IDH3 mediates apoptosis of alveolar epithelial cells type 2 due to mitochondrial Ca\(^{2+}\) uptake during hypocapnia

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In adult respiratory distress syndrome (ARDS) pulmonary perfusion failure increases physiologic dead-space (\(V_D/V_T\)) correlating with mortality. High \(V_D/V_T\) results in alveolar hypocapnia, which has been demonstrated to cause edema formation, atelectasis, and surfactant depletion, evoked, at least in part, by apoptosis of alveolar epithelial cells (AEC). However, the mechanism underlying the hypocapnia-induced AEC apoptosis is unknown. Here, using fluorescent live-cell imaging of cultured AEC type 2 we could show that in terms of CO\(_2\) sensing the tricarboxylic acid cycle enzyme isocitrate dehydrogenase (IDH) 3 seems to be an important player because hypocapnia resulted independently from pH in an elevation of IDH3 activity and subsequently in an increase of NADH, the substrate of the respiratory chain. As a consequence, the mitochondrial transmembrane potential (\(\Delta\Psi\)) rose causing a Ca\(^{2+}\) shift from cytosol into mitochondria, whereas the IDH3 knockdown inhibited these responses. Furthermore, the hypocapnia-induced mitochondrial Ca\(^{2+}\) uptake resulted in reactive oxygen species (ROS) production, and both the mitochondrial Ca\(^{2+}\) uptake and ROS production induced apoptosis. Accordingly, we provide evidence that in AEC type 2 hypocapnia induces elevation of IDH3 activity leading to apoptosis. This finding might give new insight into the pathogenesis of ARDS and may help to develop novel strategies to reduce tissue injury in ARDS.

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The adult respiratory distress syndrome (ARDS) is a typical complication of ventilated intensive care patients with an incidence of almost 10% and a mortality rate of ~40%. There are many predisposing factors known, including pneumonia, sepsis, or trauma. Inflammation, alveolar-capillary barrier dysfunction, development of lung edema, and formation of atelectasis typically in the dorsal regions of both lungs are the hallmarks of ARDS. Accordingly, opening of atelectatic lung regions, avoiding their re-collapse by applying a positive end-expiratory pressure, reducing the tidal volume during ventilation, and, if necessary, extracorporeal oxygenation and CO\(_2\) removal are the treatment approaches in ARDS.

Although the pathogenesis of ARDS has been extensively studied over the last decades, a causal therapy has not been found yet. During the last years it became clear that in both the ARDS and its experimental equivalent, the acute lung injury, intravascular coagulation, and perfusion disorders lead to an increase of the physiologic dead-space fraction (\(V_D/V_T\)), which is associated with increased mortality.\(^1\) In contrast to the CO\(_2\) accumulation in the arterial blood, the CO\(_2\) concentration in alveoli may drop, especially in lung regions with high \(V_D/V_T\). It has been shown that alveolar hypocapnia may contribute to tissue injury, including depletion of surfactant, which is produced by alveolar epithelial cells (AEC) type 2 and normally opposes the alveolar-collapsing tendency by lowering the air–liquid surface tension. Furthermore, reduced secretion of surfactant is commonly seen in ARDS patients and is associated with worse outcomes especially in critically-ill patients (which has been excellently reviewed by J Laffey et al.\(^2–6\)). Thus, the alveolar hypocapnia-induced AEC type 2 damage seems to be likely in ARDS. However, the underlying mechanism has not been elucidated yet.

Apoptosis of AEC potentially has an important role in ARDS. Decreased size and condensation of the chromatin of AEC type 2 have been shown early in human ARDS.\(^7\) Moreover, caspase-cleaved cytokeratin-18, a marker for epithelial cell apoptosis, increases in bronchoalveolar lavage fluids during ARDS.\(^8\)

Several pathways of apoptosis induction are known, also in the context of ARDS. One important trigger for apoptosis is the mitochondrial Ca\(^{2+}\) overload, especially in combination with increased mitochondrial production of reactive oxygen species (ROS).\(^9\) Importantly, mitochondrial Ca\(^{2+}\) uptake and ROS production can mutually be dependent. The driving force for mitochondrial Ca\(^{2+}\) uptake is the negative transmembrane potential gradient (\(\Delta\Psi\)) across the mitochondrial inner membrane.\(^10,11\) There are reports that \(\Delta\Psi\) can be increased by stimulation of the respiratory chain with its substrate NADH, which is produced in the tricarboxylic acid (TCA) cycle by the enzymes isocitrate dehydrogenase (IDH) 3 and \(\alpha\)-ketoglutarate dehydrogenase\(^{12–15}\) together with CO\(_2\).
According to the law of mass action the accumulation of the product may inhibit, whereas a rapid clearance of the product may accelerate its enzymatic reaction rate. We therefore hypothesize that a decrease of the alveolar partial pressure of CO₂ (pCO₂), which occurs in lung regions with high Vₒ/Vₜ, increases IDH3 activity, NADH production, and thus ΔΨ in AEC.

Using real-time fluorescence microscopy of AEC type 2 we could identify that mitochondrial IDH3 is a key player in CO₂ sensing. Accordingly, under hypocapnic conditions the resulting increase of IDH3 activity elicits an increase of ΔΨ and subsequently a shift of Ca²⁺ from the cytosol into mitochondria. In consequence, the hypocapnia-induced mitochondrial Ca²⁺ uptake leads to elevation of mitochondrial ROS production, and thus apoptosis. The understanding of this context could help to develop strategies to reduce AEC type 2 injury, including formation of atelectasis, in ARDS patients with elevated Vₒ/Vₜ. Furthermore, the opening of atelectasis, a central element of ADRS therapies, should be re-evaluated in non-perfused lung regions to prevent hypocapnic conditions and thus apoptosis of AEC.

Results

Hypocapnia increases IDH3 activity and thus NADH production. Using extracts of primary isolated AEC type 2 we were able to show that hypocapnia at constant pH of 7.4 did not induce a detectible increase of IDH3 activity (Figure 1a). We assumed that the IDH3 product NADH, which was measured for IDH3 activity determination, was oxidized by the complex 1 of the respiratory chain before its colorimetric assay detection. Therefore, we inhibited the complex 1 by rotenone at pCO₂ of 40 mm Hg in native cells, as indicated. Mean IDH3 protein in native (lane 2 and 3), in scRNA- (lane 4 and 5), and IDH3-siRNA-treated (lane 6 and 7) A549 cells. Lane 1 and 8: molecular weight ladder.

