Metformin effectively treats Tsc1 deletion-caused kidney pathology by upregulating AMPK phosphorylation

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Abstract
Tuberous sclerosis complex (TSC) is characterized by hamartomatous lesions in multiple organs, with most patients developing polycystic kidney disease and leading to a decline of renal function. TSC is caused by loss-of-function mutations in either Tsc1 or Tsc2 gene, but currently, there is no effective treatment for aberrant kidney growth in TSC patients. By generating a renal proximal tubule-specific Tsc1 gene-knockout (Tsc1−/−) mouse model, we observed that Tsc1−/− mice developed aberrantly enlarged kidneys primarily due to hypertrophy and proliferation of proximal tubule cells, along with some cystogenesis, interstitial inflammation, and fibrosis. Mechanistic studies revealed inhibition of AMP-activated protein kinase (AMPK) phosphorylation at Thr-172 and activation of Akt phosphorylation at Ser-473 and Thr-308. We therefore treated Tsc1−/− mice with the AMPK activator, metformin, by daily intraperitoneal injection. Our results indicated that metformin increased the AMPK phosphorylation, but decreased the Akt phosphorylation. These signaling modulations resulted in inhibition of proliferation and induction of apoptosis in the renal proximal tubule cells of Tsc1−/− mice. Importantly, metformin treatment effectively prevented aberrant kidney enlargement and cyst growth, inhibited inflammatory response, attenuated interstitial fibrosis, and protected renal function. The effects of metformin were further confirmed by in vitro experiments. In conclusion, this study indicates a potential therapeutic effect of metformin on Tsc1 deletion-induced kidney pathology, although currently metformin is primarily prescribed to treat patients with type 2 diabetes.

Introduction
Tuberous sclerosis complex (TSC), a genetic disorder affecting multiple organ systems (including the brain, skin, heart, lungs, and kidneys), is caused by loss-of-function mutations in either TSC1 or TSC2 gene1. In normal human cells, the proteins TSC1 (also known as hamartin) and TSC2 (also called tuberin), which are encoded by TSC1 and TSC2 genes, respectively2,3, interact with TBC1D7 to form a functional ternary complex in which TSC2 acts as a selective GTPase-activating protein for the small GTPase, Rheb (Ras homolog enriched in brain)4. When TSC1 or TSC2 gene mutations inactivate the ternary TSC complex, Rheb is released to activate the mechanistic target of rapamycin complex 1 (mTORC1), causing dysregulated cell proliferation. In total, 60–80% of TSC patients suffer from renal manifestations of aberrant cell growth and dysregulated cell proliferation, resulting in renal cysts and hamartomatous renal tumors such as angiomyolipomas1,5,6. Such renal lesions can cause kidney enlargement and tubulointerstitial fibrosis, leading to impaired renal function and/or angiomyolipoma-associated hemorrhage7,8. Although more than half of
TSC patients fail to be diagnosed at the early stage of this disease, with innovative detection technologies, more and more TSC patients are diagnosed with kidney pathology.

Currently, mTORC1 is considered a pharmacological target for the treatment of TSC. Unfortunately, mTORC1 inhibitors currently used in the clinic seemed to only have a limited cytostatic effect, as the TSC tumors grew again after mTORC1 inhibitors were discontinued, and long-term mTORC1 inhibition could cause detrimental side effects. It is noteworthy that deletion of human TSC2 and PKD1 genes results in severe infantile polycystic kidney disease (PKD)—a contiguous gene syndrome. Neither sirolimus (rapamycin) nor everolimus (sirolimus analog) is effective in treating PKD, and these drugs have been reported to cause albuminuria (caused by sirolimus) and leukopenia (caused by everolimus). Moreover, our previous study demonstrated that either pharmacologic inhibition of mTORC1 or genetic deletion of rpS6 phosphorylation exacerbated structural and functional damage of the kidneys in Tsc1 gene-deleted mice, suggesting that alternative strategies are needed for effective treatment of TSC-associated kidney disease.

AMP-activated protein kinase (AMPK) is involved in multiple physiological regulations. AMPK regulates cell proliferation, apoptosis, and inflammation, in addition to regulation of energy metabolism. For instance, modulating AMPK has been demonstrated to be an effective maneuver for treating primary breast cancer, ovarian cancer, and inflammatory bowel disease. TSC-associated kidney pathology includes tubulointerstitial fibrosis and decreased apoptosis. Interestingly, pharmacological activation of AMPK has been demonstrated to be an effective strategy for the treatment of mouse Pkd1 deletion-induced PKD. Metformin is a potent AMPK activator and well-tolerated drug currently used for the treatment of type 2 diabetes. Metformin has been reported to increase AMPK activity in Pkd1−/− cells and prevent renal interstitial fibrosis in a mouse model of unilateral ureteral obstruction (UUO), in addition to ameliorating renal injuries in PKD. Enhancing AMPK phosphorylation with metformin can inhibit the growth and survival of cancer cells by inhibiting protein synthesis and angiogenesis, and has been reported for the treatment of cancer. Furthermore, metformin suppresses macrophage infiltration and inhibits expression of the inflammatory cytokine IL-6. Interestingly, metformin also alleviates lung fibrosis.

