

# HIF and HOIL-1L-mediated PKC $\zeta$ degradation stabilizes plasma membrane Na,K-ATPase to protect against hypoxia-induced lung injury

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Organisms have evolved adaptive mechanisms in response to stress for cellular survival. During acute hypoxic stress, cells down-regulate energy-consuming enzymes such as Na,K-ATPase. Within minutes of alveolar epithelial cell (AEC) exposure to hypoxia, protein kinase C zeta (PKC $\zeta$ ) phosphorylates the  $\alpha_1$ -Na,K-ATPase subunit and triggers it for endocytosis, independently of the hypoxia-inducible factor (HIF). However, the Na,K-ATPase activity is essential for cell homeostasis. HIF induces the heme-oxidized IRP2 ubiquitin ligase 1L (HOIL-1L), which leads to PKC $\zeta$  degradation. Here we report a mechanism of prosurvival adaptation of AECs to prolonged hypoxia where PKC $\zeta$  degradation allows plasma membrane Na,K-ATPase stabilization at ~50% of normoxic levels, preventing its excessive down-regulation and cell death. Mice lacking HOIL-1L in lung epithelial cells (*Cre<sup>SPC</sup>/HOIL-1L<sup>fl/fl</sup>*) were sensitized to hypoxia because they express higher levels of PKC $\zeta$  and, consequently, lower plasma membrane Na,K-ATPase levels, which increased cell death and worsened lung injury. In AECs, expression of an  $\alpha_1$ -Na,K-ATPase construct bearing an S18A ( $\alpha_1$ -S18A) mutation, which precludes PKC $\zeta$  phosphorylation, stabilized the Na,K-ATPase at the plasma membrane and prevented hypoxia-induced cell death even in the absence of HOIL-1L. Adenoviral overexpression of the  $\alpha_1$ -S18A mutant Na,K-ATPase in vivo rescued the enhanced sensitivity of *Cre<sup>SPC</sup>/HOIL-1L<sup>fl/fl</sup>* mice to hypoxic lung injury. These data suggest that stabilization of Na,K-ATPase during severe hypoxia is a HIF-dependent process involving PKC $\zeta$  degradation. Accordingly, we provide evidence of an important adaptive mechanism to severe hypoxia, whereby halting the exaggerated down-regulation of plasma membrane Na,K-ATPase prevents cell death and lung injury.

hypoxia | Na,K-ATPase | HOIL-1L | PKC $\zeta$  | alveolar epithelial cells

Short-term exposure to hypoxia leads to inhibition of Na,K-ATPase, a high-ATP-consuming enzyme, independently of hypoxia-inducible factor (HIF) (1). However, during prolonged hypoxia, HIF plays an important role in maintaining cell homeostasis (2–5). HIF regulates genes that increase energy generation via anaerobic glycolysis and those that decrease ATP-consuming enzymes, thereby preserving cell metabolism during hypoxia (2–4, 6). The Na,K-ATPase utilizes ~30% of the cell's ATP under basal conditions to maintain the Na<sup>+</sup> and K<sup>+</sup> concentration gradients across the cell membrane necessary for cellular homeostasis (1, 7, 8). Hypoxia occurs in individuals with normal respiratory function during ascent to high altitude and in patients with pulmonary edema due to heart failure and acute lung injury (9–11). Hypoxia has been reported to inhibit edema reabsorption from the alveolar spaces by inhibiting the sodium channels, which are responsible for the apical sodium entry, and basolateral membrane Na,K-ATPase, which is responsible for Na<sup>+</sup> extrusion (12–14). The hypoxia-mediated down-regulation of the Na,K-ATPase at the

alveolar epithelial cell (AEC) basolateral membrane is mediated by protein kinase C zeta (PKC $\zeta$ ) phosphorylation of the Na,K-ATPase  $\alpha_1$  catalytic subunit at Ser-18, which in turn triggers Na,K-ATPase endocytosis (1, 15–17).

PKC isoenzymes play a role in the cellular adaptation to stress by regulating survival, proliferation, migration, and apoptosis (18–21). PKC $\zeta$  is a member of the atypical class of PKC isoforms. Unlike the conventional and novel isoforms, atypical PKCs do not respond to the second messenger diacylglycerol or calcium, but they are activated by stimuli-dependent phosphorylation (22, 23). In the basal state, PKCs are auto-inhibited by their pseudosubstrates and converted into catalytically competent enzymes by a series of phosphorylations (18, 22, 24). However, the mechanisms that regulate termination of PKC signaling are incompletely understood. We reported that, in cancer cells, tumor growth is promoted via

## Significance

Exposure to hypoxia requires adaptive mechanisms for survival. During acute hypoxia, Na,K-ATPase endocytosis in alveolar epithelial cells occurs via protein kinase C zeta (PKC $\zeta$ ) phosphorylation of  $\alpha_1$ -Na,K-ATPase independently of the hypoxia-inducible factor (HIF). However, exaggerated Na,K-ATPase down-regulation leads to cell death. Here we report that during prolonged hypoxia plasma membrane Na,K-ATPase levels were maintained at ~50% of normoxic values due to HIF-mediated up-regulation of HOIL-1L, which targets PKC $\zeta$  for degradation. Silencing HOIL-1L in the lung epithelium prevented PKC $\zeta$  degradation, causing Na,K-ATPase downregulation. Accordingly, HIF regulation of HOIL-1L targets the phosphorylated PKC $\zeta$  for degradation and serves as an hypoxia-adaptive mechanism to stabilize the Na,K-ATPase, avoiding significant lung injury.

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the transcription of heme-oxidized IRP2 ubiquitin ligase 1L (HOIL-1L), which acts as the E3 ubiquitin ligase for PKC $\zeta$ , targeting it for proteasomal degradation (20, 25). HOIL-1L together with HOIL-1-interacting protein (HOIP) and Shank-associated RH-domain-interacting protein (SHARPIN) form the linear ubiquitination assembly complex (LUBAC) (26–29). We and others have found that, when acting independently of LUBAC, HOIL-1L adds Lys-48-linked chains and serves as an ubiquitin E3 ligase (20, 30).

