



Original Contribution

Primary role of mitochondrial Rieske iron–sulfur protein in hypoxic ROS production in pulmonary artery myocytes

Amit S. Korde, Vishal R. Yadav, Yun-Min Zheng, Yong-Xiao Wang *

Center for Cardiovascular Sciences, Albany Medical College, Albany, NY 12208, USA

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ABSTRACT

This study was designed to determine whether: (1) hypoxia could directly affect ROS production in isolated mitochondria and mitochondrial complex III from pulmonary artery smooth muscle cells (PASMCs) and (2) Rieske iron–sulfur protein in complex III might mediate hypoxic ROS production, leading to hypoxic pulmonary vasoconstriction (HPV). Our data, for the first time, demonstrate that hypoxia significantly enhances ROS production, measured by the standard ROS indicator dichlorodihydrofluorescein/diacetate, in isolated mitochondria from PASMCs. Studies using the newly developed, specific ROS biosensor pHyPer have found that hypoxia increases mitochondrial ROS generation in isolated PASMCs as well. Hypoxic ROS production has also been observed in isolated complex III. Rieske iron–sulfur protein silencing using siRNA abolishes the hypoxic ROS formation in isolated PASM complex III, mitochondria, and cells, whereas Rieske iron–sulfur protein overexpression produces the opposite effect. Rieske iron–sulfur protein silencing inhibits the hypoxic increase in $[Ca^{2+}]_i$ in PASMCs and hypoxic vasoconstriction in isolated PAs. These findings together provide novel evidence that mitochondria are the direct hypoxic targets in PASMCs, in which Rieske iron–sulfur protein in complex III may serve as an essential, primary molecule that mediates the hypoxic ROS generation, leading to an increase in intracellular Ca^{2+} in PASMCs and HPV.

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Pulmonary arteries (PAs) contract in response to hypoxia, delineated as hypoxic pulmonary vasoconstriction (HPV). This unique response is an important physiological process that optimizes the matching of regional alveolar ventilation to pulmonary perfusion in the lungs, but may also serve as a major pathological factor in the development of pulmonary hypertension. A crucial factor for HPV is an increase in intracellular Ca^{2+} concentration ($[Ca^{2+}]_i$) in PA smooth muscle cells (PASMCs) [1].

Pharmacological and genetic studies in PASM cells and tissues indicate that the hypoxic increase in $[Ca^{2+}]_i$ in PASMCs and associated HPV are mediated by ROS signaling, which results from mitochondria and NADPH oxidase (NOX) [1–6]. The NOX-mediated, hypoxic ROS formation is secondary to mitochondrial ROS; this newly identified ROS-induced ROS generation provides a positive feedback mechanism, contributing to the hypoxic increase in $[Ca^{2+}]_i$ in PASMCs and HPV [7–9]. These findings further demonstrate the importance of mitochon-

drial ROS in hypoxic cellular responses in PASMCs. Evidence of the essential role of mitochondrial ROS in hypoxic Ca^{2+} and associated contractile responses, together with the fact that mitochondria are the most voracious intracellular oxygen consumers, would seem to indicate that mitochondria are the primary hypoxic sensor in PASMCs [1–6]. However, direct experimental evidence to support this view is lacking.

Numerous studies, particularly using isolated mitochondria, have revealed that under physiological conditions, mitochondria may generate ROS predominantly in the electron transport chain (ETC) complex I in most types of cells because of a higher NADH/NAD⁺ ratio in the matrix and/or a higher proton-motive force [10]. Up to now, there have been no reports showing ROS formation in isolated mitochondria from PASMCs. Studies in intact cells and tissues using various complex inhibitors suggest that complex III is likely to be the primary site of the hypoxic ROS generation in PASMCs [1–6]. Further research is needed to verify the critical, initial role of complex III in PASMCs. Equally importantly, it is necessary to determine which molecule in complex III may serve as a critical factor in the initiation of hypoxic ROS generation in PASMCs.

In this study, thus, we sought to determine whether hypoxia could directly affect ROS generation in isolated mitochondria and complex III from PASMCs. As Rieske iron–sulfur protein is a major functional component of complex III [11], we also wanted to test whether this molecule could play an important role in the hypoxic ROS generation in PASM complex III, mitochondria, and cells, mediating the hypoxic increase in $[Ca^{2+}]_i$ in PASMCs and hypoxic vasoconstriction in pulmonary arteries.

Abbreviations: BSA, bovine serum albumin; ETC, electron transport chain; H₂DCE/DA, 5,6-chloromethyl-2,7-dichlorodihydrofluorescein diacetate; DMEM, modified Dulbecco's minimal essential medium; EGTA, ethylene glycol tetraacetic acid; GAPDH, glyceraldehyde-3-phosphate dehydrogenase; Hepes, 4-(2-hydroxyethyl)-1-piperazineethanesulfonic acid; HPV, hypoxic pulmonary vasoconstriction; HRP, horseradish peroxidase; NOX, NADPH oxidase; PA, pulmonary artery; PBSS, phosphate-buffered saline solution; SDS, sodium dodecyl sulfate; SMC, smooth muscle cell; ROS, reactive oxygen species.

* Corresponding author. Fax: +1 518 262 8101.

E-mail address: wangy@mail.amc.edu (Y.-X. Wang).

Materials and methods

Electron microscopic examination of isolated mitochondria

All the animals in this study were used in accordance with the approved Animal Care and Use Protocol of Albany Medical College. Resistance (third and smaller intralobar) PASM tissues and cells were prepared from Swiss Webster mice, as we reported previously [12,13].

Mitochondria were isolated from PASM tissues by differential centrifugation [14]. All following the steps were performed at 4 °C. Isolated PAs were exposed to papain (0.5 mg/ml) for 25 min, placed in an all-glass Dounce homogenizer that contained 1 ml isolation buffer consisting of 215 mM mannitol, 75 mM sucrose, 20 mM 4-(2-hydroxyethyl)-1-piperazineethanesulfonic acid (Hepes), 1 mM ethylene glycol tetraacetic acid (EGTA), and 0.1% bovine serum albumin (BSA; pH 7.2), and homogenized. The homogenate was spun twice at 1300g in an Eppendorf microcentrifuge for 3 min. The resulting supernatant was topped off with isolation buffer and then spun at 13,000g for 10 min. The pellet was resuspended in 500 µl isolation buffer and placed inside the nitrogen cell disruption bomb (Parr Instrument) under a pressure of up to 1200 psi for 10 min. The samples were brought up to 2 ml with isolation buffer and centrifuged at 13,000g for 10 min. The pellet was resuspended in isolation buffer and centrifuged at 10,000g for 10 min. The collected pellet was resuspended in isolation buffer to yield the mitochondrial suspension (10 mg protein/ml).