Figure 1  IDH3 activity and NADH concentration in AEC type 2. (a) Group data of IDH3 activity in primary isolated AEC type 2 measured every 5 min under normocapnic (pCO₂: 40 mm Hg) or hypocapnic (pCO₂: 0 mm Hg) conditions without or with rotenone (1 μM), respectively, at constant extracellular pH of 7.4. 50 000 cells/well. Mean ± S.E., 1P<0.05 versus pCO₂ 0 mm Hg (hypocapnia) alone. Each measurement of a single experiment was repeated twice. In total, 2–10 cells were analyzed per picture. Each experiment was repeated six times. (b) Color-coded images and (c) quantification of changes in τbound in primary isolated AEC type 2 under normocapnic (pCO₂: 40 mm Hg) conditions alone or in combination with rotenone or after hypocapnia (pCO₂: 0 mm Hg) in combination with rotenone. Mean ± S.E., repeated four times. (d) Images of primary isolated AEC type 2 show the pseudocolor-coded NAD(P)H. Upper panel, left group: left picture – normocapnic condition (baseline), right picture – normocapnic condition (control intervention); upper panel, right group: left picture – normocapnic condition (baseline), right picture – hypocapnic condition alone (intervention); lower panel, left group: left picture – normocapnic condition (baseline), right picture – normocapnic condition (intervention); lower panel, right group: left picture – normocapnic condition (baseline), right picture – hypocapnic condition+rotenone (intervention); pH was constant at 7.4. A total of 2–10 cells were analyzed per picture. Experiments were repeated four times. (e) Δ percentage change of the NAD(P)H fluorescence in primary isolated AEC type 2 under normocapnic baseline conditions, after rotenone (1 μM) or vehicle application, and again under normo- or hypocapnic conditions. As indicated, black line represents normocapnic condition, red line represents hypocapnic condition. Experiments were conducted at constant pH of 7.4 throughout. Mean ± S.E., 1P<0.05 versus rotenone at pCO₂ of 40 mm Hg (normocapnia). In total, 2–10 cells were analyzed per image. Experiments were repeated four times. (f) Rate of NAD(P)H fluorescence change during baseline conditions, hypocapnia alone, normocapnia in combination with rotenone or hypocapnia in combination with rotenone at constant pH of 7.4, as indicated. Mean ± S.E., 1P<0.05 versus pCO₂ 0 mm Hg alone (control), 1P<0.05 versus rotenone at pCO₂ of 40 mm Hg (normocapnia). In total, 3–10 cells were analyzed per image. Experiments were repeated four times. (g) Images showing NADH autofluorescence (measured at ex.: 360 nm and em: 510 nm) of native or IDH3-siRNA-treated A549 cells superfused with normocapnic (pCO₂: 40 mm Hg) or hypocapnic (pCO₂: 0 mm Hg) buffer without or with rotenone, as indicated. (h) Tracing of the percentage change of the NADH auto-fluorescence in native or IDH3-siRNA-treated A549 cells superfused with normo- or hypocapnic buffer, as indicated, at constant pH of 7.4. Rotenone (1 μM) was applied as indicated. (i) Group data of maximal NADH auto-fluorescence response 10 min following rotenone (1 μM) under hypocapnic conditions or 10 min following additionally switching to hypocapnic conditions in native or IDH3-siRNA-treated A549 cells in comparison with normocapnic baseline conditions. Experiments were performed at constant extracellular pH of 7.4. Mean ± S.E., 1P<0.05 versus baseline (pCO₂: 40 mm Hg), 1P<0.05 versus rotenone at pCO₂ of 40 mm Hg in native cells, 1P<0.05 versus rotenone at pCO₂ of 0 mm Hg in native cells. In total, 4–11 cells were analyzed per picture. Experiments were repeated six times. (j) Left) Gels show RT-PCR products for IDH3- and GAPDH-mRNA in native (lane 2 and 3), in IDH3-siRNA- (lane 4 and 5), and scRNA-treated (lane 6 and 7) A549 cells. Lane 1 and 8: molecular weight ladder. (Right) Group data of relative IDH3-mRNA expression, 1P<0.05 versus scRNA (control), n=4–5 each. (k) Western blot show IDH3 protein in native (lane 2 and 3), in scRNA- (lane 4 and 5), and IDH3-siRNA-treated (lane 6 and 7) A549 cells. Lane 1 and 8: molecular weight ladder.
(P)H auto-fluorescence increase was higher under hypo- than normocapnic conditions (Figures 1g–i). Importantly, the NAD (P)H auto-fluorescence in A549 cells was confined to mitochondrial localization, indicating again the mitochondrial origin of the NAD(P)H production (Figure 1g). As auto-fluorescence is not specific to NADH we further measured NADH concentration in A549 cells extracts by using an enzymatic assay. We revealed that NADH concentration 15 min upon onset of hypocapnia was only slightly increased in comparison to normocapnia (Supplementary Figure 2). However, after rotenone hypocapnia doubled the NADH concentration within 30 min (Supplementary Figure 2), indicating again that without rotenone treatment the NADH produced under hypocapnia is immediately consumed by the complex 1 of the respiratory chain.