To date, however, the effect of AMPK modulation on TSC-associated kidney disease remains unknown. In the present study, we established an animal model of TSC by knocking out mouse Tsc1 gene in the renal proximal tubule (Tsc1PrtKO), and investigated the effect of metformin on renal pathology and functional changes to evaluate the potential of metformin for the treatment of TSC-associate kidney disease. Utilizing this Tsc1PrtKO mouse model, we also examined the cellular and molecular mechanisms of action of metformin in treating renal pathology and preserving renal function in TSC.

Materials and methods
Reagents and antibodies
Antibodies against β-actin (#60009-1-lg), IL-1β (#16806-1-AP), IL-6 (#66146-1-lg), and Fibronectin (#15613-1-AP), as well as goat anti-rabbit (#SA00001-1) and goat anti-mouse (#SA00001-15) secondary antibodies were purchased from ProteinTech (Rosemont, IL, USA). Antibodies against TSC1 (#6935), p-AMPK Thr-172 (#2535), AMPK (#5832), p-Akt Ser-473 (#4060), p-Akt Thr-308 (#13038), Akt (#2920), PCNA (#2586), Caspase-3 and Cleaved caspase-3 (#9664), BAD (#9239), BAX (#5023), F4/80 (#30325), CD3 (#99940), CD4 (#25229), and α-SMA (#19245) were purchased from Cell Signaling Technology (Danvers, MA, USA). Rabbit anti-Ki-67 (#15580) was purchased from ABCAM. Fluorescein-labeled LTL (#FL-1321) and Dylight 594 anti-rabbit IgG (#15580) were purchased from Vector Labs. Peroxidase-Technology (Danvers, MA, USA). Rabbit anti-Ki-67 (#15580) was purchased from ABCAM. Fluorescein-labeled LTL (#FL-1321) and Dylight 594 anti-rabbit IgG (#15580) were purchased from Vector Labs. Peroxidase marker and DAB + Substrate (#5007) were purchased from Dako.

Generation of renal proximal tubule-specific Tsc1 gene-knockout mice
Tsc1-flxed mice (Strain Name: STOCK Tsc1tm1Djk/J, Stock Number: J005680) were purchased from Nanjing Biomedical Research Institute of Nanjing University. The γGT-Cre mice (Strain Name: STOCK Tg (gGT1-CRE) M3Egn/J, Stock Number: 012841) were purchased from The Jackson Laboratory. Animals were housed at the veterinary facility (West 24th building), Shanghai Medical School, Fudan University. Animal care and all experimental procedures were approved by Fudan University Basic Medical School’s Experimental Animal Ethical Committee. Renal proximal tubule cell-specific homozygous Tsc1-knockout (Tsc1PrtKO) mice were generated by crossing Tsc1PrtKO mice with a transgenic mouse line expressing Cre recombinase under the control of the gamma-glutamyl transpeptidase promoter (γGT-Cre). The γGT-Cre mouse line has been used to successfully delete different genes of interest selectively in the renal proximal tubule. Tsc1PrtKO mice were used for all the Tsc1PrtKO/Tsc1PrtKO mice were included and the rest were excluded), which was not described before. For each group used in experiments, at least five mice were guaranteed for the adequate power to detect a
prespecified effect size. All mice were maintained and handled according to protocols approved by the NIH Animal Care and Use Committee. Block randomization was applied for the animal studies during the experiment and assessing the outcome.

Cell culture

NRK cells were from Prof. Lu Limin (Department of Physiology, Fudan University), which was obtained from the cell bank of Chinese Academy of Sciences (China). The NRK cells were cultured in high-glucose DMEM media (Gibco) containing 10% fetal bovine serum (Gibco), maintained at 37°C in a 5% CO₂ cell incubator (Thermofisher).

**Tsc1** RNA interference and metformin and dorsomorphin treatment of NRK cells

After the NRK cells were seeded at 3 x 10⁵ cells per well in 12-well plates for 24 h, **Tsc1** siRNA (25 nM) was transfected for 42 h. The siRNA sequence was 5'-CGCCTTTATGGAAATGACCCTTGTA-3', purchased from Gene Pharma, China. The cells were then treated with 3 mM metformin (Sangon Biotech) with or without 10 μM dorsomorphin (Selleck Chemicals) for 2 h. The NRK cells were lysed with radioimmunoprecipitation assay (RIPA) buffer (Sangon Biotech) containing protease inhibitors on ice for 40 min before centrifugation at 12,500 rpm for 20 min for collection of cell lysates. Equal amounts of protein samples were loaded onto 10% sodium dodecyl sulfate polyacrylamide gel electrophoresis (SDS-PAGE) for immunoblotting analysis as we described previously.

