whereas DAG activates protein kinase C (PKC), which affects the sensitivity of the contractile apparatus to Ca^{2+} . However, recent identification of receptors such as Type II taste receptors (TAS2Rs), proton-sensing receptor (OGR-1) and gamma-aminobutyric acid (GABA) receptors on ASM cells has renewed interest in establishing the causal relationship between calcium elevation and contraction in ASM. Paradoxically, elevation of intracellular calcium by agonists belonging to TAS2Rs, OGR-1 and GABARs leads to relaxation of ASM cells [14–16]. In general, these functional diversities of calcium in ASM suggest signaling and functional compartmentalization. What role different compartments such as mitochondria play in calcium signaling compartmentalization in ASM cells needs further investigation.

The intracellular calcium response elicited by the contractile agonists is biphasic. The initial phase is a rapid and transient rise of calcium concentration from the baseline levels, which is responsible for ASM tension development. The rapid rise in ASM $[\text{Ca}^{2+}]_i$ then declines to a lower, sustained steady-state concentration slightly above the baseline level, which is associated with tension maintenance [17–20]. The initial calcium spike is due to a release of calcium from intracellular stores, the SR, induced by IP_3 , while the sustained phase of the

response originates from the calcium influx from the extracellular space [18, 19, 21–23]. The initial increase in calcium is coupled to calcium uptake into mitochondria and is necessary in order to drive ATP synthesis [24–27].

As a major player in the regulation of intracellular calcium, mitochondria are strategically localized either in close proximity of the IP₃R calcium release sites of the SR, or the calcium influx channels on the plasma membrane [25, 28]. IP₃-mediated rise of intracellular calcium results in large, rapid increases in [Ca²⁺]_m in parallel with cytosolic calcium ([Ca²⁺]_{cyto}) rises evoked by the contractile agonist [25]. Other mechanisms of elevating intracellular calcium such as calcium influx from the extracellular space has much less influence on [Ca²⁺]_m compared to the calcium rise mediated by IP₃ [25]. Furthermore, caffeine-induced calcium release from the SR (mediated primarily via ryanodine-sensitive SR stores) increases [Ca²⁺]_m [29] suggesting possible involvement of the ryanodine receptor in the regulation of mitochondrial calcium. Interestingly, kinetics analysis demonstrated that the increase in [Ca²⁺]_m coincides with the increase in [Ca²⁺]_i and remains elevated after [Ca²⁺]_i returns to resting levels. Earlier studies in various types of smooth muscle showed that agonist-induced [Ca²⁺]_i increase is almost fully inhibited when mitochondrial calcium uptake is prevented, suggesting an indispensable role for mitochondria in the regulation of intracellular calcium signaling [30–33]. The direct evidence for the role of mitochondria in calcium handling comes from studies using protonophore carbonyl cyanide *p*-trifluoromethoxy phenylhydrazone (FCCP), which is known to depolarize mitochondria. FCCP pretreatment attenuates caffeine-induced increase in [Ca²⁺]_m and increases the half-time for recovery of [Ca²⁺]_i to resting values [29]. More direct evidence was provided by using a system in which calcium release was evoked by photolysis of caged IP₃ instead of agonist stimulation. IP₃-induced calcium release was significantly reduced by inhibiting mitochondrial calcium uptake under conditions where ATP levels, PH, and plasma membrane potential were all controlled [30, 34–36]. Collectively, multiple lines of evidence suggest the critical role of mitochondria in calcium regulation in smooth muscle cells.

It has become clear that the mitochondrial calcium uptake is not simply a passive process. Mitochondria contain their own specific calcium transport systems. The uptake of calcium into the mitochondria is mediated by the MCU, the molecular identity of which has recently been unraveled. Calcium release from mitochondria is mediated by either the Na⁺/Ca²⁺ or H⁺/Ca²⁺ exchanger [37–41]. Recent studies suggest that the MCU has a higher threshold of activation in response to Ca²⁺ higher than 10 μM, whereas the [Ca²⁺]_{cyto} in response to agonist stimulation is around 2 μM [42]. One logical explanation is that MCUs are in close proximity to IP₃ gated channels on the SR, where [Ca²⁺]_{cyto} is high enough to be sensed by mitochondria. Perhaps this geographical orientation of mitochondria in proximity to calcium release channels on the SR can be called calcium “microdomains” or hotspots [25, 43–47]. In this theory, mitochondria-SR communications are strictly dependent on the spatial localization of mitochondria within the cells. Spatial and temporal organization of mitochondria in microdomains and the molecular mechanisms by which mitochondria sense the high concentrations of calcium remain to be determined in ASM cells.

Whether mitochondria act purely as a calcium sink or as a calcium relay mechanism is still not clear. Recent studies in ASM cells determining the effect of [Ca²⁺]_i clearance by either

the sarcoplasmic/endoplasmic reticulum calcium ATPase (SERCA) or mitochondria further support an important role for mitochondria in calcium handling [48]. The mathematical models of calcium regulation in ASM indicate that calcium is first sequestered by mitochondria before it is pumped back into the SR. Furthermore, the mitochondria uncoupler, carbonyl cyanide 4-(trifluoromethoxy) phenylhydrazone, but not SERCA inhibitor cyclopiazonic acid, significantly affected calcium decay after calcium release from the SR [48]. Taken together, by localizing to the calcium release site of the SR and rapidly taking up calcium, mitochondria could potentially modulate the rate of calcium rise as well as the magnitude of the calcium spike in response to a contractile agonist. On the other hand, the release of calcium from mitochondria back to the cytosol during the decay phase could delay the recovery of cytosolic calcium to baseline levels [49–53]. Therefore, mitochondrial calcium handling could contribute not only to the local calcium buffering but also to propagating and synchronizing calcium signals in ASM cells.