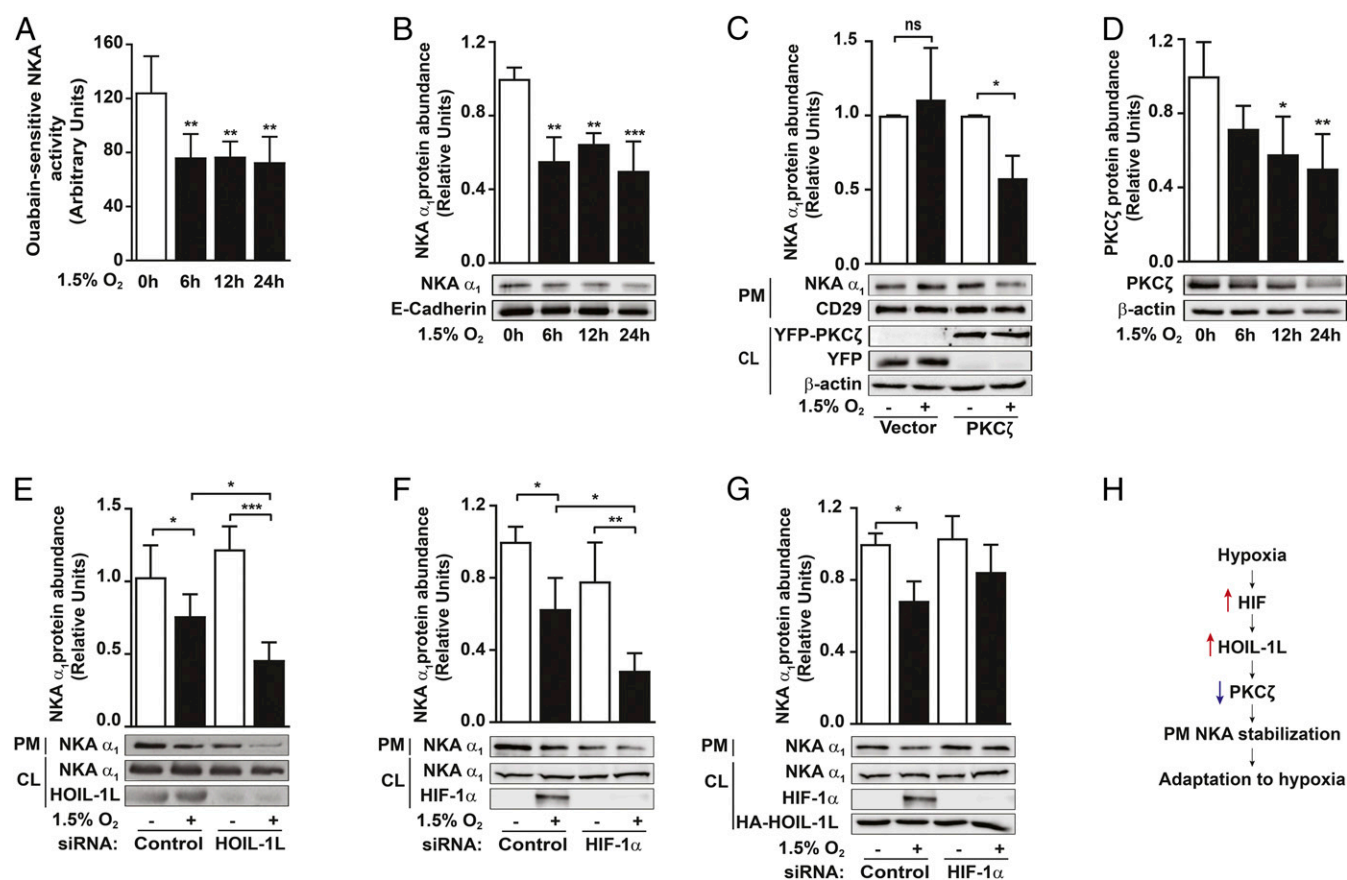
Here we report that, in lung epithelial cells exposed to prolonged hypoxia in vitro, the Na,K-ATPase is stabilized at a plateau lower than levels in normoxic conditions via a HIF-mediated up-regulation of HOIL-1L. Hypoxia promotes the translocation of phosphorylated PKC $\zeta$  to the plasma membrane where it interacts with HOIL-1L, which targets it for degradation. This PKC $\zeta$  degradation limits Na,K-ATPase down-regulation and safeguards alveolar epithelial function. To examine this pathway in vivo, we generated mice with lung epithelial-specific deletion of HOIL-1L (*Cre<sup>SPC</sup>/HOIL-1L<sup>fl/fl</sup>*). Exposure of these mice to prolonged hypoxia resulted in higher levels of PKC $\zeta$ , excessive depletion of plasma membrane Na,K-ATPase, and more severe lung injury.

This sensitivity was rescued by overexpression of a phosphorylation-resistant mutant, Na,K-ATPase. Collectively, these data suggest that HOIL-1L-mediated ubiquitination and degradation of PKC $\zeta$  results in Na,K-ATPase stabilization at the plasma membrane as an important adaptive mechanism to preserve the alveolar epithelium from significant injury.

## Results

### PKC $\zeta$ Degradation by HOIL-1L Stabilizes Plasma Membrane Na,K-ATPase During Prolonged Hypoxia.

During prolonged exposure of primary rat alveolar type II (ATII) cells to 1.5% O<sub>2</sub> (hypoxia), Na,K-ATPase activity and protein abundance at the plasma membrane decreased by ~50%, compared with cells exposed to 21% O<sub>2</sub> (normoxia), and then stabilized (Fig. 1*A* and *B*). As observed in *SI Appendix*, Fig. S1*A–C*, mitochondrial ATP production decreased after 6 h of hypoxia compared with the normoxic values. However, all hypoxia time points showed similar intracellular ATP content (*SI Appendix*, Fig. S1*D*). In pVHL-deficient renal clear carcinoma cells (RCC4), where HIF is stabilized, HOIL-1L levels increased while PKC $\zeta$  levels decreased (20, 31) and hypoxia-induced  $\alpha_1$ -Na,K-ATPase endocytosis was prevented (Fig. 1*C*). Overexpression of

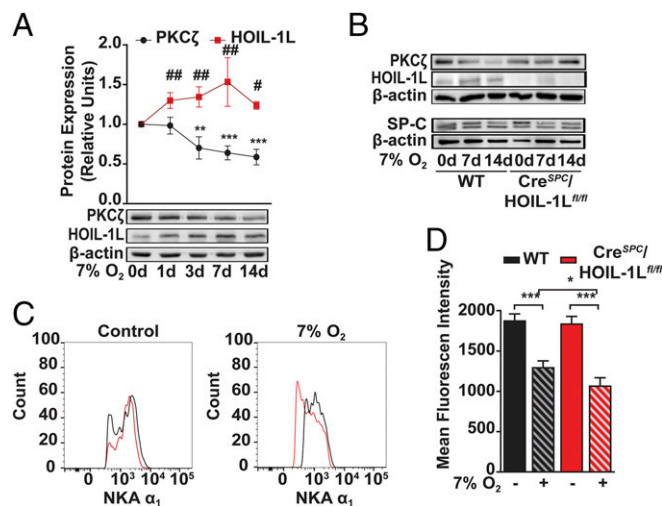


**Fig. 1.** PKC $\zeta$  degradation by HOIL-1L leads to the stabilization of plasma membrane Na,K-ATPase during hypoxia. (*A* and *B*) Rat ATII (r-ATII) cells were exposed to normoxia (21% O<sub>2</sub>) or hypoxia (1.5% O<sub>2</sub>) for up to 24 h. (*A*) Na,K-ATPase activity was measured in isolated membranes by a colorimetric assay as described in *Experimental Procedures* ( $n = 6$ ). (*B*) Plasma membrane  $\alpha_1$ -Na,K-ATPase (NKA  $\alpha_1$ ) was determined after cell-surface labeling with biotin in cells exposed to hypoxia or normoxia by immunoblot with a NKA  $\alpha_1$ -specific antibody ( $n = 3$ ). (*C*) RCC4 cells were transfected with empty vector (Vector) or YFP-PKC $\zeta$  (PKC $\zeta$ ). Forty-eight hours after transfection, cells were exposed to normoxia or hypoxia for 2 h, and then the expression of plasma membrane (PM)  $\alpha_1$ -Na,K-ATPase was determined after cell-surface labeling with biotin and determined by Western blot. Transfection efficiency was determined by performing an immunoblot of cell lysate (CL) proteins using YFP-specific antibody and  $\beta$ -actin antibody as a loading control ( $n = 6$ ). (*D*) PKC $\zeta$  levels were assessed in the cell lysates of r-ATII cells exposed to normoxia or hypoxia for up to 24 h by Western blot ( $n = 5$ ). (*E–G*) A549 cells were transfected with the indicated siRNA and exposed to normoxia or hypoxia for 24 h. Expression of  $\alpha_1$ -Na,K-ATPase at the PM was determined by cell-surface biotinylation and Western blot ( $n = 3$ ). (*H*) Schematic representation of the hypoxia-induced pathway for PM Na,K-ATPase stabilization. Representative immunoblots and bar graphs showing the densitometry quantifications of immunoblots in relation to the corresponding loading controls are shown. Data are expressed as mean  $\pm$  SD. Statistical significance was calculated using one-way ANOVA and the Tukey multiple comparisons test (\* $P < 0.05$ , \*\* $P < 0.01$ , \*\*\* $P < 0.001$ ).

