

# Knockout Mice Reveal a Major Role for Alveolar Epithelial Type I Cells in Alveolar Fluid Clearance

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## Abstract

Active ion transport by basolateral Na-K-ATPase (Na pump) creates an Na<sup>+</sup> gradient that drives fluid absorption across lung alveolar epithelium. The  $\alpha 1$  and  $\beta 1$  subunits are the most highly expressed Na pump subunits in alveolar epithelial cells (AEC). The specific contribution of the  $\beta 1$  subunit and the relative contributions of alveolar epithelial type II (AT2) versus type I (AT1) cells to alveolar fluid clearance (AFC) were investigated using two cell type-specific mouse knockout lines in which the  $\beta 1$  subunit was knocked out in either AT1 cells or both AT1 and AT2 cells. AFC was markedly decreased in both knockout lines, revealing, we believe for the first time, that AT1 cells play a major role in AFC and providing insights into AEC-specific roles in alveolar homeostasis. AEC monolayers derived from knockout mice demonstrated decreased short-circuit current and active Na<sup>+</sup> absorption, consistent with *in vivo* observations. Neither hyperoxia nor ventilator-induced lung injury increased wet-to-dry lung weight ratios in knockout lungs relative to control lungs. Knockout mice showed increases in Na pump  $\beta 3$  subunit expression and  $\beta 2$ -adrenergic receptor expression. These results demonstrate a

crucial role for the Na pump  $\beta 1$  subunit in alveolar ion and fluid transport and indicate that both AT1 and AT2 cells make major contributions to these processes and to AFC. Furthermore, they support the feasibility of a general approach to altering alveolar epithelial function in a cell-specific manner that allows direct insights into AT1 versus AT2 cell-specific roles in the lung.

**Keywords:** lung; ion transport; Na-K-ATPase;  $\beta 1$  subunit;  $\beta 2$ -adrenergic receptor

## Clinical Relevance

This study demonstrates an important role for the Na-K-ATPase  $\beta 1$  subunit in ion transport and alveolar fluid clearance and that AT1 cells contribute a major portion of alveolar fluid clearance in the lung. Compensatory mechanisms involving increased  $\beta 3$  subunit levels as well as  $\beta 2$ -adrenergic receptor protein expression can be activated in response to  $\beta 1$  subunit deletion.

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Lung alveolar epithelium, composed of alveolar epithelial type I (AT1) and type II (AT2) cells, forms a tight barrier that limits leakage of solutes and water from the interstitial and vascular compartments into alveolar air spaces. Regulation of alveolar fluid homeostasis is based on active ion transport across the alveolar epithelium (1). Polarized localization and function of the basolateral Na-K-ATPase (Na pump) and apical sodium channels (2) are of crucial importance in transepithelial ion transport and the accompanying fluid clearance across the alveolar epithelial barrier (3, 4).

AT2 cells, despite covering only ~2.5% of the alveolar surface (5), have been assumed to contribute the major portion of transport activity in alveolar epithelium, on the basis of high channel and pump density in this cell type (6). However, AT1 cells express both epithelial sodium channel (ENaC) and Na-K-ATPase subunit proteins (7–9) and, in addition to covering a large surface area, have very high water permeability (10). It is therefore likely that AT1 cells make an important contribution to alveolar fluid clearance (AFC) (11, 12), although the relative contributions of AT1 and AT2 cells to ion transport and AFC in the lung are entirely unknown.

The Na pump catalyzes active transport of cytoplasmic  $\text{Na}^+$  in exchange for extracellular  $\text{K}^+$  at the basolateral cell surface (13, 14). Na-K-ATPase is a heterotrimer composed of one  $\alpha$ , one  $\beta$ , and one  $\gamma$  subunit. The  $\alpha$  subunit harbors the catalytic function of the Na pump, whereas the  $\beta$  subunit is important in the maturation of the structure and the function of the holoenzyme and for its transport and localization to the cell membrane (15). The function of the  $\gamma$  subunit is less clear, but experiments have demonstrated a modulatory role of Na pump activity and ion affinity that is tissue specific, on the basis of the differential expression of various  $\gamma$  subunits (FXD proteins) in different tissues (16). The  $\beta 1$  subunit is highly expressed in both AT1 and AT2 cells (17), and Na pumps composed of  $\alpha 1$  and  $\beta 1$  subunits are thought to be the predominant isozyme expressed in both cell types (18, 19), although expression of the  $\alpha 2$  subunit in lung and AT1 cells has been reported (12). A number of *in vivo*

studies based on Na pump  $\beta 1$  subunit overexpression in rodent lungs provide evidence to support an important role of this subunit in Na pump function and AFC, both at baseline and during lung injury (19–24). Expression of the  $\beta 3$  subunit has been demonstrated in rat lung (25), but the role of this subunit in the lung is largely unknown. Clinically, in patients with acute lung injury/acute respiratory distress syndrome, the capacity for higher levels of AFC is associated with better outcomes (26, 27).

The primary goals of this study were to elucidate the roles of Na pump  $\beta 1$  subunit in active transepithelial ion transport and AFC and to determine the relative contributions of AT1 versus AT2 cells to these processes. We generated two mouse lines with conditional knockout of the  $\beta 1$  subunit in either AT1 cells or in both AT1 and AT2 cells in mouse lung. This approach allowed us to directly assess the contribution of the  $\beta 1$  subunit *in vivo* and, we believe for the first time, to investigate specifically the roles of the two different types of alveolar epithelial cells (AEC) in alveolar function. Some of the results of these studies have been reported previously in abstract form (28–30).

## Materials and Methods

### Generation of Knockout Mice

Mice with a floxed allele of the Na pump  $\beta 1$  subunit gene (*Atp1b1*<sup>F/F</sup>) were generated (see online supplement). Briefly, *Atp1b1*<sup>F/F</sup> mice were crossed to *Aqp5-cre* mice (31) to generate *Atp1b1*<sup>Aqp5-cre</sup> mice deficient in the  $\beta 1$  subunit in AT1 cells, and to *Sftpc-cre* mice (32) to generate *Atp1b1*<sup>Sftpc-cre</sup> mice deficient in the  $\beta 1$  subunit in the entire alveolar epithelium (i.e., both AT1 and AT2 cells). All animal protocols were approved by the Institutional Animal Care and Use Committee at the University of Southern California.

### Alveolar Fluid Clearance

Phosphate-buffered saline with 5% bovine serum albumin (BSA) (Sigma-Aldrich, St. Louis, MO) and 0.25 milligrams per milliliter BSA-Alexa Fluor 594 conjugate (Life Technologies, Carlsbad, CA) as tracer were instilled intratracheally into anesthetized mice. Alveolar fluid was aspirated after 30 minutes. AFC was

calculated from fluorescence measured in instillate and aspirate.

### In Vivo Lung Permeability

Permeability *in vivo* was calculated after jugular vein injection of 10 milligrams of fluorescein-BSA (Life Technologies) per kilogram of body weight from fluorescence measured 2 hours later in bronchoalveolar lavage fluid and serum.

### Wet-to-Dry Lung Weight Ratios

Lungs were removed surgically, weighed, and then dried at 65°C for 48 hours. Dry weight was recorded and wet-to-dry lung weight ratios were calculated.

### Isolation of AT2 Cells and Primary Culture of Mouse Alveolar Epithelial Cell Monolayers

AT2 cells were isolated from mouse lungs, and mouse alveolar epithelial cell monolayers (MAECM) were cultured as described previously (33).

### Bioelectric Properties of MAECM

Transepithelial electrical resistance ( $R_T$ ) ( $\text{k}\Omega \cdot \text{cm}^2$ ) and spontaneous potential difference ( $PD$ ) (mV) of MAECM were measured using a Millicell-ERS device (Millipore, Bedford, MA) on Day 6 after AT2 cells had transdifferentiated to an AT1 cell-like phenotype. Equivalent short-circuit current ( $I_{EQ}$ ) ( $\mu\text{A}/\text{cm}^2$ ) was calculated as  $PD/R_T$ .  $PD$  and short-circuit current ( $I_{SC}$ ) ( $\mu\text{A}/\text{cm}^2$ ) were measured in modified Ussing chambers in the presence or absence of terbutaline, amiloride, and pimozide.

### Unidirectional Flux of $\text{Na}^+$

Unidirectional flux of  $\text{Na}^+$  was measured across MAECM at 37°C using <sup>22</sup>NaCl (American Radiolabeled Chemicals, St. Louis, MO).