Isolated mitochondria were fixed with 4% paraformaldehyde and 0.25% glutaraldehyde in Sorenson's buffer and infiltrated with LR-White resin using standard procedures. Sections of 15-nm thickness were cut on a Leitz Ultramicrotome and collected on nickel grids and counterstained with uranyl acetate and lead citrate. The grids were examined and photographed on a Hitachi H7100 transmission electron microscope.

Measurement of cytochrome c release in isolated mitochondria

Fifty microliters of isolated mitochondrial suspension (10 µg of protein) was added to microplate wells that contained 75 µl of cytochrome c conjugate (monoclonal antibody against cytochrome c conjugated to horseradish peroxidase). After incubation for 2 h at ~22 °C, 100 µl of a H₂O₂ (100 µM) and tetrabenazine (1 mM) mixture was added and incubated for 30 min at ~22 °C. The reaction was stopped by adding 100 µl of stop solution with 2 N hydrochloric acid. A change in the difference in the absorbance at 450 and 570 nm ($A_{450} - A_{570}$), measured at 37 °C using a FlexStation-III spectrophotometer (Molecular Devices), was taken as the amount of cytochrome c released [14].

Detection of ROS production

Isolated mitochondrial samples (50 µg protein) were added to microplate wells containing respiration buffer that included 5 mM pyruvate, 2.5 mM malate, 10 µM dichlorodihydrofluorescein diacetate (H₂DCF/DA; Molecular Probes), and 5 µM horseradish peroxidase (HRP). After incubation for 10 min, fluorescence was measured at 37 °C using the FlexStation-III spectrophotometer with an excitation wavelength of 485 nm and emission wavelength of 532 nm. ROS production was determined by subtracting the fluorescence intensity measured in control wells that contained assay buffer without mitochondria [14].

Mitochondrial intermembrane space ROS generation in isolated PSMCs was assessed using the newly developed, specific ROS biosensor pHyPer-dMito (exclusively targeted to mitochondrial inner membrane) [15]. Freshly isolated PSMCs were cultured at 37 °C in modified Dulbecco's minimal essential medium (DMEM) containing 10% fetal bovine serum and 1% penicillin/streptomycin. After culture for 48 h, by which time the cells became ~80% confluent, the cells were transfected with the pHyPer-dMito construct. After transfection for

72 h, an equal number of cells were added to microplate wells with normal physiological saline solution. HyPer was alternatively excited at 420 and 500 nm. Emitted fluorescence at 510 nm was measured at 37 °C using the FlexStation-III spectrophotometer.

Measurement of ROS formation in isolated complex III was similar to that in isolated mitochondria, except that 40 µM reduced decylubiquinone, 2 mM potassium cyanide, and 50 µM cytochrome c were used as substrates instead of pyruvate and malate. Isolated complex III was obtained from isolated PASM mitochondria using immunocapture kits (Mitoscience) according to the manufacturer's instructions. Isolated mitochondria were incubated in phosphate-buffered saline solution (PBSS) with 20 mM lauryl maltoside and protease inhibitor mixture on ice for 30 min and centrifuged at 72,000g at 4 °C for 10 min. The resulting supernatant was incubated at ~22 °C for 3 h with 10 µl of beads that were attached with specific complex III antibodies. The beads were collected by gentle centrifugation at 500g for 30 s and washed using PBSS for 5 min three times. Complex III samples were eluted by adding 50 µl of 1% sodium dodecyl sulfate (SDS). Collected complex III samples were added in respiration buffer containing H₂DCF/DA and HRP. After incubation for 10 min, fluorescence was measured at 37 °C using the FlexStation-III spectrophotometer.

Measurement of mitochondrial ATP production

ATP production was measured using an ATP Determination Kit (Molecular Probes) according to the manufacturer's instructions. Isolated mitochondrial samples (10 µg) were added in 500 µl ATP detection buffer. After incubation for 15 min, emitted luminescence at 560 nm was measured at 37 °C using the FlexStation-III spectrophotometer. The background luminescence was read before the addition of samples. The amount of ATP in the samples was calculated from the standard curve.

Determination of mitochondrial respiration activity

Mitochondrial respiration activity was determined using a miniature Clark electrode (Hansatech Instruments) in a sealed, thermostated (37 °C), and continuously stirred chamber [14]. Isolated mitochondria (50 µg protein) were added to the chamber with 500 µl respiration buffer containing 215 mM mannitol, 75 mM sucrose, 2 mM MgCl₂, 2.5 mM inorganic phosphate, 20 mM Hepes, and 0.1% fatty-acid-free BSA (pH 7.2). State 1 respiration was determined before addition of any substrate, State 2 respiration after addition of 5 mM pyruvate and 2.5 mM malate, State 3 respiration after addition of 1 mM adenosine diphosphate, State 4 respiration after addition of 1 µg/ml oligomycin, and State 5 respiration after addition of 1 µM carbonyl cyanide-*p*-trifluoromethoxyphenylhydrazone. Respiration activity was determined by the respiration control ratio between State 3 and State 4.

Measurement of complex III activity

Isolated complex III samples (5 µg protein) were added to 96-well microplates that contained 100 µl assay buffer (37 °C) with 40 µM reduced decylubiquinone and 2 mM potassium cyanide per well. The assay was initiated by adding 50 µM cytochrome c. The reduction of cytochrome c, taken as the activity of complex III, was measured at 550 nm using the FlexStation-III spectrophotometer.

Western blot analysis

Western blot analysis was performed as described in previous reports [12,13]. Samples (10 µg protein) were loaded onto 12% SDS gels and then transferred onto nitrocellulose membrane. The membrane was incubated with primary antibodies (Invitrogen) for the complex III subunit core 1 protein, complex IV subunit COX IV protein, or Rieske iron-sulfur protein at a dilution of 1:2000 followed

by appropriate secondary antibodies (Invitrogen) at a dilution of 1:4000. The targeted proteins were probed with an enhanced chemiluminescence detection kit (Santa Cruz Biotechnology) and quantified using Multi Gauge software (Fujifilm).