In ASM cells, acetylcholine (ACh)-induced contractile response is associated with regenerative and propagating Ca^{2+} oscillations [54]. The rise and sustainment of ACh-induced Ca^{2+} -oscillations involving IP_3 -induced Ca^{2+} release were initially observed in fluo3-loaded porcine ASM cells [55, 56]. Real-time spatial and temporal confocal imaging analysis indicated that within a small region of the cell, ACh-induced calcium oscillations are biphasic and are propagating throughout the cells. The oscillations originate from either one end or in the middle of the cell, and propagate to the other or both ends of the cell. Increases in ACh concentration increase frequency and propagation velocity of calcium oscillations without affecting the peaks [54]. In contrast, the global spatial and temporal integration of the regional oscillation responses indicate an ACh concentration–dependent increase in both peak and mean cellular calcium concentrations. This pattern of calcium elevation in ASM cells involves the coordinated actions of calcium influx from the extracellular space via plasma membrane ion channels and release of calcium from the intracellular calcium store. What role, if any, mitochondria have in the regulation of calcium oscillations in ASM cells is not known. Previous studies indicate that initiation of calcium oscillations by ACh requires activation of the IP_3 Rs in the SR [57]. The increase in $[\text{Ca}^{2+}]_i$ promotes the binding of Ca^{2+} to the IP_3 Rs. This enhances the open probability of IP_3 Rs leading to a continuous rise of the $[\text{Ca}^{2+}]_i$, which will stimulate neighboring IP_3 Rs and further propagate a Ca^{2+} wave. Although refilling the SR with Ca^{2+} reestablishes the Ca^{2+} driving force, the SR $[\text{Ca}^{2+}]$ may also be critical for the regulation of the SR release channel. When Ca^{2+} binds to a second site on the IP_3 Rs, the additional binding of calcium reduces open probability of calcium release channels on SR. The localized decrease in $[\text{Ca}^{2+}]$ of the SR and decreased open probability together terminate Ca^{2+} release from the SR.

While IP_3 -based Ca^{2+} oscillations require an IP_3 -induced Ca^{2+} release pathway, caffeine-induced Ca^{2+} release involves a mechanism independent of the IP_3 Rs [55, 56]. ACh-induced repetitive, propagating Ca^{2+} oscillations in isolated porcine tracheal SMCs were attenuated by ryanodine and caffeine [58] suggesting the involvement of ryanodine receptors (RyRs) in calcium oscillations. IP_3 appears to be indispensable in the initiation of the Ca^{2+} oscillations. However, IP_3 alone could not maintain Ca^{2+} oscillations. The maintenance of the Ca^{2+} oscillations requires the RyRs [58] and this involves activation of RyRs via a calcium-induced calcium release (CICR) mechanism [59]. Ryanodine pretreatment not only inhibited

ACh-induced Ca^{2+} oscillations in muscle bundles from porcine or human airways, but also attenuated carbachol-induced contraction of mouse tracheal rings [60–62] suggesting the functional role of RyRs in the regulation of ASM contraction. Similar results were found using other less specific inhibitors of the RyR, such as procaine and tetracaine. Although in other cell types, Ca^{2+} oscillations are thought to occur via the IP_3R , in ASM cells, both IP_3R and RyRs appear to play role in the regulation of Ca^{2+} oscillations. Additionally, in ASM cells, NAD metabolites such as cyclic ADP-ribose (cADPR) and nicotinic acid dinucleotide phosphate (NAADP) are known to regulate calcium release from the SR [63].

Although mitochondria can buffer calcium, it is puzzling that the mitochondrial calcium uptake fails to decrease cytosolic calcium during agonist-induced calcium oscillations [64, 25, 65]. The possible explanation is that calcium taken up by mitochondria may limit the chance of negative feedback regulation, which prevents the accumulation of high enough levels of local calcium or locations that would inhibit the open probabilities of IP_3R . Nevertheless, mitochondria have the capability to modulate the activity of intracellular calcium release channel and store-operated, calcium influx channels [66, 67]. In colonic smooth muscle cells, inhibition of mitochondrial calcium uptake but not efflux impeded IP_3R sensitive calcium release [68]. It is not clear whether mitochondria play a role in the regulation of calcium homeostasis mechanisms involving CICR or via NAD metabolites in ASM cells.

In order to sustain agonist-induced contraction of ASM, influx of extracellular Ca^{2+} is required. Ca^{2+} oscillations arise from intracellular stores and are sustained by calcium influx from the extracellular space [69]. Calcium influx can occur through a variety of mechanisms including voltage- and receptor-operated membrane channels, and store-operated calcium entry (SOCE) [17, 23, 70–73]. In striated muscle, excitation-contraction coupling is often dependent on membrane depolarization via opening of voltage-dependent Ca^{2+} channels. However, voltage- dependent Ca^{2+} channels do not appear to strongly influence membrane depolarization and refilling of internal stores in ASM cells [74] suggesting other mechanisms for Ca^{2+} entry to maintain ASM contraction. Store depletion-induced Ca^{2+} entry via store-operated Ca^{2+} channels (SOCs) is a common mechanism to activate Ca^{2+} influx via plasma membrane of ASM [75, 76]. The mechanistic link between store depletion and influx involves stromal interacting molecule (STIM) 1, and its communication with a cell membrane protein or channel (Orai1) [77, 78]. STIM1 is expressed and plays a role in agonist-induced Ca^{2+} influx in human ASM [76]. STIM1 is a potential Ca^{2+} sensor in the SR and has a SR luminal Ca^{2+} binding domain. Upon store emptying, STIM1 relocates and forms clusters adjacent to the plasma membrane. However, the mechanism by which STIM1 is linked to the membrane channel activation remains unclear [76]. Another major route for Ca^{2+} influx involves calcium release-activated calcium channels (CRAC) that are activated when the Ca^{2+} content within the SR/ER falls. Originally, SERCA pump activity was considered one of the major determinants of CRAC activities because when SERCA pump activity is decreased, CRAC current develops. Recent studies in T cells indicate that mitochondria also play a key role in the regulation of this Ca^{2+} influx pathway. Following Ca^{2+} release from the store (ER), mitochondria rapidly take up calcium, which results in further Ca^{2+} depletion of the store with concomitant activation of CRAC channels [79–82]. Ca^{2+} uptake by mitochondria increases the sensitivity of store-operated Ca^{2+} entry [81]. The

mechanisms remain unclear, but it appears that mitochondria reduce Ca^{2+} -dependent inactivation of CRAC channels via microtubule-dependent translocation of mitochondria to the plasma membrane [79, 80]. Inhibiting mitochondrial movement abolishes mitochondrial plasma membrane translocation and reduces sustained Ca^{2+} signals [79]. The physiological significance of these phenomena is still unclear particularly in ASM cells, but they all support a role for mitochondria in Ca^{2+} signaling by modulating Ca^{2+} transport pathways.