### Western Analysis

Details regarding methods and antibodies used are provided in the online supplement.

### Lung Histology

Standard methods as described in online supplement were used.

### Antibody Staining

Staining methods and antibody information are described in the online supplement.

**Table 1.** Quantitative Reverse Transcriptase–Polymerase Chain Reaction Primers

Gene	GenBank Accession No.	Primer Sequence (5' → 3')	Product Size (bp)
<i>Atp1a1</i> (Na <sup>+</sup> ,K <sup>+</sup> -ATPase α1)	NM_144900	TCAAGTCTTGGAGCTCGGAACT (FP) ACGTCTGCATCCCCACATG (RP)	66
<i>Atp1a2</i> (Na <sup>+</sup> ,K <sup>+</sup> -ATPase α2)	NM_178405	TGAGCTGGGCCGAAAATACC (FP) GGTCCATCTCTAGCCAGAAT (RP)	85
<i>Atp1a3</i> (Na <sup>+</sup> ,K <sup>+</sup> -ATPase α3)	BC034645	TCTGCCCTGCTTAAGTGCATT (FP) TCCGTTCTCGCATCAGCTT (RP)	61
<i>Atp1a4</i> (Na <sup>+</sup> ,K <sup>+</sup> -ATPase α4)	NM_013734	TCAGGAGTCTGTTCCCATAGCTAA (FP) GGAGAGCTGACTCGGAAGCA (RP)	62
<i>Atp1b1</i> (Na <sup>+</sup> ,K <sup>+</sup> -ATPase β1)	NM_009721	TTCATCGGGACCATCCAAGT (FP) TCCTGGTATGTGGGCTTCAGT (RP)	62
<i>Atp1b2</i> (Na <sup>+</sup> ,K <sup>+</sup> -ATPase β2)	NM_013415	TGCCCACACAATTTCCAACAT (FP) ACTCCCAGACTCCTCTCTGTCTCT (RP)	64
<i>Atp1b3</i> (Na <sup>+</sup> ,K <sup>+</sup> -ATPase β3)	NM_007502	GCCGAGTGGAAAGCTGTTTCAT (FP) GGTGCGCCCCAGAACT (RP)	56
<i>Atp1b4</i> (Na <sup>+</sup> ,K <sup>+</sup> -ATPase β4)	NM_133690	TGAAAATGACATTCGATCCATCA (FP) TAAGGGTAGTAGCGGAGGTCAA (RP)	69
<i>Polr2a</i>	NM_009089	GGCAAGGTCCCACAACCA (FP) ACAATTGATGTGTCCAGGTATGATG (RP)	93

Definition of abbreviations: bp, base pair; FP, forward primer; RP, reverse primer.

### RNA Isolation, Reverse Transcription and Quantitative Polymerase Chain Reaction

Details regarding methods are provided in the online supplement. Primer sequences are listed in Table 1.

### Hyperoxia Exposure

Mice were housed in cages inside a hyperoxia chamber with 95% oxygen for 65 hours.

### Ventilator-Induced Lung Injury

Anesthetized mice were ventilated with an Inspira ASV ventilator (Harvard Apparatus, Holliston, MA) under either noninjurious or injurious ventilation conditions (see online supplement).

### Data Analysis

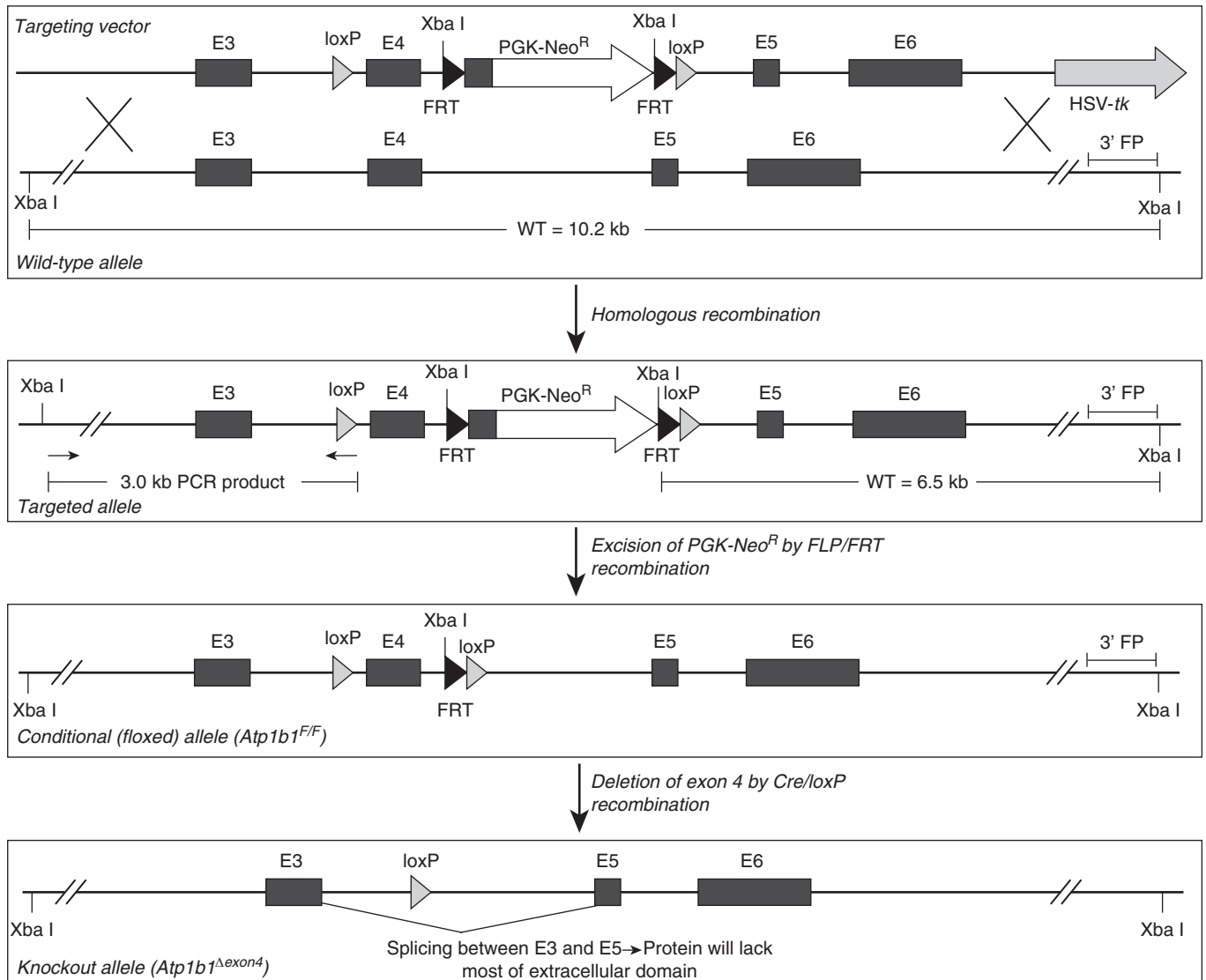
Data are shown as mean ± SEM. Unpaired Student's *t* test was used for comparisons of two group means. Multiple (three or more) group means were analyzed by one-way analysis of variance with post hoc tests based on Student-Newman-Keuls approaches. *P* < 0.05 is considered statistically significant.

