

LETTERS

Angiotensin-converting enzyme 2 protects from severe acute lung failure

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Acute respiratory distress syndrome (ARDS), the most severe form of acute lung injury, is a devastating clinical syndrome with a high mortality rate (30–60%) (refs 1–3). Predisposing factors for ARDS are diverse^{1,3} and include sepsis, aspiration, pneumonias and infections with the severe acute respiratory syndrome (SARS) coronavirus^{4,5}. At present, there are no effective drugs for improving the clinical outcome of ARDS^{1–3}. Angiotensin-converting enzyme (ACE) and ACE2 are homologues with different key functions in the renin–angiotensin system^{6–8}. ACE cleaves angiotensin I to generate angiotensin II, whereas ACE2 inactivates angiotensin II and is a negative regulator of the system. ACE2 has also recently been identified as a potential SARS virus receptor and is expressed in lungs^{9,10}. Here we report that ACE2 and the angiotensin II type 2 receptor (AT₂) protect mice from severe acute lung injury induced by acid aspiration or sepsis. However, other components of the renin–angiotensin system, including ACE, angiotensin II and the angiotensin II type 1a receptor (AT_{1a}), promote disease pathogenesis, induce lung oedemas and impair lung function. We show that mice deficient for *Ace2* show markedly improved disease, and also that recombinant ACE2 can protect mice from severe acute lung injury. Our data identify a critical function for ACE2 in acute lung injury, pointing to a possible therapy for a syndrome affecting millions of people worldwide every year.

The renin–angiotensin system has an important role in maintaining blood pressure homeostasis, as well as fluid and salt balance^{11–13}. ACE2 is a homologue of ACE, and functions a negative regulator of the renin–angiotensin system^{6–8}. Although ACE2 is expressed in the lungs of humans¹⁰ and mice (see Supplementary Fig. 1a, b), nothing is known about its function in the lungs. However, mortality following SARS coronavirus infections approaches almost 10% owing to the development of ARDS^{14–16}. To elucidate the role of ACE2 in acute lung injury, we examined the effect of *Ace2* gene deficiency in mouse experimental models that mimic the common lung failure pathology observed in several human diseases, including sepsis, acid aspiration and pneumonias such as SARS and avian influenza A¹⁷.

Aspiration of gastric contents with a low pH is a frequent cause of acute lung injury/ARDS^{1–3}. Acid aspiration in wild-type mice, which mimics human acute lung injury^{18,19}, resulted in rapid impairment of lung functions assessed by increased lung elastance (a measure of the change in pressure achieved per unit change in volume, representing the stiffness of the lungs) (Fig. 1a), decreased blood oxygenation

(Fig. 1b) and the development of pulmonary oedema (Fig. 1c). Acid aspiration resulted in increased alveolar wall thickness, oedema, bleeding, inflammatory cell infiltrates and formation of hyaline membranes (Fig. 1d). Notably, acid-treated *Ace2* knockout mice⁸ showed significantly greater lung elastance compared with control wild-type mice, but there were no differences in lung elastance between saline-treated *Ace2* knockout and wild-type mice (Fig. 1a). Moreover, loss of *Ace2* resulted in worsened oxygenation (Fig. 1b), massive lung oedema (Fig. 1c), increased inflammatory cell infiltration and hyaline membrane formations (Fig. 1d) in response to acid aspiration. It should be noted that ACE2 protein expression is typically downregulated in wild-type mice following acid challenge (Fig. 1e).

Sepsis is the most common cause of acute lung injury/ARDS^{1–3}. We therefore examined the effect of *Ace2* gene deficiency on sepsis-induced acute lung injury using caecal ligation and perforation (CLP)²⁰. CLP causes lethal peritonitis and sepsis due to a polymicrobial infection that is accompanied by acute lung failure²⁰. Whereas all CLP-treated wild-type mice survived, only two out of ten CLP-treated *Ace2* knockout mice survived the 6 h experimental observation period (Fig. 2a). CLP resulted in lung failure defined by increased elastance (Fig. 2a), pulmonary oedema (Fig. 2b) and leukocyte accumulation (Fig. 2c) in wild-type mice. CLP-treated *Ace2* knockout mice had a marked worsening of lung functions (Fig. 2a), increased oedema (Fig. 2b) and leukocyte accumulation (Fig. 2c) compared with wild-type mice. In addition, *Ace2* knockout mice also developed markedly enhanced acute lung injury after endotoxin challenge¹⁸ (see Supplementary Fig. 2a–c). *Ace2* maps to the X chromosome, and it should be noted that loss of ACE2 expression resulted in equally severe acute lung injury phenotypes in male (*Ace2*^{−/y}) and female (*Ace2*^{−/−}) mice. Our data from three different acute lung injury models show that loss of *Ace2* expression precipitates severe acute lung failure.