Mitochondria in the regulation of ASM proliferation

Increase of bronchial smooth muscle mass characterized by hyperplasia and hypertrophy of ASM is a key feature of remodeling in asthma. The underlying mechanisms are complicated but mitochondria play an important role. Under pathological conditions, elevation of intracellular Ca^{2+} levels are associated with subsequent activation of Ca^{2+} -calmodulin-calcineurin-NFAT signaling cascade leading to gene expression and increased cell proliferation [83–85]. Nuclear factor of activated T cells (NFATs) are a family of transcription factors that play a central role in the proliferation of a variety of cell types and functions of organ systems. Calcium activates calmodulin and calcineurin, resulting in dephosphorylation and translocation of NFAT from the cytosol to the nucleus, where it functions as the transcription factor to induce gene expression. Interestingly, NFAT translocation is exquisitely sensitive to Ca^{2+} microdomains near open store-operated CRAC channels [83]. Kinetics linking CRAC channel activation to NFAT nuclear localization indicate that the rate-limiting step in calcium-mediated gene expression is the exit of NFAT from the nucleus (not the sustained calcium elevation in the cytosol). These findings suggest a novel mechanism by which Ca^{2+} -dependent responses can be sustained despite the termination of the initial Ca^{2+} signal to allow gene expression to continue after the primary stimulus has been removed [83]. In ASM cells, activation of vanilloid transient receptor potential channel (TRPV4) by synthetic and endogenous TRPV4 agonists increase proliferation of ASM cells. Additional studies suggest that NFAT is activated upon stimulation of TRPV4 and this presumably involves colocalization of TRPV4 and calcineurin in calcium microdomains. Translocation of activated NFAT leads to induction of ASM cell proliferation [84]. Recent studies also suggest a role for microRNA miR-143–3p in the regulation of ASM cell proliferation in an NFAT-dependent manner [85]. Levels of miR-143–3p were found to be significantly decreased in ASM cells from asthmatic patients compared to non-asthmatic individuals. As a negative regulator of NFATc1, miR-143–3p overexpression significantly decreased transforming growth factor beta (TGF- β)-induced ASM proliferation [85].

While the rise of intracellular Ca^{2+} concentration in response to agonist stimulation is essential for ASM contraction, restoration of normal basal Ca^{2+} concentration and preventing long-lasting rise of Ca^{2+} could be critical for ASM proliferation. Excessive cytoplasmic Ca^{2+} is taken up into the SR by SERCA or extruded to the extracellular space by plasma membrane Ca^{2+} -ATPase (PMCA) and NCX [86–89]. Recent studies indicate that PMCA is likely involved in the regulation of ASM proliferation via a calcium-dependent mechanism [90]. Inhibiting PMCA decreased the decay rate of the bradykinin-induced Ca^{2+} transient. This impaired Ca^{2+} clearance caused sustained elevation of intracellular Ca^{2+} and ASM proliferation suggesting the role of PMCA in ASM cell proliferation [90]. Like in

many other cell types, SERCA is the major mechanism in human ASM for replenishing calcium stores. In human ASM cells, SERCA is regulated by calcium/calmodulin-dependent protein kinase II (CaMKII)-related mechanisms and Inflammation maintains intracellular Ca^{2+} levels by decreasing SERCA expression and attenuating Ca^{2+} reuptake into SR. Because Ca^{2+} levels in ASM are increased in asthma, these studies suggest that decreased SERCA expression and function could be the mechanism for the increase in intracellular Ca^{2+} [91, 92]. Recent studies also suggest a role for SERCA in the regulation of ASM proliferation and remodeling. Both SERCA2 mRNA and protein levels are reduced in ASM cells of asthmatic patients [93]. The decrease in SERCA2 levels is associated with attenuation of agonist-induced rise of intracellular Ca^{2+} and a delay in returning to the baseline level of Ca^{2+} . Knocking down SERCA2 by siRNA approaches increased cell spreading and more importantly ASM proliferation [93]. These findings suggest that the decrease in SERCA2 expression may happen during early stage of asthma. This decrease in SERCA2 expression and function decreases Ca^{2+} reuptake into SR, leading to increased intracellular Ca^{2+} levels, which may contribute to the ASM phenotypic switch and progression of asthma.

As the key regulator of intracellular calcium, mitochondrial dysfunction contributes to ASM cell proliferation and remodeling. There is increasing evidence that cross talk between mitochondrial calcium buffering and SR calcium homeostasis plays an important role in ASM proliferation. Inflammation and cigarette smoke cause mitochondrial dysfunction which impair the calcium buffering capacity of mitochondria [94, 95]. The decrease in Ca^{2+} buffering leads to an increase in intracellular Ca^{2+} level and presumably ASM proliferation. Mitochondrial dysfunction also increases mitochondrial ROS generation, resulting in ER/SR stress and breaks in Ca^{2+} homeostasis in ASM. While the majority of the studies involving patients of different disease severity indicated that elevated intracellular calcium concentration contributes to the hyperproliferative phenotype of ASM cells in asthma, disrupted calcium balance changes mitochondrial biogenesis, potentially leading to excessive ASM proliferation. In support of this hypothesis recent studies have demonstrated that ASM cells from asthmatic patients have an increased mitochondrial number and oxygen consumption, indicating increased mitochondrial biogenesis [96]. Furthermore, in severe asthma, altered calcium handling involves phosphorylation of calcium/calmodulin-dependent protein kinase IV (CaMK-IV) and subsequent activation of peroxisome proliferator-activated receptor γ coactivator-1 α (PGC1- α), nuclear respiratory factor-1 (NRF-1) and mitochondrial transcription factor A (mtTFA) [97]. mtTFA is a transcription factor playing a key role in the regulation of mitochondrial DNA replication. These findings further suggest increased biogenesis of mitochondria in asthmatic ASM cells. In addition, mitochondrial dynamics and mitochondrial dynamics-related proteins are closely related to the intracellular Ca^{2+} levels. Increased intracellular and mitochondrial Ca^{2+} enhance mitochondrial fission [98] and are associated with changes in the expression of mitochondrial fission and fusion proteins. In fact, mitofusin (Mfn) 2, also called hyperplasia suppressor gene, is implicated not only in mitochondria fusion and tethering of mitochondrial and ER/SR membranes, but also in cell proliferation [99, 100]. These observations reiterate the hypothesis that mitochondrial mechanisms play a central role in the regulation of ASM cell proliferation in airway diseases.