To further elucidate the specificity of IDH3 in the hypocapnia-induced NADH production process we downregulated this enzyme in A549 cells. The mitochondrial IDH3 knockdown...
blocked the hypocapnia-induced increase of the NAD(P)H auto-fluorescence (Figures 1g–i). The IDH3 knockdown was verified on mRNA and protein level (Figures 1j and k, respectively). Taken together, we conclude first that, independent from pH, hypocapnia increases the mitochondrial IDH3 activity and thus the NADH production rate and second that, directly after its synthesis, NADH is totally consumed by the complex 1 of the respiratory chain. The hypocapnia-induced acceleration of IDH3 activity and thus NADH production subsequently increases mitochondrial ΔΨ. We next determined whether the hypocapnia-induced increase of IDH3 activity, NADH production rate and subsequent respiratory chain NADH consumption elevated ΔΨ. The characterization of the primary isolated AEC type 2 was revealed by the detection of LTG-stained lamellar bodies (Figure 2a). In these cells, hypocapnia at constant pH of 7.4 led to an increase of the ratio between the mitochondrial and cytosolic tetramethylrhodamine methyl ester (TMRM) fluorescence intensity equivalent for an increase of ΔΨ (Figures 2b,c and e). Under normocapnic conditions, rotenone decreased, as expected, the TMRM ratio (Figures 2d and e) verifying the correctness of the method. In these rotenone-treated cells, hypocapnia did not increase the TMRM ratio and thus ΔΨ (Figures 2d and e). All these results could be reproduced in A549 cells (Figures 2f–i).
Figure 2. ΔΨ in alveolar epithelial cells in vitro. (a) Images show primary isolated AEC type 2. For ΔΨ quantification cells were loaded with TMRM. For characterization identical cells were additionally stained with the AEC type 2-specific lamellar body marker LTG (middle). (b) Images showing fluorescence of TMRM-loaded primary isolated AEC type 2 superfused with normocapnic (pCO₂: 40 mm Hg) or hypocapnic (pCO₂: 0 mm Hg) buffer without or with rotenone (1 μM), as indicated. (c, d) Tracing of the ratio between mitochondrial and cytosolic TMRM fluorescence intensity in primary isolated AEC type 2 superfused with normo- or hypocapnic buffer without or with rotenone (1 μM), as indicated, at constant pH of 7.4. (e) Group data of maximal TMRM ratio response 15 min after switching from normocapnic (pCO₂: 40 mm Hg) baseline to hypocapnic (pCO₂: 0 mm Hg) conditions in primary isolated AEC type 2 at constant extracellular pH of 7.4. Mean ± S.E., †P<0.05 versus baseline (pCO₂: 40 mm Hg), ‡P<0.05 versus pCO₂: 0 mm Hg alone. In total, 3–9 cells were analyzed per picture. Experiments were repeated 4–6 times. (f) Images show rhod-2 fluorescence of native, scRNA, or IDH3-siRNA-treated A549 cells under normocapnic conditions alone or in combination with rotenone and under hypocapnic condition alone or in combination with rotenone, as indicated. (g) Tracing of the ratio between mitochondrial and cytosolic TMRM fluorescence intensity of native, scRNA- or IDH3-siRNA-treated A549 cells superfused with normo- or hypocapnic buffer, as indicated, at constant pH of 7.4. (h) Tracing of the ratio between mitochondrial and cytosolic TMRM fluorescence intensity of native A549 cells superfused with normocapnic buffer alone (pCO₂: 40 mm Hg) or in combination with rotenone, or superfused with hypocapnic buffer (pCO₂: 0 mm Hg) in combination with rotenone, as indicated, at constant pH of 7.4. (i) Group data of maximal TMRM ratio response 15 min after switching from normocapnic (pCO₂: 40 mm Hg) baseline condition to hypoxia alone, to normocapnia plus rotenone, or to hypoxia (pCO₂: 0 mm Hg) plus rotenone in native, scRNA-, IDH3-siRNA-, or MCU-siRNA-treated A549 cells at constant extracellular pH of 7.4, as indicated. Mean ± S.E., †P<0.05 versus baseline (pCO₂: 40 mm Hg), ‡P<0.05 versus pCO₂: 0 mm Hg alone in native or scRNA-treated A549 cells. In total, 4–12 cells were analyzed per picture. Experiments were repeated 4–6 times. (j) (Left) Gels show RT-PCR products for MCU- and UBC-mRNA in native (lane 1–4), in scRNA- (lane 5–8), and MCU-siRNA-treated (lane 9–12) A549 cells. Lane 13: no template control, lane 14: negative control, lane 15 molecular weight ladder. (Right) Group data of relative MCU-mRNA expression, *P<0.05 versus scRNA (control), n = 4 each. (k) Western blot show MCU protein in native (lane 1 and 4), in scRNA- (lane 2 and 5), and MCU-siRNA-treated (lane 3 and 6) A549 cells. Lane 7: molecular weight ladder.
Figure 2 Continued

f. 
| pCO₂ [mmHg]: | 40 | 0 | 40 | 40 | 0 |
|----------------|----|---|----|----|---|
| rotenone:      | -  | + | -  | -  | - |
| nativ          |    |   |    |    |   |
| scRNA          |    |   |    |    |   |
| IDH3-siRNA     |    |   |    |    |   |


h. 
| pCO₂ [mmHg]: | 40 | 0 |
|----------------|----|---|
| rotenone      |    |   |


i. 
| pCO₂ [mmHg]: | 0 | 40 | 0 | 0 | 0 |
|----------------|---|----|---|---|---|
| rotenone:      | - | + | - | - | - |
| scRNA          | - | - | + | - | - |
| IDH3G-siRNA    | - | - | - | + | - |
| MCU-siRNA      | - | - | - | - | + |


j. 
| MCU (151bp)  |
|---------------|
| 1 2 3 4 5 6 7 |
| 8 9 10 11 12 13 14 15 |
| UBC (90bp)    |
| 1 2 3 4 5 6 7 |

k. 
| Relative expression |
|---------------------|
| scRNA | MCU-siRNA |
| 0.0   | 1.5       |

30 kD
Furthermore, in these cells the IDH3 knockdown inhibited the hypocapnia-induced increase of ΔΨ (Figures 2f, g and i). Summarizing, the hypocapnia-induced increase of IDH3 activity, NADH production rate, and subsequent respiratory chain NADH consumption caused an elevation of mitochondrial ΔΨ.

As mentioned before, ΔΨ can be the driving force for mitochondrial Ca\(^{2+}\) uptake but it is also well known that mitochondrial Ca\(^{2+}\) regulates IDH3 and respiratory chain activity. Accordingly, we conclude that mitochondrial Ca\(^{2+}\) is not responsible for the observed ΔΨ changes. We could demonstrate that the silencing of the mitochondrial Ca\(^{2+}\) uniporter (MCU) did not inhibit the hypocapnia-induced ΔΨ increase (Figure 2i). The MCU knockdown was verified on mRNA and protein level (Figures 2j and k, respectively).

Hypocapnia induces mitochondrial Ca\(^{2+}\) uptake from the cytosol. We next determined in primary isolated AEC type 2 the hypocapnia-induced mitochondrial Ca\(^{2+}\) uptake. The characterization of these cells was again revealed by the detection of LTG-stained lamellar bodies (Figure 3a). Switching from normo- to hypocapnic conditions induced a rhod-2 fluorescence intensification, equivalent to an increase of the mitochondrial Ca\(^{2+}\) concentration [Ca\(^{2+}\)]\(_{\text{mito}}\) (Figures 3b,c,f). This response was completely reversible after switching back to normocapnic buffer. All experiments were performed at a constant extracellular pH of 7.4. The mitochondrial localization of the rhod-2 fluorescence was proofed by its colocalization with the mitochondrial marker MitoTracker green (MTG) (data not shown) and by the observation that in AEC type 2 the hypocapnia-induced Ca\(^{2+}\) uptake. Furthermore, in these cells the IDH3 knockdown inhibited the hypocapnia-induced increase (Figure 3d). We therefore assumed that hypocapnia causes an overshooting increase of [Ca\(^{2+}\)]\(_{\text{cyt}}\) assuming that cells compensate the hypocapnia-induced [Ca\(^{2+}\)]\(_{\text{cyt}}\) decrease by mechanisms, as e.g., decrease of cell membrane Ca\(^{2+}\) pump activity. The pretreatment of AEC type 2 cells with ruthenium red or rotenone inhibited the hypocapnia-induced [Ca\(^{2+}\)]\(_{\text{cyt}}\) decrease (Figures 4d and e). According to these findings, in A549 cells hypocapnia reduced [Ca\(^{2+}\)]\(_{\text{cyt}}\) 20 ± 2%, which was inhibited by rotenone (Figures 4f,g,i). In addition, both the MCU and IDH3 silencing inhibited the hypocapnia-induced decrease of [Ca\(^{2+}\)]\(_{\text{cyt}}\) (Figure 4h).

