

No Amelioration of Uromodulin Maturation and Trafficking Defect by Sodium-4-Phenylbutyrate *in vivo*: Studies in Mouse Models of Uromodulin-associated Kidney Disease

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Background: *In-vitro* studies suggested the chemical chaperone 4-PBA as treatment of Uromodulin-associated kidney disease (UAKD).

Results: 4-PBA did not ameliorate uromodulin maturation defects in two UAKD mouse models. Renal abundance/phosphorylation of several NF-kappaB pathway components was increased.

Conclusion: 4-PBA failed to cure UAKD *in-vivo* but deteriorated kidney function.

Significance: NF-kappaB pathway was revealed as potential target for UAKD treatment.

ABSTRACT

Uromodulin (UMOD)-associated kidney disease (UAKD) belongs to the hereditary progressive ER storage diseases caused by maturation defects of mutant UMOD protein. Current treatments of UAKD patients are symptomatic and cannot prevent disease progression. Two *in vitro* studies reported a positive effect of the chemical chaperone sodium-4-phenylbutyrate (4-PBA) on mutant UMOD maturation. Thus, 4-PBA was suggested as potential treatment for UAKD. This study evaluated the effects of 4-PBA in two mouse models of UAKD. In contrast to previous *in vitro* studies, treatment with 4-PBA did not increase HSP70 expression or improve maturation and trafficking

of mutant UMOD *in vivo*. Kidney function of UAKD mice was actually deteriorated by 4-PBA-treatment. In transfected tubular epithelial cells, 4-PBA did not improve maturation, but increased the expression level of both mutant and wild-type UMOD protein. Activation of NF- κ B pathway in thick ascending limb of Henle's loop cells of UAKD mice was detected by increased abundance of RelB and phospho-IKK α/β , an indirect activator of NF- κ B. Further, the abundance of NF- κ B1 p105/p50, NF- κ B2 p100/p52 and TRAF2 was increased in UAKD. NF- κ B activation was identified as a novel disease mechanism of UAKD and might be a target for therapeutic intervention.

INTRODUCTION

Uromodulin-associated kidney disease (UAKD) is a dominant hereditary disorder caused by mutations of the uromodulin (UMOD) gene and comprises medullary cystic kidney disease type 2 (MCKD2; OMIM #603860), familial juvenile hyperuricemic nephropathy (FJHN; OMIM #162000), and glomerulocystic kidney disease (GCKD; OMIM #609886) (1-3). In about 80% of UAKD cases, earliest symptoms are hypouricemic hyperuricemia, frequently associated with gout, commonly diagnosed in the late teenage years or in young adults (1,4,5).

Further, mild defect in urinary concentration ability and a usually strong decrease of urinary uromodulin excretion are present. UAKD is progressive and can lead to end-stage renal disease at an age ranging between 25 and more than 70 years with the need of renal replacement therapy (4,6,7). Histomorphological kidney alterations usually comprise diffuse tubulointerstitial fibrosis with moderate inflammatory cell infiltration, tubular atrophy, and intracytoplasmic accumulation of uromodulin in tubular cells (4,8). In about one-third of the patients, uni- or bilateral renal cysts are diagnosed (7).

Allelic *UMOD* variants causing UAKD are rare and have to be differentiated from common allelic variants of the *UMOD* gene. In various GWAS studies, common allelic *UMOD* promoter variants were identified to be associated with increased risk of chronic kidney disease (CKD) and other complex traits, such as hypertension and kidney stones (reviewed in 4). These risk variants of the *UMOD* promoter were recently identified to increase *UMOD* expression and secretion which leads, by influencing salt reabsorption in the kidney, to increased risk of developing hypertension and CKD (9).

Uromodulin is selectively expressed in cells of the thick ascending limb of Henle's loop (TALH) and of the early distal convoluted tubules (10). It is a highly glycosylated protein processed in the endoplasmic reticulum (ER) and Golgi apparatus, translocated to the luminal cell membrane and released in the urine by proteolytic cleavage. So far, more than 70 different UAKD-causing *UMOD* mutations have been published, of which all but four (in-frame deletions) lead to amino acid substitutions (4,7,11). Mutant uromodulin isoforms exhibit a protein maturation defect and are retained in the ER (12,13). ER retention of mutated *UMOD* is considered as a key step in the pathogenesis of UAKD (4). Analyses of kidney biopsies of UAKD patients revealed a hyperplastic ER in TALH cells with overexpression of BiP and PDI, indicative of ER stress (8,14,15). ER hyperplasia and activation of ER-resident molecular chaperones (e.g. BiP) and foldases represent cellular counteractions of unfolded protein response (UPR) to increase folding capacity of the ER (16). Gain-of-toxic function of uromodulin mutations may lead to disturbance of ER homeostasis in tubular cells, to kidney dysfunction and finally to UAKD (17).

Therapies of UAKD other than the symptomatic corrections of hyperuricemia and of water and electrolyte imbalance are not available (4). Treatment

of hyperuricemia for prevention of gout can be efficiently done by xanthine oxidase inhibitors like allopurinol and febuxostat, by uricosuric drugs like probenecid or by benzbromarone (5,18). However, treatment of hyperuricemia does not prevent disease progression and deterioration of kidney function in UAKD (5). *UMOD* maturation retardation in the ER and ER overload represent the central pathomechanism in UAKD, a member of the ER storage diseases (4). Thus, novel therapeutic strategies for UAKD focus on improvement of ER capacity and function, and this approach might delay disease progression. Two independent *in vitro* studies postulated a positive effect of the chemical chaperone sodium 4-phenylbutyrate (4-PBA) on mutant *UMOD* protein maturation, suggesting that 4-PBA might reduce ER retention of *UMOD* and might ameliorate UAKD (19,20).

The present study evaluated the effects of 4-PBA in *Umod*^{A227T} and *Umod*^{C93F} mutant mice, two recently described models of UAKD (17,21).

EXPERIMENTAL PROCEDURES

Animals. *Umod*^{A227T} and *Umod*^{C93F} mutant mice were generated by ENU mutagenesis as described previously (17,21). Both *Umod* mutant mouse lines exhibit the key features of UAKD like maturation defect and retention of uromodulin in the hyperplastic endoplasmic reticulum of TALH cells and impaired kidney function with mild defect in urinary concentration ability, reduced fractional excretion of uric acid, and reduced uromodulin excretion. Both mouse lines were maintained on the C3HeB/FeJ (C3H) genetic background and housed in a specific pathogen-free mouse facility as previously described (17,21). All mice received food (V1124-3, Ssniff, Soest, Germany) and water *ad libitum* and mouse husbandry was performed under standard environment conditions (22 ± 2°C, 40-50% relative humidity, 12-h light/dark cycle). All animal experiments were approved and carried out in accordance with the German Animal Welfare Act with permission from the responsible veterinary authority.