Although enhanced airway contractility and airway remodeling are the two main pathological features of asthma, the role of oxygen (hypoxia and hyperoxia) in modulating ASM structure and function of the airway during these processes is not known. Recent studies demonstrate that human fetal ASM cells respond to oxygen in a dose-dependent manner, which could contribute to childhood airway disease [101]. In fetal ASM cells, both acetylcholine and histamine induced calcium mobilization and agonist-induced calcium response was augmented upon exposure to increased oxygen up to 60%. However, exposure of fetal ASM cells to more than 60% oxygen diminished histamine- and acetylcholine-induced calcium mobilization. Further, fetal ASM showed enhanced proliferation at baseline compared to adult ASM cells. Proliferation of fetal ASM was increased by hyperoxia up to 60% whereas exposure to more than 60% oxygen resulted in diminished proliferation coupled with enhanced apoptosis. Interestingly, mitochondrial fusion was increased in response to higher than 60% of oxygen, supporting a role for fetal ASM mitochondria in not only sensing the oxygen but also assessing the risks of supplemental oxygen in prematurity [101].

Mitochondria in the regulation of ROS-mediated effects in ASM

ROS regulates many aspects of ASM biology including contraction, proliferation and migration. As in many other cell types, mitochondria are one of the major sites of ROS production in ASM. Electrons leaked from ETC during electron transfer react with oxygen to generate ROS in the form of superoxide (O_2^-). Mitochondrial ROS (mtROS) level is regulated by antioxidant enzymes such as superoxide dismutase and catalase that can eliminate ROS by converting O_2^- to hydrogen peroxide (H_2O_2) and ultimately to harmless H_2O . Other major determinants of mtROS are the redox status of ETC, and electrical and chemical gradients [102–104]. However, the mechanism by which electrical/chemical gradients regulate ROS generation is not well understood. Although mtROS can be biologically important in a variety of physiological processes including cell signaling and autophagy, mtROS overproduction is detrimental.

ROS can cause cellular damage and impair ASM contraction [105] (Pan unpublished data). The consequence of ROS on ASM contractility is dependent on the dose, the duration of exposure and the type of species of ROS [105–107]. For example, treating canine distal airway strips or bovine tracheal strips with hydrogen peroxide (10^{-5} – 10^{-3} M) resulted in ASM contractions that were partially reversed by oxygen radical scavengers, suggesting the contraction is mediated by ROS [106]. In a separate study, it was demonstrated that H_2O_2 treatment contracts trachealis muscle in a concentration-dependent manner beginning at 1 mM, which is not attributable to the release of prostaglandins, acetylcholine, or other neurotransmitters. On the contrary, 0.1 mM or more of H_2O_2 attenuated the responses to electrical field stimulator, acetylcholine, and KCl in a concentration-dependent manner in another study [108]. It appears that at high concentrations, H_2O_2 affected the responses of ASM by actions on neurotransmission, muscarinic receptors, and downstream components. However, electric field induced-contractile responses of isolated bronchi are enhanced after exposure to H_2O_2 in rats [108]. Exposure of rabbits or cats to ROS-generating enzyme systems leads to enhanced contraction to acetylcholine *in vitro* and *in vivo* [109, 110]. In addition, ozone and H_2O_2 are able to induce increases in airway responsiveness to

acetylcholine and bradykinin [111, 112]. Interestingly, ROS can decrease function and number of β -adrenergic receptors present in lung tissue (presumably decreasing the ability of ASM to relax), which can be reversed by superoxide dismutase [113–117]. In asthmatic ASM *in vivo* and in primary ASM culture, the generation of ROS is increased with an associated increase in the expression of adenine dinucleotide phosphate oxidase type 4 (NOX4) enzyme, resulting in oxidative stress in ASM cells. This increase in oxidative stress is correlated to the degree of airflow obstruction and airway hyperresponsiveness (AHR) [118, 119]. More importantly, the antioxidant mitubiquinone mesylate (MitoQ) reverses ozone-induced AHR and inflammation, suggesting a key role for mitochondrial ROS in the development of these pathological conditions [11]. Increased ROS generation damages cellular DNA, carbohydrates, proteins and lipids leading to increased inflammatory responses [120, 121]. Furthermore, increased oxidative stress damages mitochondria such as in patients with COPD in which mitochondrial dysfunction drives inflammation and ASM remodeling [11]. Mice exposed to oxidative stress develop lung inflammation and AHR, similar to that observed in patients with COPD, which is associated with decreased mitochondrial membrane potential and increased mitochondria ROS generation. All these studies clearly demonstrate the role of mitochondria in the regulation of oxidative stress and ROS generation in airway resident cells such as ASM cells leading to increased airway inflammation, remodeling and AHR (Figure 3).

Exposure of cultured ASM cells to mitogenic stimuli switches ASM cells from a contractile, hypoproliferative to a proliferative, hypocontractile phenotype, as indicated by changes in molecular phenotypic markers [122, 123]. Many studies suggest that ROS can regulate cell proliferation [124–126]. Alternatively, in other cell types such as in tumors, although proliferating phenotypes are associated with low levels of mtROS, the role for mtROS in ASM cell proliferation is not well characterized. Cell proliferation and migration begin with stimulation of cell surface receptors and the transduction of the external signals to a series of cellular responses. ERK and PI3K activation appear to be the dominant signal transduction pathways for RTK-, GPCR-, or cytokine-stimulated ASM growth [127]. Accumulating evidence indicates that ROS are important modulators of mitogenic signaling via several different mechanisms. First, ROS can function as the second messenger in signal transduction [128]. Secondly, ROS regulates cell cycle progression by directly modulating the phosphorylation and ubiquitination of cell cycle-related enzymes [128]. Finally, ROS can activate growth factor receptors in the absence of the growth factor ligands leading to cell proliferation [129]. Antioxidants such as N-acetylcysteine, catalase and probucol significantly reduce ASM proliferation induced by fetal bovine serum or platelet-derived growth factor suggesting that ROS is involved in mitogenic stimulation of ASM cell [130].