## Results

### *Atp1b1* Gene Targeting and Generation/Verification of β1 Knockout Mice

We generated a mouse line with a conditional (floxed) allele of the *Atp1b1* gene (encoding the Na-K-ATPase β1 subunit) to enable cell-specific gene knockout by Cre/loxP recombination (Figure 1). Mouse *Atp1b1* contains six exons; exon 1 codes for the cytoplasmic domain of the Na-K-ATPase β1 subunit protein, and the first part of exon 2 codes for the transmembrane domain. The remainder of the gene codes for the large extracellular domain that makes up 243 of the 305 amino acids in this protein. We flanked exon 4 of *Atp1b1* with loxP sites to create a conditional allele of this gene (*Atp1b1*<sup>F/F</sup>) (Figure 1). Cre/loxP-mediated deletion of exon 4 results in a frame-shift mutation after splicing of exon 3 to exon 5. To elucidate the importance of the β1 subunit in adult mouse lung fluid homeostasis and its relative importance in AT1 versus AT2 cells, we generated two separate lines, one harboring a conditional β1

knockout specifically in AT1 cells using *Aqp5-cre* mice (31) and one in which β1 was inactivated in both AT1 and AT2 cells (alveolar epithelium-specific knockout) using *Sftpc-cre* mice (32). These two knockout lines are referred to as *Atp1b1*<sup>Aqp5-cre</sup> and *Atp1b1*<sup>Sftpc-cre</sup>, respectively. Knockout lines were analyzed to verify deletion of the Na pump β1 subunit gene (see online supplement). Genomic PCR (see Figure E1A in the online supplement) confirmed that the floxed *Atp1b1* allele could be deleted correctly by Cre/loxP recombination. Western analysis (Figure E1B) confirmed that Na pump β1 protein was absent in AT2 cells isolated from *Atp1b1*<sup>Sftpc-cre</sup> knockout mice. We confirmed (Figure E1C) that *Aqp5-cre* activates green fluorescent protein (GFP) expression from a *ROSA*<sup>mT/mG</sup> reporter transgene specifically in AT1 cells in the alveolar epithelium and that efficiency of this Cre/loxP-mediated reporter activation is very high (>90% of AT1 cells) (31). Finally, we confirmed that *Sftpc-cre* activates GFP expression in the *ROSA*<sup>mT/mG</sup> reporter transgene in both AT1 and AT2 cells, although the GFP signal



**Figure 1.** Gene targeting strategy. An *Atp1b1* gene targeting vector was made by floxing exon 4, introducing a PGK-Neo<sup>R</sup> positive selection marker upstream of the 3' loxP site, and placing an HSV-*tk* negative selection marker in a 3'-flanking position in the genomic fragment. The *Atp1b1* gene targeting vector was introduced into W2 embryonic stem (ES) cells, and clones that had undergone homologous recombination were identified by Southern blot using a flanking probe (3' FP) binding a 6.5-kb Xba I fragment from the targeted allele and a 10.2-kb fragment from the wild-type allele. Positive clones were then verified by PCR, with primers amplifying a 3.0-kb portion on the 5' side of the targeted allele. After karyotyping of positive ES cell clones, chimeric mice were produced. Germline-transmitting chimeras were bred to FLPeR mice to remove the FRT-flanked PGK-Neo<sup>R</sup> selection marker, generating mice harboring the final floxed *Atp1b1* allele (*Atp1b1*<sup>F/F</sup>). Cre/loxP-mediated deletion of exon 4 generates a deleted allele (*Atp1b1*<sup>Δexon4</sup>) with an out-of-frame mutation that disrupts the coding sequence in exons 5 and 6 (E5 and E6). FRT, flippase recognition target; PGK, phosphoglycerate kinase promoter; WT, wild type.

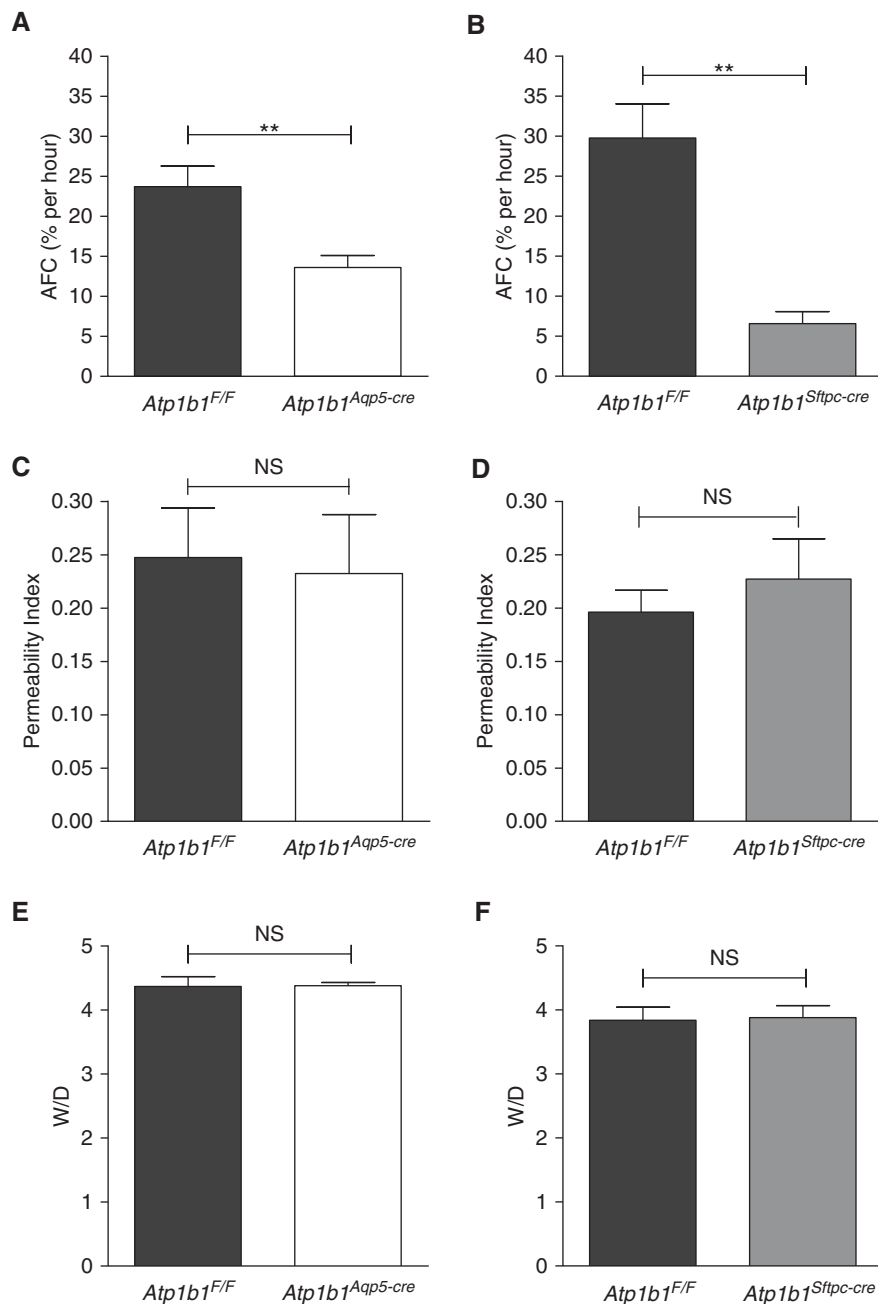
is considerably stronger in AT2 cells, in part reflecting the very thin AT1 cell architecture *in vivo* (Figure E1C).

#### AFC Is Markedly Reduced in Mice Deficient in the Na Pump $\beta$ 1 Subunit

We measured AFC in the two  $\beta$ 1 subunit knockout lines (*Atp1b1*<sup>Aqp5-cre</sup> and *Atp1b1*<sup>Sftpc-cre</sup>) and compared them to respective floxed litter mate control mice (*Atp1b1*<sup>F/F</sup>). AFC in *Atp1b1*<sup>Aqp5-cre</sup>

knockout mice, deficient in the  $\beta$ 1 subunit specifically in AT1 cells, demonstrated a highly significant ( $P = 0.006$ ) reduction (43%) in AFC compared with *Atp1b1*<sup>F/F</sup> mice (Figure 2A). *Atp1b1*<sup>Sftpc-cre</sup> knockout mice, which lack the  $\beta$ 1 subunit in both AT1 and AT2 cells, revealed even lower AFC, which was significantly ( $P = 0.002$ ) reduced by 78% compared with control mice (Figure 2B). These data demonstrate that the Na pump  $\beta$ 1

subunit is of crucial importance in AFC after fluid instillation and that both AT1 and AT2 cells make major contributions to AFC in the adult mouse lung. The portion of AFC attributable to AT1 and AT2 cells can be estimated as  $\sim 55\%$  (i.e.,  $43/78 = 0.55$ ) and  $\sim 45\%$  (i.e.,  $[78-43]/78 = 0.45$ ), respectively, assuming that AT1/AT2 cell number ratios are similar in control and knockout mice on the basis of histological examination of lungs from floxed control