PCR primers and genotyping

Genomic DNA was isolated from mouse ear or toe biopsy samples for polymerase chain reaction (PCR) genotyping. PCR primers used for the floxed **Tsc1** allele are 5'-GTCAGACGAGTGGATAAGGC-3' and 5'-GAA TCAACCCCACAGAGCAT-3', while γGT-Cre transgene-detecting primers are 5'-AGGTGTAGAGAAGGCA CTTAGC-3' and 5'-CTATCGCCATCTTCCAGCAGG-3'. PCR conditions for both floxed-**Tsc1** and γGT-Cre are 94°C for 3 min followed by 94°C for 30 s, 65°C for 60 s, and 72°C for 60 s for 36 cycles, with an additional 2-min extension at 72°C.

Histologic examination, immunohistochemistry, immunofluorescence, TUNEL stain, and immunoblotting analysis

Mouse kidneys were fixed in 4% paraformaldehyde for paraffin-embedded kidney sections (5 μm), which were then deparaffinized and rehydrated for the following staining techniques. For histologic examination, H&E staining was performed using the standard methods. Immunohistochemistry and immunofluorescence staining were performed as described previously. Briefly, rehydrated kidney sections were subjected to antigen retrieval using the Antigen Unmasking Solution purchased from Vector, followed by blocking with 5% normal goat serum. The sections were then incubated with primary antibodies (indicated in the respective figures) at 4°C overnight, washed three times in PBS. For immunohistochemistry, the sections were subsequently incubated in a secondary antibody (peroxidase marker, Dako) at 37°C for 1 h. Then the signals were visualized using 1:50 diluted DAB (Dako), followed by counterstaining with hematoxylin and capturing images using a Nikon microscope. For immunofluorescence staining, after incubation with the primary antibodies indicated and washing with PBS, the sections were incubated with Dylight 594-conjugated secondary antibodies and LTL for 1 h, and images were captured using the Zeiss microscope system running on the ZEN software. For terminal deoxynucleotidyl transferase dUTP nick end labeling (TUNEL) stain, sections were processed as described above for the IHC and labeled with TdT using the In-Situ Cell Death Detection Kit (Roche). Image J was applied for the calculation of the immunohistochemical-positive area. Staining was visualized using 1:50 diluted DAB (Dako). Immunoblotting analyses were performed as we described previously.

Measurement of kidney function

Serum creatinine, albumin, and blood urea nitrogen (BUN) levels were measured as previously described. As for lactic acid in tissue levels, the same region of cortical tissues of mice was homogenized in RIPA cell lysis buffer, centrifuged at 12,500 rpm at 4°C for 20 min, and the supernatant was collected as the samples. Blood samples were collected from mice at different ages indicated in the corresponding figures. Serum creatinine, albumin, and BUN levels were immediately measured according to the instruction of Creatinine, Albumin, and Blood Urea Nitrogen Reagent Set (Pointe Scientific). Urine samples were collected simultaneously from mice at different ages and monitored by SDS-PAGE and Coomassie stain.

ELISA measurement

Culture media were collected from NRK cells treated as indicated above. The collected media were centrifuged at 2000–3000 rpm for 20 min, and the supernatants were collected for detection of IL-1β secretion according to the manufacturer’s instructions of the IL-1 beta Rat ELISA Kit (Invitrogen).

Masson trichrome staining

Kidney sections (5 μm) were deparaffinized, rehydrated, and stained with the Masson Trichrome Stain Kit.
Fig. 1 (See legend on next page.)
according to the manufacturer’s instructions (Sigma-Aldrich).

Sirius red staining
The slides were incubated with a 0.1% sirius red solution dissolved in aqueous saturated picric acid for 1 h, washed in acidified water (0.5% hydrogen chloride), dehydrated, and mounted with DPX Mounting. Collagen and non-collagen components were stained red and orange, respectively.

Statistical analysis
Data are presented as means ± SEM for at least three separate experiments (each in triplicate). An estimate of variation within each group is being statistically compared. An unpaired \( t \) test was used for statistical analysis; ANOVA and Bonferroni \( t \) tests were used for multiple group comparisons using GraphPad Prism 7. A \( P \) value < 0.05 (*\( P \)) compared with control was considered statistically significant.