YFP-PKC $\zeta$  in these cells restored the Na,K-ATPase endocytosis (Fig. 1C). Hypoxia does not affect PKC $\zeta$  mRNA levels or protein synthesis (20). However, hypoxia caused a time-dependent decrease of PKC $\zeta$  levels in rat A111 cells (Fig. 1D). To assess the role of LUBAC during the hypoxia-induced Na,K-ATPase down-regulation, we performed loss-of-function experiments using A549 cells. Even though A549 cells are tumor cells, we and many others have reported that they have many characteristics of A111 cells (32, 33), including the regulation of Na,K-ATPase (1, 33, 34). HOIL-1L in A549 cells led to a more significant decrease of plasma membrane  $\alpha_1$ -Na,K-ATPase abundance during hypoxia compared with cells transfected with control siRNA (Fig. 1E). As with HOIL-1L silencing, HIF1 $\alpha$  silencing resulted in fewer Na,K-ATPase molecules observed at the plasma membrane of AECs exposed to hypoxia compared with cells transfected with a control siRNA and exposed to hypoxia (Fig. 1F). Overexpression of HOIL-1L rescues the effects of HIF1 $\alpha$  silencing, preventing the Na,K-ATPase endocytosis (Fig. 1G). However, silencing of HOIP did not further alter the  $\alpha_1$ -Na,K-ATPase abundance at the plasma membrane after 24 h of hypoxia (SI Appendix, Fig. S1E). Collectively, these data suggest that, in AECs exposed to hypoxia, HIF regulates HOIL-1L, which, by promoting PKC $\zeta$  degradation, leads to plasma membrane Na,K-ATPase stabilization (Fig. 1H).

**HOIL-1L Silencing Leads to Exaggerated  $\alpha_1$ -Na,K-ATPase Down-Regulation in Lung Epithelium During Hypoxia.** Analysis of peripheral lung tissue cell lysates from C57BL/6 (WT) mice exposed to 7% O<sub>2</sub> (hypoxia) for up to 14 d showed a significant increase in HOIL-1L in parallel with a decrease in PKC $\zeta$  protein abundance (Fig. 2A). We generated *Cre<sup>SPC</sup>/HOIL-1L<sup>fl/fl</sup>* mice, which bear a lung epithelial-specific deletion of the *Rbck1* (HOIL-1L) gene (as described in Experimental Procedures). We analyzed, by RNA sequencing, the transcriptome of A111 cells isolated from WT and *Cre<sup>SPC</sup>/HOIL-1L<sup>fl/fl</sup>* mice in basal conditions. Only 20 of 13,617 detected genes were differentially expressed, suggesting that the deletion of HOIL-1L in the alveolar epithelium did not cause major changes in the epithelium (SI Appendix, Table S1). In contrast to the observed decrease in PKC $\zeta$  levels in A111 cells from control mice exposed to hypoxia, PKC $\zeta$  levels did not change in A111 cells isolated from *Cre<sup>SPC</sup>/HOIL-1L<sup>fl/fl</sup>* mice (Fig. 2B). The expression levels of neither HOIP nor SHARPIN, the other components of LUBAC, changed in lung tissue of mice exposed to hypoxia (SI Appendix, Fig. S2A). We assessed, by flow cytometry, the effects of HOIL-1L deletion on  $\alpha_1$ -Na,K-ATPase abundance at the plasma membrane of A111 cells from mice exposed to 7% O<sub>2</sub> for 7 d (SI Appendix, Fig. S2B, for gating strategy) and found no differences between the  $\alpha_1$ -Na,K-ATPase abundance in A111 cells from WT (black) or *Cre<sup>SPC</sup>/HOIL-1L<sup>fl/fl</sup>* mice (red) kept in room air (control) (Fig. 2C and D, solid bars), whereas A111 cells from *Cre<sup>SPC</sup>/HOIL-1L<sup>fl/fl</sup>* mice exposed to 7% O<sub>2</sub> had lower  $\alpha_1$ -Na,K-ATPase abundance compared with cells from WT mice exposed to hypoxia (Fig. 2C and D, striped bars). These data suggest that in AECs from mice exposed to hypoxia, the absence of HOIL-1L leads to increased PKC $\zeta$  levels and decreased plasma membrane  $\alpha_1$ -Na,K-ATPase protein abundance.

**Hypoxia Causes Lung Injury in Mice with HOIL-1L Deletion in the Lung Epithelium.** To assess whether deletion of HOIL-1L in the alveolar epithelium affects lung function, we exposed mice to 7% O<sub>2</sub> for 7 d and found that the epithelial permeability to small solutes increased (Fig. 3A), as well as the total number of cells and protein concentration in the bronchoalveolar lavage fluid (BALF) (Fig. 3B and C) in *Cre<sup>SPC</sup>/HOIL-1L<sup>fl/fl</sup>* mice compared with WT mice. H&E staining of lung tissue revealed that in lungs of *Cre<sup>SPC</sup>/HOIL-1L<sup>fl/fl</sup>* mice exposed to low O<sub>2</sub> conditions there was increased thickening in the alveolar septa, more noticeable in the peribronchiolar area, compared with WT mice lungs exposed to hypoxia (Fig. 3D and E). Moreover, in lungs of *Cre<sup>SPC</sup>/HOIL-1L<sup>fl/fl</sup>* mice exposed to hypoxia



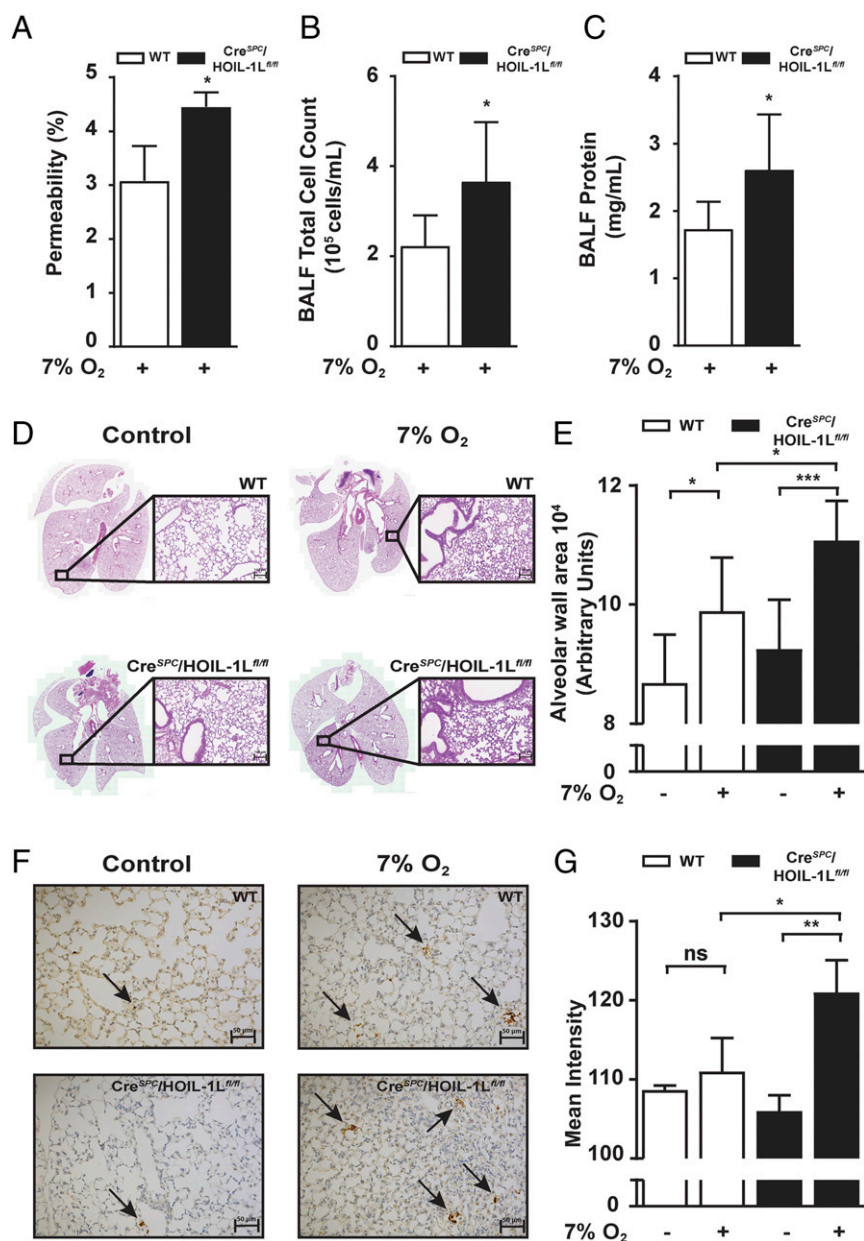
**Fig. 2.** Silencing of HOIL-1L in the alveolar epithelium decreases Na,K-ATPase protein abundance at the plasma membrane in vivo. (A) WT mice were kept in room air (0 d) or exposed to 7% O<sub>2</sub> for up to 14 d, and PKC $\zeta$  (black) and HOIL-1L (red) levels were analyzed by Western blot in lung peripheral tissue homogenates. Representative immunoblots and graph show the densitometry quantifications of proteins in relation to  $\beta$ -actin. Graph represents mean  $\pm$  SD ( $n = 4$ ). (B) Mouse A111 cells were isolated from WT and *Cre<sup>SPC</sup>/HOIL-1L<sup>fl/fl</sup>* mice treated as in A. Levels of PKC $\zeta$ , HOIL-1L, SP-C, and  $\beta$ -actin were determined in cell lysates by Western blot. SP-C was used as a marker of A111 cells. Representative immunoblots are shown ( $n = 3$ ). (C and D) The  $\alpha_1$ -Na,K-ATPase (NKA  $\alpha_1$ ) levels at the plasma membrane were assessed by flow cytometry of EpCAM<sup>+</sup> epithelial cells from lung homogenates of WT (black) and *Cre<sup>SPC</sup>/HOIL-1L<sup>fl/fl</sup>* (red) mice kept in room air or exposed to 7% O<sub>2</sub> for 7 d. (C) Representative histograms from three independent experiments are shown. (Left) Room air (control). (Right) 7% O<sub>2</sub>. (D) Quantification of  $\alpha_1$ -Na,K-ATPase levels comparing  $\alpha_1$ -Na,K-ATPase mean fluorescence intensity from room air (solid bars) and 7% O<sub>2</sub> for 7 d (striped bars;  $n = 3$ ). Graph bars represent mean  $\pm$  SD. Statistical significance was calculated using one-way ANOVA and the Tukey multiple comparisons test (\* $P < 0.05$ , \*\* $P < 0.01$ , \*\*\* $P < 0.001$ ).