In the pulmonary artery, smooth muscle cells switch from OXPHOS to glycolysis and activation of mitochondrial fission protein dynamin-like protein 1 (DLP1) is associated with the hyperproliferative phenotype in smooth muscle cells in pulmonary arterial hypertension [131]. However, little is known about the role of metabolic shift in ASM cells during airway inflammatory conditions.

Increased ROS generation increases mitochondria fragmentation and mitochondrial dysfunction. When this happens, damaged mitochondria are removed by autophagy.

shift of human ASM cells into a synthetic state, which is characterized by increased mitochondrial fragmentation and cellular proliferation [3].

Defects in mitochondrial function have been identified as the cause for onset of smoking related diseases like COPD [143]. Cigarette smoke introduces over 5000 chemical compounds, which contain ROS such as H_2O_2 , O_2^- , and hydroxyl free radicals. Smokers have also been shown to have elevated levels of endogenous ROS production in addition to exogenous sources of ROS in cigarette smoke. Direct evidence for cigarette smoke-induced structural and functional changes in ASM came from a recent study that demonstrated cigarette smoke-induced mitochondrial fragmentation and decreased oxygen consumption rate in ASM cells [95].

Mitochondrial DNA (mtDNA) is separate and unique from nuclear DNA. Many mtDNA encoded proteins are involved in the repair of oxidative damage of cellular proteins from ROS and reactive nitrogen species (RNS), which are byproducts of oxidative phosphorylation.¹³ Proteins involved in the maintenance of mtDNA, such as superoxide dismutases, are encoded by nuclear genes. Disruption of the dynamic interaction and crosstalk between mitochondria and the SR can lead to defects in mtDNA repair mechanisms, elevated levels of $[\text{Ca}^{2+}]_{\text{cyto}}$ and increased ROS production [3, 144]. Furthermore, elevated levels of ROS can overwhelm cellular antioxidant responses and increase the activation of transcription factors involved in inflammation such as nuclear factor kappa-light-chain-enhancer of activated B cells (NF- κ B) and NRF-2¹³². These observations suggest that ablation of the interplay between mitochondria and the nucleus plays a key role in the disruption of mitochondrial energetics, cell fate determination and pathogenesis of many human disorders including asthma, COPD, neurodegenerative diseases and cancer¹³².

Mitochondria in hypoxia-mediated effects on ASM

ASM cells are dependent upon the diffusion of oxygen across the basement membrane or microvessels within the connective tissue of the airway wall [145]. Asthmatic lung tissue is susceptible to hypoxia due to the thickening of the basement membrane [145, 146]. Increased levels of ROS during hypoxic stress following bronchoconstriction can lead to expression of transcription factors belonging to the hypoxia inducible factor (HIF) family [145]. HIF-1 regulates ATP production by suppressing the ETC and driving mitochondria to switch from OXPHOS to β -oxidation of fatty acids for energy during hypoxia [144]. Under normal conditions, HIF1- α is hydroxylated at two proline residues and targeted for proteasomal degradation via ubiquitinylation by Von Hippel-Lindau protein (VHL) and elongin C (Eloc) [145]. Under hypoxic conditions such as those witnessed in asthmatic ASM, HIF1- α hydroxylation is impaired, allowing HIF1- α to form a heterodimeric complex with HIF1- β and bind to hypoxia response elements in the nucleus [145]. The currently known gene targets downstream of these response elements are known to play a role in angiogenesis, erythropoiesis, ventilation, glycolysis and autophagy [145] (Figure 4).

Consistently elevated HIF-1 levels are known to contribute to hyperresponsiveness in asthmatic ASM [147]. ASM proliferation, and the subsequent rise in number of

mitochondria, increases the oxygen demand necessary to fuel ATP production. Oxygen deprivation of ASM due to thickening of the basement membrane leads to HIF-mediated angiogenesis of microvessels within the submucosa of the airway wall [145, 146]. The airway submucosa of asthmatic patients has been observed to have higher levels of angiogenic factors such as VEGF, EGF and angiopoietin in response to pro-inflammatory cytokines like IL-13 or TGF- β [148]. However, the structural integrity in the endothelium of these newly formed microvessels is often poor, as similarly observed in the hypoxic environment of tumors [149]. Degradation of tight junctions within the endothelium of these vessels allows excess fluid and immune cells to leak into the submucosa of the airway wall, resulting in edema and localized inflammation [145]. Mitochondria are pivotal in the regulation of multiple pathogenic processes emanating from the physiological response of mitochondria to hypoxia and energy demand of the cell.

Recent literature suggests that platelets may also play a key role in inflammation and airway remodeling due to increased platelet reactivity in asthmatic individuals [150]. Immunologic and nonimmunologic depletion of platelets in ovalbumin (OVA)-sensitized mice resulted in significant attenuation of ASM thickening and extracellular matrix deposition [150]. Thrombi have been shown to release growth factors that facilitate the smooth muscle growth. Furthermore, chronically challenged OVA-mice suffered from airway remodeling in the absence of eosinophils, suggesting that air remodeling may persist even following the attenuation of an inflammatory immune response [150]. Platelets are able to directly interact with eosinophils that buildup in airway tissue following hypoxic angiogenesis, further reinforcing the immune-mediated inflammatory response [149]. Although mitochondrial number and morphology remained unchanged in platelets of asthmatics, expression of citric acid cycle enzymes aconitase, succinate dehydrogenase and citrate synthase are upregulated in platelet mitochondria in asthmatics compared to healthy individuals under hypoxic conditions [149]. This dynamic difference in mitochondrial function allows for increased ATP and ROS production despite restricted oxygen intake in individuals with asthma. Despite these findings, the mechanisms by which immunological cell-derived ATP, growth factor and cytokine production affect ASM proliferation remain to be elucidated [150]. Further investigation is also necessary to determine the importance of platelets and immune cell crosstalk on ASM mitochondrial energy dynamics and proliferation in the pathogenesis diseases such as asthma and COPD.