Rieske iron–sulfur protein silencing and overexpression

Specific small interfering RNAs (siRNAs) for Rieske iron–sulfur protein with the sequence AAGGUGCCUGACUUCUGAA were obtained from Dharmacon. One microgram of siRNAs and 6 μ l of Lipofectamine 2000 were mixed in 3 ml of Opti-MEM (Invitrogen) for 15 min at $\sim 22^\circ\text{C}$ (which helps the siRNAs enter the cells) and then applied to primary cultured PSMCs [16]. The cells were cultured in a 5% CO_2 incubator at 37°C for 72 h. As a control, cells were treated identically, but either untransfected or transfected with scrambled siRNA with the sequence GCUUACUGCGUAUAGGUCACA.

To silence Rieske iron–sulfur protein expression in isolated PAs, siRNAs for Rieske iron–sulfur protein were introduced into tissues using reverse permeabilization, as described earlier [17]. Isolated PAs were transferred to culture dishes and exposed to the following three successive solutions (4°C): (1) 10 mM EGTA, 120 mM KCl, 5 mM ATP, 2 mM MgCl_2 , 20 mM TES (pH 6.80) for 20 min; (2) 120 mM KCl, 5 mM ATP, 2 mM MgCl_2 , 20 mM TES, and 20 nM siRNAs (pH 6.8) for 3 h; and (3) 120 mM KCl, 5 mM ATP, 10 mM MgCl_2 , 20 mM TES, and 20 nM siRNAs (pH 6.8) for 30 min. Subsequently, the PAs were bathed in a fourth solution containing 140 mM NaCl, 5 mM KCl, 10 mM MgCl_2 , 5 mM glucose, and 2 mM Mops (pH 7.1, 22°C), in which the extracellular Ca^{2+} concentration was gradually increased from 0.01 to 0.1 to 1.8 mM every 15 min. After this, the PAs were placed in DMEM/F-12 medium (supplemented with 1 mM L-glutamine, 50 U/ml penicillin, and 50 $\mu\text{g}/\text{ml}$ streptomycin) and cultured at 37°C for 72 h. As a control, PAs were treated similarly, but not treated with siRNAs for Rieske iron–sulfur protein or treated with scrambled siRNAs.

In the Rieske iron–sulfur protein overexpression studies, primary cultured cells were transfected with the pCMV-XL4 vector containing the Rieske iron–sulfur gene (Origene), not transfected, or transfected with the expression vector alone for 72 h.

The efficiency of Rieske iron–sulfur protein silencing and overexpression was determined using Western blot analysis, as described above.

Measurement of $[\text{Ca}^{2+}]_i$

As described in our previous reports [12,18], cells were incubated with 10 μM Fura-2/AM at 37°C for 30 min. Fura-2 was excited at 340 and 380 nm with the FlexStation-III spectrophotometer, and emitted fluorescence was detected at 510 nm.

Muscle tension measurement

Muscle tension in isolated PA rings was measured as described earlier [12,13]. PA rings 3 mm in length were placed in 2 ml tissue bath (Radnoti) at 37°C . All the PAs used were devoid of the endothelium. The absence of the endothelium was confirmed by the lack of 10 μM methacholine-evoked relaxation. Passive tension of 200 mg was applied to each of the PA rings. The viability of the PAs was tested with 10 μM phenylephrine. Experiments started after an equilibration period for 90 min. Contraction recordings were made using a highly sensitive force transducer (Harvard Apparatus) with a Powerlab/4SP recording system (AD Instruments).

Hypoxia

To induce hypoxic responses in isolated complex III, mitochondria, and cells, the medium used was first aerated with a 20% O_2 , 5% CO_2 , and 75% N_2 mixture (normoxic gas) for 10 min and then with a 1% O_2 ,

5% CO_2 , and 96% N_2 mixture (hypoxic gas) for 5 min to induce hypoxic responses, as we reported previously [7–9,12,13,18–20]. As a control, samples were treated with normoxic gas alone. To assess HPV, isolated PA rings were equilibrated with the above-mentioned normoxic gas for 90 min and then exposed to hypoxia for up to 45 min. Oxygen tension in normoxic and hypoxic bath solutions was determined using an OXEL-1 oxygen electrode connected to an ISO₂ isolated dissolved oxygen meter (World Precision Instruments).

Statistical analysis

Data are expressed as means $\pm$ standard error of the mean obtained from at least three independent experiments. Student's *t* test or one-way ANOVA with Bonferroni post hoc test was used to determine the significance of differences between comparisons. A $P < 0.05$ was accepted as statistically significant.

Results

Hypoxia causes an increase in ROS generation in isolated mitochondria from PSMCs

Mitochondria in PSMCs are considered to be implicated in hypoxic sensing to enhance ROS generation, leading to HPV [1–6]. To provide direct experimental evidence for this view, we sought to determine whether hypoxia could directly affect ROS production in isolated mitochondria from PSMCs. Electron microscopic studies showed that isolated mitochondria had well-preserved structures (Fig. 1A). Application of Triton X-100 (1 mM) caused a massive cytochrome *c* release (Fig. 1B). These data indicate that the isolated mitochondria were intact and viable.

We were excited to find, for the first time, that acute hypoxic exposure for 5 min significantly enhanced ROS production, as determined using the classic ROS detection probe $\text{H}_2\text{DCF}/\text{DA}$, in isolated mitochondria (Fig. 1C). However, hypoxia did not affect mitochondrial ATP production (Fig. 1D). Mitochondrial oxygen consumption was not affected, either (Fig. 1E). In these experiments, the mean O_2 tension in the normoxic and hypoxic solutions in the recording tube was 160.1 ± 9.0 and 12.8 ± 3.4 mm Hg, respectively.

Using pHyPer-dMito, an expression vector that encodes the newly developed, specific ROS detection biosensor HyPer and is exclusively targeted to the mitochondrial inner membrane [15], we further revealed that hypoxia causes a large increase in HyPer-derived fluorescence ($[\text{ROS}]_i$) in PSMCs (Fig. 1F). These findings are consistent with our previous reports using the traditional ROS detection approaches [7–9,18]. As control experiments, we also assessed the effect of exogenous H_2O_2 on HyPer-derived fluorescence in PSMCs. The results indicated that H_2O_2 could result in a concentration-dependent increase in HyPer signals (Fig. 1G).

Hypoxia enhances ROS formation in isolated complex III from PSMCs

Pharmacological studies imply that the hypoxic ROS production in PSMCs predominantly occurs at the mitochondrial ETC complex III [1]. Consistent with previous data, here we unveiled that acute hypoxia largely enhanced ROS generation in isolated complex III from PSMCs (Fig. 2A). The isolated complex III had a high purity, as Western blot analysis showed the presence of the complex III subunit core 1 protein, but not the complex IV subunit COX IV protein, in isolated complex III preparations (Fig. 2B).