**Figure 2.** Alveolar fluid clearance (AFC), lung permeability, and wet-to-dry lung weight ratio (W/D) in  $\beta 1$  knockout mice. (A) AFC (% per h) in  $Atp1b1^{Aqp5-cre}$  mice ( $13.6 \pm 1.5$ ) ( $n = 7$ ) was significantly reduced compared with  $Atp1b1^{F/F}$  mice ( $23.7 \pm 2.6$ ) ( $n = 7$ ). (B) AFC (% per h) in  $Atp1b1^{Sftpc-cre}$  mice ( $6.6 \pm 1.5$ ) ( $n = 4$ ) was significantly reduced compared with  $Atp1b1^{F/F}$  mice ( $29.8 \pm 4.2$ ) ( $n = 4$ ). (C and D) There was no significant difference in *in vivo* permeability to fluorescein-bovine serum albumin between  $Atp1b1^{F/F}$  ( $0.248 \pm 0.047$ ) ( $n = 6$ ) and  $Atp1b1^{Aqp5-cre}$  mice ( $0.232 \pm 0.055$ ) ( $n = 7$ ) (C) or between  $Atp1b1^{F/F}$  ( $0.196 \pm 0.020$ ) ( $n = 6$ ) and  $Atp1b1^{Sftpc-cre}$  mice ( $0.227 \pm 0.038$ ) ( $n = 6$ ) (D). (E and F) At baseline, W/D of  $Atp1b1^{F/F}$  ( $4.37 \pm 0.15$ ) ( $n = 8$ ) versus  $Atp1b1^{Aqp5-cre}$  ( $4.38 \pm 0.05$ ) ( $n = 12$ ) mice (E) and of  $Atp1b1^{F/F}$  ( $3.84 \pm 0.08$ ) ( $n = 6$ ) versus  $Atp1b1^{Sftpc-cre}$  ( $3.88 \pm 0.08$ ) ( $n = 6$ ) mice (F) were not significantly different.  $**P < 0.01$ . NS, not significantly different from each other ( $P > 0.05$ ).

mice and both knockout lines that showed no obvious morphologic differences (Figure E2). AFC in  $Atp1b1^{Aqp5-cre}$  and  $Atp1b1^{Sftpc-cre}$  knockout mice (57% and

22%, respectively, relative to floxed control mice) are likely based on active sodium transport because of residual Na pump activity.

### Unchanged Lung Permeability and Absence of Lung Edema in $\beta 1$ Knockout Mice

To investigate if  $\beta 1$  knockout lungs have altered paracellular permeability that might contribute to reductions in AFC, we injected fluorescein-BSA into the jugular vein of floxed and knockout mice. As shown in Figures 2C and 2D, lung permeability to fluorescein-BSA was not different in  $\beta 1$  knockout mice compared with floxed control mice. Consistent with these findings, lungs of both  $Atp1b1^{Aqp5-cre}$  and  $Atp1b1^{Sftpc-cre}$  mice had wet-to-dry lung weight ratios that were not different from their respective floxed control mice (Figures 2E and 2F), suggesting that, in both  $\beta 1$  subunit knockout lines, residual Na pump activity is sufficient to maintain lung fluid homeostasis.

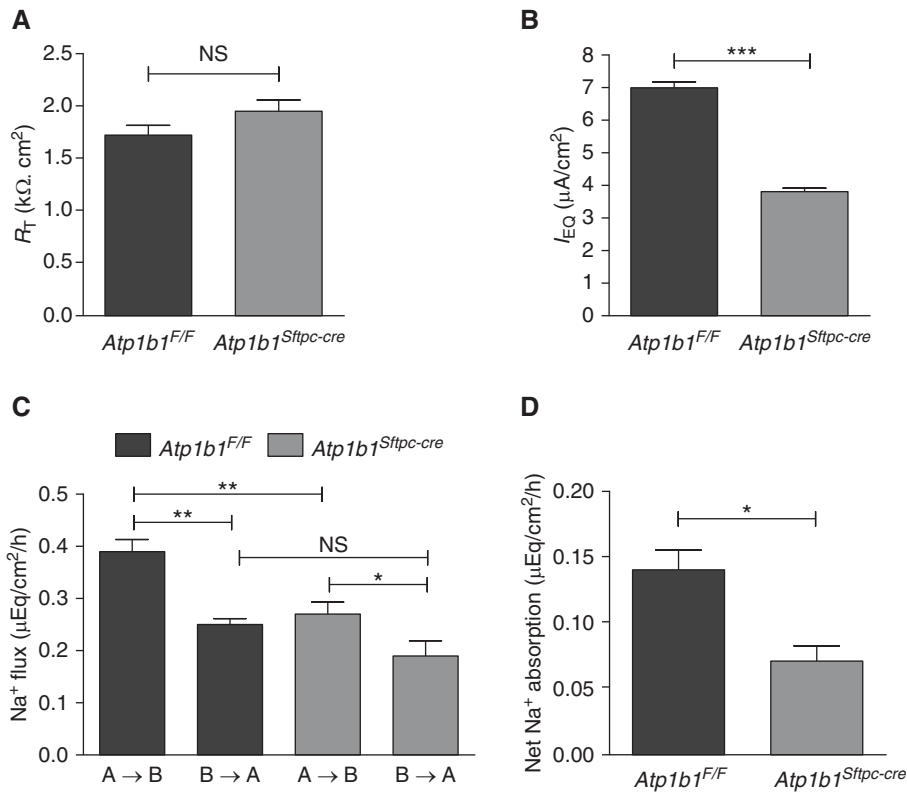
### Decreased $I_{EQ}$ and Unchanged $R_T$ Across MAECM Deficient in $\beta 1$ Subunit

Bioelectric properties ( $R_T$  and  $I_{EQ}$ ) were evaluated in mouse alveolar epithelial cell monolayers (MAECM) prepared by cultivation of freshly isolated AT2 cells derived from  $Atp1b1^{F/F}$  control and  $Atp1b1^{Sftpc-cre}$  knockout mice. As shown in Figure 3A,  $R_T$  of  $Atp1b1^{F/F}$  control and  $Atp1b1^{Sftpc-cre}$  knockout monolayers on Day 6 were not significantly different. In contrast,  $I_{EQ}$  of  $Atp1b1^{Sftpc-cre}$  knockout monolayers were significantly reduced compared with  $Atp1b1^{F/F}$  control monolayers, consistent with results *in vivo* (Figure 3B) and supporting the observation that the Na pump  $\beta 1$  subunit plays a crucial role in active ion transport across MAECM.

### Decreased $Na^+$ Absorption in MAECM Deficient in $\beta 1$ Subunit

To more directly evaluate active ion transport across  $Atp1b1^{F/F}$  control and  $Atp1b1^{Sftpc-cre}$  knockout MAECM, unidirectional  $Na^+$  flux and net  $Na^+$  absorption were determined under zero electrical gradient (i.e., short-circuit) conditions in Ussing chambers. Figure 3C shows the unidirectional  $Na^+$  flux in each direction (i.e., apical to basolateral [A $\rightarrow$ B] and basolateral to apical [B $\rightarrow$ A]). In both control and knockout monolayers, unidirectional  $Na^+$  flux in the A $\rightarrow$ B direction was significantly greater than that in the B $\rightarrow$ A direction. Knockout monolayers had significantly lower unidirectional  $Na^+$  flux in the A $\rightarrow$ B direction ( $P < 0.01$ ) and unchanged





**Figure 3.** Bioelectric properties and  $\text{Na}^+$  absorption of  $\beta 1$  knockout mouse alveolar epithelial cell monolayers (MAECM). (A) Transepithelial electrical resistance ( $R_T$ ) across  $Atp1b1^{F/F}$  MAECM ( $1.72 \pm 0.09 \text{ k}\Omega \cdot \text{cm}^2$ ) ( $n = 42$ ) was not significantly different compared with  $Atp1b1^{Sftpc-cre}$  MAECM ( $1.95 \pm 0.11 \text{ k}\Omega \cdot \text{cm}^2$ ) ( $n = 45$ ). (B) Equivalent short-circuit current ( $I_{EQ}$ ) was significantly lower in  $Atp1b1^{Sftpc-cre}$  MAECM ( $3.81 \pm 0.11 \mu\text{A}/\text{cm}^2$ ) ( $n = 45$ ) compared with  $Atp1b1^{F/F}$  MAECM ( $7.00 \pm 0.18 \mu\text{A}/\text{cm}^2$ ) ( $n = 42$ ). (C) Unidirectional  $\text{Na}^+$  fluxes in the apical-to-basolateral (A $\rightarrow$ B) direction were significantly lower in  $Atp1b1^{Sftpc-cre}$  MAECM, whereas fluxes in the basolateral-to-apical (B $\rightarrow$ A) direction were unchanged ( $n = 3$ ). (D) Net  $\text{Na}^+$  absorption of  $Atp1b1^{F/F}$  and  $Atp1b1^{Sftpc-cre}$  MAECM ( $n = 3$ ) were significantly different, with lower absorption in knockout monolayers. \* $P < 0.05$ ; \*\* $P < 0.01$ ; \*\*\* $P < 0.001$ .

unidirectional  $\text{Na}^+$  flux in the B $\rightarrow$ A direction, compared with control monolayers. Baseline net  $\text{Na}^+$  absorption of knockout monolayers was significantly lower than that of control monolayers (Figure 3D). Net  $\text{Na}^+$  absorption in control and knockout monolayers was not significantly different from the observed active ion transport rate (i.e.,  $I_{SC}$ ) during the  $\text{Na}^+$  flux measurements (data not shown), indicating that estimated net  $\text{Na}^+$  absorption accounts for most of the observed active ion transport across MAECM.