Taken together studies to date suggest that healthy mitochondria are essential for normal structure and function of the airway cells, which is central to lung physiology. Mitochondrial dysfunction associated with aberrant mitochondrial dynamics and excessive mitochondrial ROS generation can be used as markers, as well as targets in the pathogenesis of lung diseases. One such example is that the administration of mitochondrial targeted antioxidant, MitoTEMPO (2-(2,2,6,6-Tetramethylpiperidin-1-oxyl-4-ylamino)-2-oxoethyl) triphenylphosphonium chloride) in mouse attenuated neonatal hyperoxia-induced lung injury [151]. Similarly, MitoTEMPO significantly decreased mitochondrial ROS generation, TGF- β , and collagen deposition in OVA-challenged mice [152]. In a mouse model of COPD, both a mitochondrial antioxidant MitoQ, and an inhibitor of DLP1 which blocks mitochondrial fission, reversed the adverse effect of cigarette smoking on mitophagy [153]. However, the direct evidence connecting various lung diseases and mitochondria abnormalities is missing.

Whether mitochondrial changes are cause or effect of lung diseases needs to be established. Additional studies on targeting drugs precisely to mitochondria may provide novel therapeutic strategies for lung diseases.

Summary

Chronic inflammation in the lungs leads to progressive structural and functional changes in airway cells including ASM cells resulting in bronchoconstriction. The cellular and molecular pathogenesis of airway diseases includes the progression of ASM from a mature, contractile phenotype into a synthetic and hyperproliferative phenotype. The importance of changes in calcium mobilization, calcium sensitivity and cellular bioenergetics in asthmatic ASM has been well established in the literature. However, there is ever increasing evidence implicating the role of mitochondria in the regulation of these cellular dynamics.

Inflammation, hypoxia, cigarette smoke are major causes of mitochondrial dysfunction including mitochondrial fragmentation, uncoupling from the SR and change in mitochondrial membrane potential all of which lead to increased ROS production and a greater amount of stress in ASM cells. These phenomena play significant roles in ASM cell proliferation (fragmentation) and calcium hypersensitivity (mitochondrial uncoupling).

A deeper understanding of the importance of mitochondria in asthma and COPD open new avenues for drug discovery. Mitochondria-specific antioxidants such as MitoQ and MitoTEMPO have been shown to be effective in reversing mitochondrial dysfunction, suppressing ROS and preventing oxidative damage in *in vitro* studies and animal models [154]. Despite the need to further elucidate the specific role of mitochondrial dysfunction in the pathology of AHR, remodeling and inflammation, mitochondrial-specific therapeutics represent a potentially effective option to eventually be used in place of or in tandem with the current standard of care.

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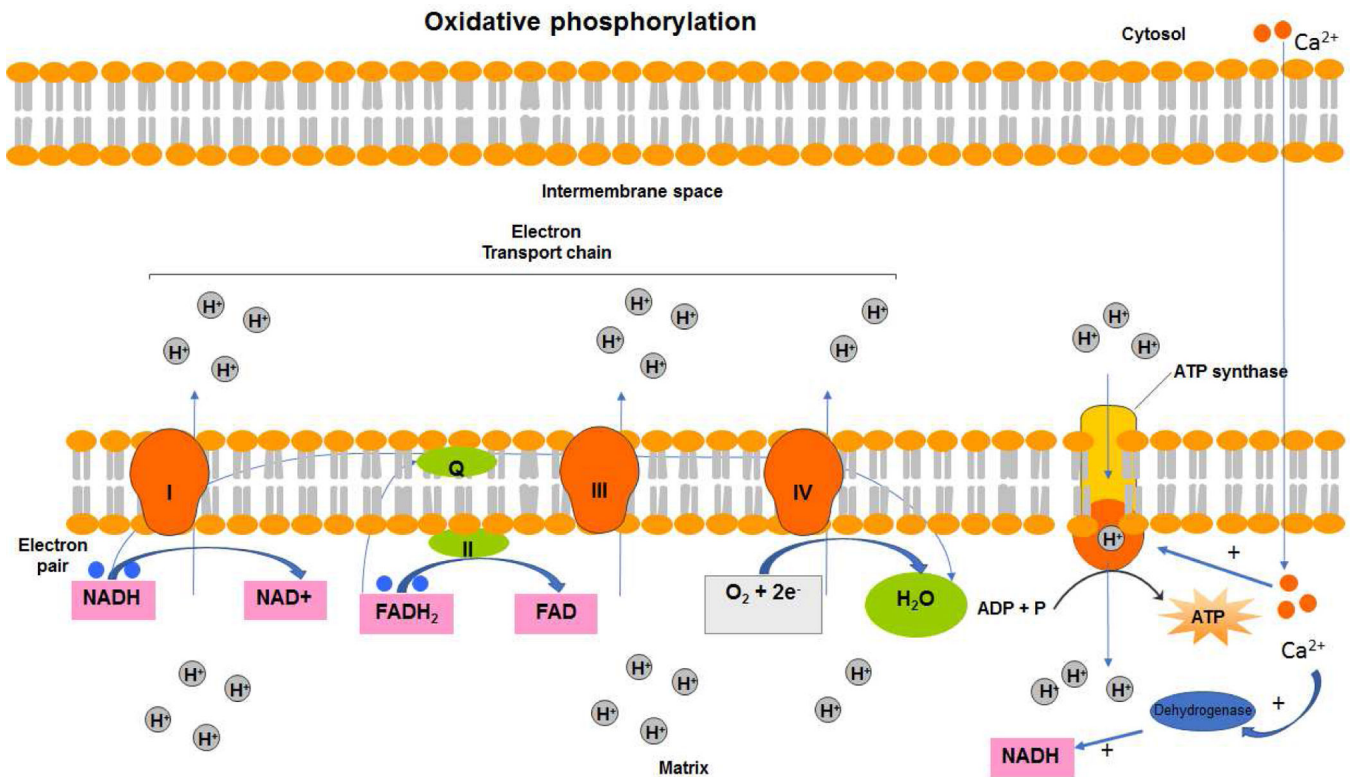


Figure 1: Oxidative phosphorylation.