Interestingly, the activity of isolated complex III was remarkably enhanced as well (Fig. 2C). These data provide the novel evidence that hypoxia may enhance the functional activity of complex III and then its ROS formation, leading to an increase in $[\text{ROS}]_i$ in PSMCs.

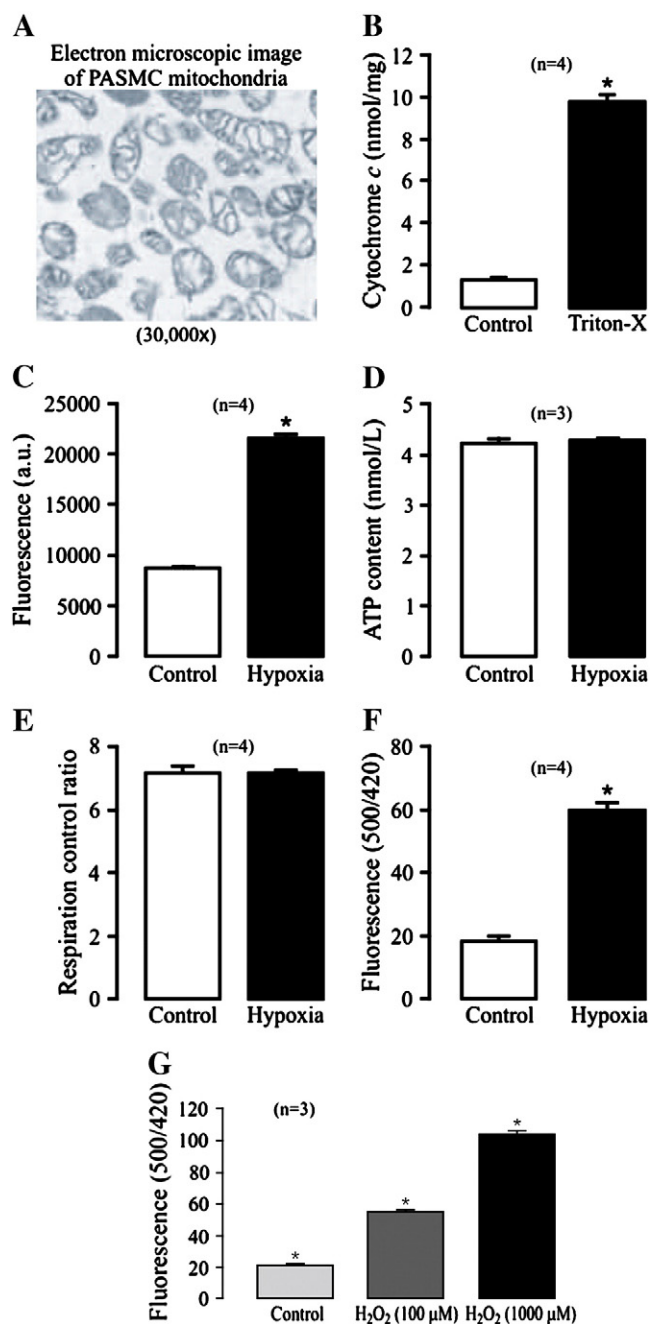


Fig. 1. Hypoxia causes a large increase in ROS production in isolated mitochondria from PSMCs. (A) Representative electron microscopic image of isolated mitochondria. (B) Cytochrome c release in isolated mitochondria before and after application of Triton X-100 (1 mM) for 5 min. Numbers in parentheses refer to the number of independent experiments performed. * $P < 0.05$ compared with control (before application of Triton X-100). (C) Effects of acute hypoxic exposure for 5 min on ROS production, determined by assessing the fluorescence of the ROS detection dye H₂DCF, in isolated mitochondria. (D) Effects of acute hypoxia on ATP production in isolated mitochondria. (E) Effects of hypoxia on respiration activity in isolated mitochondria. (F) Effects of hypoxia on mitochondrial intermembrane space ROS production, examined by measuring the fluorescence of the specific ROS biosensor pHyPer, in isolated PSMCs. (G) Effects of H₂O₂ on HyPer signals in isolated PSMCs.

Rieske iron–sulfur protein silencing blocks the hypoxic ROS generation in isolated PASM complex III, mitochondria, and cells

Rieske iron–sulfur protein is an important functional component in complex III [11]. Thus, we wondered whether this molecule is required for the hypoxic ROS generation in complex III of PSMCs. Our data indicate that transfection with specific siRNAs for Rieske iron–

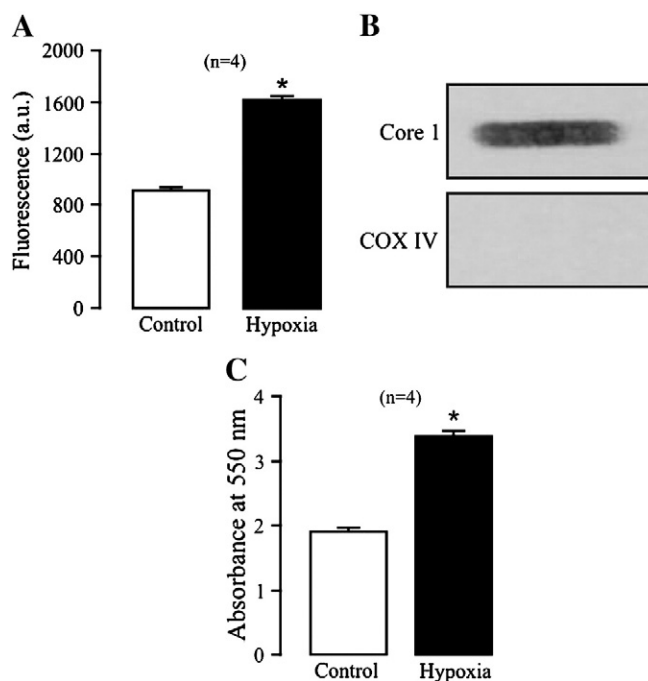


Fig. 2. Hypoxia significantly increases ROS production in isolated complex III from PSMCs. (A) Effects of hypoxia on ROS formation (H₂DCF fluorescence) in isolated complex III. (B) Representative Western blots show the presence of the complex III subunit core 1 protein, but not the complex IV subunit COX IV protein, in isolated complex III samples. (C) Effects of hypoxia on isolated complex III activity, gauged by assessing the absorbance of cytochrome c reduction at 550 nm.

sulfur protein suppressed its expression in isolated PSMCs by 90% (Fig. 3A). However, transfection with scrambled siRNAs had no effect.