To test whether loss of ACE2 is essential for disease pathogenesis, we performed a rescue experiment using recombinant human ACE2 protein (rhuACE2) (see Supplementary Fig. 3a, b). Injection of rhuACE2 into acid-treated *Ace2* knockout mice decreased the degree of acute lung injury, as assessed by lung elastance (Fig. 2d) and pulmonary oedema formation (Fig. 2e). When we injected rhuACE2 protein into acid-treated wild-type mice, lung function (Fig. 2f) and oedema formation (see Supplementary Fig. 3c) were also rescued. In saline-treated wild-type or *Ace2* knockout mice, injections of

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rhuACE2 did not affect pulmonary functions (Fig. 2d–f). Catalytically inactive ACE2 protein (mut-rhuACE2) (see Supplementary Fig. 3a, b) did not rescue the severe lung phenotype in *Ace2* knockout mice (Fig. 2d, e) and had no effect on the severity of acute lung injury in wild-type animals (Fig. 2f and Supplementary Fig. 3c). These results show that the catalytic activity of ACE2 can directly protect lungs from acute lung injury.

ACE2 is a homologue of ACE, both of which are central enzymes in the renin–angiotensin system^{6–8}. ACE cleaves the decapeptide angiotensin (Ang)I into the octapeptide AngII (refs 11, 12). ACE2 cleaves a single residue from AngI to generate Ang1–9 (refs 6, 7), and a residue from AngII to generate Ang1–7 (ref. 6). In this way, ACE2 negatively regulates the renin–angiotensin system by inactivating AngII (Fig. 3a). Acid aspiration in wild-type mice resulted in marked downregulation of ACE2 protein, but ACE levels remained constant (Fig. 1e). Moreover, only catalytically active ACE2 improved the acute lung injury phenotype in mutant and wild-type mice (Fig. 2d–f). To clarify whether acute lung injury shifts the functional equilibrium between ACE and ACE2, we measured AngII levels in acid-treated and control mice. Acid aspiration markedly increased AngII levels in lungs (Fig. 3b) and plasma (see Supplementary Fig. 4a) of wild-type mice. We observed a further, significant increase in AngII levels in the lungs (Fig. 3b) and plasma (see Supplementary Fig. 4a) of acid-treated *Ace2* knockout mice. Thus, acute lung injury results in decreased ACE2 expression and increased production of AngII.

On the basis of these results we speculated that, in contrast to ACE2, ACE promotes disease pathogenesis through increased AngII production (Fig. 3a). Indeed, genetic inactivation of *Ace* on an *Ace2* wild-type or *Ace2* knockout background markedly decreased AngII levels in lung and plasma in our acid injury model (see Supplementary Fig. 4b, c). Moreover, treatment with rhuACE2 protein attenuated lung injury (Fig. 2d–f) and further reduced AngII levels in the lungs of acid-treated mice (Supplementary Fig. 4d). In contrast to *Ace2* knockout mice, *Ace*^{-/-} mice²¹ were partly protected against acute lung injury induced by acid-aspiration (Fig. 3c and Supplementary Fig. 5). These effects were dependent on gene dosage and were observed to a lesser extent in *Ace*^{+/-} mice. In addition,

inactivation of *Ace* on an *Ace2* knockout background rescued the severe lung failure (Fig. 3d), oedema formation (Fig. 3e) and histological changes (Fig. 3f) compared with *Ace2* knockout mice. Similarly, in endotoxin-induced acute lung injury, the severe lung impairments in *Ace2* knockout mice were reversed by additional *Ace* gene deficiency (see Supplementary Fig. 6). Thus, ACE promotes acute lung injury pathology and ACE2 alleviates it.