Experimental schedule of administration of sodium-4-phenylbutyrate (4-PBA) in vivo. *In vivo* administration of the chemical chaperone 4-PBA with the drinking water in male homozygous *Umod*^{A227T} mutants, male homozygous *Umod*^{C93F} mutants, and their male wild-type littermates started at an age of two months. At this age, the UAKD phenotype is

already clinically present in *Umod* mutant mice of both lines but is in an early disease stage where no progressed morphological kidney alterations like interstitial fibrosis, tubular atrophy or infiltrates of inflammatory cells are present. Drug administration was performed for 60±4 days. Pharmaceutical-grade 4-phenylbutyric acid sodium salt (Scandinavian Formulas, Pennsylvania, USA, water solubility: 0.62 g/ml) was dissolved in drinking water and administered at a daily dosage of 1 g/kg body weight. 4-PBA solutions were renewed every third day. Mice of the placebo groups received drinking water without supplement. Body weights of mice were measured weekly, and water intake was measured per cage after each solution change. Blood sampling was performed at day 0 prior to the start of the experiment, and at day 30±4 and 60±4 at the mid and the end of the experiment. On day 56±4, the mice were individually transferred to metabolic cages (Tecniplast, No. 3600M021, Italy). After a conditioning period of two days to the metabolic cages, water consumption and urine excretion was assessed for two days. On day 60±4, the animals were anaesthetized for retro-orbital blood collection, followed by necropsy.

Clinical chemical analyses. Analyses of blood and urine parameters were carried out as described previously (17,22). Plasma creatinine values were determined enzymatically (Biomed), while urine creatinine concentration was determined by the Jaffé method.

Analyses of urine and kidney function. Urine osmolality was determined by freezing point depression analysis. Daily excretion of solutes was measured. Calculations of creatinine clearance and of fractional excretion of a solute x (FEx) was performed as described previously (17,23).

Administration of 4-PBA in vitro. MTC (immortalized murine proximal tubular epithelial cells) (24) were friendly donated by G. Wolf, Jena and were cultured on collagen-coated plates in DMEM high glucose medium containing 10% fetal calf serum (FKS), 1% penicillin/streptomycin, and 1% insulin/transferrin/selenium. Transfection of MTC cells (passage 6) was performed by Nucleofection™ (Lonza, program U12) with the constructs pcDNA3-*Umod*^{wt}, pcDNA3-*Umod*^{A227T}, and pcDNA3-*Umod*^{C93F}, respectively, and stable clones were generated by G418 selection (2 mg/ml). Two stable clones per

construct (with high and with medium UMOD expression) were selected for further analyses.

Isolation of primary kidney cells was performed by digest with 0.1% (w/v) collagenase II (Life Technology, Germany) for 30 min at 37°C after mincing kidneys of homozygous *Umod*^{A227T} mutant, homozygous *Umod*^{C93F} mutant, or wild-type mice. After gently pressing the digested renal tissue through a 100-µm cell strainer, a second digest with 0.1% (w/v) collagenase II and 7.5 U/ml DNase I for 30 min at 37°C was performed. Finally, undigested tubular fragments and glomeruli were removed by filtration through a 40-µm cell strainer, and cell suspension was seeded on collagen-coated plates or 8-well slides (BD Bioscience, Germany) in DMEM/Ham's F12 supplemented with 20% FKS, 1% penicillin/streptomycin, 1% sodium pyruvate, 1% non-essential amino acids, and 1% insulin/transferrin/selenium.

Treatment of MTC clones with 4-PBA was performed at passages 12 – 19. The doses of drug supplement were tested in pilot studies. Seeding of cells on collagen-coated 12-well plates, volume of culture medium and duration of treatment were standardized. Treatment of MTC clones was performed on two different stably transfected *Umod*^{wt}-MTC clones and on each three different MTC clones stable transfected with *Umod*^{A227T} or *Umod*^{C93F}, respectively. Treatment of primary kidney cells was performed on 8-well slides.

Morphological analysis. Histological analyses of kidneys were performed as described previously (21). Paraffin sections were stained with hematoxylin and eosin (H&E), Masson-Trichrom, and PAS-silver. Immunohistochemistry was performed using the following primary antibodies: rabbit polyclonal antibody against human THP (H-135; Santa Cruz Biotechnology), rat monoclonal antibody against mouse uromodulin (clone 774056; R&D Systems), rabbit polyclonal antibody against HSP70 (Cell Signaling), rabbit monoclonal antibody against RelB (C1E4, no. 4922, Cell Signaling), and rabbit monoclonal antibody against phospho-IKKα/β (Ser176/180) (no. 2697, Cell Signaling). Immunoreactivity was visualized using 3,3'-diaminobenzidine tetrahydrochloride dihydrate (DAB) (brown color) or using Vector Red Alkaline Phosphatase Substrate Kit I (red color). Nuclear counterstaining was done with haemalum.

For immunofluorescence analyses of cells, the following primary antibodies were used: polyclonal

goat antiserum against human uromucoid (MP Biomedicals) and polyclonal rabbit anti GRP78/BiP IgG (Abcam). Fluorescence analyses were performed using a confocal laser scanning microscope (LSM 710, Zeiss).

Western blot analyses. Twenty-four-hour urine samples, standardized for equal creatinine levels, were analyzed for uromodulin content as described previously (21). MTC cells were lysed in Laemmli extraction buffer, protein concentration was determined by BCA assay and equal protein amounts (10 µg) of denatured cell lysate per lane was separated in 8% SDS-polyacrylamide minigels. Equal amounts of supernatant of cells cultured under standardized conditions were used for analysis of their UMOD content.

TALH segment density in the kidney is highest in the outer medullar region. The outer medulla was obtained after cutting each fresh kidney of four-month-old mice along the midsagittal plane into two halves and further into transverse sections, and by removing the cortex and inner medulla. After homogenization of the outer medulla in Laemmli extraction buffer and determination of protein concentration by BCA assay, 15 µg of denatured tissue lysate per lane was separated in 10% or 12% SDS-polyacrylamide minigels and blotted on PVDF membranes. Equal loading was controlled by Ponceau staining.

The following primary antibodies were used: rabbit polyclonal antibody against human THP (H-135; Santa Cruz Biotechnology), rabbit polyclonal antibody against BiP (no. 3183), rabbit monoclonal antibody against GAPDH (no. 2118), rabbit polyclonal antibody against IKKα (no. 2682), rabbit monoclonal antibody against phospho-IKKα/β (Ser176/177) (C84E11, no: 2078), rabbit polyclonal antibody against NF-κB1 p105/p50 (no. 12540), rabbit polyclonal antibody against NF-κB2 p100/p52 (no. 4882), rabbit monoclonal antibody against NF-κB p65 (no. 4764), rabbit monoclonal antibody against RelB (C1E4, no. 4922), rabbit polyclonal antibody against TRAF2 (no. 4712), and rabbit polyclonal antibody against TRAF3 (no. 4729) (all antibodies except for THP were bought from Cell Signaling). Bound antibodies were visualized using ECL reagent (GE Healthcare Amersham Biosciences). Signal intensities were quantified using ImageQuant (GE Healthcare). Standardization of equal loading was referred to the

signal intensities of GAPDH of the corresponding PVDF membrane.

Statistical analysis. Data are shown as means ± SD as indicated. Data were analyzed by using unpaired Student's *t*-test or 2-way ANOVA as indicated using Sigma Plot 12.0. Holm-Bonferroni correction for multiple comparisons was performed where indicated.