Transfection of siRNAs for Rieske iron–sulfur protein could decrease the basal activity of isolated complex III from PSMCs (Fig. 3B). Importantly, siRNAs for Rieske iron–sulfur protein, but not scrambled siRNAs, completely prevented acute hypoxia from enhancing the activity of isolated complex III. Rieske iron–sulfur protein silencing also moderately inhibited the basal ROS generation and fully abolished the hypoxic ROS formation in isolated complex III (Fig. 3C). Consistent with its direct effect, hypoxia did not alter Rieske iron–sulfur protein expression in PSMCs (Fig. 3D).

Rieske iron–sulfur protein overexpression augments the hypoxic ROS generation in isolated PASM complex III, mitochondria, and cells

Transfection with Rieske iron–sulfur gene, but not expression vector alone, significantly augmented its protein overexpression in isolated PSMCs (Fig. 4A).

In contrast to its silencing, Rieske iron–sulfur protein overexpression could increase the basal activity as well as the hypoxic increase in the activity of isolated complex III (Fig. 4B). Moreover, Rieske iron–sulfur protein overexpression also enhanced the hypoxic ROS generation in isolated complex III (Fig. 4C).

Rieske iron–sulfur protein silencing prevents, whereas Rieske iron–sulfur protein overexpression enhances, the hypoxic ROS generation in isolated PASM mitochondria and cells

Compatible with the effect in isolated complex III, Rieske iron–sulfur protein silencing reduced the basal ROS generation and blocked the hypoxic ROS generation in isolated mitochondria (Fig. 5A). The hypoxic increase in mitochondrial ROS generation, determined using HyPer, was inhibited as well in PSMCs (Fig. 5B).

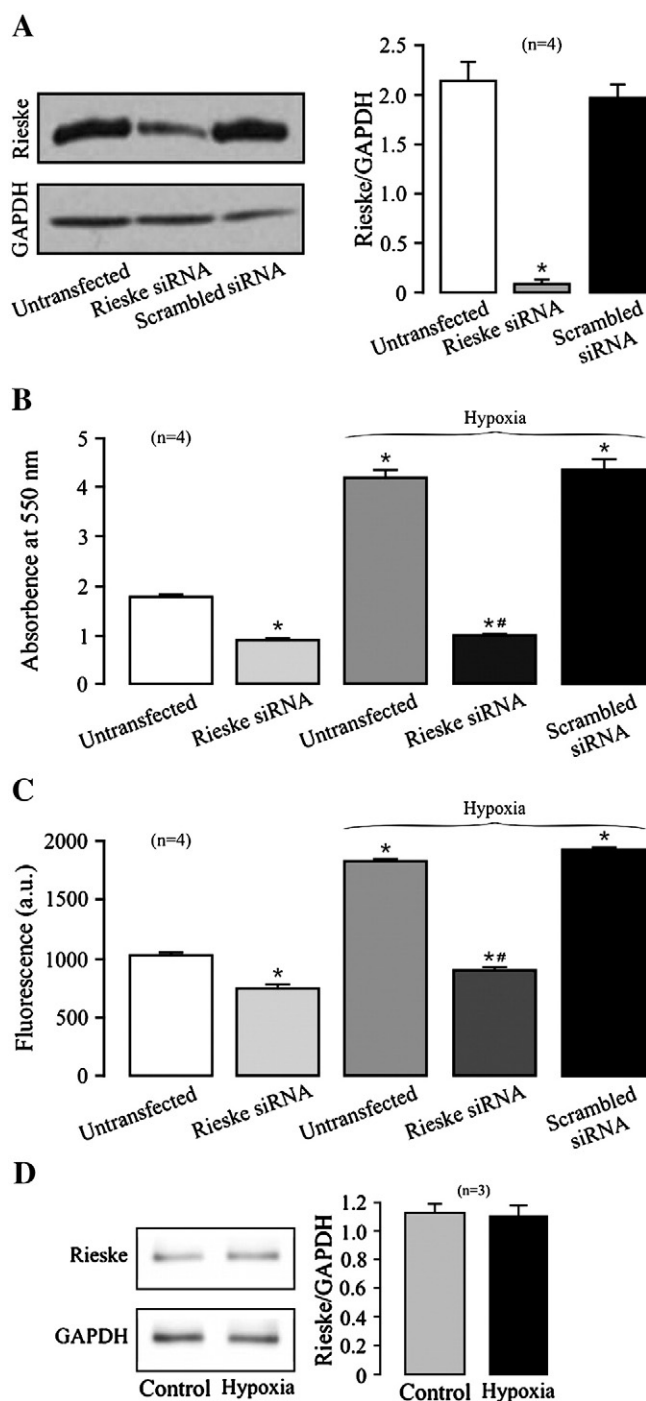


Fig. 3. Rieske iron-sulfur protein silencing blocks the hypoxic ROS production in isolated complex III from PASM cells. (A) Effects of siRNA-mediated Rieske iron-sulfur protein silencing on its protein expression in PASM cells. Representative Western blots illustrate Rieske iron-sulfur protein and glyceraldehyde-3-phosphate dehydrogenase (GAPDH) protein expression in untransfected PASM cells (control) and PASM cells transfected with Rieske iron-sulfur protein siRNAs or scrambled siRNAs. Bar graph summarizes the effects of Rieske iron-sulfur protein siRNAs on Rieske iron-sulfur protein expression levels. $*P<0.05$ compared with untransfected cells (control). (B) Effects of Rieske iron-sulfur protein silencing on the acute hypoxic increase in the activity of isolated complex III. $*P<0.05$ compared with normoxic, untransfected samples; $**P<0.05$ compared with hypoxic, untransfected samples. (C) Effects of Rieske iron-sulfur protein silencing on the hypoxic increase in ROS formation in isolated complex III. (D) Effects of acute hypoxia on Rieske iron-sulfur protein expression in PASM cells.

As expected, Rieske iron-sulfur protein overexpression was able to enhance the basal ROS generation as well as the hypoxic ROS generation in isolated PASM mitochondria (Fig. 5C) and cells (Fig. 5D).