#### Sodium Channel Function Unaffected by $\beta 1$ Knockout

AEC harbor apical amiloride-sensitive ENaC and pimozide-sensitive cyclic nucleotide-gated (CNG) nonselective

cation channels (4). To determine if knockout of the Na pump  $\beta 1$  subunit had any effects on these ion channels, MAECM were treated with ENaC or CNG channel inhibitors, and bioelectric properties were evaluated. The addition of amiloride to apical fluid caused a rapid  $\sim 70\%$  decrease in  $I_{SC}$  of both control and knockout MAECM (Figures 4A and 4B). Subsequent treatment with apical pimozide ( $10 \mu\text{M}$ ) at 60 minutes led to a slight additional decrease in  $I_{SC}$  of both monolayers. When the CNG inhibitor pimozide was added first,  $I_{SC}$  of both control and knockout monolayers gradually decreased by  $\sim 45\%$  (Figures 4C and 4D). When amiloride ( $10 \mu\text{M}$ ) was subsequently added to apical fluid,  $I_{SC}$  of both monolayers rapidly decreased further by  $\sim 35\%$ . These data suggest that the

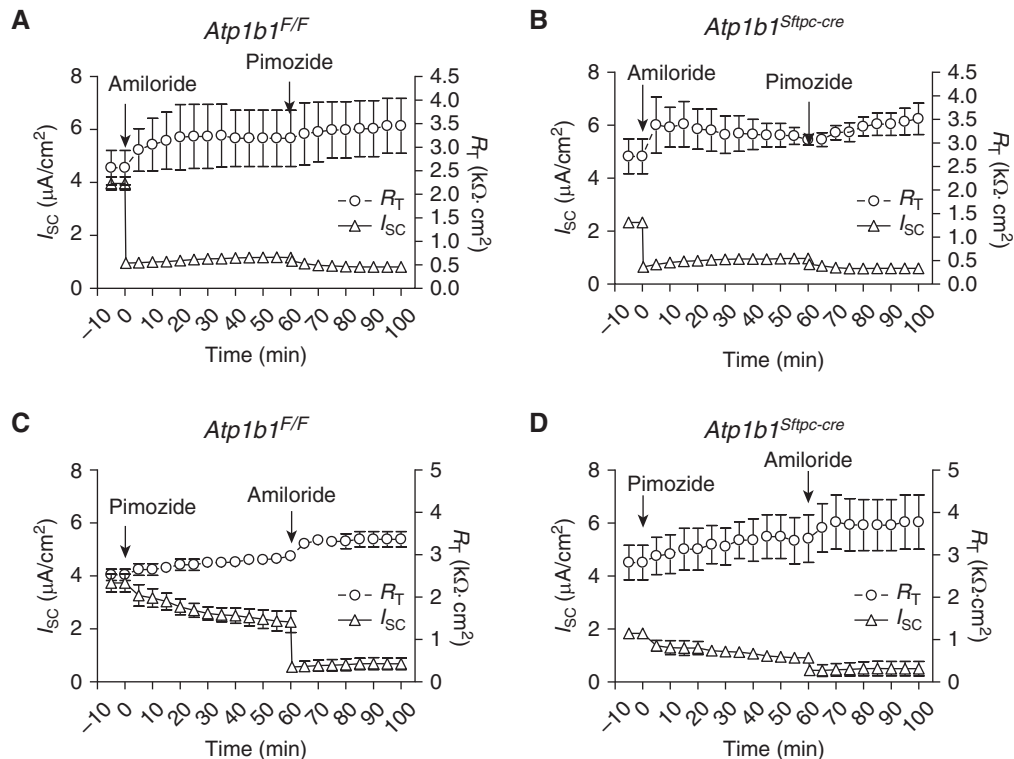
function of apical sodium channels remains unchanged in  $\beta 1$  knockout MAECM.

#### Wet-to-Dry Lung Weight Ratios after Hyperoxia or Ventilator-induced Lung Injury

Although residual AFC in both  $\beta 1$  knockout lines was sufficient to maintain a normal lung fluid balance in the unchallenged lung, we assessed whether or not decreased AFC in mice deficient in the  $\beta 1$  subunit affected their susceptibility to lung injury from hyperoxia or ventilator-induced lung injury (VILI). All animals developed increased wet-to-dry lung weight ratios after exposure to  $>95\%$  oxygen for 65 hours (Figure 5A), but neither  $Atp1b1^{Aqp5-cre}$  nor  $Atp1b1^{Sftpc-cre}$  knockout mice exhibited ratios different from  $Atp1b1^{F/F}$  control mice subjected to the same experimental conditions. Mice of both  $Atp1b1^{F/F}$  and  $Atp1b1^{Sftpc-cre}$  genotypes mechanically ventilated for 3 hours with a peak inspiratory pressure of 40 cm  $\text{H}_2\text{O}$  and without positive-end expiratory pressure had significantly higher wet-to-dry lung weight ratios compared with mice ventilated under noninjurious conditions at 20 cm  $\text{H}_2\text{O}$ , also without positive-end expiratory pressure (Figure 5B), but there was no significant difference in wet-to-dry weight ratios between control and knockout mice after VILI.

#### Increased Expression of Na Pump $\beta 3$ Subunit Protein in $Atp1b1^{Sftpc-cre}$ Knockout Mice

Both  $Atp1b1^{Aqp5-cre}$  and  $Atp1b1^{Sftpc-cre}$  knockout mice had normal wet-to-dry lung weight ratios at baseline (Figures 2E and 2F), suggesting that residual Na-K-ATPase activity was sufficient to maintain fluid homeostasis. The Na pump  $\beta 3$  subunit is the second highest expressed  $\beta$  subunit in AEC. Western analysis revealed increased  $\beta 3$  protein expression in freshly isolated AT2 cells derived from  $Atp1b1^{Sftpc-cre}$  mice (Figures 6A and 6B), and antibody staining of MAECM from knockout mice showed increased expression of  $\beta 3$  subunit protein compared with control MAECM (Figures 6C and 6D), although  $\beta 3$  subunit expression at the mRNA level in whole lung, freshly isolated AT2 cells, or MAECM did not change (Figures 6E–6H). Relative mRNA expression levels of other  $\alpha$  and  $\beta$  subunit



**Figure 4.** Short-circuit current ( $I_{SC}$ ) response to amiloride and pimoziide. (A and B) At  $t=0$ , amiloride ( $10 \mu\text{M}$ ) was added to apical fluid of (A)  $Atp1b1^{F/F}$  MAECM ( $n=3$ ) and (B)  $Atp1b1^{Sftpc-cre}$  MAECM ( $n=3$ ). After 60 minutes, pimoziide ( $10 \mu\text{M}$ ) was added to apical fluid. (C and D) At  $t=0$ , pimoziide ( $10 \mu\text{M}$ ) was added to apical fluid of (C)  $Atp1b1^{F/F}$  MAECM ( $n=3$ ) and (D)  $Atp1b1^{Sftpc-cre}$  MAECM ( $n=3$ ). After 60 minutes, amiloride ( $10 \mu\text{M}$ ) was added to apical fluid.

genes in whole lung of  $Atp1b1^{Aqp5-cre}$  knockout mice were either unchanged or only moderately different from  $Atp1b1^{E/F}$  control lungs, whereas in  $Atp1b1^{Sftpc-cre}$  knockout lungs, a more substantial reduction in  $\beta 1$  subunit RNA levels was observed (Figure 6F), likely because the  $\beta 1$  gene is deleted in the entire epithelium. Similarly,  $Atp1b1^{Sftpc-cre}$  knockout AT2 cells and MAECM demonstrated significantly lower expression of  $\beta 1$  subunit (Figures 6G and 6H). Expression of Na pump  $\alpha 1$  subunit protein in AT2 cells obtained from  $Atp1b1^{Sftpc-cre}$  knockout mice was unchanged, whereas  $\beta 1$  subunit protein was absent (Figure 6A). Increased expression of  $\beta 3$  subunit protein may contribute to residual AFC in  $Atp1b1^{Sftpc-cre}$  knockout mice.