We could demonstrate that in native A549 cells ATP alone caused an increase of [Ca\(^{2+}\)]\(_{\text{cyt}}\), which was comparable to that after ATP in combination with rotenone. Further, the ATP response in native A549 cells did not differ from the ATP responses in scRNA- or IDH3-siRNA-treated cells. This indicates that both rotenone and IDH3 knockdown did not affect cell viability at least in terms of receptor-mediated Ca\(^{2+}\)-signaling cascades (Supplementary figure 5).

Furthermore, we were also interested whether pH may play a role in Ca\(^{2+}\) response induced by hypocapnia. Without additional buffering, the removing of the dissolved CO\(_2\) increased the pH of the superfusion buffer from 7.4 to 7.8. Under these conditions hypocapnia induced a similar reduction of [Ca\(^{2+}\)]\(_{\text{cyt}}\) in A549 cells as at constant extracellular pH of 7.4 revealed by buffering (Supplementary figure 6A). In contrast, an alkalinization of 7.8 under normocapnic conditions did not induce an alteration of [Ca\(^{2+}\)]\(_{\text{cyt}}\) (Supplementary figure 6A). Furthermore, the cytosolic pH was not affected in unbuffered cells (Supplementary figure 6B). We therefore conclude that the hypocapnia-induced [Ca\(^{2+}\)]\(_{\text{cyt}}\) responses were pH independent.

Since Ca\(^{2+}\) can be shifted directly from the endoplasmatic reticulum (ER) to mitochondria, we measured Ca\(^{2+}\) in the ER [Ca\(^{2+}\)]\(_{\text{ER}}\). In contrast to [Ca\(^{2+}\)]\(_{\text{cyt}}\) and [Ca\(^{2+}\)]\(_{\text{mito}}\), hypocapnia did not change [Ca\(^{2+}\)]\(_{\text{ER}}\) (Supplementary figure 7A–C). However, this ratio was clearly elevated after ATP application indicating the correctness of the method (Supplementary figure 7A–C). Taken these results together, we conclude that hypocapnia and thus acceleration of IDH3 activity induces a Ca\(^{2+}\) shift from the cytosol to the mitochondria, whereas, the ER seems to have a minor role in this scenario.

Both, the hypocapnia-induced increase of [Ca\(^{2+}\)]\(_{\text{mito}}\) and associated mitochondrial ROS production evoke apoptosis of AEC type 2. To investigate whether hypocapnia causes mitochondrial ROS production in primary isolated AEC type 2 cells we used the fluorescent MitoSOX Red mitochondrial superoxide indicator for live-cell imaging. We could show that under hypocapnia the ROS production, expressed as delta increase of the dye, was enhanced in comparison with normocapnia (Figure 5). To evaluate the respiratory chain as potential source of ROS we used rotenone under both normo- and hypercapnia. Rotenone almost diminished the hypocapnia-induced delta increase, whereas rotenone with normocapnia did not change ROS production in comparison with normocapnia alone (Figure 5).

In addition, ROS production was also inhibited after ruthenium red incubation, which blocked the mitochondrial
Figure 3  \([\text{Ca}^{2+}]_{\text{mito}}\) in AEC type 2. (a) Images show primary isolated AEC type 2. For characterization cells were stained with the cytosolic dye fura 2 and with the AEC type 2-specific lamellar body marker LTG (middle). (b) For measurement of \([\text{Ca}^{2+}]_{\text{mito}}\), AEC type 2 cells were loaded with rhod-2. Images showing fluorescence of rhod-2-loaded primary isolated AEC type 2 under normocapnic (pCO2: 40 mm Hg) baseline condition and after hypocapnia (pCO2: 0 mm Hg) alone or in combination with rotenone (1 μM), as indicated. (c–e) Tracing of rhod-2 fluorescence in primary isolated AEC type 2 superfused with normocapnic or hypocapnic buffer alone or in combination rotenone (1 μM) or ruthenium red (10 μM), as indicated. Experiments were performed at constant extracellular pH of 7.4, respectively. (f) Group data of maximal rhod-2 fluorescence responses in primary isolated AEC type 2 10 min following switching from normocapnic (pCO2: 40 mm Hg) baseline to hypocapnic (pCO2: 0 mm Hg) conditions without or with rotenone (1 μM) or ruthenium red (10 μM) pretreatment, as indicated. Experiments were performed at constant extracellular pH of 7.4 throughout. Mean ± S.E., \(^*P < 0.05\) versus baseline (pCO2: 40 mm Hg). 5–10 cells were analyzed per picture. Experiments were repeated 6 times. (g) Images showing fluorescence of rhod-2-loaded A549 cells under normocapnic (pCO2: 40 mm Hg) baseline or hypocapnic (pCO2: 0 mm Hg) conditions. (h, i) Tracings of rhod-2 fluorescence in native, scRNA-, IDH3-siRNA-, or MCU-siRNA-treated A549 cells. Cells were superfused with normo- or hypocapnic buffer, as indicated. Experiments were performed at constant extracellular pH of 7.4. (j) Group data of maximal rhod-2 fluorescence responses in native, scRNA-, IDH3-siRNA-, or MCU-siRNA-treated A549 cells 10 min after switching from normocapnic (pCO2: 40 mm Hg) baseline to hypocapnic (pCO2: 0 mm Hg) conditions. Experiments were performed at constant extracellular pH of 7.4 throughout. Mean ± S.E., \(^*P < 0.05\) versus baseline (pCO2: 40 mm Hg), \(^{†}P < 0.05\) versus pCO2: 0 mm Hg alone in native or scRNA-transfected cells. 5–10 cells were analyzed per picture. Experiments were repeated 6 times.
calcium uniporter and subsequently the mitochondrial 
Ca\textsuperscript{2+}-uptake. Mitochondrial Ca\textsuperscript{2+} and ROS are potential 
trigger for apoptosis. Therefore, we used a CellEvent 
Caspase-3/7 fluorogenic substrate to detect apoptotic cells 
upon hypercapnia in a live-cell image setting. As depicted in 
Figure 6a, the percent of apoptotic cells increased from 
20.1 ± 5.9 under normocapnia to 48.8 ± 6.6 under hypercapnia. Treatment with rotenone or ruthenium red prevented the 
initiation of apoptosis (23.8 ± 8.1% apoptotic cells and 
16.4 ± 4.6% apoptotic cells, respectively) (Figure 6a). In 
order to validate these results we measured as an additional 
apoptosis marker the expression of cleaved poly-(ADP- 
ribose) polymerase (PARP) in primary isolated AECs type 2. 
Western blot analysis of this protein revealed that only cells 
cultured under hypocapnic conditions express cleaved PARP, 
which was inhibited by rotenone or ruthenium red (Figure 6b). Under normocapnic condition alone or in combination with 
rotenone or ruthenium red cleaved PARP was not detectable 
(Figure 6b).