RESULTS

1. Clinical and clinical-chemical findings in placebo and 4-PBA treated UAKD mouse models and wild-type controls

1.1. Umod mutant versus wild-type mice after placebo treatment. Four-month-old homozygous mutants (*Umod*^{A227T} and *Umod*^{C93F}) receiving drinking water without supplement (placebo treatment group) had significantly increased plasma urea levels and plasma lipase activities compared to corresponding wild-type control animals, whereas plasma triglyceride concentrations were significantly decreased (Table 1, Figure 1A). Additionally, plasma calcium level was mildly increased in *Umod*^{C93F} mutant mice compared to wild-type mice. *Umod* mutants of both lines exhibited distinctly reduced body weight, significantly reduced urine osmolality combined with slightly increased urine volume, significantly increased urinary calcium excretion and fractional excretion (FE) of calcium as well as reduced urinary excretion of phosphate and reduced FE_P and FE_{urea}, when compared to wild-type mice (Table 2, Figure 1B). Further, *Umod*^{C93F} mutants exhibited increased 24-hour urinary potassium excretion vs. wild-type mice. These findings are in line with earlier published effects of mutant UMOD expression on clinical-chemical parameters of plasma and urine in both *Umod*^{A227T} and *Umod*^{C93F} mutant mice representing models for UAKD (17,21). *Umod*^{C93F} mutants exhibited a more severe UAKD phenotype than mutants of the *Umod*^{A227T} mouse line (21).

1.2. Umod mutant versus wild-type mice after 4-PBA treatment. After two months of administration of 4-PBA, at an age of 4 months, *Umod*^{A227T} mutants and *Umod*^{C93F} mutants displayed distinctly higher plasma levels of urea, creatinine and calcium as well as higher plasma lipase activities and decreased plasma triglyceride levels, compared to wild-type controls (Table 1, Figure 1A). Further, 4-PBA treated *Umod*^{C93F} mutant mice exhibited a significantly increased plasma chloride concentration compared to

wild-type controls. A mild increase of plasma potassium concentration was detected in 4-PBA treated *Umod*^{A227T} mutants compared to corresponding wild-type controls.

Most of the urinary parameters of 4-PBA treated *Umod* mutants were similarly altered as in the placebo treatment group (Table 2, Figure 1B). 4-PBA treated *Umod* mutants displayed reduced body weight, reduced urine osmolality combined with slightly increased 24-hour urine volume, increased urinary excretion and FE of calcium, reduced urinary excretion of phosphate and reduced FE_{urea} in comparison to 4-PBA treated wild-type mice. Further, daily excretions of sodium and potassium as well as FE_K were increased in *Umod* mutants of both mouse lines. Additionally, *Umod*^{C93F} mutants, which exhibited a more severe UAKD phenotype than mutants of the *Umod*^{A227T} mouse line, excreted more urea and creatinine in their 24-hour urine and had a strongly decreased FE_P when compared to 4-PBA treated wild-type controls.

1.3. *Umod* mutant mice after 4-PBA versus placebo treatment. Additional clinical-chemical plasma parameters showed trends to be altered by 4-PBA treatment of *Umod* mutants compared to their genotype-matched placebo treated counterparts (Table 1&2, Figure 1). In *Umod* mutant mice of both lines, 4-PBA treatment led to a mild but significant increase of plasma chloride levels and to a distinct decrease of the plasma concentrations of inorganic phosphate. Further, 4-PBA treated *Umod*^{C93F} mutant mice exhibited a significantly higher plasma urea concentration compared to placebo treated mice of the same genotype, indicating worsening of renal function by 4-PBA in the more severely UAKD affected mutant mouse line *Umod*^{C93F}.

In the setting of UAKD, long-term administration of 4-PBA led to significantly increased 24-hour urinary excretion of sodium in *Umod* mutants from both lines when compared to the placebo treated *Umod* mutants of the same genotype. Urine osmolality was significantly reduced and urine volume excretion was increased in 4-PBA treated vs. placebo treated *Umod*^{A227T} mutant mice. In contrast, urine osmolality of *Umod*^{C93F} mutants receiving 4-PBA was significantly increased when compared to placebo treated mice of the same genotype. However, as the UAKD phenotype is more pronounced in *Umod*^{C93F} mutants compared to *Umod*^{A227T} mutants, urine excretion volume was still distinctly higher and urine

osmolality distinctly lower in *Umod*^{C93F} mutants when compared to *Umod*^{A227T} mutants after placebo treatment, and this genotype difference was attenuated after 4-PBA treatment. Further striking alterations of kidney function of *Umod*^{C93F} mutants due to 4-PBA treatment were a significant reduction of urinary excretion of phosphate and of FE_P. Additionally, urinary excretion of potassium was increased in 4-PBA-treated *Umod*^{A227T} mutants. In 4-PBA treated *Umod*^{C93F} mutants, 24-hour urinary excretion of calcium, FE_{Ca} and FE_{Na} were increased when compared to the values of the genotype-matched placebo treated mice.

1.4. Wild-type mice after 4-PBA versus placebo treatment. In 4-PBA treated wild-type mice, plasma calcium and cholesterol levels tended to be lower than in wild-type mice after placebo treatment (Table 1, Figure 1A). Further, long-term oral administration of 4-PBA in young-adult wild-type mice resulted in a slight but significant decrease of urine osmolality and in a distinct decrease of 24-hour urinary excretion of phosphate and of FE_{urea} (Table 2, Figure 1B).

1.5. 4-PBA treated *Umod*^{C93F} versus 4-PBA treated *Umod*^{A227T} mutant mice. *Umod*^{C93F} mutants of the 4-PBA treatment group exhibited, besides the more prominent reduction of body weight also present in mice of the placebo group, significantly stronger reductions of FE_P and FE_{urea} when compared to 4-PBA treated *Umod*^{A227T} mutant mice (Table 2, Figure 1B). These results indicate the more pronounced UAKD phenotype in *Umod*^{C93F} mutants and a stronger effect of 4-PBA treatment in this line.

1.6. Progression of UAKD during 4-PBA and placebo treatment. At an age of two months, before testing the putative therapeutic effect of 4-PBA, renal dysfunction of homozygous mutants (*Umod*^{A227T} and *Umod*^{C93F}) was already clinically evident by distinctly increased plasma urea concentrations in mutants compared to age-matched wild-type controls (Table 3). With increasing age, mutant mice of both lines exhibited a continuous increase of plasma urea concentrations irrespective of treatment group (Figure 2). However, the relative increase of plasma urea levels between two and four months of age was significantly more pronounced in 4-PBA treated versus placebo treated *Umod*^{C93F} mutant mice. Of note, plasma urea concentrations of *Umod*^{C93F} mutants were distinctly higher than those of *Umod*^{A227T} mutants in

both the 4-PBA and the placebo treatment groups. This reflects the more severe UAKD phenotype in the *Umod*^{C93F} mutant mouse line, which is in line with previous results indicating that renal dysfunction is more pronounced in *Umod*^{C93F} mutant mice than in age-matched *Umod*^{A227T} mutants.

2. Effect of 4-PBA administration on urinary uromodulin excretion and TALH morphology. One characteristic of UAKD represents the reduced urinary uromodulin excretion due to the dominant negative effects of amino acid-changing *UMOD* mutations on protein maturation and secretion. As shown earlier, urinary *UMOD* excretion was strongly decreased in *Umod* mutant mice, with the extent of decrease being more pronounced in *Umod*^{C93F} mutant mice than in *Umod*^{A227T} mutant mice of the same allelic status (17,21). After 4-PBA treatment for 2 months, homozygous mutant mice had a reduced *UMOD* excretion compared to wild-type mice, and the reduction of *UMOD* excretion was more pronounced in mutants of the *Umod*^{C93F} mouse line than in the *Umod*^{A227T} line (not shown). A putative beneficial therapeutic effect through a molecular chaperone is expected to attenuate the protein maturation defect and to increase the urinary *UMOD* excretion. However, *Umod* mutant mice of both lines treated with the molecular chaperone 4-PBA had similar urinary uromodulin contents as genotype-matched mutant mice of the placebo group (Figure 3A).