Results

\( Tsc1 \) deletion in renal proximal tubules caused striking kidney growth

\( Tsc1 \) gene-floxed mice (\( Tsc1^{floxed/floxed} \)) were crossed with transgenic mice expressing the Cre recombinase under the control of the gamma-glutamyl transpeptidase (\( \gamma \)GT) promoter to generate a proximal tubule-specific \( Tsc1 \) gene-knockout (\( Tsc1^{ptKO} \)) mouse line. Sex-matched littermates with a genotype of \( Tsc1^{ptKO/ptKO} \) were used as control (called \( Tsc1^{Ctrl} \) mice hereafter) (Fig. S1a). \( Tsc1^{floxed/floxed} \) mice were identified by PCR as those carrying only the \( Tsc1 \)-floxed allele (230 bp), while \( Tsc1^{ptKO/wt} \) mice were those carrying both the \( Tsc1 \)-floxed allele (230 bp) and a wild-type allele (192 bp) of the \( Tsc1 \) gene. The \( \gamma \)GT-Cre gene was detected by PCR as a 370-bp band in \( Tsc1^{ptKO} \) mice but not in \( Tsc1^{Ctrl} \) mice (Fig. S1b). By 4 weeks of age, \( Tsc1^{ptKO} \) mice exhibited a smaller body size (Fig. S1c), but had strikingly larger kidneys (Fig. S1d), compared with \( Tsc1^{Ctrl} \) littermates.

As shown in Fig. 1, compared with \( Tsc1^{Ctrl} \) littermates, \( Tsc1^{ptKO} \) mice developed markedly enlarged kidneys by 2 weeks of age, which continued to enlarge aberrantly by 3 and 4 weeks of age (Fig. 1a), with significant increases in kidney weight (Fig. 1b) and kidney-to-body weight ratios (Fig. 1c). Renal histology revealed hypertrophy of renal tubules in the kidneys of \( Tsc1^{ptKO} \) mice by 2 weeks of age, and some enlarged renal tubules became dilated after 3 weeks of age (Fig. 1d). By 4 weeks of age, \( Tsc1^{ptKO} \) mice developed some microscopic renal cysts (Fig. 1d, e). More detailed examination of the renal pathology under higher magnification revealed that the large kidney phenotype of \( Tsc1^{ptKO} \) mice was caused by both cellular hypertrophy and proliferation, leading to enlargement of renal tubules with occasional microscopic hamartoma formation (Fig. 1e).

\( Tsc1 \) deletion in renal proximal tubules caused moderate albuminuria

To determine whether the structural damage of the kidneys shown in Fig. 1 has caused functional impairment of the kidneys in \( Tsc1^{ptKO} \) mice, we measured urinary protein excretion, serum creatinine, and BUN levels. As shown in Fig. 2a, urinalysis by SDS-PAGE revealed that \( Tsc1^{ptKO} \) mice developed mild albuminuria by 2 weeks of age that became markedly more apparent by 4 weeks of age (Fig. 1f). The proteinuria was confirmed by statistically significant increases in urinary albumin-to-creatinine ratios (Fig. 1g). The increased excretion of albumin in the urine may be explained by the data shown in Fig. 2, indicating that the renal proximal tubule cells in \( Tsc1^{ptKO} \) mice were activated to undergo dedifferentiation and proliferation, so their normal function of reabsorbing albumin from the glomerular filtrate might have been compromised. However, \( Tsc1^{ptKO} \) mice had normal serum creatinine levels (Fig. 2h) and BUN levels (Fig. 2i), indicating that the excretion of serum creatine and BUN was not decreased.
Tsc1 deletion in renal proximal tubules induced cell proliferation, suppressed apoptosis, and promoted inflammation and fibrosis

Immunohistochemistry with an antibody specific for Ki-67, a marker of proliferating cells, revealed aberrant proliferation of renal tubular epithelial cells in Tsc1^ptKO mice (Fig. 2a), which was confirmed by quantitation of Ki-67-positive cells in the kidney sections of Tsc1^ptKO mice, compared with that of Tsc1^Ctrl littermates (Fig. 2b). Double-immunofluorescence staining with Lotus Tetragonolobus
Lectin (LTL, a marker specific for proximal tubules) and the Ki-67 antibody indicated that increased proliferating cells are largely in the proximal tubules of Tsc1\(^{ptKO}\) mice; the increases in Ki-67-positive cells lining the renal proximal tubules were demonstrated to be statistically significant (Fig. 2d). Furthermore, immunoblotting revealed down-regulation of cleaved caspase-3, Bcl-2-associated X protein (BAX), and Bcl-2-associated death promoter (BAD) in Tsc1\(^{ptKO}\) mice (Fig. 2c), leading to decreased apoptosis in the kidneys of Tsc1\(^{ptKO}\) mice, compared with that of Tsc1\(^{Ctrl}\) mice, as indicated by TUNEL assays and quantitative data (Fig. 2e). The decreased apoptosis was consistent with decreases in cleaved caspase-3-positive cells in Tsc1\(^{ptKO}\) mice, indicated by immunofluorescence staining of kidney sections with LTL and an antibody specific for cleaved caspase-3 and the corresponding quantitative data (Fig. 2f).