**Discussion**

We provide evidence that in AEC type 2 hypcapnia increases 
IDH3 activity and NADH production, ΔΨ, [Ca\textsuperscript{2+}]\textsubscript{mito}, and 
mitochondrial ROS concentration, which in turn induces 
apoptosis (Figure 7).

For the first time, at least to our knowledge, we were able to 
measure an increase of the IDH3 activity in primary isolated 
AEC type 2 in response to hypcapnia. In this context, we 
have to critically emphasize that we could not see any 
increase of NADH concentration in AEC type 2 subjected 
solely to hypocapnia. Instead, after rotenone we could 
measure a steady rise of NADH concentration, which could 
father be increased in combination with hypcapnia. We 
concluded that NADH produced under hypcapnic conditions 
is immediately consumed by complex 1 and can thus not been 
detected. Inhibition of complex I however blocked the NADH 
consumption and after this intervention the hypcapnia-inducd NADH production was detectable.
Figure 4  

[Ca^{2+}]_cyt in alveolar epithelial cells in vitro. (a) Images show primary isolated AEC type 2. For [Ca^{2+}]_cyt measurement cells were loaded with fura 2 (left). For characterization cells were additionally stained with LTG (middle), a marker for type 2-specific lamellar bodies. (b) Images show the 340:380 ratio in pseudocolor of the identical fura 2-loaded primary isolated AEC type 2 under normocapnic (pCO₂: 40 mm Hg) baseline or hypocapnic (pCO₂: 0 mm Hg) conditions at constant pH of 7.4. (c) Tracing of [Ca^{2+}] from an identical epithelial cell superfused with normo- (pCO₂: 40 mm Hg) or hypocapnic (pCO₂: 0 mm Hg) buffer, as indicated. (d) Tracings of [Ca^{2+}] from an identical epithelial cell superfused with normo- (pCO₂: 40 mm Hg) or hypocapnic (pCO₂: 0 mm Hg) buffer, as indicated. Cells were pretreated with rotenone or ruthenium red, as indicated. (e) Bars show group data of maximal [Ca^{2+}]_cyt response in primary isolated AEC type 2. 15 min after intervention. Cells were pretreated with vehicle, rotenone (1 μM), or ruthenium red (10 μM), as indicated. Under baseline conditions cells were superfused with buffer at following conditions: pCO₂: 40 mm Hg and pH 7.4 for each group. Buffer conditions after intervention were: pCO₂: 0 mm Hg and pH 7.4 for each group. Mean ± S.E., *P<0.05 versus baseline (pCO₂ of 40 mm Hg), †P<0.05 versus pCO₂ of 0 mm Hg alone. In total, 3–9 cells were analyzed per picture. Experiments were repeated six times each. (f) Images show the pseudocolor-coded 340:380 ratio for fura 2-loaded A549 cells under normocapnic (pCO₂: 40 mm Hg) baseline or hypocapnic (pCO₂: 0 mm Hg) conditions at constant pH of 7.4. (g) Tracings of [Ca^{2+}] from an identical epithelial cell superfused with normo- (pCO₂: 40 mm Hg) or hypocapnic (pCO₂: 0 mm Hg) buffer, as indicated. pH was constant at 7.4. (h) Tracings of [Ca^{2+}] from an identical epithelial cell superfused with normo- (pCO₂: 40 mm Hg) or hypocapnic (pCO₂: 0 mm Hg) buffer, as indicated. Cells were untreated or treated with scRNA, IDH3-siRNA, or MCU-siRNA, as indicated. (i) Group data of maximal [Ca^{2+}]_cyt responses 10 min following switching from normocapnic (pCO₂: 40 mm Hg) baseline to hypocapnic conditions (pCO₂: 0 mm Hg) at constant extracellular pH of 7.4 in native, scRNA-, IDH3-siRNA-, or MCU-siRNA-treated A549 cells. Mean ± S.E., *P<0.05 versus baseline (pCO₂: 40 mm Hg). †P<0.05 versus pCO₂: 0 mm Hg in native or scRNA-treated A549 cells. In total, 3–10 cells were analyzed per picture. Experiments were repeated 6 times.
The dependency of the respiratory chain activity from NADH has been demonstrated by Kuznetsov et al.\textsuperscript{19} Hence, as a consequence of hypocapnia-induced elevation of NADH production, $\Delta\psi$ and thus the driving force for the mitochondrial Ca$^{2+}$ uptake increased. This hypothesis was supported by the IDH3 knockdown. However, the IDH3 knockdown did not completely blocked the hypocapnia-induced NADH and $\Delta\psi$ responses. Reason for this result could be either that the IDH knockdown was not efficient enough or, more importantly, other mechanisms for the hypocapnia-induced NADH and $\Delta\psi$ responses are involved. Hypothetically, low pCO$_2$ could also act on mitochondrial Ca$^{2+}$ handling, e.g., on increase of mitochondrial Ca$^{2+}$ influx or decrease of mitochondrial Ca$^{2+}$ efflux. In this context another aspect has to be discussed: because IDH3 is one of the first enzymes of the TCA cycle and therefore upstream of all other TCA cycle enzymes, e.g., the CO$_2$ producing alpha-ketoglutarate dehydrogenase, the IDH3 knockdown would affect these enzyme activities, which could also contribute to the hypocapnia-induced changes of in NADH and $\Delta\psi$. This might be the reason why the IDH knockdown was highly efficient. In contrast, it could be assumed that the IDH3 knockdown would affect the overall flux through the TCA cycle. Unfortunately, we cannot exclude this hypothesis. However, we measured cytosolic Ca$^{2+}$ responses to the purinergic receptor agonist ATP in IDH3-siRNA-treated AEC and the revealed Ca$^{2+}$ responses were comparable to native and even to rotenone-treated AEC indicating that, at least, G-protein coupled receptor Ca$^{2+}$ signaling is intact.