A morphological characteristic of UAKD is the intracytoplasmic accumulation of uromodulin in the hyperplastic endoplasmic reticulum in TALH cells. Uromodulin accumulation in the perinuclear compartment of TALH cells was demonstrated in placebo treated *Umod* mutant mice of both lines, but also after 4-PBA treatment (Figure 3B). Further, ultrastructural alterations of TALH cells such as perinuclear stacked lamellar structures were similar in 4-PBA and placebo treated *Umod* mutant mice (not shown). In earlier analyses, these perinuclear lamellar structures were identified as hyperplastic endoplasmic reticulum where intracytoplasmic uromodulin retention and accumulation takes place (21).

3. *In vitro* assays

3.1. Effect of 4-PBA on uromodulin secretion and expression in stably transfected MTC cells. In view of published *in vitro* effects of 4-PBA (19,20), stably transfected clones of the immortalized murine kidney cell line MTC (24) expressing wild-type or

mutant *UMOD* were generated. Mature uromodulin protein was detected in supernatant of MTC cells stably transfected with wild-type, A227T mutant, or C93F mutant uromodulin expression vectors. There were differences of the uromodulin content in the supernatant between different stably transfected cell clones irrespective of the type of transfected *Umod* variant (not shown). 4-PBA administration (1 mM – 5 mM) to *UMOD* expressing MTC cells for 48 hours led to an increase of uromodulin content in supernatant compared to control containing no 4-PBA in the culture medium (Figure 4A). There was a correlation between concentration of 4-PBA and amount of uromodulin in the culture medium with higher concentrations of up to 5 mM 4-PBA resulting in a higher uromodulin content in the supernatant. However, the increased uromodulin secretion was not due to improved protein maturation efficiency but was paralleled by an increase of uromodulin expression as demonstrated by increased uromodulin content in cell lysates. This effect was observed for stably transfected MTC clones irrespective of the type of transfected *Umod* variant. The intracellular content of both mature and immature uromodulin was increased and this was particularly seen in *Umod* mutant transfected MTC cells.

3.2. Effect of 4-PBA on uromodulin expression in kidney cells of *Umod* mutant mice *in vitro* and on HSP70 expression in TALH cells *in vivo*. To further elucidate the putative effect of 4-PBA to ameliorate mutant uromodulin maturation and to resolve ER hyperplasia, primary kidney cells of *Umod* mutant mice were cultured for 48 hours in the presence of either 2 mM 4-PBA, 5 mM 4-PBA or without drug supplementation. Double-immunofluorescence analyses were performed to identify *UMOD*-positive TALH cells, to localize *UMOD* and to visualize the ER by the ER chaperone BiP (Figure 4B). Of note, prominent BiP signals localizing to the enlarged ER were present in TALH cells of *Umod* mutant mice. No obvious effect of 4-PBA on uromodulin retention and BiP signal in TALH cells of *Umod* mutant mice was observed.

Ma et al. (20) reported an impact of the heat shock protein HSP70 in amelioration of protein maturation of mutant uromodulin in transfected MDCK cells by 4-PBA *in vitro*. Therefore, we analyzed the protein abundance of HSP70 in kidneys of UAKD-affected *Umod* mutant and wild-type mice. HSP70 was predominantly expressed in collecting

duct cells of the renal papilla, but its abundance was only very weak or absent in TALH cells irrespective of genotype (Figure 5). After administration of 4-PBA for 2 months, HSP70 immunostaining pattern and intensity were similar irrespective of genotype when compared to placebo control mice.

4. Analysis of unfolded protein response pathway in TALH cells of *Umod* mutant mice.

To elucidate the mechanism of unfolded protein response (UPR) pathway in TALH cells in the setting of UAKD, protein abundances of factors of the NF- κ B pathway were analyzed. Therefore, the renal segment with predominance of TALH cells, the outer medulla, was prepared (Figure 6A). The lysates contained high amount of TALH segments which was confirmed by the more than two-fold increase of the ER protein BiP in *Umod* mutant mice of both lines compared to wild-type controls. In *Umod* mutant mice, protein abundances of p105 and p50 of NF- κ B1 were significantly increased in the outer medulla compared to that of wild-type mice (Figure 6B). A two-fold increased abundance of p100 and p52 of NF- κ B2 and an at least three-fold increased abundance of RelB was detected in UAKD affected mice compared to wild-type controls, whereas minor alterations of protein abundance of NF- κ B p65 were observed. Analyses of upstream factors of the NF- κ B pathway revealed increased abundances of phospho-IKK α/β and TRAF2 in *Umod* mutant mice of both lines compared to wild-type mice. Protein abundances of IKK α and TRAF3 were similar in all genotype groups.

To have a more detailed view on the role of NF- κ B pathway in UAKD, the abundance of phospho-IKK α/β (Ser176/180), representing the active form of IKK α and IKK β , was analyzed immunohistochemically (Figure 7A). In wild-type mice, phospho-IKK α/β was predominantly and strongly detected in the basolateral cytoplasmic compartment of collecting duct cells, but its abundance in TALH cells was low. In *Umod* mutant mice, the abundance of phospho-IKK α/β in TALH cells appeared similar as in collecting duct cells. Further, the staining pattern in TALH cells differed between *Umod* mutant mice and wild-type controls. A strong cytoplasmic basal and paranuclear staining pattern of phospho-IKK α/β was present in TALH cells of *Umod* mutant mice, while in wild-type mice the staining was weak and apical accentuated. RelB was immunohistochemically localized with variable staining intensity in the

cytoplasm and the nucleus of tubular cells (Figure 7B). Whereas only few tubular cells were detected containing RelB in wild-type mice, the majority of tubular cells of the outer medulla of *Umod* mutant mice were immunopositive for RelB and the staining intensity was stronger than in the corresponding region of wild-type mice. Identification of TALH segment was enabled by detection of UMOD.

DISCUSSION

Specific curative therapies for UAKD are not available. Symptomatic treatment addresses correction of water and electrolyte imbalances and of hyperuricemia but does not prevent disease progression. The central disease mechanism of UAKD is the protein maturation defect caused by mutant uromodulin leading to UMOD retention in the ER and ER overload. Thus, amelioration of protein maturation capacity by molecular chaperons might represent a promising therapeutic approach to reduce UMOD retention in ER of TALH cells and consequently to reduce TALH dysfunction in UAKD. *In vitro* studies demonstrated a positive effect of the chemical chaperone 4-PBA on mutant uromodulin maturation (19,20).