In addition, immunoblotting analysis of renal cortical homogenates unveiled increases in the expression levels of TNF-α, IL-1β, and IL-6 in Tsc1\(^{ptKO}\) mice, compared with those in Tsc1\(^{Ctrl}\) mice (Fig. 3a). These findings may be explained by increases in inflammatory cell infiltrates, including F4/80-positive macrophages as well as CD3- or CD4-positive lymphocytes, in the interstitial areas of Tsc1\(^{ptKO}\) mice, compared with those of Tsc1\(^{Ctrl}\) mice, revealed by immunohistochemical staining with the indicated antibodies (Fig. 3b). Moreover, Masson’s trichrome and sirius red staining revealed renal interstitial fibrosis in Tsc1\(^{ptKO}\) mice (Fig. 3c).

**Daily metformin treatment reduced aberrant kidney growth and attenuated albuminuria in Tsc1\(^{ptKO}\) mice**

Previous studies have shown that metformin inhibits renal fibrosis in a mouse model of UUO\(^{27,28}\) and attenuates renal injury in PKD\(^{21}\). To test the effect of metformin on Tsc1 deletion-induced abnormal kidney growth, we treated Tsc1\(^{ptKO}\) mice with metformin by intraperitoneal injection once a day starting from 1 week of age for 3 consecutive weeks. Compared with vehicle-treated Tsc1\(^{ptKO}\) mice, metformin treatment markedly reduced kidney size (Fig. 4a, b), kidney weight (Fig. 4c), and kidney/body weight ratios (Fig. 4d) in Tsc1\(^{ptKO}\) mice. There was a 22.7% reduction in kidney weight and a 32.2% reduction in kidney-to-body weight ratios (Fig. 4c, d). Compared with the vehicle-treated group, metformin treatment also markedly attenuated cystic renal lesions (Fig. 4b, e). Importantly, metformin treatment effectively preserved renal function, evidenced by marked decreases in proteinuria (Fig. 4f) and urinary albumin-to-creatinine ratios (Fig. 4g), with no significant alterations in the normal serum creatinine (Fig. 4h) and BUN levels in Tsc1\(^{ptKO}\) mice (Fig. 4i), compared with vehicle-treated Tsc1\(^{ptKO}\) mice. Thus, metformin effectively reduced the abnormal enlargement of kidneys and mitigated the albuminuria in Tsc1\(^{ptKO}\) mice.

Metformin treatment inhibited excessive proliferation, induced apoptosis, suppressed inflammation, and attenuated fibrosis in the kidneys of Tsc1\(^{ptKO}\) mice

Previous studies have demonstrated that metformin inhibits the survival and growth of cancer cells\(^{29,30}\). To further investigate the beneficial effects of metformin in our Tsc1-knockout mouse model, we analyzed cell proliferation, apoptosis, interstitial inflammation, and fibrosis in the kidneys of Tsc1\(^{ptKO}\) mice treated with metformin, compared with vehicle-treated Tsc1\(^{ptKO}\) mice. As shown in Fig. 5a, b, immunohistochemical staining with antibodies against Ki-67, a marker for proliferating cells, revealed a significant reduction in Ki-67-positive cells in the kidneys of Tsc1\(^{ptKO}\) mice. Double-immunofluorescence staining with LTL (Green) and Ki-67 (Red) with statistical analysis demonstrated that the proliferating cells were primarily...
among the epithelial cells lining renal proximal tubules and renal cysts in $Tsc1^{ptKO}$ mice, and were markedly reduced by treatment with metformin (Fig. 5d).

In addition, we observed marked upregulation of several apoptosis mediators, including cleaved caspase-3, BAD, and BAX in metformin-treated $Tsc1^{ptKO}$ mice, compared with vehicle-treated $Tsc1^{CTRL}$ mice (Fig. 5c). Such molecular signaling alterations were accompanied by increases in TUNEL-positive cells in metformin-treated $Tsc1^{ptKO}$ mice relative to vehicle-treated $Tsc1^{ptKO}$ mice, which were also indicated in the statistical analysis of TUNEL-positive cells (Fig. 5e). Furthermore, double-immunofluorescence staining with LTL (green) and cleaved caspase-3 and the corresponding quantitative data confirmed that metformin treatment induced marked apoptosis of the LTL-positive renal proximal tubular epithelial cells, with some apoptotic cells having sloughed off the basement membrane and fragmented within the lumen, resulting in shrunken renal cysts (Fig. 5f). These results suggest that metformin has both antiproliferative and proapoptotic effects on the renal proximal tubular epithelial cells in $Tsc1^{ptKO}$ mice.