Both ACE and ACE2 are non-specific proteases that cleave additional substrates^{11,12}. Thus, although increased levels of AngII have been correlated with *Ace2* deficiency, it has not been shown that upregulation of the AngII pathway accounts for the observed phenotypes of *Ace2* knockout mice *in vivo*. The receptors for AngII in mice are angiotensin II type 1a (AT_{1a}) receptor²², the type 1b (AT_{1b}) receptor and the type 2 (AT₂) receptor²³. AT_{1a} and AT₂, but not AT_{1b} receptor expression is found in the lungs²⁵. We therefore explored which AngII receptor subtypes are responsible for ACE/ACE2 regulated acute lung injury, and whether AngII signalling through its receptors is responsible for ACE2-regulated lung pathology (Fig. 3a). Compared with wild-type mice, genetic loss of AT_{1a} receptor expression in *Agtr1a*^{-/-} mice²⁴ markedly improved lung function (Fig. 4a) and reduced oedema formation (see Supplementary Fig. 7a). In contrast, inactivation of the AT₂ receptor (*Agtr2*^{-/-})²⁵ aggravated acute lung injury (Fig. 4a and Supplementary Fig. 7a). AngII levels induced by acid aspiration in both *Agtr1a*^{-/-} and *Agtr2*^{-/-} mice were comparable to those in wild-type controls (not shown).

We next attempted to rescue acute lung injury in *Ace2* knockout mice using specific AT₁ and AT₂ receptor blockers. Pharmacological inhibition of AT₁ attenuated the severity of acid-induced lung injury in *Ace2* knockout mice (Fig. 4b and Supplementary Fig. 7b). Inhibition of AT₂ had no apparent effect on the acute lung injury phenotypes of *Ace2* knockout mice (Fig. 4b). These data show that the AT_{1a} and AT₂ receptors have opposite functions in controlling the severity of acute lung injury, and that actions of AngII through the AT_{1a} receptor have a causative role in acute lung failure.

Pulmonary oedema could arise from increased hydrostatic pressure (due to pulmonary vascular constriction) and/or enhanced