4-PBA is an orally bioavailable aromatic short-chain fatty acid and is used as U.S. Food and Drug Administration-approved drug (BUPHENYL[®], Ucylyd Pharma Inc.; AMMONAPS[®], Orphan Europe of Immeuble "Le Guillaumet") for clinical use as an ammonia scavenger in long-term treatment of hyperammonemia in urea cycle disorders (25,26). 4-PBA reduces ammonia levels in blood as it offers itself as an alternative to urea synthesis to excrete waste nitrogen from the body. It is converted to 2-phenylacetate and subsequently excreted with urine as nitrogen-containing phenylacetylglutamine in humans. Further, a clinical effect of 4-PBA in treatment of homozygous mutants suffering from beta thalassemia was demonstrated (27) and also a putative curative effect of 4-PBA in spinal muscular atrophy is currently under critical discussion (28,29). Various studies demonstrated the effect of the chemical chaperone 4-PBA in stabilizing protein conformation, improving ER folding capacity, facilitating the trafficking of mutant proteins, reducing ER stress, modulating the unfolded protein response and thus improving protein homeostasis in ER storage diseases (25,30,31). For instance, 4-PBA has been shown to enhance *in vitro* and *in vivo* the adaptive capacity of the ER and to improve systemic insulin action in the setting of

obesity-induced insulin resistance in type 2 diabetes (30). Further, trafficking of the cystic fibrosis transmembrane regulator with the mutation $\Delta F508$ (CFTR $\Delta F508$) was increased and secretion of the mutant $\alpha 1$ -ATZ protein in $\alpha 1$ -antitrypsin deficiency was enhanced by 4-PBA (32,33). Regarding the putative therapeutic effect of 4-PBA in the setting of UAKD, two independent *in vitro* studies demonstrated that mutant *UMOD* transfected cells exhibited an increased secretion of uromodulin in the supernatant and had an enforced immunofluorescent uromodulin staining signal on their cell membrane (19,20). In the present study, however, we were not able to demonstrate a positive effect of 4-PBA to improve mutant uromodulin maturation *in vivo* in our UAKD mouse models *Umod*^{A227T} and *Umod*^{C93F}. Intracellular accumulation of uromodulin in TALH cells was still present in *Umod* mutant mice after two months treatment with water soluble and orally bioavailable 4-PBA, and uromodulin excretion with urine remained low at a similar range as in genotype-matched placebo-treated mice. 4-PBA is rapidly metabolized to phenylacetate, and both components have short elimination half-lives of 0.76 hours and 1.29 hours in humans, respectively. Therefore, missing bioavailability of 4-PBA to TALH cells has to be taken into account as a potential explanation for the lack of a therapeutic effect in our UAKD mouse models. However, this is unlikely since administration with the drinking water ensured continuous intake of 4-PBA throughout the day. Further, effects of 4-PBA on some kidney parameters, like urine osmolality and urinary excretion of phosphate, were clearly detected. Basseri et al. (31) demonstrated an *in vivo* effect of 4-PBA on modulating unfolded protein response and attenuating ER stress in adipocytes using similar preparation, dosage, and oral administration of 4-PBA like we used in our study, but changed drinking water only weekly in contrast to half-weekly fresh preparation made in our study. To further exclude the possibility that the missing effect of 4-PBA in our *in vivo* study was due to insufficient bioavailability of the drug at the site of action, the TALH cells, 4-PBA was administered *in vitro* to culture medium of freshly isolated kidney cells of *Umod* mutant mice for 48 hours. Again, 4-PBA was ineffective to attenuate uromodulin retention and ER hyperplasia in TALH cells isolated from kidneys of *Umod* mutant mice.

To evaluate the effect of 4-PBA on secretion of murine *UMOD* A227T and *UMOD* C93F mutant protein and to repeat published experiments indicating

a positive effect of 4-PBA on secretion of human uromodulin *in vitro*, we performed assays on stably transfected murine proximal tubular epithelial (MTC) cells. MTC cells do not express endogenous uromodulin protein but provide the maturation capacity for GPI-linked secreted proteins thereby facilitating the interpretation of maturation ability of the mutant protein. We obtained various stable *Umod* transfected MTC clones for each construct (*Umod*^{wt}, *Umod*^{A227T} and *Umod*^{C93F}) secreting variable amounts of uromodulin into the supernatant. There was no relationship between the type of *Umod* mutation and maturation efficiency of stably transfected cells. There were numerous *in vitro* studies published analyzing the maturation time of uromodulin until secretion and the amount of protein secreted into the supernatant by transfected cells. A common finding of these studies was that all *UMOD* mutations analyzed needed a longer time period for protein maturation and that the efficiency of protein maturation was different between different *UMOD* mutations (3,12-14,34). However, these *in vitro* models were described to be not suitable to analyze potential disease mechanisms downstream of mutant uromodulin retention as the cellular models reported so far did not reproduce key cellular hallmarks of UAKD, which are mutant uromodulin aggregation, ER membrane expansion, and ER dilation (4). *In vitro* 4-PBA administration to supernatant was reported to increase secretion of both wild-type and mutant uromodulin, which was also associated with stronger membrane labeling of uromodulin of stably transfected cells (19,20). In our *in vitro* studies, we also detected a strong increase of uromodulin secretion of both wild-type and mutant protein after 4-PBA stimulation. But in parallel, the amount of intracellular uromodulin in MTC cells was also increased indicating an increase of uromodulin synthesis rather than changes in maturation efficiency of uromodulin. There are numerous reports of 4-PBA as a well-known histone deacetylase (HDAC) inhibitor, which increases transcription of a number of genes (25,35,36). For instance, beneficial therapeutic effects of 4-PBA due to its HDAC inhibitor function were demonstrated for motor neuron survival in mice with amyotrophic lateral sclerosis (35) or for treatment of different types of tumors (25). The therapeutic effect of 4-PBA to correct trafficking of mutant CFTR $\Delta F508$ in cystic fibrosis epithelial cells could be attributed to increased expression of some cytosolic molecular chaperones like heat shock protein 70 (HSP70) (37). In consequence, increased *UMOD* content in supernatant

as well as in cell lysate of UMOD-expressing MTC clones in our study might be caused by the HDAC inhibition due to 4-PBA administration increasing UMOD expression in a dose-dependent manner.

The *in vitro* results obtained by 4-PBA testing on UMOD transfected cells might be highly dependent on the cell type used for *in vitro* analyses. Ma et al. (20) reported a similar steady-state expression of the molecular chaperone HSP70 in mock-transfected and wild-type UMOD transfected MDCK cells, an immortalized collecting duct cell line, but a decrease of HSP70 expression in UMOD mutant transfected MDCK cells, and an increase of HSP70 expression in mutant UMOD expressing cells up to the level of wild-type UMOD expressing cells after incubation with 10 mM 4-PBA. The heat shock protein HSP70 is predominantly expressed in the kidney in papillary collecting duct cells to protect these cells from hypertonic stress present in the hypertonic environment of the renal papilla (38). HSP70 protein abundance was low or absent in TALH cells of both wild-type mice and *Umod* mutant mice indicating a minor relevance of this molecular chaperone in uromodulin protein maturation in TALH cells *in vivo*. Therefore, *in vitro* models of UMOD transfected cells might be unsuitable for testing putative therapeutic effects of drugs, as Rampoldi et al. (4) already described that these cell models are not useful to analyze the molecular intracellular pathways in UAKD.