there was increased apoptosis (as measured by TUNEL assay) compared with lungs of WT mice (Fig. 3F and G).

It has been reported that hypoxia induces an inflammatory response with recruitment of immune cells into the alveolar spaces (35). To assess whether the increased injury observed in *Cre<sup>SPC</sup>/HOIL-1L<sup>fl/fl</sup>* mice in response to hypoxia was due to an exaggerated immune response, we performed flow cytometric analysis of the immune cells in the lungs. While the numbers of neutrophils and interstitial macrophages in mice exposed to hypoxia were elevated, there was no difference in immune cell composition between WT and *Cre<sup>SPC</sup>/HOIL-1L<sup>fl/fl</sup>* mice either in basal state or after being exposed to hypoxia (SI Appendix, Fig. S3).

To determine whether the HOIL-1L-mediated cell adaptation to hypoxia was due to the stabilization of Na,K-ATPase at the plasma membrane, we used A549 cells expressing GFP- $\alpha_1$ -Na,K-ATPase-WT ( $\alpha_1$ -WT) or GFP- $\alpha_1$ -Na,K-ATPase-S18A ( $\alpha_1$ -S18A). The Na,K-ATPase bearing an S18A mutation is resistant to PKC $\zeta$  phosphorylation and thus fails to internalize during hypoxia (1, 33, 36). In  $\alpha_1$ -WT cells transfected with HOIL-1L-specific siRNA, exposure to hypoxia significantly increased cell death compared with  $\alpha_1$ -WT cells transfected with a scrambled siRNA (Fig. 4A). In contrast, in  $\alpha_1$ -S18A A549 cells neither the silencing of HOIL-1L nor the exposure to hypoxia increased cell death (Fig. 4A), supporting the hypothesis that during prolonged hypoxia HOIL-1L exerts a protective effect by promoting PKC $\zeta$  degradation, which stabilizes the  $\alpha_1$ -Na,K-ATPase at the AEC plasma membrane.