#### Increased $\beta 2$ -Adrenergic Responsiveness *In Vivo* in $\beta 1$ Subunit Knockout Mice

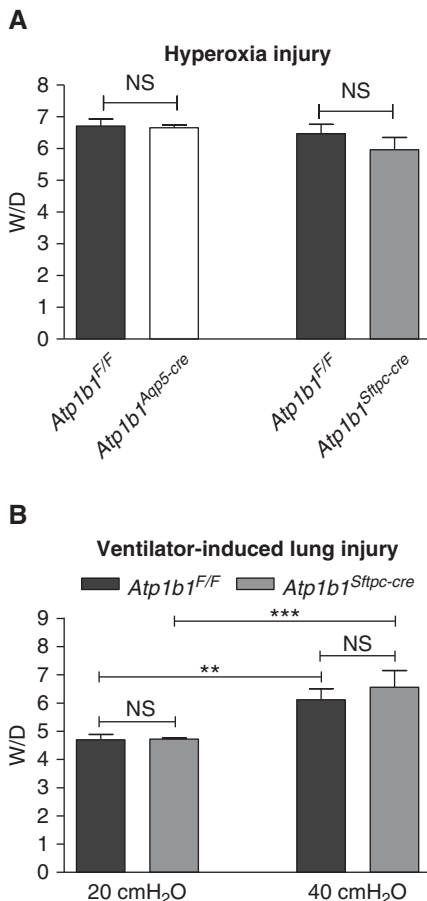
When terbutaline was added to the instillate in AFC experiments, AFC increased markedly in both  $Atp1b1^{Aqp5-cre}$  and  $Atp1b1^{Sftpc-cre}$  knockout mice,

reaching levels close to those measured in  $Atp1b1^{E/F}$  control mice (+Terbutaline in Figures 7A and 7B). This was in contrast to AFC in the absence of terbutaline, which showed significantly lower AFC in both knockout mice (−Terbutaline in Figures 7A and 7B). To investigate the possibility that these observations could be caused by higher expression of the  $\beta 2$ -adrenergic receptor ( $\beta 2\text{AR}$ ) in knockout mice, we performed Western analysis of the whole lung. As shown in Figures 7C and 7D,  $\beta 2\text{AR}$  protein levels were significantly higher in the lungs of mice from both knockout lines compared with those of control mice, which may contribute to increased responsiveness to terbutaline stimulation and residual AFC in  $Atp1b1^{Aqp5-cre}$  and  $Atp1b1^{Sftpc-cre}$  knockout mice.

#### Discussion

We generated and analyzed two mouse lines harboring knockout of the

Na-K-ATPase  $\beta 1$  subunit, either in AT1 cells only ( $Atp1b1^{Aqp5-cre}$  mice) or in both AT1 and AT2 cells ( $Atp1b1^{Sftpc-cre}$  mice) in the lung. AFC was reduced by 78% in  $Atp1b1^{Sftpc-cre}$  mice, providing evidence for an important role of the  $\beta 1$  subunit under conditions of excess alveolar fluid volume. AFC was also reduced substantially in  $Atp1b1^{Aqp5-cre}$  mice (by 43%), indicating that AT1 cells are major contributors to active ion transport and associated fluid clearance (~55% of overall AFC) in mouse lung. There were no changes in lung permeability in mice deficient in the Na pump  $\beta 1$  subunit. Despite lower active ion transport and decreased AFC, both knockout lines had normal wet-to-dry lung weight ratios at baseline, and no differences in wet-to-dry lung weight ratios were observed among genotypes after hyperoxia or VILI. Analysis of the bioelectric properties of cultured AEC monolayers demonstrated that AT1-like cells deficient in the Na pump  $\beta 1$  subunit featured significantly decreased active  $\text{Na}^+$  absorption and  $I_{EQ}$  (and  $I_{SC}$ ), consistent with the



**Figure 5.** W/D in mice exposed to hyperoxia or subjected to ventilator-induced lung injury (VILI). (A) W/D were not significantly different between floxed and knockout mice after hyperoxic injury (65 h in >95% oxygen). W/D in *Atp1b1<sup>F/F</sup>* and *Atp1b1<sup>Aqp5-cre</sup>* mice were  $6.70 \pm 0.22$  versus  $6.66 \pm 0.09$  ( $n = 3$ ) and in *Atp1b1<sup>F/F</sup>* and *Atp1b1<sup>Sftpc-cre</sup>* mice were  $6.47 \pm 0.30$  versus  $5.96 \pm 0.40$  ( $n = 3$ ). (B) No significant difference in W/D between genotypes was observed after either noninjurious (PIP = 20 cm H<sub>2</sub>O) or injurious (PIP = 40 cm H<sub>2</sub>O) ventilation. W/D in *Atp1b1<sup>F/F</sup>* and *Atp1b1<sup>Sftpc-cre</sup>* were  $4.70 \pm 0.11$  versus  $4.72 \pm 0.02$  ( $n = 3$ ) after noninjurious ventilation and  $6.12 \pm 0.17$  versus  $6.57 \pm 0.26$  ( $n = 5$ ) after VILI. PIP, peak inspiratory pressure. \*\* $P < 0.01$ ; \*\*\* $P < 0.001$ .

observation that knockout mice were unable to clear fluid at the same rate as floxed control mice and demonstrating that the  $\beta 1$  subunit is crucial to AFC. Finally, *in vitro* analysis revealed that  $R_T$  was unaffected in MAECM deficient in the  $\beta 1$  subunit, consistent with unchanged permeability in knockout lungs.

### Cell-Specific Deletion of *Atp1b1* and Effects on Na Pump Subunit Expression

In Figure 6E, which shows subunit expression levels in whole lung in AT1 cell-specific knockout mice, the decrease in  $\beta 1$  mRNA did not reach statistical significance. Given the fact that AT2 cells are present in larger numbers than are AT1 cells in the lung ( $\sim 2:1$ ) (5), a relatively limited effect on whole lung  $\beta 1$  expression levels might be expected in AT1 cell-specific knockout lungs. It is also possible that the  $\beta 1$  expression level per cell differs between wild type AT1 and AT2 cells. If the level is higher in AT2 cells, this would also contribute to the apparent limited decrease in  $\beta 1$  mRNA in whole lungs of AT1 cell-specific knockout mice. A compensatory increase in  $\beta 1$  expression in AT2 cells in AT1 cell-specific knockout mice would have the same effect of masking the decreased mRNA level in AT1 cells when analyzing expression in whole lung. The decreased mRNA levels of the  $\alpha 1$ ,  $\alpha 2$ , and  $\beta 2$  subunits in freshly isolated AT2 cells shown in Figure 6G were relatively small but nevertheless significant. At the protein level, we were able to detect  $\alpha 1$ , which did not change significantly (Figure 6A), whereas  $\alpha 2$  and  $\beta 2$  proteins were undetectable. From other studies involving sodium pump subunits, it is known that changes in mRNA level are not always reflected at the protein level (e.g.,  $\beta 1$  subunit is increased at the mRNA level in rat lung exposed to hyperoxia, whereas no corresponding increase in  $\beta 1$  protein was found [34]).