Long-term administration of 4-PBA had, besides changes in phosphate levels (lower plasma phosphate levels and lower 24-hour phosphate excretion with urine) and mild decrease of urine osmolality, only minor effects on renal function of healthy wild-type mice compared to untreated wild-type controls. However, in the setting of UAKD, long-term treatment with 4-PBA changed further parameters of renal function, like blood chloride and phosphate levels, urine osmolality and renal handling of some electrolytes like sodium. The most prominent changes caused by 4-PBA treatment represented the strongly reduced fractional excretion of phosphate in homozygous *Umod*^{C93F} mutant mice, exhibiting a more severe uromodulin maturation defect and UAKD phenotype as compared to *Umod*^{A227T} homozygotes. Although a known function of 4-PBA is to act as nitrogen scavenger and thus to be able to decrease plasma ammonia and urea levels, plasma urea concentrations of 4-PBA treated mice were similar to those of placebo-treated genotype-matched controls or

were even increased like in homozygous *Umod*^{C93F} mutant mice. Thus, 4-PBA treatment rather resulted in a further deterioration of renal function in both mouse models of UAKD than improving it.

To elucidate further therapeutic approaches to perform curative treatment of UAKD, one open question is the analysis of the predominant stress pathways that are activated by mutant uromodulin maturation retardation and ER retention. In our mouse models of UAKD, numerous factors of the NF-κB pathway were increased abundant and/or activated and the activation of NF-κB pathway was localized in TALH cells of *Umod* mutant mice. RelB was identified as the factor with the highest increase of protein abundance in *Umod* mutant mice compared to wild-type controls. Together with the increased abundance of NF-κB2 p52 in *Umod* mutant mice, our data indicate that the non-canonical NF-κB pathway, that predominantly targets activation of the p52/RelB NF-κB complex (39,40), might be of special importance as disease mechanism in UAKD.

In conclusion, our study demonstrated that – in contrast to the reported amelioration of the maturation and trafficking defect of mutant uromodulin *in vitro* – the chemical chaperone 4-PBA was ineffective to improve mutant uromodulin maturation and TALH function in two mouse models of UAKD *in vivo*. Activation of NF-κB pathway in thick ascending limb of Henle's loop cells of UAKD mice was demonstrated, identifying a novel disease mechanism of UAKD and a potential target for therapeutic intervention.

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AUTHOR CONTRIBUTIONS

EK, BA and RW designed the experiments, EK and SS performed the experiments, EK, SS, BR,

MHdA analyzed the data, EK, EW and RW wrote the paper.

CONFLICT OF INTEREST STATEMENT

All the authors have no competing financial interest to declare.

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LEGENDS TO FIGURES

Figure 1: Effect of placebo and 4-PBA treatment on plasma and urine parameters in wild-type and *Umod* mutant mice. (A) Clinical-chemical plasma parameters. (B) Urinary parameters of kidney function. Data presented in Tukey's box plot (whiskers go 1.5 times the interquartile range or to the highest or lowest point, whichever is shorter), with outliers represented as dots. n=6-19 per genotype and treatment group. 2-way ANOVA: ^{a, b, c}, $p < 0.05$; ^{aa, bb, cc}, $p < 0.01$; ^{aaa, bbb, ccc}, $p < 0.001$; ^a, 4-PBA vs. placebo of the same genotype; ^b, homozygous mutant vs. wild-type of the same treatment group; ^c, homozygous *Umod*^{C93F} mutant vs. homozygous *Umod*^{A227T} mutant of the same treatment group.

Figure 2: Development of plasma urea levels during 4-PBA treatment. Comparison of the plasma urea concentrations of mice at the beginning of the trial (age of mice: 2 months), after one month of treatment (age of mice: 3 months) and at the end of the 2-month-duration of therapeutic testing of 4-PBA (age of mice: 4 months). Data presented in Tukey's box plot (whiskers go 1.5 times the interquartile range or to the highest or lowest point, whichever is shorter), with outliers represented as dots. n=11-19 per genotype, treatment group and age group, except of n=8 of placebo group of homozygous *Umod*^{C93F} mutants (age 2 and 3 months), n=2-3 of placebo group of 2- and 3-month-old homozygous *Umod*^{A227T} mutants and n=6 of placebo group of 4-month-old homozygous *Umod*^{A227T} mutants. The relative increases of plasma urea concentrations with age compared to that of an age of 2 months are indicated. Student's *t*-test with Holm-Bonferroni correction: ^{a, b}, $p < 0.05$; ^{aa, bb}, $p < 0.01$; ^{aaa, bbb}, $p < 0.001$; ^a, plasma urea concentration of 3-month-old vs. 2-month-old mice of the same genotype and treatment group; ^b, plasma urea concentration of 4-month-old vs. 2-month-old mice of the same genotype and treatment group.

Figure 3: Evaluation of the *in vivo* effect of 4-PBA treatment on mutant uromodulin maturation and trafficking defect. (A) On Western Blot analysis, uromodulin abundance in urine, standardized on equal creatinine, of six mice per *Umod* mutant mouse line and treatment group were analyzed and compared to the UMOD abundance of a wild-type mouse of the placebo treatment group. Urinary uromodulin protein excretion was reduced in 4-PBA and placebo treated *Umod* mutant mice compared to placebo-treated wild-type mice. Further, the uromodulin excretion with urine was much more decreased in homozygous *Umod*^{C93F} mutant mice than in homozygous mutant mice of the *Umod*^{A227T} mutant mouse line. Compared to genotype-matched mice of the placebo control group, *Umod* mutant mice of both lines receiving 4-PBA for two months excreted similar amounts of UMOD protein with urine as placebo control mice. (B) Staining patterns of uromodulin in TALH cells were similar in placebo control mice and 4-PBA treated mice of the same genotype group. TALH cells of wild-type mice exhibited a diffuse cytoplasmic uromodulin immunopositivity with enforcement of the luminal membrane. In contrast, TALH cells of *Umod* mutant mice of both lines displayed a strong paranuclear immunopositivity for UMOD irrespective of kind of treatment. Chromogen: DAB, nuclear staining: haemalum.

Figure 4: Evaluation of the *in vitro* effect of 4-PBA on mutant uromodulin expressing cells. (A) An increased concentration of administered 4-PBA to UMOD expressing MTC cells for 48 hours led to an increased uromodulin content in supernatant compared to control containing no 4-PBA in the culture medium irrespective of the type of UMOD protein (native or mutant) expressed. In parallel to the increased UMOD content in supernatant by 4-PBA administration, the abundance of UMOD protein in cell lysate also increased irrespective of the type of UMOD protein (native or mutant) expressed. Further, immature UMOD signal appeared to be stronger in the mutant protein expressing MTC cells than in the wild-type UMOD protein expressing cells and was increased after administration of higher 4-PBA concentrations. (B) 4-PBA administration (5 mM) for 48 hours on primary kidney cells of a homozygous *Umod*^{A227T} mutant mouse did not lead to obvious changes of prominent perinuclear UMOD (green color) and BiP (orange color) immunofluorescent signals in TALH cells compared to placebo control cells. Thus, no obvious effect of 4-PBA on uromodulin retention and BiP signal intensity in TALH cells of *Umod* mutant mice was seen *in vitro*.

Figure 5: Effect of 4-PBA on HSP70 protein abundance in TALH cells of *Umod* mutant mice. HSP70 protein immunopositivity was predominantly detected in collecting duct cells of the renal papilla. In the outer

medulla, the HSP70 protein was predominantly detected in collecting duct cells, but (nearly) no expression of HSP70 could be observed in uromodulin-expressing TALH cells. There were no obvious differences in HSP70 immunostaining pattern and intensity between different *Umod* genotypes or treatment groups (placebo control group vs. 4-PBA group). Chromogen: DAB for HSP70, Vector RED for UMOD; nuclear staining: haemalum. Age of mice analyzed: four months.