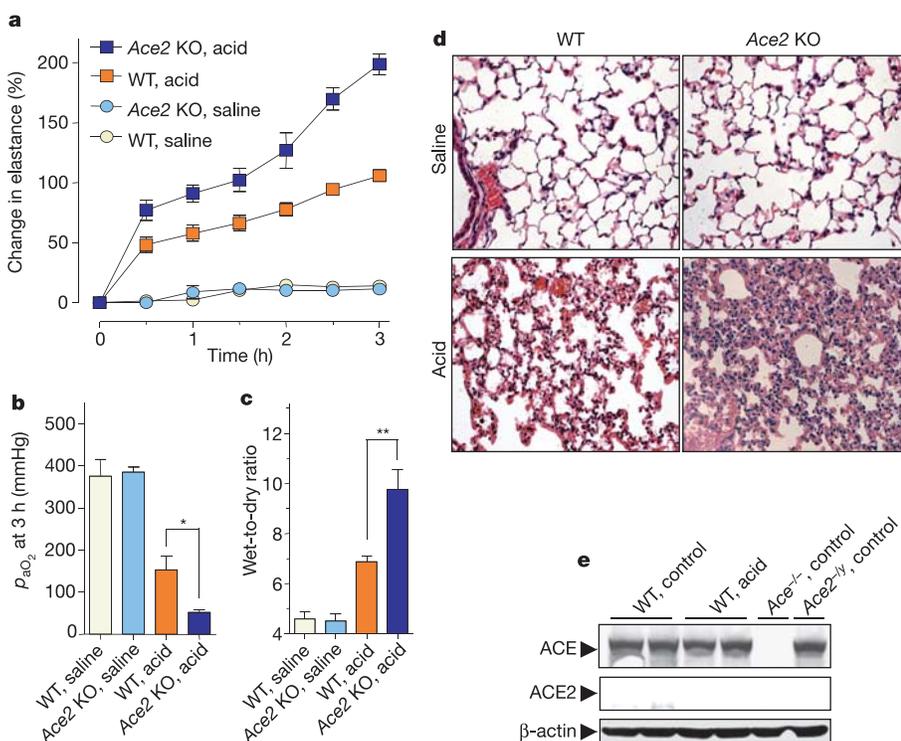


Figure 1 | Loss of ACE2 worsens acid aspiration-induced acute lung injury. **a**, Lung elastance after acid or saline treatment in wild type (WT) and *Ace2* knockout (*Ace2* KO) mice ($n = 10$ for acid-treated groups, $n = 6$ for saline-treated groups). $P < 0.05$ for the whole time course comparing acid-treated WT and *Ace2* knockout mice. **b**, Partial pressure of oxygen in arterial blood (p_{aO_2}) in acid-induced acute lung injury. **c**, Wet-to-dry weight ratios of lungs 3 h after acid injury. Single asterisk, $P < 0.05$; double asterisk, $P < 0.01$. **d**, Lung histopathology. Note the enhanced hyaline membrane formation, inflammatory cell infiltration and lung oedema in acid-treated *Ace2* knockout mice (H&E staining, $\times 200$). **e**, ACE and ACE2 protein expression in total lysates from control lungs and lungs 3 h after acid injury. Error bars indicate s.e.m.