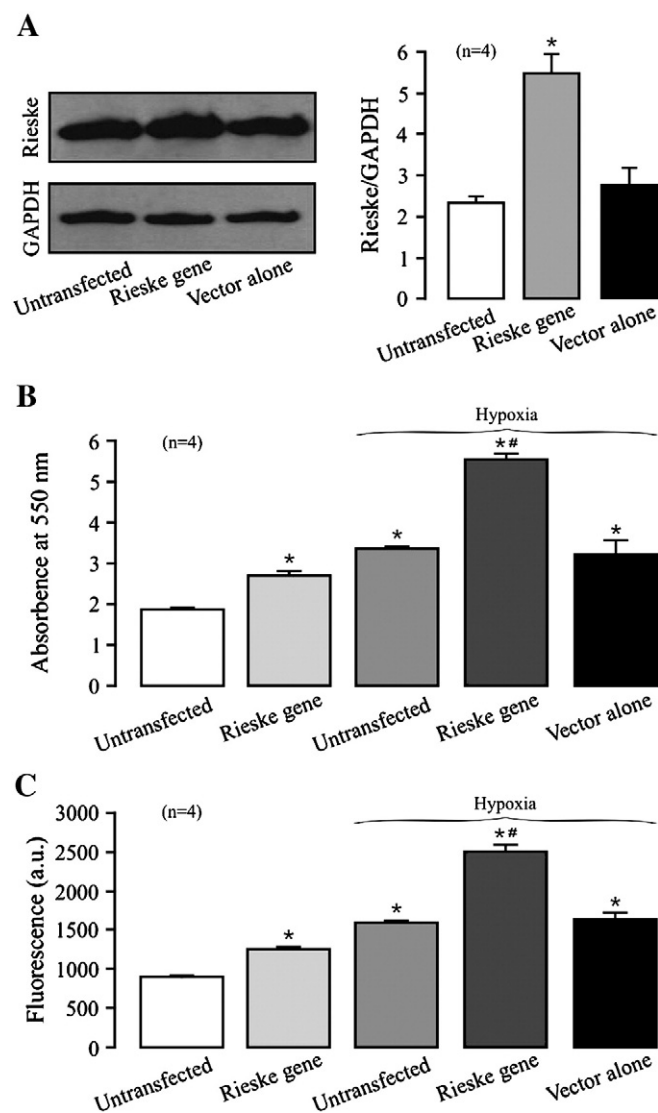


Fig. 4. Rieske iron-sulfur protein overexpression augments the hypoxic ROS generation in complex III from PASM cells. (A) Western blotting images show Rieske iron-sulfur protein and GAPDH protein expression in untransfected PASM cells (control) and PASM cells transfected with Rieske iron-sulfur gene or expression vector alone. Graph depicts the effects of Rieske iron-sulfur protein overexpression on its protein expression in PASM cells. (B) Effects of Rieske iron-sulfur protein overexpression on the hypoxic increase in isolated complex III activity. (C) Effects of Rieske iron-sulfur protein overexpression on the hypoxic ROS production in isolated complex III.

Rieske iron-sulfur protein silencing inhibits the hypoxic increase in $[Ca^{2+}]_i$ in PASM cells and HPV

Introduction of siRNAs for Rieske iron-sulfur protein into isolated PAs using reverse permeabilization [17] largely suppressed its protein expression (Fig. 6A). In line with this result, introduction of siRNAs for Rieske iron-sulfur protein markedly reduced HPV (Fig. 6B). On the other hand, scrambled siRNAs had no effect. In these studies, the mean O_2 tension in the normoxic and hypoxic solutions in the tissue bath was 155.5 ± 11.5 and 14.2 ± 4.1 mm Hg, respectively.

Similar to our previous findings [7–9,12,13,18–20], hypoxia caused a large increase in $[Ca^{2+}]_i$ in PASM cells. The hypoxic increase in $[Ca^{2+}]_i$ was mostly blocked by Rieske iron-sulfur protein silencing. The mean increase in $[Ca^{2+}]_i$ was decreased from 671.0 ± 35.5 nM in control (untransfected) cells to 173.3 ± 9.7 nM in cells transfected with siRNAs for Rieske iron-sulfur protein (Fig. 6C). In control experiments, transfection with scrambled siRNAs did not produce an effect.