To determine whether the hypocapnia-induced increase of NADH and $\Delta\psi$ elicits a Ca$^{2+}$ shift into mitochondria from the cytosol we measured both $[\text{Ca}^{2+}]_{\text{mito}}$ and $[\text{Ca}^{2+}]_{\text{cyt}}$. Accordingly, we could demonstrate on both primary isolated AEC type 2 and A549 cells that hypocapnia was accompanied not only by a $[\text{Ca}^{2+}]_{\text{mito}}$ increase but also by a $[\text{Ca}^{2+}]_{\text{cyt}}$ decrease. In order to differentiate whether the Ca$^{2+}$ responses in AEC...
type 2 were CO₂ rather than pH mediated, we additionally measured [Ca²⁺]_{cyt} primarily in A549 cells under several pH conditions and found that in AEC type 2 [Ca²⁺]_{cyt} response to CO₂ is independent from pH.

Moreover, we could show that the hypocapnia-induced [Ca²⁺]_{cyt} response was inhibited by MCU knockdown or pharmacological blockade of the MCU with ruthenium red confirming a Ca²⁺ shift from the cytosol into mitochondria. From these experiments we further conclude that the driving force for the mitochondrial Ca²⁺ uptake came from the mitochondria and not from the cytosol since in response to hypocapnia [Ca²⁺]_{cyt} further decreased. According to the literature, the [Ca²⁺]_{mito} increase can also be evoked by a Ca²⁺ shift from the ER directly into mitochondria. In our scenario this pathway seems to be unlikely because [Ca²⁺]_{ER} was unaffected by hypocapnia. However, we cannot exclude the possibility that the ER was synchronously re-filled with cytosolic Ca²⁺, whereas Ca²⁺ was shifted from the ER into mitochondria. It has also been described that mitochondria can uptake Ca²⁺ from the extracellular space. Indeed, this possibility, at least in part, cannot be excluded by our experiments, too.

The inhibition of the CO₂-induced Ca²⁺ shift by rotenone and the unaffected [Ca²⁺]_{ER} suggests that the driving force for the mitochondrial Ca²⁺ uptake was an increase of the respiratory chain activity and hence ΔΨ rather than ER. This hypothesis was confirmed by our finding that the fluorescence of TMRM, which accumulates in mitochondria in a potential-dependent manner, was enhanced in both primary isolated AEC type 2 and in A549 cells under hypo-compared with normocapnic conditions. Addition of rotenone revealed, as expected, a quick decrease of the TMRM fluorescence indicating that the respiratory chain activity is the important determinant of ΔΨ. Further, hypocapnia induced an increase of the respiratory chain substrate NADH and the knockdown of the NADH-producing IDH3 inhibited NADH, ΔΨ, [Ca²⁺]_{mito}, and [Ca²⁺]_{cytosolic} response under low CO₂ levels.

Taken together, we suggested that CO₂, a by-product of IDH3, inhibits IDH3 activity via a negative feedback loop. Under hypocapnic conditions, this inhibitory effect of CO₂ on the IDH3 activity is reduced which subsequently increases NADH production and in turn ΔΨ which, at least in part, lead to a Ca²⁺ shift into mitochondria.

The hypocapnia-induced mitochondrial Ca²⁺ uptake was associated with AEC type 2 apoptosis. Ca²⁺ enters mitochondria via the Ca²⁺ uniporter. As its pharmacological inhibition by ruthenium red prevented both Ca²⁺ uptake and apoptosis we conclude that the hypocapnia-induced Ca²⁺ uptake was, at least in part, responsible for the apoptosis of AEC type 2. We can only speculate about the underlying mechanism. Increase in [Ca²⁺]_{mito} has been shown to induce apoptosis by alterations in mitochondrial membrane permeability, formation of the permeability transition pore, cytochrome c release, and activation of caspase-9 and -3 and disruption of ΔΨ. We could demonstrate that immediately after onset of hypocapnia ΔΨ increases. However, we have no data on ΔΨ 24 h after hypocapnia because TMRM measurements are useful only for a short time period. There are reports that especially Ca²⁺ coming from the ER might be one trigger for apoptosis. As stated above, this way of apoptosis induction might be
induced mitochondrial Ca\textsuperscript{2+} uptake leads to mitochondrial ROS production (VI). Further, the hypocapnia-give any proof for the mechanism of the Ca\textsuperscript{2+}-dependent mitochondrial ROS production is well established we cannot probability that the electron will instead be donated to O\textsubscript{2}, the respiratory chain. This might statistically increases the in a higher amount of ubisemiquinone on complex III of ROS production. Theoretically, it could be the case that the perfusion failure in ventilated lung regions, equivalent to a VD/VT increase, leads to negatively regulated by its product CO\textsubscript{2} we conclude that under hypocapnic transport may occur under hypocapnia and be responsible generating superoxide. Furthermore, a reverse electron hypothesis.

**Materials and Methods**

**Animals.** Male Sprague-Dawley rats (350 g) were purchased from Charles River (Sulzfeld, Germany). After review and approval by the local ethics committee and the government of Hamburg, all animals were treated humanely in accordance with ‘principles of laboratory animal care’ (NIH Publication No. 86-23, revised 1985) as well as with the German legislation on protection of laboratory animals.

**Materials.** Fura 2-AM, LysoTracker green (LTG), MTG, rhod-2, 2\textsuperscript{,} 7\textsuperscript{,} bis-(2-carboxyethyl)-5-(and 6)-carboxyfluorescin-AM and TMRM were purchased from Molecular probes (Thermo Fisher Scientific, Waltham, MS, USA). Both rotenone and ruthenium red were purchased from Sigma-Aldrich (St. Louis, MO, USA). Vehicle for dyes and other agents was HBSS buffer (150 mmol/l Na\textsuperscript{+}, 5 mmol/l K\textsuperscript{+}, 1.0 mmol/l Ca\textsuperscript{2+}, 1 mmol/l Mg\textsuperscript{2+}, and 20 mmol/l HEPES at pH 7.4). All esterified fluorescent probes were prepared as stock solutions in dimethylsulfoxide.