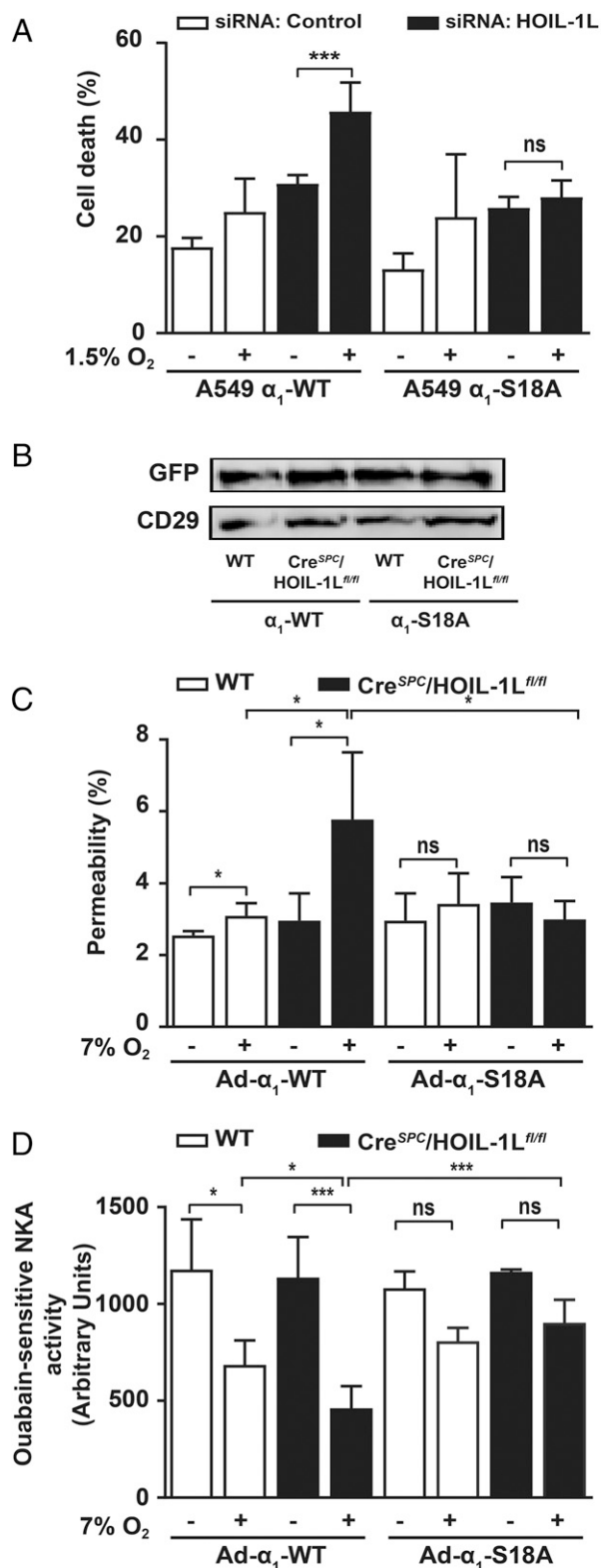


**Fig. 3.** HOIL-1L deletion in lung epithelial cells causes lung injury in mice exposed to hypoxia. WT and *Cre<sup>SPC</sup>/HOIL-1L<sup>fl/fl</sup>* mice were kept in room air (21% O<sub>2</sub>) or exposed to hypoxia (7% O<sub>2</sub>) for 7 d. (A) Alveolar epithelial permeability is expressed as the ratio of the fluorescence present in BALF to the plasma ( $n = 3$ ). (B) Total cell count in BALF ( $n = 6$ ). (C) Total protein concentration in BALF. Graph bars represent mean  $\pm$  SD ( $n = 6$ ). Statistical significance was calculated for A–C using the Student's *t* test ( $*P < 0.05$ ). (D) Lung sections were stained with H&E, and brightfield serial images were obtained. Images show representative montages using 20 $\times$  magnification images and higher magnification views of the areas indicated by squares. (Scale bar, 50  $\mu$ m.) (E) Quantification of the alveolar wall area using ImageJ/FIJI 1.46 from WT (white bars) and *Cre<sup>SPC</sup>/HOIL-1L<sup>fl/fl</sup>* (black bars) mouse lungs. Graph bars represent mean  $\pm$  SD ( $n = 3$ ). (F) Lung sections were stained with TUNEL, and brightfield serial 10 $\times$  magnification images were obtained. Arrows point to TUNEL-positive nuclei. (Scale bar, 50  $\mu$ m.) (G) Quantification of TUNEL-positive nuclei mean intensity per lung section montage from WT (white bars) and *Cre<sup>SPC</sup>/HOIL-1L<sup>fl/fl</sup>* (black bars) mouse lungs. Graph bars represent mean  $\pm$  SD ( $n = 3$ ). Statistical significance for E and G was calculated using one-way ANOVA and the Tukey multiple comparisons test ( $*P < 0.05$ ,  $**P < 0.01$ ,  $***P < 0.001$ ).

We also infected WT and *Cre<sup>SPC</sup>/HOIL-1L<sup>fl/fl</sup>* mice before exposure to 7% O<sub>2</sub> with an adenoviral construct coding for GFP- $\alpha_1$ -Na,K-ATPase -WT (Ad  $\alpha_1$ -WT) or GFP- $\alpha_1$ -Na,K-ATPase-S18A (Ad  $\alpha_1$ -S18A). The infection efficiency was assessed by measuring the expression of GFP- $\alpha_1$ -Na,K-ATPase in membranes from lung peripheral tissue by immunoblotting with an anti-GFP antibody (Fig. 4B). As observed in noninfected mice (Fig. 3A), lungs of *Cre<sup>SPC</sup>/HOIL-1L<sup>fl/fl</sup>* mice expressing Ad  $\alpha_1$ -WT exposed to hypoxia had increased permeability and total cell counts in the BALF compared with WT mice (Fig. 4C and *SI Appendix, Fig. S4*).

However, *Cre<sup>SPC</sup>/HOIL-1L<sup>fl/fl</sup>* mice expressing  $\alpha_1$ -S18A exposed to hypoxia did not have significant changes in lung permeability or total cell counts compared with mice breathing room air (Fig. 4C and *SI Appendix, Fig. S4*). Moreover, the decrease in Na,K-ATPase activity observed in *Cre<sup>SPC</sup>/HOIL-1L<sup>fl/fl</sup>* mice exposed to hypoxia was rescued in mice infected with Ad- $\alpha_1$ -S18A (Fig. 4D).

**Phosphorylation at Thr-410 Is Required for PKC $\zeta$  Ubiquitination and Degradation.** Phosphorylation in the activation loop of the PKC $\zeta$  kinase domain at Thr-410 exposes it to further phosphorylation



**Fig. 4.** Alveolar epithelial function in *Cre<sup>SPC</sup>/HOIL-1L<sup>fl/fl</sup>* mice exposed to hypoxia is rescued by Na,K-ATPase  $\alpha_1$ -S18A overexpression. (A) Cell death was measured using a lactate dehydrogenase (LDH) assay in incubation medium from A549 cells stably expressing rat Na,K-ATPase  $\alpha_1$ -WT (A549  $\alpha_1$ -WT) or  $\alpha_1$ -S18A (A549  $\alpha_1$ -S18A) that were transfected with scrambled (control, white bars) or HOIL-1L (HOIL-1L, black bars) siRNA and exposed to normoxia or 1.5% O<sub>2</sub> for 24 h ( $n = 3$ ). (B–D) WT (white bars in C and D) and *Cre<sup>SPC</sup>/HOIL-1L<sup>fl/fl</sup>* (black bars in C and D) mice were infected with either adenovirus expressing GFP-NaK-ATPase  $\alpha_1$ -WT (Ad- $\alpha_1$ -WT) or GFP-NaK-ATPase  $\alpha_1$ -S18A (Ad- $\alpha_1$ -S18A)

and is considered the key event in stabilizing PKC $\zeta$  in the active conformation at the plasma membrane (37). In COS-7 cells transfected with FLAG-PKC $\zeta$ -WT, hypoxia-induced PKC $\zeta$  phosphorylation increased after 4 h and returned to basal levels after 24 h of exposure (Fig. 5A). To assess whether AMPK-induced phosphorylation is necessary for PKC $\zeta$  down-regulation during hypoxia, we infected rat A10 cells with a dominant-negative AMPK  $\alpha_1$  subunit adenovirus (DN-AMPK) or an empty adenovirus (Null). The lack of AMPK activity was sufficient to prevent the hypoxia-induced PKC $\zeta$  down-regulation (Fig. 5B). To further confirm the role of phosphorylation in PKC $\zeta$  degradation, we used loss- and gain-of-function analysis. Expression of PKC $\zeta$  bearing a T410A mutation, FLAG-PKC $\zeta$ -T410A (T410A), prevented the hypoxia-induced degradation of PKC $\zeta$ , whereas a PKC $\zeta$  phosphorylation mimic, FLAG-PKC $\zeta$ -T410E (T410E), was degraded during hypoxia similarly to PKC $\zeta$ -WT (Fig. 5C). The degradation of FLAG-PKC $\zeta$ -T410E during hypoxia was prevented in HOIL-1L-silenced cells (Fig. 5D).