Figure 6: Analysis of unfolded protein response pathway in the outer medulla of *Umod* mutant mice. (A) The high density of TALH segments in the outer medulla was demonstrated by protein abundances of mature and immature UMOD and the increased abundance of BiP in *Umod* mutant mice compared to wild-type mice. (B) Protein abundances of NF- κ B1 p105/p50, NF- κ B2 p100/p52, NF- κ B p65, RelB, phospho-IKK α / β , IKK α , TRAF2 and TRAF3. Signal intensities of the factors of NF- κ B pathway were corrected for GAPDH signal intensities of the same PVDF-membrane, which was stripped several times to facilitate the detection of multiple proteins. Mean of protein abundance of wild-type mice was set on a value of 1 [mean (wild-type) = 1]. Student's *t*-test with Holm-Bonferroni correction: *p* vs. wild-type, *, *p* < 0.05; **, *p* < 0.01; ***, *p* < 0.001.

Figure 7: Immunohistochemical localization of factors of NF- κ B pathway in TALH cells of *Umod* mutant mice. (A) Phospho-IKK α / β was predominantly detected in the basolateral cytoplasmic compartment of collecting duct cells. In TALH cells of wild-type mice, a weak diffuse cytoplasmic immunopositivity for phospho-IKK α / β was observed. In contrast, TALH cells of *Umod* mutant mice exhibited a strong signal of phospho-IKK α / β in the cytoplasmic basal and paranuclear compartment which signal intensity was in a similar range as that of collecting duct cells. Chromogen: DAB for phospho-IKK α / β , Vector RED for UMOD; nuclear staining: haemalum. Age of mice analyzed: four months. CD: collecting duct profiles. Wild-type: *Umod*^{wt} mouse; UAKD: homozygous *Umod*^{A227T} mutant mouse. (B) In wild-type mice, RelB was detected in few tubular cells in the cytoplasm with a variable but predominantly weak staining intensity and in some nuclei. In contrast to wild-type mice, tubular cells of *Umod* mutant mice exhibited a stronger cytoplasmic and nuclear RelB staining intensity and more tubular cells were stained. UMOD immunohistochemistry enabled identification of TALH segments. Serial kidney sections were used for RelB and UMOD immunohistochemistry and corresponding kidney regions are shown. DAB; nuclear staining: haemalum. Age of mice analyzed: four months. Wild-type: *Umod*^{wt} mouse; UAKD: homozygous *Umod*^{C93F} mutant mouse.

Table 1: Plasma data at an age of 4 months of placebo mice or after 2 months of 4-PBA administration.

	<i>Wild-type</i>		<i>Umod^{A227T} homozygote</i>		<i>Umod^{C93F} homozygote</i>	
	<i>Placebo</i>	<i>4-PBA</i>	<i>Placebo</i>	<i>4-PBA</i>	<i>Placebo</i>	<i>4-PBA</i>
Na ⁺ (mmol/l)	155.9 ± 3.1	154.3 ± 3.6	151.7 ± 6.6 ^b	156.4 ± 2.3 ^{aa, b}	155.5 ± 4.3 ^c	156.1 ± 1.9
K ⁺ (mmol/l)	4.07 ± 0.29	3.94 ± 0.19	4.15 ± 0.40	4.21 ± 0.32 ^{bb}	4.22 ± 0.59	3.98 ± 0.26
Cl ⁻ (mmol/l)	114.6 ± 1.9	113.5 ± 2.8	111.8 ± 4.1 ^b	114.5 ± 1.5 ^a	113.9 ± 3.7	116.2 ± 1.1 ^{a, bb}
Ca ²⁺ (mmol/l)	2.30 ± 0.07	2.21 ± 0.08 ^a	2.32 ± 0.15	2.34 ± 0.09 ^{bbb}	2.37 ± 0.07 ^{bbb}	2.36 ± 0.11 ^{bbb}
P _i (mmol/l)	2.39 ± 0.44	2.11 ± 0.31	2.64 ± 0.70	1.94 ± 0.36 ^{aaa}	2.30 ± 0.56	1.92 ± 0.26 ^a
Total protein (g/l)	51.1 ± 1.8	51.3 ± 2.7	49.6 ± 3.8	52.3 ± 1.7 ^a	52.1 ± 2.4 ^c	52.6 ± 1.7
Creatinine (μmol/l)	9.00 ± 1.92	8.00 ± 1.32	9.86 ± 0.96	9.93 ± 1.30 ^{bbb}	9.46 ± 1.83	9.45 ± 1.11 ^b
Urea (mmol/l)	10.15 ± 1.16	11.02 ± 1.29	20.54 ± 2.36 ^{bbb}	20.49 ± 2.21 ^{bbb}	23.08 ± 2.53 ^{bbb, c}	25.99 ± 2.95 ^{aaa, bbb, ccc}
Cholesterol (mmol/l)	3.56 ± 0.42	3.20 ± 0.42 ^a	3.41 ± 0.16	3.31 ± 0.24	3.76 ± 0.25 ^{cc}	3.45 ± 0.25 ^a
Triglycerides (mmol/l)	1.72 ± 0.87	1.67 ± 0.60	1.04 ± 0.27 ^b	1.25 ± 0.51 ^b	0.80 ± 0.24 ^{bbb}	0.93 ± 0.23 ^{bb, c}
ALAT (U/l)	32.1 ± 14.2	38.7 ± 24.0	29.3 ± 8.7	43.1 ± 44.8	28.1 ± 6.4	31.4 ± 5.2
ASAT (U/l)	56.1 ± 14.5	68.7 ± 48.9	48.5 ± 5.4	72.3 ± 55.2	65.8 ± 29.5	58.3 ± 8.1
Lipase (U/l)	45.6 ± 7.5	43.9 ± 7.8	54.2 ± 5.5 ^{bb}	49.8 ± 8.2 ^b	56.1 ± 6.0 ^{bbb}	55.9 ± 4.7 ^{bbb, c}

Age of mice analyzed: 4 months. Data are means ± SD. n=6-19 per genotype. P_i, inorganic phosphate; ALAT, alanine aminotransferase (EC 2.6.1.2); ASAT, aspartate aminotransferase (EC 2.6.1.1). 2-way ANOVA: ^{a, b, c}, $p < 0.05$; ^{aa, bb, cc}, $p < 0.01$; ^{aaa, bbb, ccc}, $p < 0.001$; ^a, 4-PBA vs. placebo of the same genotype; ^b, homozygous mutant vs. wild-type of the same treatment group; ^c, homozygous *Umod^{C93F}* mutant vs. homozygous *Umod^{A227T}* mutant of the same treatment group.

Table 2: Urine parameters at an age of 4 months of placebo mice or after 2 months of 4-PBA administration.