Mitochondria are responsible for energy production through the process of cellular respiration. The reduction of each complex in the electron transport chain allows for protons to be pumped into the intermembrane space. The buildup of protons in its space eventually creates a concentration gradient known as the proton motive force. The influx of protons back into the mitochondrial matrix through the proton pump, F₁F₀ ATP synthase, drives the conversion of ADP and inorganic phosphate into ATP, which is an important source of energy for smooth muscle contraction. Activation of multiple mitochondrial enzymes is dependent upon calcium concentration in the mitochondria. Uptake of calcium from cytosol into mitochondrial matrix positively regulate ATP synthesis.

G protein-coupled receptor mediated Ca^{2+} signaling

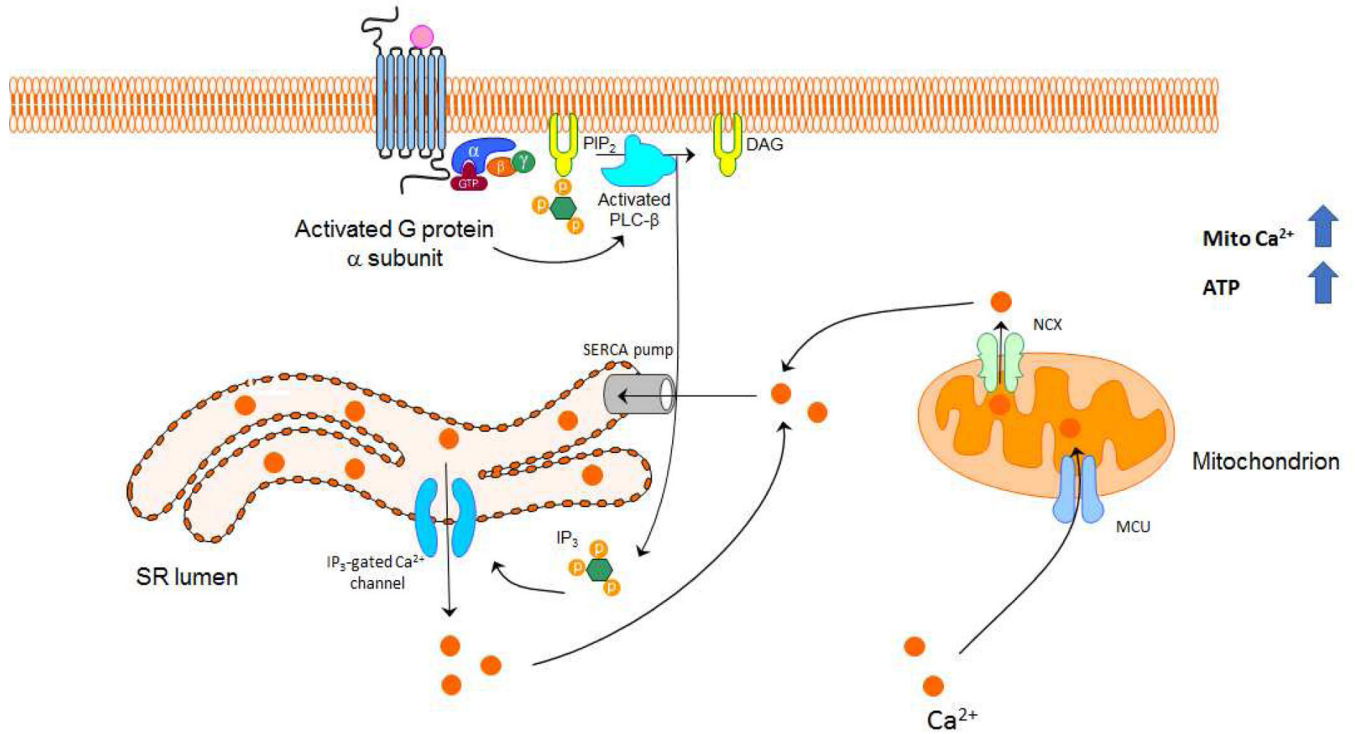


Figure 2: G-protein mediated Ca^{2+} mobilization.

ASM contraction and relaxation are dependent on changes in cytosolic and mitochondrial calcium. GPCR stimulation and subsequent G protein activation leads to the activation of PLC- β . PIP₂ is cleaved by PLC- β into IP₃ and DAG. IP₃ binds to IP₃-gated Ca^{2+} channels on the SERCA membrane, causing these channels to open and allow Ca^{2+} to be released into the cytosol. Mitochondria which are in close proximity to the SR are able to uptake Ca^{2+} through the MCU, where it can be used by calcium-dependent dehydrogenases in the TCA cycle. Transient mitochondrial calcium fluxes thus lead to the increases in ATP production necessary for both temporary and prolonged ASM contraction.