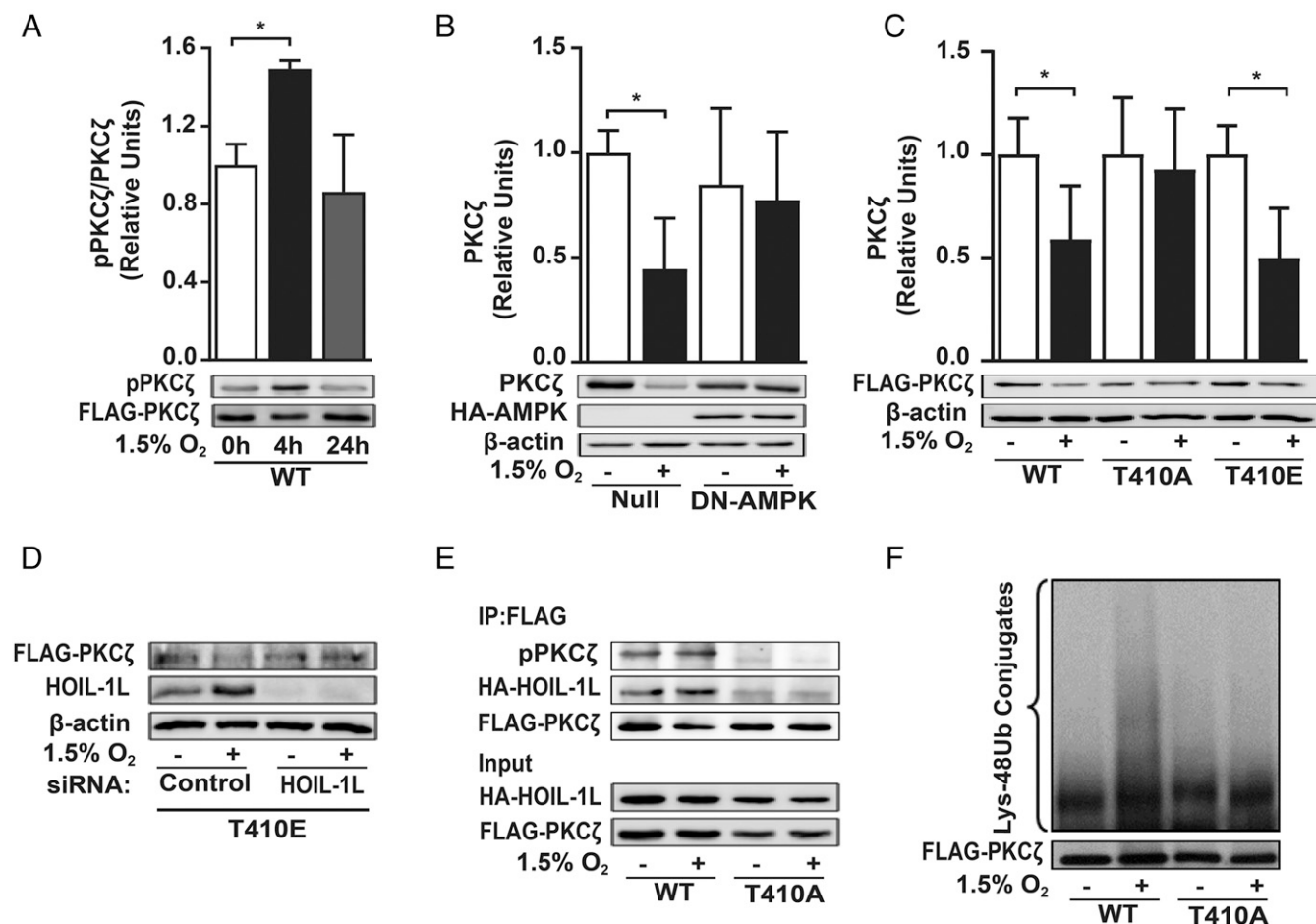
To determine whether the interaction between PKC $\zeta$  and HOIL-1L during hypoxia requires PKC $\zeta$  phosphorylation, cells were cotransfected with HA-HOIL-1L and either FLAG-PKC $\zeta$ -WT or FLAG-PKC $\zeta$ -T410 and exposed to hypoxia, and then proteins in the cell lysate were coimmunoprecipitated by use of an anti-FLAG antibody. Exposure to hypoxia increased PKC $\zeta$ -WT phosphorylation and the interaction between HOIL-1L and PKC $\zeta$ -WT (Fig. 5E, lanes 1 and 2). Overexpression of PKC $\zeta$ -T410A prevented not only the hypoxia-induced PKC $\zeta$  phosphorylation but also the interaction between HOIL-1L and PKC $\zeta$  (Fig. 5E, lanes 3 and 4). Similarly, transfection with FLAG-PKC $\zeta$ -T410A prevented the hypoxia-induced PKC $\zeta$  ubiquitylation (Fig. 5F).

**PKC $\zeta$  Degradation Is Initiated in the Plasma Membrane.** Hypoxia-induced PKC $\zeta$  phosphorylation is required for its translocation to the plasma membrane, as translocation was prevented in cells overexpressing FLAG-PKC $\zeta$ -T410A (Fig. 6A). PKC $\zeta$  remained phosphorylated for at least 4 h (Fig. 5A), and its removal from the plasma membrane had a significant effect after 6 h of hypoxia exposure (Fig. 6B). These data suggest that the interaction between HOIL-1L and the phosphorylated PKC $\zeta$  occurs at the plasma membrane, which was confirmed by transfecting COS-7 cells with FLAG-PKC $\zeta$ -WT and HA-HOIL-1L, exposing the cells to hypoxia for 4 h, isolating the membranes, and coimmunoprecipitating the solubilized proteins with an anti-FLAG antibody (Fig. 6C). Moreover, the hypoxia-induced PKC $\zeta$  ubiquitylation occurs at the plasma membrane (Fig. 6D).

## Discussion

Adaptation to hypoxia requires coordination between energy demand and supply to promote cell homeostasis. The Na,K-ATPase is a major energy consumer, accounting for 20–70% of metabolic ATP depending of the tissue (38, 39). To conserve energy during acute hypoxia, the Na,K-ATPase  $\alpha_1$  subunit is phosphorylated by PKC $\zeta$ , which triggers its endocytosis, independently of HIF (1, 36). Here we describe that AEC exposure to prolonged hypoxia results in decreased Na,K-ATPase levels; however, the cells survive with the Na,K-ATPase level stabilized at a lower plateau (~50% of

and kept in room air or exposed to 7% O<sub>2</sub> for 7 d. (B) Adenovirus infection efficiency was determined by measuring GFP expression in lung peripheral tissue by Western blot. CD29 was used as a loading control. (C) Alveolar epithelial permeability is expressed as the ratio of the fluorescence present in BALF to the plasma ( $n = 3$ ). (D) Na,K-ATPase activity measured in isolated plasma membranes from lung peripheral tissue by a colorimetric assay as described in *Experimental Procedures*. Graph bars represent mean  $\pm$  SD. Statistical significance was calculated for E and G using one-way ANOVA and the Tukey multiple comparisons test (\* $P < 0.05$ , \*\*\* $P < 0.001$ ).



**Fig. 5.** PKC $\zeta$  phosphorylation is required for its ubiquitylation and degradation. Cells were exposed to normoxia or 1.5% O<sub>2</sub> for up to 24 h. (A) COS-7 cells were transfected with FLAG-PKC $\zeta$ -WT, and 4 h after treatment cells were lysed and proteins were immunoprecipitated with anti-FLAG antibody. Densitometry quantification of immunoblots of phospho-PKC $\zeta$  (pPKC $\zeta$ ) in relation to FLAG is shown ( $n = 3$ ). (B) Rat ATII cells were infected with null adenovirus (Null) or HA-tagged adenovirus carrying a dominant-negative, kinase-dead (K45R) variant of AMPK (DN-AMPK) after 24 h of normoxia or 1.5% O<sub>2</sub>. PKC $\zeta$  expression was evaluated by Western blot;  $\beta$ -actin was used as a loading control ( $n = 3$ ). Statistical significance was calculated for A and B using one-way ANOVA and the Tukey multiple comparisons test ( $*P < 0.05$ ). (C) PKC $\zeta$  degradation was assessed after 24 h of treatment in COS-7 cells that were transfected with FLAG-WT-PKC $\zeta$  (WT), FLAG-T410A-PKC $\zeta$  (T410A), or FLAG-T410E-PKC $\zeta$  (T410E) by densitometry quantification of immunoblots of FLAG in relation to  $\beta$ -actin ( $n = 3$ ). Statistical significance was calculated using unpaired Student's  $t$  test ( $*P < 0.05$ ). (D) COS-7 cells were transfected at the same time with FLAG-T410E-PKC $\zeta$  (T410E) and scrambled (control) or HOIL-1L siRNA before 24 h of normoxia or 1.5% O<sub>2</sub> exposure. The pPKC $\zeta$  and HOIL-1L expression levels were detected by Western blot analysis;  $\beta$ -actin was used as a loading control. Representative immunoblots from three independent experiments are shown. (E) COS-7 cells were cotransfected with PKC $\zeta$ -WT (WT) or PKC $\zeta$ -T410A (T410A) and HA-HOIL-1L, exposed to normoxia or 1.5% O<sub>2</sub> for 4 h, and lysed; cell lysate was used for coimmunoprecipitation using anti-FLAG antibody. Phospho-PKC $\zeta$  and HOIL-1L were immunoblotted to assess interaction between proteins. FLAG was used as a loading control. Representative immunoblots from three independent experiments are shown. (F) PKC $\zeta$  ubiquitination was evaluated in COS-7 cells transfected with PKC $\zeta$ -WT (WT) or PKC $\zeta$ -T410A (T410A). After 6 h, proteins in the cell lysate were immunoprecipitated using an anti-FLAG antibody and immunoblotted for Lys-48 polyubiquitination. Representative immunoblots from three independent experiments are shown.

normoxia levels) (Fig. 1 A and B) to conserve energy. This is achieved via HIF-induced up-regulation of the E3 ubiquitin ligase HOIL-1L and phosphorylation of PKC $\zeta$  by AMPK. HOIL-1L then interacts with and ubiquitylates the phosphorylated PKC $\zeta$  at the plasma membrane, triggering its degradation.