### Relative Contributions of AT1 and AT2 Cells to Alveolar Ion and Water Transport

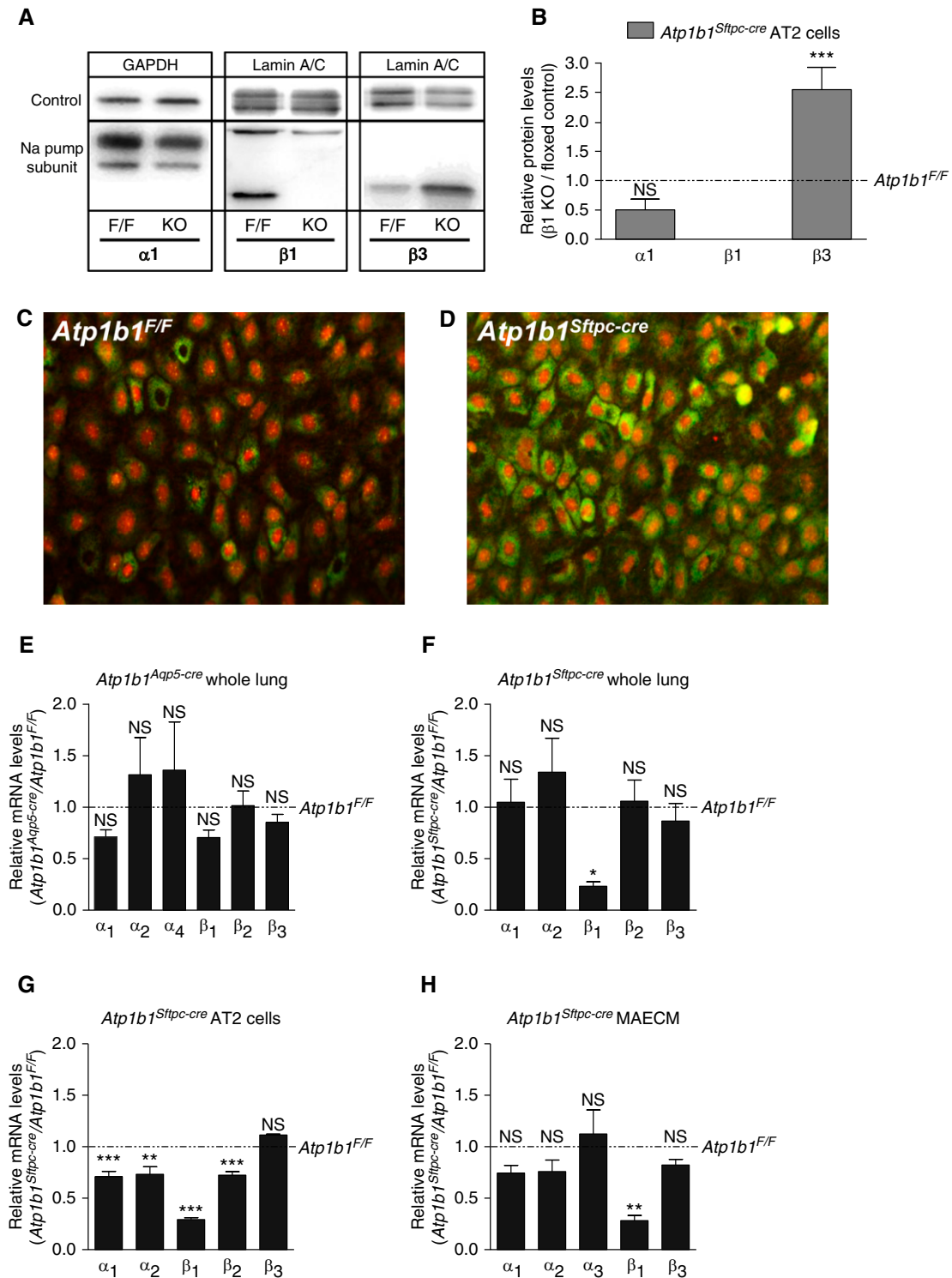
AT1 cell-specific  $\beta 1$  knockout mice showed  $\sim 43\%$  reduction in AFC, whereas animals lacking  $\beta 1$  in both AT1 and AT2 cells demonstrated  $\sim 78\%$  reduction in AFC. As noted above, these results indicate that a major portion ( $\sim 55\%$ ) of AFC in the lung is normally driven by AT1 cells, providing, we believe, the first direct evidence to support a major contribution of AT1 cells to AFC in the lung. Considering that the number of AT2 cells in the lung is greater than the number of AT1 cells (5), AFC contributed per AT1 cell is greater than that per AT2 cell. This finding contrasts with the

earlier presumption that AT2 cells are responsible for most of the ion (and accompanying fluid) transport in the lung (6). However, because the total surface area of AT1 cells is much larger than that of AT2 cells (73-fold difference in rat lung [5]), the density of Na pumps per AT2 cell is likely much higher than that per AT1 cell. In these calculations of relative contribution to AFC from AT1 and AT2 cells, we assumed that  $\beta 1$  is efficiently and similarly knocked out in both AT2 and AT1 cells in the *Atp1b1<sup>Sftpc-cre</sup>* knockout line, which is likely because the *Sftpc* promoter driving the *cre* transgene starts to be expressed early during lung development (35) and the *Atp1b1* gene will thus be deleted in progenitors of both AT2 and AT1 cells (36). Figure E1C (right panel) shows efficient Cre-mediated reporter activation, resulting in GFP expression in both AT1 and AT2 cells, although expression is considerably weaker in AT1 cells. Because the myristoylated GFP reporter protein encoded by the *ROSA<sup>mT/mG</sup>* transgene is inserted into the plasma membrane, an equal amount of myristoylated GFP expressed per cell in AT2 and AT1 cells would be expected to result in a weaker signal in AT1 cells given their considerably larger cell surface area. When comparing the two knockout lines and calculating the relative contributions of AT1 and AT2 cells, we have assumed equal knockout efficiency in AT1 cells in the two lines on the basis of efficient reporter activation in AT1 cells in both *Aqp5<sup>cre</sup>*; *ROSA<sup>mT/mG</sup>* and *Sftpc<sup>cre</sup>*; *ROSA<sup>mT/mG</sup>* (Figure E1C, left and right panel, respectively).

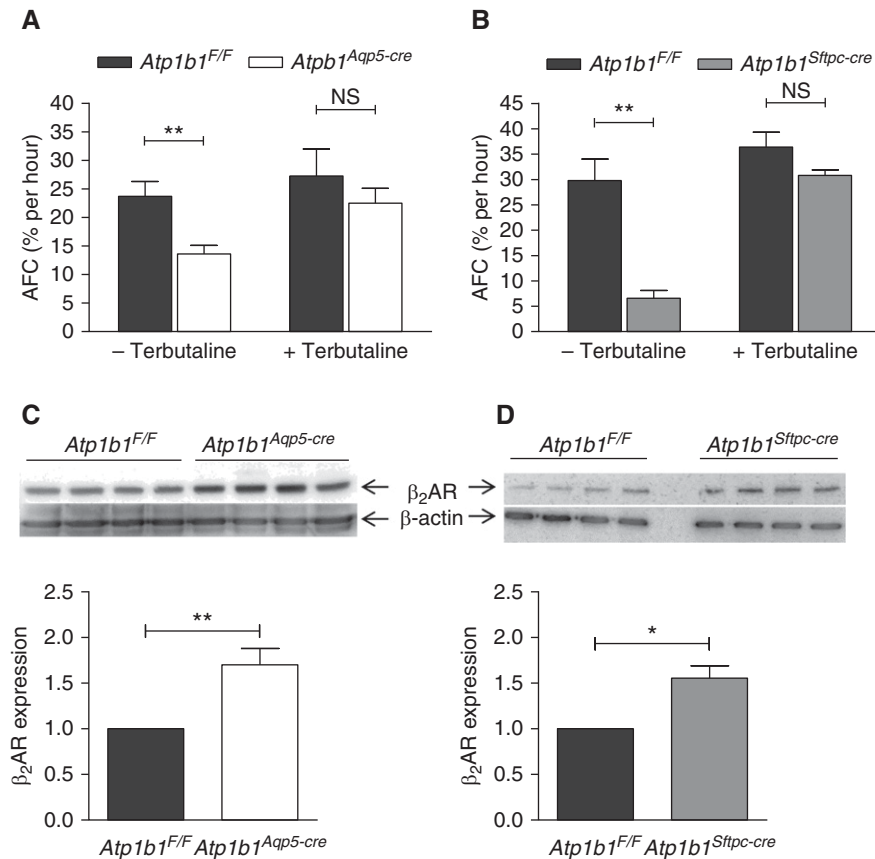
### Changes in Transport Properties of $\beta 1$ Knockout MAECM

Isolated mouse AT2 cells cultured on polycarbonate filters transdifferentiate into AT1-like cells, forming functional monolayers with high  $R_T$  that are suitable for characterization of ion transport and bioelectric properties (33). Given that Na pump  $\beta 1$  subunit is the most highly expressed  $\beta$  subunit in the lung, we hypothesized that knockout of *Atp1b1* would result in decreased active  $\text{Na}^+$  transport. MAECM deficient in  $\beta 1$  subunit exhibited lower net  $\text{Na}^+$  absorption compared with MAECM from floxed mice. These results, demonstrating