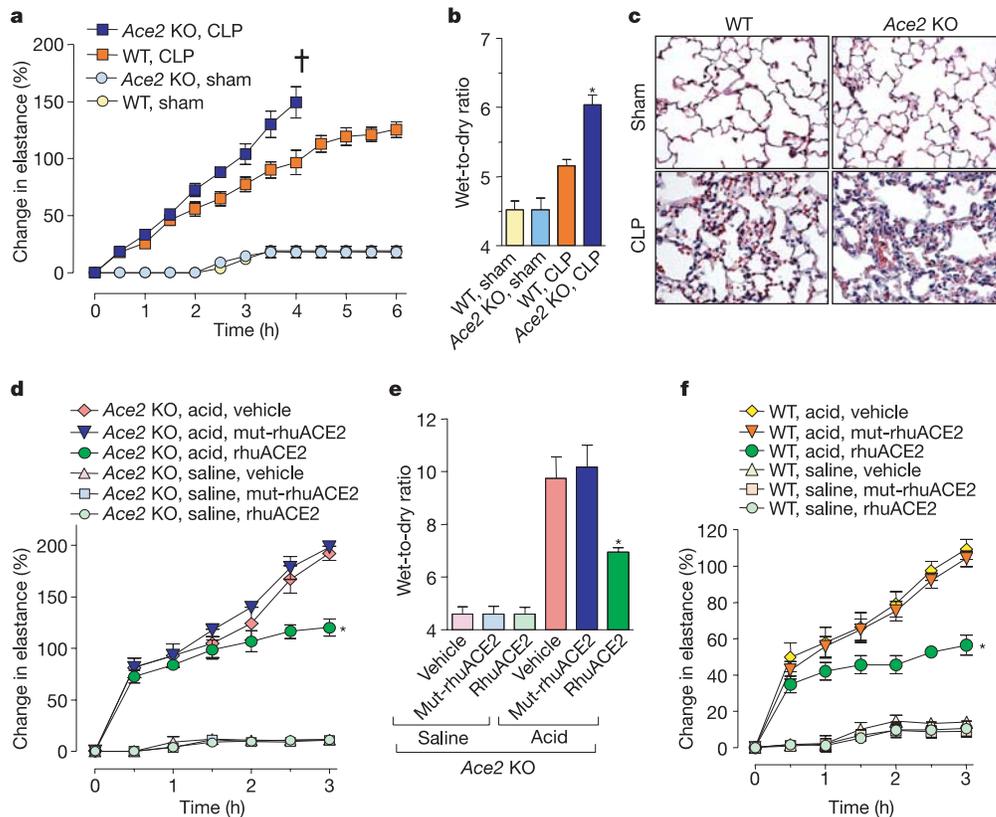


Figure 2 | ACE2 controls acute lung failure. **a**, Lung elastance after acute lung injury in WT and *Ace2* knockout (KO) mice induced by caecal ligation perforation (CLP). Eighteen hours after sham or CLP surgery, animals received mechanical ventilation for 6 h ($n = 10$ in CLP-treated groups, $n = 6$ in sham-treated groups). As 8/10 CLP-treated *Ace2* knockout mice died at 4–4.5 h, only data up to 4 h are shown. CLP-treated *Ace2* knockout mice had significantly higher elastance than CLP-treated WT mice ($P < 0.01$). **b**, **c**, Wet-to-dry weight ratios of lungs (**b**) and lung histopathology (**c**) in sham or CLP-treated WT and *Ace2* knockout mice determined after 4 h of ventilation. Asterisk denotes a significant difference ($P < 0.05$) between CLP-treated WT and *Ace2* knockout mice. Note the enhanced lung oedema and inflammatory infiltrates in *Ace2* knockout mice

(H&E staining, $\times 200$). **d**, **e**, Lung elastance (**d**) and wet-to-dry weight ratios (**e**) after acid or saline instillation of *Ace2* knockout mice injected intraperitoneally with recombinant human ACE2 protein (rhuACE2; 0.1 mg kg^{-1}), mutant rhuACE2 (mut-rhuACE2; 0.1 mg kg^{-1}) or vehicle ($n = 6$ per group). Asterisk denotes a significant difference ($P < 0.05$) comparing rhuACE2-treated *Ace2* knockout mice with mut-rhuACE2-treated and vehicle-treated *Ace2* knockout mice at 3 h. **f**, Lung elastance after acid instillation in WT mice treated with rhuACE2 protein (0.1 mg kg^{-1}), mut-rhuACE2 protein (0.1 mg kg^{-1}) or vehicle ($n = 6$ –8 per group). Asterisk denotes a significant difference ($P < 0.05$) between WT mice treated with rhuACE2 and mut-rhuACE2 or with vehicle at 3 h. Errors bars indicate s.e.m.

microvascular permeability²⁶. We first tested whether AngII can increase hydrostatic pressure using isolated, perfused murine lungs *ex vivo*²⁷. In this system, pulmonary perfusion pressures were comparable between wild-type and *Ace2* knockout mice under baseline control conditions (wild-type $3.0 \pm 1.9 \text{ cm H}_2\text{O}$, $n = 6$ versus *Ace2* knockout $1.8 \pm 1.6 \text{ cm H}_2\text{O}$, $n = 9$; mean \pm s.e.m.), and these values were not changed by either acid-treatment or continuous perfusion of the bacterial endotoxin lipopolysaccharide (LPS). Pulmonary perfusion pressures generated by AngI or AngII injection into lungs of acid-instilled animals or into lungs perfused with LPS were also similar between wild-type and *Ace2* knockout mice (see Supplementary Fig. 8a, b). Moreover, fractional shortening using echocardiography (an indicator of left ventricular systolic function) and mean arterial pressures were comparable between *Ace2* knockout and wild-type mice during the experimental period (see Supplementary Fig. 9a, b). Thus, the severe lung oedemas in *Ace2* knockout mice do not seem to be secondary to systemic haemodynamic alterations.

As enhanced pulmonary vascular permeability is a hallmark of acute lung injury/ARDS in humans², we examined whether loss of *Ace2* results in increased vascular permeability using Evans Blue dye injections as an *in vivo* indicator of albumin leakage²⁸. Acid aspiration increased vascular permeability in wild-type mice. In *Ace2*

knockout mice, pulmonary Evans Blue accumulation was greatly increased after acid aspiration (Fig. 4c, d). These results were confirmed using fluorescein isothiocyanate (FITC)-conjugated dextran (40 kDa) as another marker to assess vascular leakage of macromolecules (data not shown). Vascular permeability was significantly attenuated in the lungs of *Agtr1a*^{-/-} mice (Fig. 4e). We suggest that loss of ACE2 expression in acute lung injury leads to leaky pulmonary blood vessels through AT₁ receptor stimulation. However, hydrostatic oedemas cannot be excluded, and the effects of local AngII production on lung blood vessels require further investigation^{27,29}.