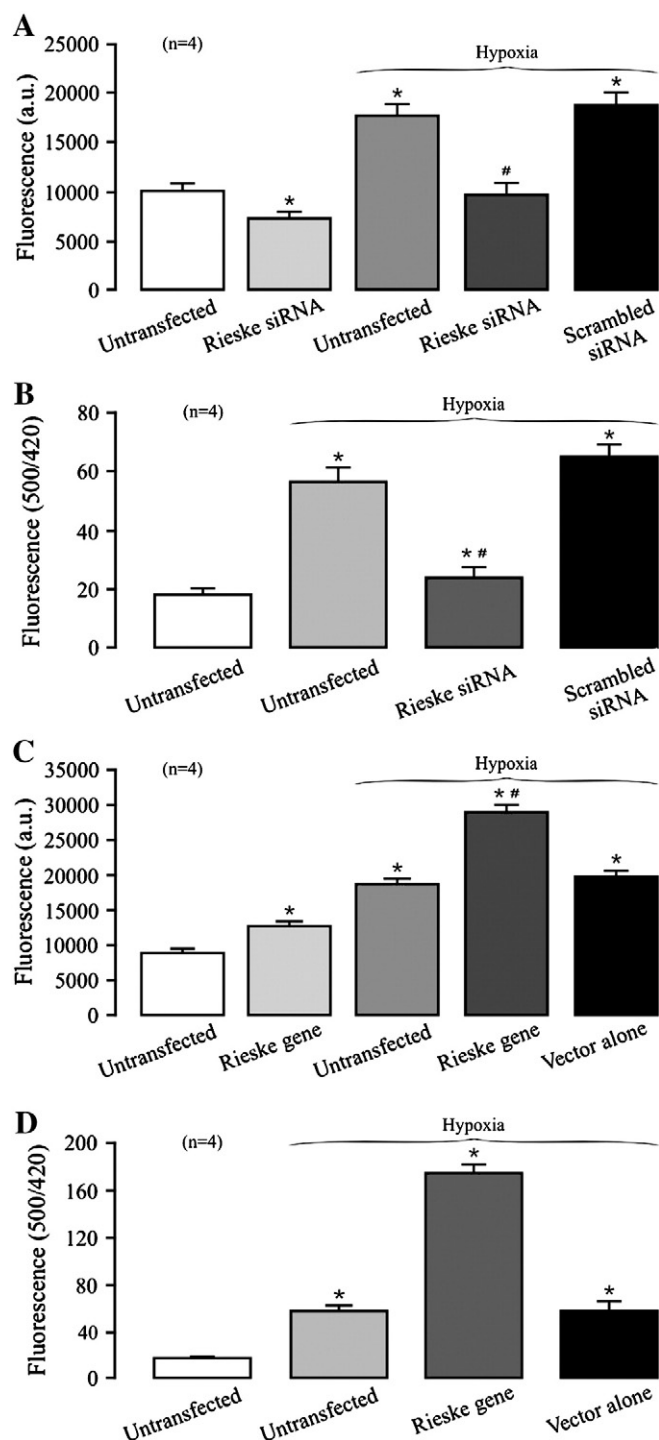


Fig. 5. Rieske iron-sulfur protein silencing prevents, whereas Rieske gene overexpression enhances, the hypoxic ROS production in isolated PASM mitochondria and cells. (A) Effects of Rieske iron-sulfur protein silencing on the hypoxic ROS production in isolated mitochondria from PSMCs. (B) Effects of Rieske iron-sulfur protein silencing on mitochondrial ROS production in PSMCs. (C) Effects of Rieske iron-sulfur protein overexpression on the hypoxic ROS production in isolated mitochondria from PSMCs. (D) Effects of Rieske iron-sulfur protein overexpression on mitochondrial ROS production in PSMCs.

Discussion

Based on previous pharmacological data in cells and tissues, we and others have proposed that mitochondria are implicated in hypoxic sensing to affect their ROS generation, mediating the hypoxic increase in $[Ca^{2+}]_i$ in PSMCs and associated HPV [1–6]. As this long-

standing conjecture lacks the support of direct experimental evidence, we have contemplated whether hypoxia may cause an increase in ROS production in isolated mitochondria from PSMCs. The findings presented in this study, for the first time, reveal that mitochondria in PSMCs can directly respond to hypoxia by producing ROS. Reports from our laboratory and many others have shown that hypoxia causes an increase in $[ROS]_i$ in isolated PSMCs, PAs, and lungs, whereas a few research groups have reported that hypoxia decreases, rather than increases, $[ROS]_i$ [1–6]. These inconsistent results have been thought to be attributed to the potentially dissimilar hypoxic responses between single cells and multiple-cell tissues and between freshly isolated and passaged cells [21,22]. The freshly isolated mitochondria are devoid of these possible problems and thus can be considered reliable preparations for studies of direct hypoxic sensing. It is also worth pointing out that by making simultaneous measurement of H_2DCF -derived fluorescence signal and staining of mitochondria, we have found that the hypoxic ROS generation in mitochondrial areas is earlier and larger than that in nonmitochondrial and whole-cell areas [8]. These data, together with the hypoxic ROS generation in isolated mitochondria (this study) and intact mitochondria of PSMCs [23], fully support the role of mitochondria in hypoxic sensing in PSMCs.

The site for the hypoxic generation of mitochondrial ROS in PSMCs is uncertain. Using mitochondrial ETC complex I, III, and IV inhibitors, earlier investigations by Archer et al. suggest that hypoxia could decrease ROS production at complex I and/or III, leading to HPV [24,25]. However, a number of later studies by several different laboratories using complex inhibitors imply that complex III is the major site of the hypoxic ROS production to cause an increase in $[Ca^{2+}]_i$ in PSMCs and HPV [8,26–29]. Moreover, it has also been reported that complexes I, II, III, and IV are all likely to be involved in HPV [30–33]. Notably, our work further indicates that the ETC molecules before the complex III ubiquinone site may act as a functional unit to mediate hypoxic ROS generation at complex III in PSMCs, as simultaneous treatment with complex I and II inhibitors, I and III preubiquinone site inhibitors, as well as II and III preubiquinone site inhibitors, neither have an additive effect on the hypoxic increase in $[ROS]_i$ nor the $[Ca^{2+}]_i$ [8]. Here, we have found that hypoxia significantly enhances ROS formation in isolated complex III. This novel result demonstrates that the mitochondrial complex III has an inherent hypoxia-sensing capability. In addition, we have revealed that hypoxia does not alter mitochondrial ATP production and respiration. These data, together with the hypoxic ROS generation in isolated complex III, indicate that mitochondrial hypoxic sensing does not necessarily depend on the reduced availability of oxygen at cytochrome c and electron flux through the ETC, at least in PSMCs. In essence, we have provided the first evidence-based proof for the original hypothesis by Guzy and Schumacker that complex III is not only the source of ROS signals, but also the oxygen sensor per se [6]. Interestingly, our data further disclose that the activity of isolated complex III, similar to ROS formation, is also significantly enhanced by hypoxia. Thus, the complex III ROS formation is likely to be dependent on its activity, supporting previous reports that pharmacological inhibitors that block complex III activity can prevent the hypoxic ROS generation in PSMCs, PAs, and lungs [1–3,5].

Rieske iron-sulfur protein is a component of complex III and critical for its functional activity [11]; accordingly, we investigated its potential role in hypoxic ROS generation in PSMCs. Using siRNA technology, we found that Rieske iron-sulfur protein silencing completely blocks the hypoxic increase in isolated complex III activity and ROS formation. Likewise, hypoxic ROS production in isolated PASM mitochondria and cells is fully abolished by Rieske iron-sulfur protein silencing. In line with its silencing, Rieske iron-sulfur protein overexpression significantly augments the hypoxic ROS formation in isolated complex III, mitochondria, and PSMCs. Moreover, Rieske iron-sulfur protein silencing reduces, and Rieske iron-sulfur protein overexpression increases, the basal activity of isolated complex III and