When we performed additional experiments to compare metformin-treated $Tsc1^{ptKO}$ mice with vehicle-treated $Tsc1^{CTRL}$ mice, we found that metformin markedly decreased the levels of TNF-α, IL-1β, and IL-6, as unveiled by immunoblotting (Fig. 6a). Metformin also markedly decreased F4/80-, CD3-, or CD4-positive inflammatory cells, revealed by immunohistochemistry (Fig. 6b). Moreover, Masson’s Trichrome and sirius red staining detected a marked reduction in collagen deposition, while immunohistochemistry revealed a marked reduction in α-SMA-positive myofibroblasts and fibronectin deposition in $Tsc1^{ptKO}$ mice in response to metformin treatment (Fig. 6c), which were also indicated by the quantitative data of fibronectin- and α-SMA-positive areas (Fig. 6d). These results suggest that in addition to an anti-inflammatory effect, metformin also has an anti-fibrotic effect in treating $Tsc1^{ptKO}$ mice.

The efficacy of metformin in treating cystic kidney disease was largely mediated by upregulation of AMPK phosphorylation

Previous studies have demonstrated that Akt mediates the survival of renal proximal tubular cells because apoptosis induced by serum withdrawal, hydrogen peroxide, etoposide, or excess free arachidonic acid is inhibited by activation of Akt. Renal proximal tubular cells make up the bulk of the renal cortex. Interestingly, our additional mechanistic studies by immunoblotting of renal cortical homogenates indicated that $Tsc1^{ptKO}$ mice exhibited discernible decreases in Thr-172-phosphorylated AMPK along with parallel increases in Ser-473- and Thr-308-phosphorylated Akt levels, compared with $Tsc1^{CTRL}$ mice.

The efficacy of metformin in treating cystic kidney disease was largely mediated by upregulation of AMPK phosphorylation.
Fig. 4 Metformin prevented Tsc1 deletion-induced kidney growth and ameliorated Tsc1 deletion-induced renal injuries by 4 weeks of age. Representative a kidney images and b kidney sections of metformin (20 mg/mL in vehicle, i.p.) and vehicle- (0.9% saline, i.p.) treated Tsc1\textsuperscript{ptKO} mice. The metformin- and vehicle-treated mice were given daily weight-adjusted i.p. injections from 1 to 4 weeks of age. Metformin prevented c kidney weights and d kidney- to-body weight ratios in Tsc1\textsuperscript{ptKO} mice, compared with vehicle-treated group. Data were from 4 weeks of age mice (n = 5–6 each per group) and expressed as means ± SEM, **P < 0.01. e Higher-magnification light microscopy demonstrated that metformin decreased renal cysts, as well as suppressed enlarged proximal tubules in Tsc1\textsuperscript{ptKO} mice, compared with vehicle-treated group. (Scale bar: 2 mm in all images of (b), 100 μm in all images of (e)). f Coomassie staining showed that metformin reduced albuminuria in Tsc1\textsuperscript{ptKO} mice, compared with vehicle-treated group. Data were shown of three independent experiments. Metformin significantly decreased g ACR in Tsc1\textsuperscript{ptKO} mice, compared with vehicle-treated group. Besides, reductions of h Scr and i BUN were detected in Tsc1\textsuperscript{ptKO} mice, although there was no statistical significance. Data were from 4 weeks of age mice (n = 5–6 each per group) and expressed as means ± SEM, **P < 0.01.
Immunohistochemical staining with antibodies specific for phospho-AMPK and phosphor-Akt, respectively, confirmed that such signaling alterations were largely localized to the epithelial cells lining the activated proximal tubules and renal cysts in Tsc1^prKO mice (Fig. 7b). Importantly, compared with vehicle-treated Tsc1^prKO mice, metformin-treated Tsc1^prKO mice showed upregulated AMPK phosphorylation and inhibited Akt phosphorylation,
revealed by immunoblotting (Fig. 7c) and confirmed by immunohistochemistry (Fig. 7d).

The in vivo animal data demonstrated above suggest that metformin treatment effectively reduces Tsc1 deletion-induced aberrant kidney growth by upregulating AMPK phosphorylation, leading to inhibition of the pro-survival factor, Akt, and activation of the proapoptotic factors, BAD and cleaved caspase-3, in the renal proximal tubular cells. To confirm such a molecular mechanism, we performed additional experiments in cultured NRK cells, a rat renal proximal tubular cell line. As shown in Fig. 8a, siRNA-induced knockdown of Tsc1 downregulated Thr-172-phosphorylated AMPK, but upregulated Ser-473- and Thr-308-phosphorylated Akt. Such effects were prevented by metformin treatment. These effects were reversed by treatment with dorsomorphin, an AMPK inhibitor.