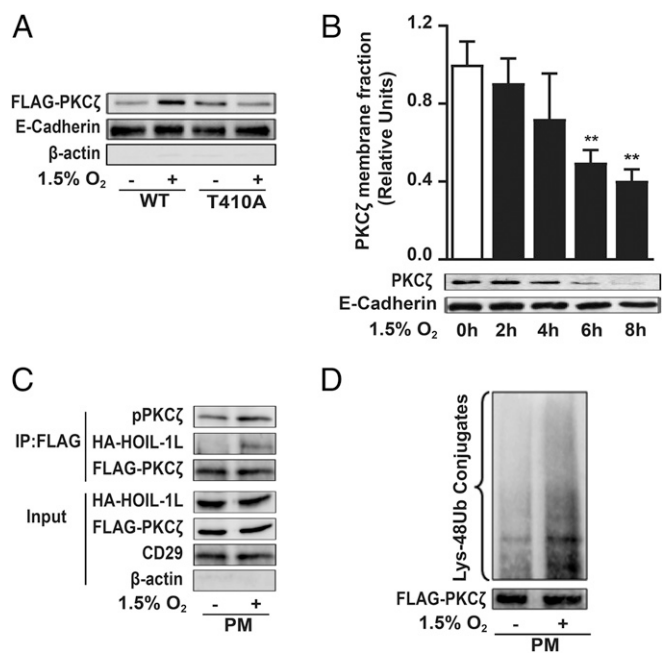
HOIL-1L is a member of LUBAC, which catalyzes the non-degradative linear ubiquitination of proteins. However, not only is the activity of HOIL-1L on PKC $\zeta$  independent of the function of HOIP (the LUBAC catalytic component), but also HOIP competes with PKC $\zeta$  for binding to HOIL-1L (20). We found that silencing of HOIP does not affect Na,K-ATPase levels at the plasma membrane during hypoxia (SI Appendix, Fig. S1E).

In RCC4 cells, which express constitutive levels of HIF1 $\alpha$ , high levels of HOIL-1L, and low levels of PKC $\zeta$ , hypoxia fails to induce Na,K-ATPase endocytosis, which is rescued by overexpression of

PKC $\zeta$  (Fig. 1D). In agreement with these data, silencing of HOIL-1L in hypoxia-exposed cells increased Na,K-ATPase down-regulation and cell death. The increase in cell death in the absence of HOIL-1L is rescued by expressing a phosphorylation-deficient Na,K-ATPase  $\alpha_1$  subunit (Fig. 4A), indicating the specificity of the effect. As we have previously reported, the phosphorylation by PKC $\zeta$  of the plasma membrane Na,K-ATPase- $\alpha_1$  subunits triggers the internalization of Na,K-ATPase molecules during hypoxia. It has been described that myristoylated alanine-rich C kinase substrate proteins (MARCKS) in the nonphosphorylated state sequester PIP<sub>2</sub>, preventing the formation of signaling complexes and thereby promoting the stability of plasma membrane proteins (40, 41). However, MARCKS is not a substrate for the atypical PKCs.

Concordant with the *in vitro* results, ATII cells isolated from WT mice exposed to 7% O<sub>2</sub> for 7 d had increased HOIL-1L and





**Fig. 6.** PKC $\zeta$  degradation is initiated at the cell plasma membrane. (A) COS-7 cells were transfected with FLAG-WT-PKC $\zeta$  (WT) or FLAG-T410A-PKC $\zeta$  (T410A) and exposed to 1.5% O<sub>2</sub> for 10 min, and pPKC $\zeta$  levels in the isolated plasma membrane fraction were determined by Western blot to assess PKC $\zeta$  translocation to the plasma membrane as described in *SI Appendix, SI Experimental Procedures*. E-cadherin was used as loading control for membrane fraction, and  $\beta$ -actin was used as a marker of cytosolic proteins. Representative immunoblots from four independent experiments are shown. (B) Isolated plasma membrane proteins from rat A1II cells exposed to normoxia or 1.5% O<sub>2</sub> were immunoblotted using an anti-PKC $\zeta$  antibody. E-cadherin was used as loading control ( $n = 3$ ). Statistical significance was calculated using one-way ANOVA and the Dunnett's multiple comparisons test (\*\* $P < 0.01$ ). (C) COS-7 cells were cotransfected with FLAG-WT-PKC $\zeta$  (WT) and HA-HOIL-1L; after 4 h of normoxia or 1.5% O<sub>2</sub> exposure, plasma membrane proteins were isolated and used for coimmunoprecipitation using anti-FLAG antibody and immunoblotted for pPKC $\zeta$  and HOIL-1L. FLAG was used as a loading control.  $\beta$ -Actin and CD29 are shown as loading controls for cytosolic and membrane fraction, respectively. Representative immunoblots from at least three independent experiments are shown. (D) PKC $\zeta$  ubiquitination was evaluated in COS-7 cells transfected with FLAG-PKC $\zeta$ -WT. After 4 h of hypoxia, an immunoprecipitation with anti-FLAG antibody was performed in isolated plasma membrane and immunoblotted for Lys-48 polyubiquitylation. Representative immunoblots from at least three independent experiments are shown.

decreased PKC $\zeta$  levels (Fig. 2A). In contrast, PKC $\zeta$  levels did not decrease in A1II cells from *Cre<sup>SPC</sup>/HOIL-1L<sup>fl/fl</sup>* mice exposed to hypoxia as they lacked HOIL-1L. We found significant  $\alpha_1$ -Na,K-ATPase down-regulation (Fig. 2E), more injury (Fig. 3), and increased apoptosis (Fig. 3) in lungs from *Cre<sup>SPC</sup>/HOIL-1L<sup>fl/fl</sup>* mice than in lungs from WT mice after low O<sub>2</sub> treatment. We have previously reported that hypoxia moderately impairs the alveolar fluid clearance (42). However, this impairment did not result in evident alveolar edema, allowing the mice to adapt and survive hypoxic conditions. Interestingly, alveolar type I cells express lower levels of LUBAC compared with A1II cells (*SI Appendix, Fig. S5*). While A1II cells express only the Na,K-ATPase  $\alpha_1$  isoform, A1I cells express both  $\alpha_1$  and  $\alpha_2$  Na,K-ATPase isoforms. The  $\alpha_2$  isoform does not have a PKC consensus phosphorylation domain (Scansite 3 beta). Therefore, the alveolar fluid clearance could be conserved in A1I cells and protect against exaggerated lung injury.