**Figure 6.** Expression of Na pump subunits in sodium  $\beta 1$  knockout mice. (A) Representative Western blots reveal that Na pump  $\alpha 1$  subunit was unchanged,  $\beta 1$  was undetectable, and  $\beta 3$  was increased in alveolar epithelial type II (AT2) cells from knockout mice. The upper band in  $\beta 1$  blot appears to be nonspecific. (B) Deletion of Na pump  $\beta 1$  subunit significantly increased  $\beta 3$  protein in AT2 cells isolated from *Atp1b1<sup>Sftpc-cre</sup>* (KO) compared with *Atp1b1<sup>F/F</sup>* (F/F) mice ( $n = 3$ ). (C and D) Antibody staining of MAECM (Day 6) for Na pump  $\beta 3$  subunit in *Atp1b1<sup>F/F</sup>* cells (C) and *Atp1b1<sup>Sftpc-cre</sup>* cells (D) shows increased expression in MAECM generated from  $\beta 1$  knockout mice. (E–H) Relative mRNA expression of  $\alpha$  and  $\beta$  subunit genes in whole lung from (E) *Atp1b1<sup>Aqp5-cre</sup>* and (F) *Atp1b1<sup>Sftpc-cre</sup>* mice ( $n = 4$ ), and (G) isolated AT2 cells ( $n = 3$ ) and (H) MAECM from *Atp1b1<sup>Sftpc-cre</sup>* mice ( $n = 3$ ). Levels are relative to expression in *Atp1b1<sup>F/F</sup>* control lung, isolated AT2 cells, or MAECM. \* $P < 0.05$ , \*\* $P < 0.01$ , \*\*\* $P < 0.001$ .



**Figure 7.** AFC in the absence or presence of terbutaline and Western analysis of  $\beta_2$ -adrenergic receptor ( $\beta_2$ AR) expression in Na pump  $\beta_1$  knockout mice. (A) AFC measurements in the absence of terbutaline demonstrated lower AFC in *Atp1b1<sup>Aqp5-cre</sup>* versus control mice. In contrast, terbutaline-stimulated AFC in *Atp1b1<sup>Aqp5-cre</sup>* mice reached  $22.5 \pm 2.6\%$  per hour ( $n = 4$ ), which was not significantly different from *Atp1b1<sup>F/F</sup>* control mice ( $27.3 \pm 4.7\%$  per hour,  $n = 4$ ). (B) AFC in the absence of terbutaline was significantly lower in *Atp1b1<sup>Sftpc-cre</sup>* versus control mice, whereas terbutaline-stimulated AFC in *Atp1b1<sup>Sftpc-cre</sup>* mice was  $30.9 \pm 1.0\%$  per hour ( $n = 4$ ), which was not significantly different from *Atp1b1<sup>F/F</sup>* control mice ( $36.4 \pm 2.9\%$  per hour,  $n = 4$ ). (C and D) Western analysis of  $\beta_2$ AR protein in whole lung in (C) *Atp1b1<sup>Aqp5-cre</sup>* and (D) *Atp1b1<sup>Sftpc-cre</sup>* mice shows increased expression in both knockout lines compared with *Atp1b1<sup>F/F</sup>* control mice. \* $P < 0.05$ ; \*\* $P < 0.01$ .

the importance of the Na pump  $\beta_1$  subunit for ion transport in AT1-like cells in culture, are consistent with our findings *in vivo* in *Atp1b1<sup>Aqp5-cre</sup>* mice in which deletion of  $\beta_1$  specifically in AT1 cells resulted in a major reduction in AFC. In experiments to evaluate responses in MAECM to an inhibitor of ENaC (amiloride) or CNG (pimozide) channels, we found proportional inhibition of active ion transport in knockout and control mice, suggesting that  $\beta_1$  subunit deficiency does not alter the relative contributions of these channels to transepithelial ion transport. However, because pimozide is known to affect  $D_2$ -dopamine receptors, we cannot completely rule out the possibility that

the observed pimozide effects on  $I_{SC}$  might have been different between the genotypes. Increased expression of  $\beta_3$  subunit and  $\beta_2$ AR in MAECM were not sufficient to restore ion transport. Although  $\beta_1$  subunits in neighboring cells have been reported to interact directly as adhesion molecules (37–39),  $R_T$  was not decreased in  $\beta_1$  knockout MAECM. Although not included in this study, evaluation of the bioelectric properties of MAECM derived from AT2 cells of *Atp1b1<sup>Aqp5-cre</sup>* mice would show the effects of *de novo*  $\beta_1$  gene deletion. Thus, the  $\beta_1$  subunit gene would be deleted gradually during the process of AT2 cell transdifferentiation into AT1 cells, because transdifferentiated cells start to

express *Aqp5* and Cre. In both knockout models, deletion of the  $\beta_1$  subunit gene is dependent on the spatiotemporal expression profile of the promoter driving Cre expression. *Sftpc-cre* is expressed already in early lung progenitors in the developing embryo, whereas *Aqp5-cre* reaches appreciable expression levels perinatally. Because all studies were performed in adult mice, compensatory mechanisms may have developed in both knockout lines.

### Response to Lung Injury in $\beta_1$ Knockout Mice

We did not detect any differences in wet-to-dry lung weight ratios among genotypes after hyperoxia or VILI, although wet-to-dry lung weight ratios are relatively insensitive to small changes in alveolar fluid volume. Although AFC data would have been more useful, technical challenges prevented accurate measurement of AFC in injured lungs. The absence of higher wet-to-dry lung weight ratios after injury in  $\beta_1$  knockout mice versus control mice might be a result, in part, of a compensatory increase in expression of sodium pump  $\beta_3$  subunit and/or elevated expression of  $\beta_2$ -adrenergic receptors. Expression of  $\beta_3$  subunit has been reported in normal rat lungs (25), but there have been no reports on its role in lung injury. Elevated expression of  $\beta_2$ -adrenergic receptors in knockout mice correlated with increased responsiveness to terbutaline, supporting a potential role of adrenergic signaling as a compensatory mechanism in  $\beta_1$  knockout lungs. A higher increase ( $\sim 4$ -fold) in AFC in response to terbutaline in *Atp1b1<sup>Sftpc-cre</sup>* knockout mice is likely a reflection of increased adrenergic signaling in both AT1 and AT2 cells, because both cell types lack  $\beta_1$  in this knockout line. In *Atp1b1<sup>Aqp5-cre</sup>* knockout mice, however,  $\beta_1$  is missing only in AT1 cells, so the adrenergic response will be less pronounced ( $\sim 2$ -fold) since increased adrenergic signaling likely only takes place in AT1 cells. This reasoning is based on the assumption that increased adrenergic signaling is an intrinsic response in cells lacking the  $\beta_1$  protein and that increased adrenergic responsiveness is found only in these cells. We did not investigate expression levels of Na pump  $\gamma$  subunits in  $\beta_1$  knockout mice, although a recent report demonstrated expression of all seven *FXFD* genes (encoding  $\gamma$  subunits) in human lung at the mRNA level (40).

This study also suggested that FXYD1 is a negative regulator of Na pump activity and that increased expression of FXYD1 in lungs from patients with ARDS may indicate a role for  $\gamma$  subunits in deficient ion transport and fluid clearance.

### Conclusions

These findings from lung epithelial cell type-specific  $\beta 1$  subunit knockout mice demonstrate a major contribution of AT1 cells, greater than that of AT2 cells, to alveolar ion transport and AFC. Residual AFC in  $\beta 1$  subunit knockout mice, possibly caused, in part, by increased

expression of  $\beta 3$  subunit and  $\beta 2AR$ , appears sufficient to maintain lung fluid homeostasis at baseline. These studies demonstrate, we believe for the first time, that the roles of lung AEC can be addressed in an AT1 versus AT2 cell-specific manner *in vivo*, leading to improved understanding of the clinical implications pertaining to specific cell types in alveolar epithelium. Elucidation of the relative contributions of AT1 and AT2 cells to alveolar function/homeostasis may help lead to the development of new therapeutic approaches to lung disease. ■

**Author disclosures** are available with the text of this article at [www.atsjournals.org](http://www.atsjournals.org).

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