ARDS is the most severe form of a wide spectrum of pathological processes designated as acute lung injury². ARDS is characterized by pulmonary oedema due to increased vascular permeability, the accumulation of inflammatory cells and severe hypoxia². Predisposing factors for ARDS include sepsis, aspiration and pneumonias (including infections with SARS coronavirus^{1–5} or avian influenza viruses¹⁷). Our data show that acute lung injury results in a marked downregulation of ACE2, a key enzyme involved in the regulation of the renin–angiotensin system.

It has been previously shown that an insertion/deletion ACE polymorphism that affects ACE activity is associated with ARDS susceptibility and outcome³⁰. Our data provide a mechanistic

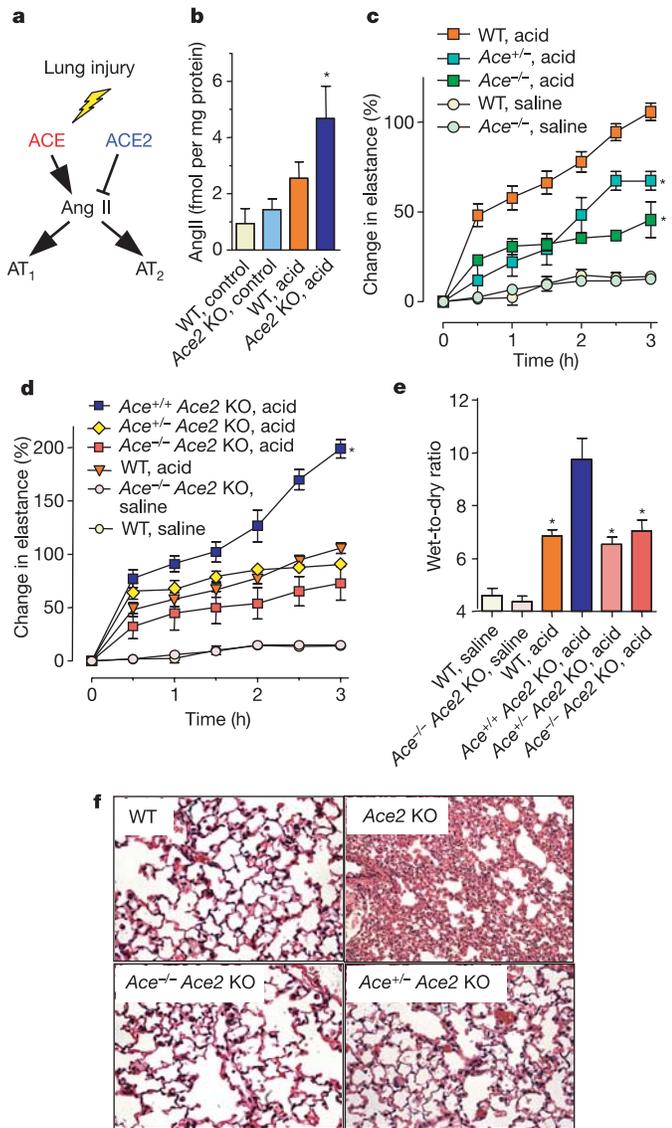


Figure 3 | ACE deficiency reduces the severity of acute lung injury. **a**, Schematic diagram of the renin–angiotensin system. **b**, Lung levels of AngII in control and acid-treated WT and *Ace2* knockout (KO) mice determined at 3 h by enzyme immunoassay ($n = 3–5$ per group). Asterisk denotes a significant difference ($P < 0.05$) between acid-treated WT and *Ace2* knockout mice. **c**, Lung elastance after acid instillation in *Ace*^{+/+} (WT), *Ace*^{+/-} and *Ace*^{-/-} mice ($n = 4–6$ mice per group). Asterisk denotes a significant difference ($P < 0.05$) comparing *Ace*^{+/+} with *Ace*^{+/-} and *Ace*^{-/-} mice at 3 h. **d, e**, Lung elastance (**d**) and wet-to-dry lung weight ratios (**e**) in acid- or saline-treated *Ace*^{+/+}*Ace2* KO, *Ace*^{+/-}*Ace2* KO, *Ace*^{-/-}*Ace2* KO and WT mice ($n = 5$ per group). Asterisk denotes a significant difference ($P < 0.05$) comparing *Ace2* KO with WT, *Ace*^{+/-}*Ace2* KO or *Ace*^{-/-}*Ace2* KO mice 3 h after acid-treatment. **f**, Lung histopathology. Severe lung interstitial oedema and leukocyte infiltration in *Ace2* KO mice are attenuated by homozygous (*Ace*^{-/-}) or heterozygous (*Ace*^{+/-}) mutations of *Ace* (H&E staining, $\times 200$). Error bars indicate s.e.m.