**Primary AEC type 2 isolation and cell culture.** Isolation and culture of AEC type 2 from adult male Sprague-Dawley rats according to procedures described previously\textsuperscript{20,21} In brief, after injection of 100 mg/kg ketamine and 20 mg/kg xylazinum (rompun), the lungs were surgically exposed and perfused with 3 \times 10 ml F-12 K medium (ATCC, Manassas, VA, USA) supplemented with 25 mM HEPES (Sigma-Aldrich) to remove blood. After multiple lavages (5 \times 10 ml) via the cannulated trachea with 5 mM ethylenediaminetetraacetic acid (EDTA, Sigma-Aldrich) and 5 mM ethylene glycol-bis (β-aminoethyl ether)-N,N,N\textsubscript{,}N\textsubscript{,} tetraacetic acid (Sigma-Aldrich) in a phosphate-buffered salt solution the lungs were removed from the body. Upon instillation of 20 ml warm F-12K medium (ATCC) enriched with 50 mM HEPES and 4.5 units/ml elastase Grade II (Worthington, Lakewood NJ, USA) the lungs were incubated at 37 °C for 30 min. After dissection of the large airways the lungs were added to 10 ml F-12K medium containing 25 mM HEPES, 20% fetal bovine serum (FBS, Biochrom AG, Berlin, Germany), and 100 μg/ml DNAse (Typ IV, Sigma-Aldrich) quickly minced and incubated at 4 °C for 10 min. Then the solution was mixed by end-over end-rotation for 4 min and filtered successively through cotton gauze (1, 2, and 4 ply, respectively) following filtration through nylon mesh (100 and 40 μm, BD Falcon, Corning, NY, USA). Upon centrifugation of the cell suspension at 150 \times g for 10 min and resuspension in 20 ml warm DMEM (Biochrom AG) the cells were applied to plates covered with rat immunoglobulin (lg) G (Sigma-Aldrich) for 1 h at 37 °C. The nonadherent cells were removed, collected in sterile tube, twice centrifuged at 150 \times g for 10 min and resuspended in 3 ml of culture medium (DMEM with 10% FBS, 1% penicillin/streptomycin and 10 ng/ml keratinocyte growth factor (Sigma-Aldrich)). After determination of cell count and viability, cells were plated on collagen-treated coverslips and incubated at 37 °C and 5% CO\textsubscript{2}.

Human lung carcinoma A549 cells were grown in F-12K Medium (ATCC) with 10% FBS, 10 000 U/ml penicillin and 10 000 μg/ml streptomycin (Biochrom AG). A549 is an alveolar epithelial cell line with type 2 cell characteristics including production of surfactant proteins. It has been widely used as a model to study the response of AEC type 2 in which the mitochondrial TCA enzyme IDH3 has constitutive expression. According to our results one could consider a novel ARDS therapy, namely CO\textsubscript{2} inhalation in order to prevent hypocapnia-induced apoptosis of AECs and tissue injury in ARDS.

**Conventional fluorescent real-time cell imaging.** For fluorescent imaging, a coverslip with fluorophore-loaded primary isolated AEC type 2 or A549 cells was mounted in an imaging/perfusion chamber on the stage of a conventional epifluorescence microscope (Olympus, Hamburg, Germany). The imaging chamber was perfused continuously with warm HEPES buffer (37 °C). Fluorophores were excited by a mercury arc lamp illumination directed through appropriate interference filters and filter sets (Semrock, Rochester, NY, USA).
Fluorescence exposure was controlled by a filter wheel (Sutter Lambda 10-C, Sutter Instrument Co). The fluorescence emission was collected using an objective lens (40 × water immersion, numerical aperture 0.8, Zeiss, Oberkochen, Germany) and captured with a charge-coupled device camera (Photometrics Coolsnaps HQ2, Tuscan, AZ, USA).

\([\text{Ca}^{2+}]_{\text{im}}\) determination. For \([\text{Ca}^{2+}]_{\text{im}}\) measurements, cells were loaded for 45 min with 10 \(\mu\text{M}\) fura 2-AM in physiological solution containing 0.02% pluronic. The cells were excited at 340 and 380 nm. The fluorescence emissions at 510 nm were recorded and \([\text{Ca}^{2+}]_{\text{im}}\) was calculated from a computer-generated 340:380 emissions ratio based on a dissociation constant of 224 nmol/l and appropriate calibration parameters.24

Fluorescence images were recorded under normo- or hypocapnic conditions in combination with an extracellular pH of 7.4 or 7.8 at 10 s intervals.

\([\text{Ca}^{2+}]_{\text{im}}\) determinations. For \([\text{Ca}^{2+}]_{\text{im}}\) measurements, cells were loaded with rhod-2-AM for 15 min followed by a Ringer’s flush. To improve the specificity of \([\text{Ca}^{2+}]_{\text{im}}\) measurement mitochondria were additionally localized by the fluorophore MTG, which binds specifically to the inner mitochondrial membrane.28 The wavelength for excitation was 560 nm, whereas monitoring the emission at 590 nm.

Mitochondrial membrane potential measurement. For mitochondrial membrane potential assessment, primary isolated AEC type 2 or A549 cells were loaded with TMRM. This indicator dye is a lipophilic cation accumulated by mitochondria in proportion to the electrical potential across their inner membrane (ΔΨm).26 The wavelength for excitation was 560 nm while monitoring the emission at 590 nm. Addition of the mitochondrial depolarizer FCCP (50 nM) resulted in a rapid loss of the mitochondrial TMRM fluorescence.

Mitochondrial NADH production. NADH concentration was quantified by measuring its auto-fluorescence intensity. Primary isolated AEC type 2 or A549 cells were excited with 360 nm while monitoring the emission at 510 nm. For determining the mitochondrial production of NADH, its auto-fluorescence was co-localized with the mitochondria-specific marker MTG.

Determination of mitochondrial ROS Production. Mitochondrial ROS production was measured using MitosOX Red (Thermo Fisher Scientific). Primary isolated AEC type 2 were stained with 10 \(\mu\text{M}\) MitosOX Red and 10 \(\mu\text{M}\) MTG (staining of mitochondria) for 10 min at 37 °C. Upon washing with normocapnia buffer, cells were then imaged with a conventional epifluorescence microscopy (Olympus) in normo- and hypocapnia puffer, respectively. Changes in the fluorescence intensity of MitosOX Red were measured at 550 nm excitation and 600 nm emission wavelengths. In an additional set of experiments we preincubated the cell with 1 \(\mu\text{M}\) rotenone to inhibit Complex I of the electron chain and/or 10 \(\mu\text{M}\) ruthenium red to inhibit the MCU. As a positive control we used Antimycin AA (2 \(\mu\text{g/ml}\)).

Determination of Caspase-3/7-positive cells. We used the Cell Event Caspase-3/7 Green Detection Reagent (Thermo Fisher Scientific) according to the manufacture’s instruction to determine apoptotic cells. In brief, upon incubation with normo- und hypocapnia puffer for 30 min at 37 °C primary isolated AEC type 2 were stained 24 h later with the caspase-3/7 reagent (10 \(\mu\text{M}\)) and counterstained with the nuclei dye Hoechst 33342 (0.5 \(\mu\text{g/ml}\)). Cells were then image with a conventional epifluorescence microscopy (Olympus) and caspase-3/7-positive cells were counted in four independent fields per well.