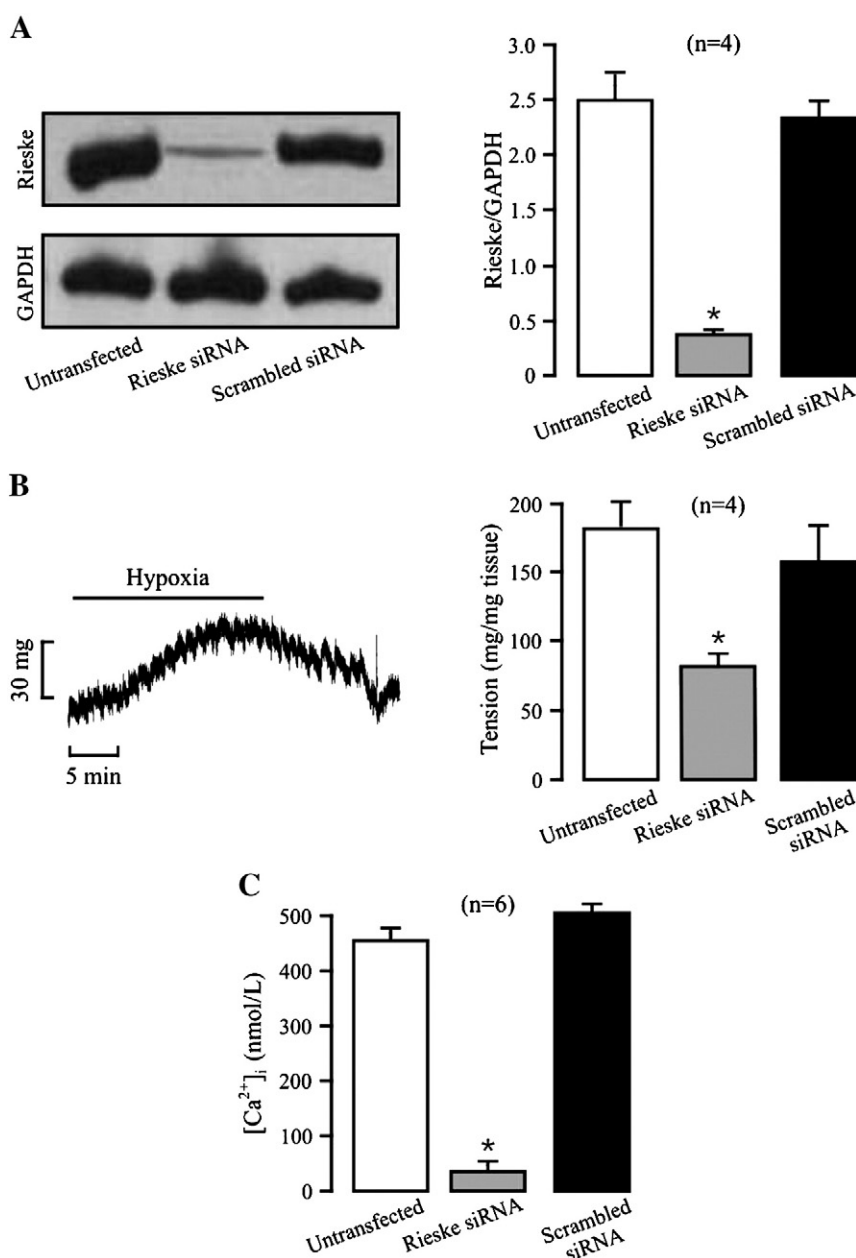


Fig. 6. Rieske iron-sulfur protein silencing inhibits HPV and hypoxic increase in $[Ca^{2+}]_i$ in PSMCs. (A) Representative Western blots show Rieske iron-sulfur protein expression in untransfected PAs (control) and PAs transfected with Rieske iron-sulfur protein siRNAs by reverse permeabilization. Bar graph summarizes the effects of Rieske iron-sulfur protein and scrambled siRNAs on Rieske iron-sulfur protein expression levels. (B) A representative recording illustrates hypoxic vasoconstriction in an isolated pulmonary artery. Graph shows the effects of Rieske iron-sulfur protein silencing on HPV. (C) Effects of Rieske iron-sulfur protein silencing on the hypoxic increase in $[Ca^{2+}]_i$ in PSMCs. The number in parentheses indicates the number of independent experiments performed from three separate preparations.

basal ROS production in isolated complex III and mitochondria. These results, together with previous publications reporting that Rieske iron-sulfur protein silencing blocks hypoxic cellular responses in mammalian cell lines and yeast [34–39], further support the view that complex III is implicated in hypoxic sensing and suggest that Rieske iron-sulfur protein plays an important role in basal and hypoxic ROS production in PSMCs. Complex III is known to carry out the ubiquinone (Q) cycle to transport electrons from complex I or II to complex IV. Ubiquinone at complex III's Qo site accepts a pair of reducing equivalents, becoming ubihydroquinone. Rieske iron-sulfur protein can seize a single electron from ubihydroquinone through its extrinsic domain and pass it to cytochrome c_1 through the movement of the extrinsic domain. Meanwhile, ubihydroquinone becomes ubisemiquinone, which transfers its electron to the cytochrome b (from b_{566} to b_{562}) and then to another ubisemiquinone at the Qi site

to regenerate ubiquinone. During this process, a small fraction of electrons may escape from ubisemiquinone to oxygen to form a superoxide anion at the Qo and also possibly at the Qi site [6,11,40,41]. As such, we speculate that hypoxia may perhaps affect the function of Rieske iron-sulfur protein in transferring electrons from ubihydroquinone to cytochrome c_1 , leading to an increase in the electron transfer rates and ROS formation in complex III. It would be very interesting to test this exciting hypothesis in future studies. A recent, elegant study has shown that ubiquinol-cytochrome c reductase binding (UQCRB) protein, an element of complex III, is involved in the hypoxic ROS generation to mediate breast cancer angiogenesis [42]. This raises the possibility that hypoxia may enhance the activity of UQCRB protein, thereby causing ROS generation in PSMCs.

As ROS mediate the hypoxic increase in $[Ca^{2+}]_i$ in PSMCs and HPV [1–6], Rieske iron-sulfur protein is possibly imperative for these

hypoxic responses as well. Indeed, Rieske iron–sulfur protein silencing significantly inhibits the hypoxic increase in $[Ca^{2+}]_i$ in PASMCs. Moreover, we have for the first time shown that Rieske iron–sulfur protein siRNAs can be introduced into isolated PAs by a reverse-permeabilization approach, causing a large reduction in HPV.

In conclusion, hypoxia can directly cause ROS generation in isolated complex III and mitochondria from PASMCs. The hypoxic ROS generation is blocked by Rieske iron–sulfur protein silencing, whereas it is augmented by Rieske iron–sulfur protein overexpression. Furthermore, Rieske iron–sulfur protein silencing inhibits the hypoxic increase in $[Ca^{2+}]_i$ in PASMCs and HPV. These findings, for the first time, provide clear evidence that Rieske iron–sulfur protein is likely to be an essential, primary molecule in hypoxic ROS generation, leading to an increase in $[Ca^{2+}]_i$ in PASMCs and ultimately pulmonary vasoconstriction.

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