explanation for these clinical findings and indicate that, in the pathogenesis of acute lung injury, AngII is upregulated by ACE and drives severe lung failure through the AT₁a receptor. On the other hand, ACE2 and the AT₂ receptor protect against lung injury. Exogenous recombinant human ACE2 attenuates acute lung failure in *Ace2* knockout as well as in wild-type mice. This combination of genetic, pharmacological and protein rescue experiments defines a new and critical role for the renin–angiotensin system in the pathogenesis of acute lung injury, and show that ACE2 is a key

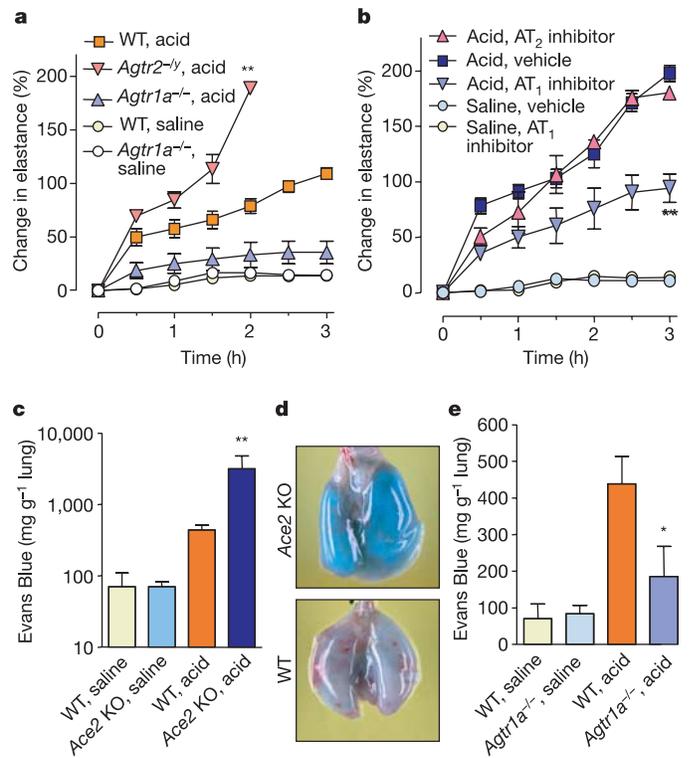


Figure 4 | The AngII receptor AT₁a controls acute lung injury severity and pulmonary vascular permeability. **a**, Lung elastance measurements in *Agtr1a*^{-/-} mice, *Agtr2*^{-/-} mice and WT mice after acid aspiration ($n = 4–6$ per group). All acid-treated *Agtr2*^{-/-} mice died after 2 h. There is a significant difference ($P < 0.01$) between acid-treated WT and acid-treated *Agtr1a*^{-/-} mice over the whole time course. Double asterisk denotes a significant difference ($P < 0.01$) between WT and *Agtr2*^{-/-} mice at 2 h. **b**, Lung elastance measurements in *Ace2* knockout mice treated with vehicle or inhibitors to AT₁ (Losartan, 15 mg kg⁻¹) or AT₂ (PD123.319, 15 mg kg⁻¹) after acid or saline instillation (see Methods, $n = 4–6$ per group). Double asterisk denotes a significant difference ($P < 0.01$) comparing *Ace2* knockout mice treated with AT₁ inhibitor with vehicle or AT₂ inhibitor treatment at 3 h. **c**, Pulmonary vascular permeability as determined by intravenous injection of Evans Blue. Extravascular Evans Blue in lungs was measured in WT and *Ace2* knockout mice 3 h after acid injury ($n = 5$ per group). Double asterisk denotes a significant difference ($P < 0.01$) between acid-treated WT and *Ace2* knockout mice. **d**, Representative images of Evans Blue-injected lungs of WT and *Ace2* knockout mice 3 h after acid aspiration. **e**, Extravascular Evans Blue in lungs of WT and *Agtr1a*^{-/-} mice 3 h after acid injury ($n = 5$ per group). Asterisk denotes a significant difference ($P < 0.05$) between acid-treated WT and *Agtr1a*^{-/-} mice at 3 h. Error bars indicate s.e.m.