2-Photon and FLIM. To determine changes of NAD+/NADH during inhibition of complex I and/or under hypocapnia treatment, FLIM was performed on a single-beam 2-photon microscope setup described elsewhere.27 In brief, the excitation wavelength of the pulsed Ti:Sa Laser (MaiTai, Spectra Physcis) was 750 nm and sent through a water-dipping lens (20 ×, NA 0.95, WD 2 mm – Olympus) to the lung surface. Emission photons passed a band-pass filter (475/47 nm) and were finally detected at the Ultra-sensitive-port by a 16-channel time-correlated single photon counter, with a maximum rate of 78 MHz (LaVision Biotec, Bielefeld, Germany). Evaluation of the multi-exponential fluorescence signal was performed by the self-developed software RINIFLIM programmed in Matlab (MathWork). For evaluation, Levenberg-Marquardt algorithms were used.

**IDH3 activity.** The NAD⁺-dependent IDH3 activity was determined using the commercial available IDH activity assay kit according to the manufactures protocol (Sigma-Aldrich). In brief, the IDH3 activity in cell extracts was measured using the substrate isocitrate and the cofactor NAD⁺ in an enzyme reaction, which results in a colorimetric (450 nm) product proportional to NADH concentration. The enzyme activity was calculated according to the standard curve. One unit of IDH3 is the amount that will generate 1.0 \(\mu\text{mol}\) NADH per minute at pH 8.0 at 37 °C.

**IDH3 and MCU knockdown.** A549 cells were transfected with 10 nM stealth siRNA of IDH3 (Thermo Fisher Scientific) and/or 10 nM siRNA of MCU (Quiagen, Hilden Germany) and scrambled siRNA (Thermo Fisher Scientific and Quiagen) by using Lipofectamine RNAiMAX (Thermo Fisher Scientific) according to the manufacturer’s protocol. In brief, siRNA was diluted in 500 \(\mu\text{l}\) Opti-MEM I medium (Quiagen) without serum in a well of the tissue culture plate. Lipofectamine RNAiMAX was added to each well containing the diluted siRNA. Cells were diluted in complete growth medium without antibodies so that the cell density after 24 h would be 30–50% confluent after plating.

**Real-time quantitative RT-PCR for IDH3 and MCU knockdown analysis.** Total RNA was isolated using the RNeasy Kit (Qiagen). A total of 2 \(\mu\text{g}\) of RNA were used for cDNA synthesis using the Maxima First Strand cDNA Synthesis Kit (Thermo Fisher Scientific) according to the manufacturer’s instructions. qRT-PCR was performed with gene- and species-specific primers. cDNA template was amplified using the QuantNova SYB GreenPCR Kit (Qiagen) on a Rotor-Gene (Qiagen) according to the manufacturer’s protocol. Quantitative analysis of data were performed by using the deltaDeltaCT method.

**Western blot for IDH3 and MCU knockdown analysis.** For immunoblotting, cells were washed twice with PBS and then snapped into RIPA buffer (Sigma-Aldrich) supplemented with the protease inhibitor cocktail complete Mini EDTA-free (Roche Diagnostics, Mannheim, Germany). Upon incubation on ice for 30 min lysed cells were centrifuged at 12 000 × g for 20 min (4 °C). The supernatants with the cytoplasmatic protein were collected, and protein concentrations were determined by using the bicinchoninic acid (BCA protein assay kit, Thermo Fisher Scientific), assay according to the manufacturer’s instructions. A total of 20 \(\mu\text{g}\) samples were reduced in sample buffer (NuPAGE sample buffer, Thermo Fisher Scientific) plus reducing agent (NuPAGE sample reducing agent, Invitrogen) at 95 °C for 5 min and run on a 4–12% Bis-Tris polyacrylamid gel (NuPage, Thermo Fisher Scientific) and transferred to nitrocellulose filters (Thermo Fisher Scientific). The filters were blocked for 1 h at room temperature in blocking puffer (PBS plus 0.1% Tween 20 containing 5% milk powder and/or 5% bovine serum albumin (BSA)) with agitation. After application to the polyclonal rabbit IDH3A antibody (1:800, GenWay Biotech Inc., San Diego, CA, USA) or to the monoclonal rabbit MCU antibody (1:1000, Cell Signaling Technology, Danvers, MA, USA) overnight at 4 °C, the filters were exposed to an ECL horseradish peroxidase-conjugated donkey anti-rabbit IgG (1:10 000 and/or 1:20 000), GE Healthcare, Little Chalfont, UK) for 1 h at room temperature with agitation. After washing with washing puffer (PBS plus 0.1% Tween 20), the filters were developed with the ECL Plus Western Blotting Detection Reagents (GE Healthcare) according the manufacturer’s protocol. For molecular weight standards, we used a MagicMark XP protein ladder (Thermo Fisher Scientific).

**Western blot analysis for PARP cleavage.** Upon incubation with normo- and hypocapnia puffer for 3 h at 37 °C primary isolated AEC type 2 were collected 24 h later and prepared for immunoblotting as described in the section ‘Western blot IDH3 and MCU knockdown analysis’. After gel electrophoresis and immunoblotting, filter was blocked for 1 h at room temperature in blocking puffer (PBS plus 0.1% Tween 20 containing 5% BSA) with agitation. Upon application of the monoclonal rabbit Anti-cleaved-PARP antibody (1:500, Abcam, Cambridge, UK) overnight at 4 °C, the filters were exposed to an ECL horseradish peroxidase-conjugated donkey anti-rabbit IgG (1:10 000 and/or 1:20 000), GE Healthcare, Little Chalfont, UK) for 1 h at room temperature with agitation. After washing washing puffer (PBS plus 0.1% Tween 20), the filters were developed with the ECL Plus Western Blotting Detection Reagents (GE Healthcare) according the manufacturer’s protocol.

**For molecular weight standards, we used a MagicMark XP protein ladder (Thermo Fisher Scientific).**

**As a positive control we used staurosporine (1 \(\mu\text{M}\)).**

**Statistics.** All data are represented as means ± S.E. Statistical data analysis was performed using SigmaStat (Systat Software, Chicago, IL, USA). Comparisons between two groups were tested using RankSumTest. Comparisons between more groups were tested by ANOVA on ranks followed by a pairwise multiple comparison.
Repeated measurements were tested using the Mann–Whitney Rank Sum Test. Statistical significance was accepted at $P<0.05$.

**Conflict of Interest**

The authors declare no conflict of interest.

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