Effects of ROS/RNS on mitochondrial function

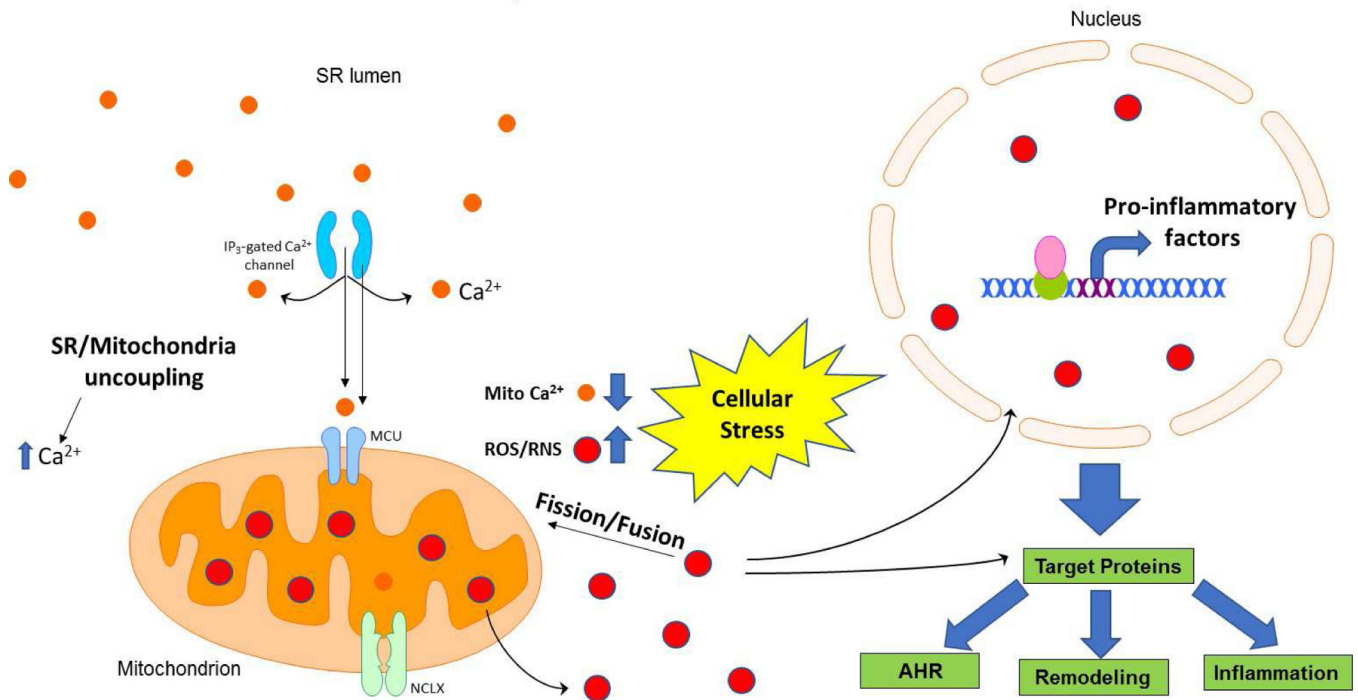


Figure 3: Role of ROS/RNS in mitochondrial dysfunction.

ASM contraction and relaxation are dependent on changes in cytosolic and mitochondrial calcium. Uncoupling of mitochondria from the ER/SR membrane leads to reduced mitochondrial Ca^{2+} flux, stress-related increases in ROS/RNS and reduced degradation of ROS. Increased ROS/RNS act as a second messenger or chemically modulate the functional ability of ASM proteins and results in increased airway inflammation, airway hyperresponsiveness (AHR) and ASM remodeling.

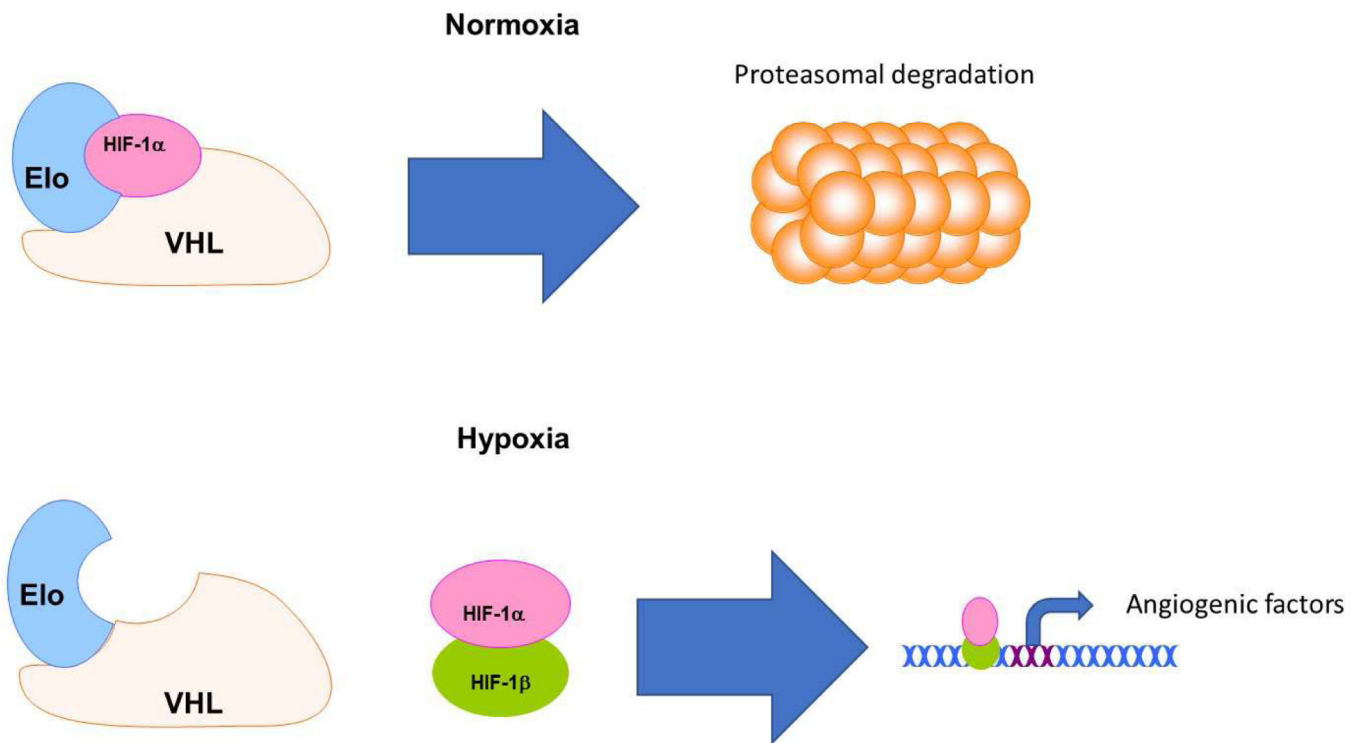


Figure 4: Hypoxia and airway smooth muscle remodeling.

Under non-hypoxic conditions, HIF-1 α is bound in a complex with VHL and Elo, hydroxylated at two proline residues and targeted for proteasomal degradation via E3 ligase-mediated ubiquitinylation. However, chronically inflamed lung tissue is susceptible to oxygen deprivation due to thickening of the basement membrane. Hypoxia allows unbound HIF-1 α to form a heterodimer with HIF- β , translocate into the nucleus and binds to hypoxia response elements. Upregulation of these response elements plays a major role in angiogenesis in the extracellular matrix of asthmatic lung tissue.