molecule involved in the development and progression of acute lung failure.

METHODS

For detailed methods please refer to the Supplementary Information. **Animals.** *Ace2*, *Ace*, *Agtr1a* and *Agtr2* mutant mice have previously been described^{2,4,5,6}. Sex-, age-, and background-matched mice were used as controls. Basal lung functions and lung structure were comparable among all the mice tested. Mice were handled in accordance with institutional guidelines. **Experimental murine models of acute lung injury.** For acid aspiration-induced acute lung injury, anaesthetized mice were intratracheally instilled with HCl (pH 1.5; 2 ml kg⁻¹) and ventilated for 3 h (refs 18, 19). To study sepsis-induced acute lung injury, we performed caecal ligation perforation (CLP)²⁰. Sham-operated mice underwent the same procedure without ligation and puncture of the caecum. Eighteen hours after sham/CLP surgery, animals were subjected to mechanical ventilation for up to 6 h. For endotoxin-induced acute lung injury, anaesthetized mice received LPS and zymosan intratracheally immediately after starting mechanical ventilation and 1 h later, respectively¹⁸. In all acute lung

injury models, total positive end expiratory pressure ($PEEP_t$) and plateau pressure (P_{plat}) were measured at the end of expiratory and inspiratory occlusion, respectively. Elastance was calculated as $(P_{plat} - PEEP_t)$ divided by tidal volume (V_T) every 30 min during the ventilation periods.

Blood oxygenation, pulmonary oedema, pulmonary vascular permeability and histology. Blood samples were obtained from the left heart ventricle and partial pressure of oxygen in arterial blood (p_{aO_2}) was measured. To assess pulmonary oedemas, the lung wet-to-dry weight ratios were calculated. Pulmonary vascular permeability was assessed by measuring the pulmonary extravasation of Evans Blue. For histological analysis, 5- μ m thick sections were cut and stained with haematoxylin and eosin (H&E).

Recombinant ACE2 and AT₁/AT₂ receptor inhibitors. Thirty minutes before acid instillation, mice received intraperitoneal injections of recombinant human ACE2 (rhuACE2) protein (0.1 mg kg⁻¹) (R&D Systems or our own rhuACE2 preparation), catalytically inactive (H374N, H378N)¹⁰ mutant recombinant human ACE2 (mut-rhuACE2) or vehicle (0.1% BSA/PBS). All animals were then ventilated for 3 h. RhuACE2 protein and mut-rhuACE2 protein were purified from transfected CHO cells by affinity chromatography. The catalytic activities of purified recombinant ACE2 proteins were measured using the fluorogenic peptide Substrate VI (R&D Systems). Mut-rhuACE2-Fc showed >95% loss of catalytic activity (see Supplementary Fig. 3a). For inhibitor studies, mice received intraperitoneal injections of the AT₁ inhibitor Losartan (15 mg kg⁻¹), the AT₂ inhibitor PD123.319 (15 mg kg⁻¹) or control vehicle 30 min before surgical procedures.

Angiotensin II peptide levels and western blotting. AngII peptide levels were measured as described⁸. For western blotting, rabbit polyclonal anti-ACE2 antibody⁸ and rabbit polyclonal anti-mouse ACE antibody (R&D Systems) were used.

Statistical analyses. All data are shown as mean \pm s.e.m.. Measurements at single time points were analysed by analysis of variance (ANOVA). Time courses were analysed by repeated measurements ANOVA with Bonferroni post-tests.

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Supplementary Information is linked to the online version of the paper at www.nature.com/nature.

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