Moreover, the knockdown of Tsc1 also elevated PCNA and reduced the expression of cleaved caspase-3 and BAD. Metformin treatment reduced PCNA, but increased cleaved caspase-3 and BAD; these effects were antagonized by treatment with the AMPK inhibitor, dorsomorphin (Fig. 8b). Our further experiments indicated that the knockdown of Tsc1 upregulated fibronectin and IL-1β. These changes were reversed by metformin treatment, but re-upregulated by dorsomorphin (Fig. 8c, d). Thus, our in vitro cell culture studies have confirmed our in vivo animal data, indicating that the restored phosphorylation level of AMPK mediates the pharmacological effect of metformin in suppressing cell proliferation, promoting apoptosis, and inhibiting the production of inflammatory cytokines as well as fibrotic factors.

**Discussion**

AMPK inactivation has been observed in multiple diseases, such as enlarged cysts in PKD, uncontrolled proliferation in breast cancer, renal inflammation induced by cyclophosphamide, and lung fibrosis after bleomycin stimulation. In our study, downregulated AMPK phosphorylation was also observed in the enlarged kidneys of Tsc1ptKO mice, which also exhibited aberrant proliferation of proximal tubule cells, renal interstitial inflammation, and fibrosis. The reduced AMPK phosphorylation in TSC is presumably caused by decreased AMP-to-ATP ratio. However, the specific mechanisms regulating AMPK phosphorylation levels still need to be further elaborated under the influence of metformin.

Our results indicate that metformin treatment inhibits cell proliferation and induces apoptosis in the renal proximal tubules of Tsc1ptKO mice by upregulating AMPK phosphorylation, in the meantime downregulating Akt phosphorylation. Our findings support the notion that AMPK phosphorylation-dependent inactivation of Akt inhibits cell proliferation and promotes apoptosis. Previous studies demonstrated that AMPK halts cycling cells in G1 phase by activating phosphorylation of p53 as well as inhibiting cycling D1, leading to cell-cycle arrest and exerting anticancer effect. Moreover, as a critical cycling procedure, mitosis is also under the regulation of AMPK by phosphorylating PPP1R12C at Ser452, which is elevated in mitotic cells. All these researches illustrate the critical role of AMPK in regulating proliferation and apoptosis, which inspires a novel idea applying AMPK activator in dealing with tumor cells.

Therefore, as potent AMPK activators, either resveratrol or metformin could slow the progress of cancers in ovary or breast through inhibiting cell proliferation and promoting apoptosis by activating AMPK phosphorylation, which demonstrates the efficacy of AMPK against tumorigenesis. Of note, due to the mutual phosphorylation regulation of AMPK and Akt, the inactivation of AMPK...
prominently facilitates the Akt phosphorylation\textsuperscript{46}, which downregulates proapoptotic factors such as BAX and BAD, leading to inhibition of cell apoptosis\textsuperscript{47}. In our study, metformin-treated \textit{Tsc1}\textsuperscript{ptKO} mice showed an evident proapoptotic effect by TUNEL assay, accompanied with immunoblotting showing downregulation of cleaved caspase-3, BAX, and BAD, as well as reduction of Ki-67. In vitro cell culture studies indicated that the effect of metformin in regulating proliferation and apoptosis was reversed by dorsomorphin, which further demonstrated the involvement of AMPK phosphorylation after metformin performance. Such a cellular mechanism may explain why metformin reduced renal cystogenesis and suppressed aberrant enlargement of the kidneys in \textit{Tsc1}\textsuperscript{ptKO} mice.

Apart from uncontrolled proliferative levels in the kidney, interstitial fibrosis is profoundly relevant to renal function as well\textsuperscript{27}. Previous studies have demonstrated the relevance between inactivation of AMPK phosphorylation and tissue