The importance of HOIL-1L in the stabilization of  $\alpha_1$ -Na,K-ATPase during prolonged hypoxia was confirmed by the rescue

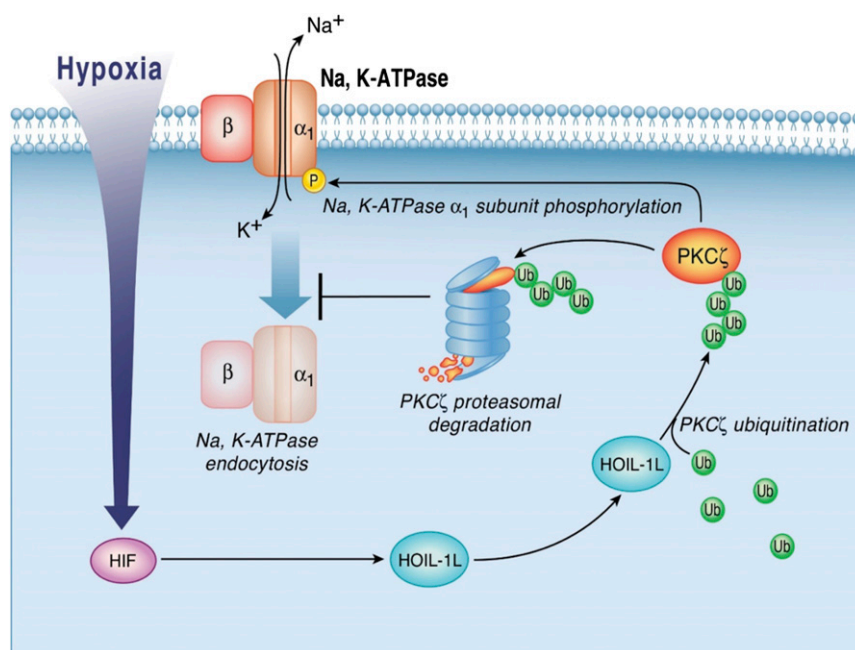
of the epithelial function in *Cre<sup>SPC</sup>/HOIL-1L<sup>fl/fl</sup>* mice overexpressing a phosphorylation-resistant Na,K-ATPase- $\alpha_1$  subunit (Fig. 4). Low O<sub>2</sub> exposure can be associated with increased immune cell recruitment and inflammatory response that has been linked with endothelial injury (43, 44). Interestingly, the increased lung injury observed in *Cre<sup>SPC</sup>/HOIL-1L<sup>fl/fl</sup>* mice during prolonged hypoxia does not appear to increase the hypoxia-induced inflammatory response in the lung but rather results from Na,K-ATPase down-regulation.

The activation of atypical PKCs is analogous to the activation of Akt, in which the loop phosphorylation is agonist-dependent and required for its translocation to the plasma membrane where the autoinhibition is removed, enabling the phosphorylation of substrates (45). In the current study, we provide evidence that the AMPK-induced phosphorylation at Thr-410 and activation are necessary for the ubiquitination and degradation of PKC $\zeta$ . Translocation of PKC $\zeta$  to the plasma membrane occurs within 10 min of hypoxia exposure, and it remains membrane-bound for at least 4 h (Fig. 5A). Similarly to PKC $\zeta$ -WT, the degradation of PKC $\zeta$ -T410E increased during hypoxia (Fig. 5C). These data suggest that both the hypoxia-induced phosphorylation of PKC $\zeta$  and the HIF-dependent HOIL-1L up-regulation are required for PKC $\zeta$  degradation to occur. Activation-dependent ubiquitylation and proteasomal degradation have been described for PKC isoforms (22, 46), suggesting that the ubiquitin proteasome system is critical for PKC regulation. Our data suggest that HOIL-1L interacts with and ubiquitylates the phosphorylated PKC $\zeta$  at the plasma membrane to trigger PKC $\zeta$  degradation (Fig. 6).

Accordingly, we provide evidence for an adaptive mechanism of cells to hypoxia, linking a HIF-regulated pathway to Na,K-ATPase stabilization at the plasma membrane (Fig. 7). During prolonged hypoxia, HIF up-regulates HOIL-1L, which targets PKC $\zeta$  for degradation, thereby preventing it from further downregulating the Na,K-ATPase to conserve ATP and promote AEC survival. This mechanism represents a noncanonical pathway in which HOIL-1L acts as an E3 ligase for PKC $\zeta$  to protect the alveolar epithelium against lung injury during severe hypoxia.

## Experimental Procedures

**Generation of *Cre<sup>SPC</sup>/HOIL-1L<sup>fl/fl</sup>* Mice.** We generated a targeting vector containing LoxP sites flanking exon 8 in the HOIL-1 genomic allele, followed by a neomycin resistance gene (*SI Appendix, Fig. S5*). T12 ES cells derived from C57BL/6 were transfected with the targeting vector, and neomycin-resistant colonies were screened for homologous recombination. Selected ES cells containing the appropriately targeted allele were injected into ICR mice blastocysts to generate chimeric mice, which then transmitted the targeted loxP-flanked HOIL-1L allele (HOIL-1L<sup>fl</sup>) to their progeny and were backcrossed with C57BL/6 mice (47). The genotypes of offspring mice were determined by PCR analysis of DNA extracted from tail-biopsy specimens. The following primer set was used to distinguish the WT-HOIL-1L allele (1,100 bp) and floxed-HOIL-1L allele (273 bp): forward primer CAGCAGGAGGGAACTACTG and reverse primer TGACAGCACTACTCATGCC. *HOIL-1L<sup>fl/fl</sup>* mice were then crossed to mice expressing a Cre recombinase driven by the SP-C promoter (SP-C-Cre<sup>+</sup>), which were a gift from B. Hogan (Duke University, Durham, NC) and are described elsewhere (48). Expression of surfactant protein C promoter begins early in development in primary lung buds and continues throughout development (49). Deletion of HOIL-1L in *Cre<sup>SPC</sup>/HOIL-1L<sup>fl/fl</sup>* mice was verified by genotyping (Transnetyx) using specific probes designed for each gene, with the following set of primers: for the probe CRE, forward primer TTAATCCATATTGGCA-GAAGGAAAACG, reverse primer CAGGCTAAGTGCCCTCTCTACA, and reporter CCTGCGGTGCTAAC; for the probe Rbck1-1 EX, forward primer CAACCCCA-TACTAGCAAGCTTAT, reverse primer CTCTCTCTGCAAACCACACT, and reporter CTCGAAGCTGACGTATAAC; for the probe Rbck1-1 FL, forward primer CAACCCCATACTAGCAAGCTTAT, reverse primer TGGAGGATGGTTAGAGGTC-AAG, and reporter CTAGATCTGCAGCCTACTC; for the probe Rbck1-1 WT, forward primer TGCTCCACCTTGACCACAAC, reverse primer GTGCTGGAGGAGTG-GTTAGAG, and reporter ACCTAGCAAGCTTCTGC. C57BL/6 mice were used as WT mice.



**Fig. 7.** Cartoon depicting the proposed mechanism for Na,K-ATPase stabilization during chronic hypoxia. Hypoxia leads to stabilization of HIF, which regulates HOIL-1L, leading to PKC $\zeta$  ubiquitylation to avoid an excessive depletion of Na,K-ATPase, and safeguarding the alveolar epithelial function as a mechanism of adaptation to hypoxic conditions.

**Alveolar Epithelial Cell Isolation and Culture.** Mice and rats were provided with food and water ad libitum, maintained on a 14-h/10-h light–dark cycle, and handled according to National Institutes of Health guidelines and the experimental protocol approved by the Northwestern University Institutional Animal Care and Use Committee.

Rat ATII cells from the lungs of male Sprague–Dawley rats (weight: 200–225 g) (29) or mouse ATII cells from WT and *Cre<sup>SPC</sup>/HOIL-1L<sup>fl/fl</sup>* mice (age: 10–12 wk) (50) were isolated by the Pulmonary Division Cell Culture and Physiology Core from Northwestern University. The day of isolation and plating of rat ATII cells was designated day 0. All experiments were conducted on day 3.

The following cell lines were used: human adenocarcinoma A549 (ATCC CCL 185), African green monkey kidney COS-7 (ATCC CRL 1651), and pVHL-deficient human RCC4 (51). A549 cells stably expressing the rat Na,K-ATPase- $\alpha_1$  subunit WT or with a Ser-18 mutation have been described elsewhere (1). A549-GFP- $\alpha_1$  cells show similar enzymatic activity and respond to stimuli similarly to WT A549 cells (36). All cells were contamination-free and grown and maintained as described elsewhere (1). After treatment, cells were washed in ice-cold PBS and, when required, solubilized in 2 $\times$  lysis buffer (Cell Signaling Technology). The lysates were cleared by centrifugation for 10 min at 14,000  $\times$  g.

Additional materials and methods can be found in *SI Appendix, SI Experimental Procedures*.

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