\begin{figure}[h]
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\includegraphics[width=\textwidth]{fig6.png}
\caption{Metformin suppressed the interstitial inflammation as well as fibrosis in \textit{Tsc1}\textsuperscript{ptKO} mice by 4 weeks of age. (a) Immunoblotting showed that metformin reduced the expression of TNF-\textalpha, IL-1\beta, and IL-6 in the renal cortex of \textit{Tsc1}\textsuperscript{ptKO} mice, compared with the vehicle-treated group. Data were shown of three independent experiments. (b) Immunohistochemistry showed that metformin decreased renal interstitial F4/80, as well as reductions of CD3- and CD4-positive staining cells in \textit{Tsc1}\textsuperscript{ptKO} mice, compared with the vehicle-treated group. (c) Decreased renal interstitial collagen formation was indicated by Masson’s trichrome and sirius red staining after metformin treatment in \textit{Tsc1}\textsuperscript{ptKO} mice, compared with the vehicle-treated group. In the meantime, decreased renal interstitial fibronectin and \alpha-SMA were demonstrated by immunohistochemistry in metformin-treated \textit{Tsc1}\textsuperscript{ptKO} mice, compared with the vehicle-treated group. (d) The drop of fibronectin and \alpha-SMA-positive areas was observed after the treatment of metformin in \textit{Tsc1}\textsuperscript{ptKO} mice, compared with the vehicle-treated group. Only fibronectin and \alpha-SMA-positive areas were analyzed for the numerator, which was divided by the total area of the \times400 magnification images and then multiplied by 100%. Data were from 4 weeks of age mice (\(n=5\) each per group) (scale bar: 100 \textmu m in all images of (b), upper four images of (c), and 50 \textmu m in lower four images of (c)).}
\end{figure}
fibrosis\(^3\); hence, activating AMPK phosphorylation has been considered a feasible way in anti-fibrosis. No matter the deposition of type IV collagen in the kidneys\(^4\), or UUO-induced renal fibrosis\(^4\), even bleomycin-induced pulmonary fibrosis\(^3\), all could be reversed by activating AMPK phosphorylation as well. Besides, metformin could suppress fibrosis not only through inhibiting the generation of myofibroblasts\(^5\), but also via downregulating the expression of HIF-1\(\alpha\)\(^5\) and TGF-\(\beta\)\(^2\) in either kidney or lung.

In our study, decreased renal fibrosis was detected in metformin-treated \(Tsc1^{ptKO}\) group, evidenced by reduced \(\alpha\)-SMA-positive fibroblasts as well as decreased fibronectin and reduced collagen levels. Our in vitro experiments further demonstrated that the suppression of fibronectin production was dependent on metformin-induced AMPK phosphorylation. The reduced renal interstitial fibrosis by metformin may contribute, at least in part, to the ameliorated renal function. Interestingly, Vesey et al.\(^5\) once showed increased secretion of \(\alpha\)-SMA and fibronectin under IL-1 stimulation in proximal tubule cells, which suggests that the suppressed fibrosis might be relevant to inhibition of inflammation as well.
Besides fibrosis, an elevated level of renal interstitial inflammation is also displayed in Tsc1<sup>ptKO</sup> mice, which could further damage renal function. AMPK phosphorylation, on the contrary, inhibited inflammation by suppressing the expression of IL-6<sup>24</sup>, TNF-α, and MCP-1 in adipocytes<sup>24</sup>, intestinal epithelial cells<sup>22</sup>, or kidney of...
Therefore, the reactivation of AMPK phosphorylation is likely to be a promising anti-inflammatory strategy for the treatment of TSC-associated renal diseases. As a potent AMPK activator, metformin was considered powerful in reducing inflammation, which not only protected renal function in DN or attenuated renal injuries induced by cyclophosphamide, but even slowed the progression of atherosclerosis as well by activating AMPK phosphorylation.

Our results demonstrated significant downregulation of inflammatory cell markers (F4/80, CD3, CD4, TNF-α, IL-1β, and IL-6) as the potential mechanism by which metformin inhibits inflammatory responses through increased AMPK phosphorylation in Tsc1flKO mice. Furthermore, siRNA downregulation of Tsc1 induced a significant increase of IL-1β production in renal proximal tubular epithelial cells in vitro. Treatment with metformin abolished the increase, which was reversed by the AMPK inhibitor, dorsomorphin. Thus, the potent anti-inflammatory effect of metformin in treating TSC-related renal pathology is dependent on AMPK phosphorylation.

In the present, we report for the first time that metformin can effectively treat the aberrant kidney growth and renal pathology caused by Tsc1 deletion in the renal proximal tubule cells of Tsc1flKO mice. Mechanistically, our results indicate that metformin treatment inhibits hyperphosphorylation of Akt at both Thr-308 and Ser-473, and thereby prevents activation and aberrant proliferation of the renal proximal tubule cells, thus preventing proteinuria and maintaining kidney function. Importantly, we also observed that metformin treatment additionally inhibits the production of pro-inflammatory cytokines, suppresses infiltration of inflammatory cells, and reduces deposition of fibrotic extracellular matrix proteins in the kidneys of Tsc1flKO mice. These multiple beneficial effects are likely what have conferred the remarkable therapeutic efficacy of metformin for the treatment of Tsc1 deletion-induced kidney pathology. Our additional mechanistic cell culture studies unveiled that these therapeutic effects of metformin are accompanied by reactivation of AMPK phosphorylation at Thr-172 that is antagonized by the AMPK inhibitor, dorsomorphin. Thus, it is our hope that this report will lead to clinical trials for the treatment of TSC-associated manifestations by repurposing the use of metformin, which is currently prescribed for the treatment of type 2 diabetic patients.
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