	<i>Wild-type</i>		<i>Umod</i> ^{A227T} <i>homozygote</i>		<i>Umod</i> ^{C93F} <i>homozygote</i>	
	<i>Placebo</i>	<i>4-PBA</i>	<i>Placebo</i>	<i>4-PBA</i>	<i>Placebo</i>	<i>4-PBA</i>
Body weight (g)	29.7 ± 2.0	30.7 ± 2.7	26.9 ± 1.2 ^{bb}	26.3 ± 1.5 ^{bbb}	24.2 ± 2.1 ^{bbb, ccc}	22.4 ± 0.7 ^{bbb, ccc}
Urine volume (ml/day)	1.2 ± 0.3	1.4 ± 0.3	1.5 ± 0.6	2.3 ± 0.5 ^{aa, bbb}	2.9 ± 0.7 ^{bbb, cc}	2.7 ± 1.0 ^{bb}
Urine osmolality (mOsm/kg H ₂ O)	3667 ± 576	3175 ± 370 ^{aa}	2919 ± 494 ^{bb}	2350 ± 390 ^{aa, bbb}	1667 ± 327 ^{bbb, ccc}	2073 ± 437 ^{a, bbb}
Na ⁺ (μmol/day)	286.3 ± 61.7	337.4 ± 84.1	289.4 ± 53.2	400.3 ± 87.8 ^{aa, b}	313.1 ± 53.5	424.6 ± 94.9 ^{aaa, bb}
K ⁺ (μmol/day)	495.3 ± 92.3	535.8 ± 121.8	508.0 ± 147.5	659.7 ± 141.6 ^{a, bb}	652.1 ± 84.4 ^{bb, c}	704.9 ± 161.1 ^{bbb}
Cl ⁻ (μmol/day)	508.8 ± 86.3	486.6 ± 114.5	453.4 ± 84.8	512.6 ± 114.3	485.0 ± 62.7	509.2 ± 98.9
Ca ²⁺ (μmol/day)	2.6 ± 0.3	3.0 ± 1.1	7.8 ± 1.5 ^{bbb}	9.5 ± 3.4 ^{bbb}	7.6 ± 2.3 ^{bbb}	10.3 ± 3.4 ^{aa, bbb}
Mg ²⁺ (μmol/day)	42.5 ± 7.8	39.7 ± 22.1	40.4 ± 14.6	38.1 ± 20.0	46.6 ± 15.4	48.7 ± 18.6
P _i (μmol/day)	160.3 ± 33.2	104.5 ± 55.5 ^{aa}	95.8 ± 64.1 ^b	58.4 ± 43.5 ^{bb}	108.4 ± 59.5 ^b	20.7 ± 31.2 ^{aaa, bbb}
Creatinine (μmol/day)	6.0 ± 1.2	5.9 ± 2.0	5.9 ± 0.8	6.8 ± 1.8	6.5 ± 0.8	7.3 ± 1.9 ^b
Urea (mmol/day)	2.7 ± 0.3	2.6 ± 0.6	2.6 ± 0.5	2.9 ± 0.6	2.9 ± 0.4	3.1 ± 0.5 ^b
Glucose (μmol/day)	3.1 ± 0.4	3.5 ± 1.1	2.3 ± 0.4	3.6 ± 1.9	3.3 ± 1.6	3.3 ± 2.8
Creatinine clearance	867.4 ± 457.1	932.8 ± 342.3	653.2 ± 113.5	725.5 ± 206.7	688.2 ± 166.8	696.7 ± 178.4
FE _{Na}	0.29 ± 0.12	0.31 ± 0.08	0.31 ± 0.03	0.40 ± 0.12 ^b	0.29 ± 0.06	0.36 ± 0.08 ^a
FE _K	18.6 ± 6.1	19.3 ± 4.7	20.3 ± 5.1	24.3 ± 6.9 ^b	22.9 ± 5.8	23.5 ± 5.2 ^b
FE _{Cl}	0.71 ± 0.26	0.61 ± 0.16	0.67 ± 0.08	0.68 ± 0.16	0.62 ± 0.14	0.58 ± 0.12
FE _{Ca}	0.18 ± 0.06	0.19 ± 0.05	0.56 ± 0.10 ^{bbb}	0.63 ± 0.26 ^{bbb}	0.46 ± 0.14 ^{bbb}	0.59 ± 0.22 ^{a, bbb}
FE _{Pi}	10.86 ± 3.56	7.57 ± 5.23	6.07 ± 3.88 ^b	5.29 ± 4.75	7.12 ± 4.34 ^b	1.42 ± 1.99 ^{aa, bbb, c}
FE _{urea}	40.9 ± 12.9	34.1 ± 8.3 ^a	21.7 ± 5.4 ^{bbb}	21.9 ± 4.6 ^{bbb}	18.5 ± 3.5 ^{cbbb}	16.2 ± 4.3 ^{bbb, cc}

Age of mice analyzed: 4 months. Data are means ± SD, standardized to 25 g body weight. n=6-19 per genotype. 2-way ANOVA: ^{a, b, c}, $p < 0.05$; ^{aa, bb, cc}, $p < 0.01$; ^{aaa, bbb, ccc}, $p < 0.001$; ^a, 4-PBA vs. placebo of the same genotype; ^b, homozygous mutant vs. wild-type of the same treatment group; ^c, homozygous *Umod*^{C93F} mutant vs. homozygous *Umod*^{A227T} mutant of the same treatment group.

Table 3: Plasma data of 2-month-old mice.

	<i>Wild-type</i>	<i>Umod</i> ^{A227T} <i>homozygote</i>	<i>Umod</i> ^{C93F} <i>homozygote</i>
Na ⁺ (mmol/l)	148.5 ± 3.8	148.5 ± 3.0	148.5 ± 4.5
K ⁺ (mmol/l)	3.93 ± 0.34	4.21 ± 0.29 ^{bb}	4.00 ± 0.33
Ca ²⁺ (mmol/l)	2.34 ± 0.07	2.34 ± 0.07	2.36 ± 0.11
Cl ⁻ (mmol/l)	109.8 ± 2.2	110.2 ± 1.9	110.8 ± 3.7
P _i (mmol/l)	2.58 ± 0.36	2.57 ± 0.36	2.33 ± 0.29 ^b
Total protein (g/l)	49.8 ± 2.6	48.3 ± 1.9 ^b	49.1 ± 2.3
Creatinine (μmol/l)	8.57 ± 1.59	9.46 ± 1.61	8.64 ± 1.53
Urea (mmol/l)	9.67 ± 1.17	14.0 ± 1.9 ^{bbb}	16.1 ± 2.6 ^{bbb, cc}
Cholesterol (mmol/l)	3.40 ± 0.28	3.25 ± 0.24	3.21 ± 0.20 ^b
Triglycerides (mmol/l)	1.31 ± 0.53	1.24 ± 0.48	1.07 ± 0.33
ALAT (U/l)	28.2 ± 10.0	30.4 ± 16.1	34.4 ± 10.0 ^b
ASAT (U/l)	41.2 ± 4.9	41.8 ± 5.1	46.3 ± 6.0 ^{bb, c}
Lipase (U/l)	52.8 ± 10.2	51.9 ± 11.0	54.3 ± 8.6

Data are means ± SD. n=19-33 animals per genotype. P_i, inorganic phosphate; ALAT, alanine aminotransferase (EC 2.6.1.2); ASAT, aspartate aminotransferase (EC 2.6.1.1). Student's *t*-test with Holm-Bonferroni correction: ^b, ^c, *p* < 0.05; ^{bb, cc}, *p* < 0.01; ^{bbb, ccc}, *p* < 0.001; ^b, homozygous mutant vs. wild-type; ^c, homozygous *Umod*^{C93F} mutant vs. homozygous *Umod*^{A227T} mutant.

Figure 1

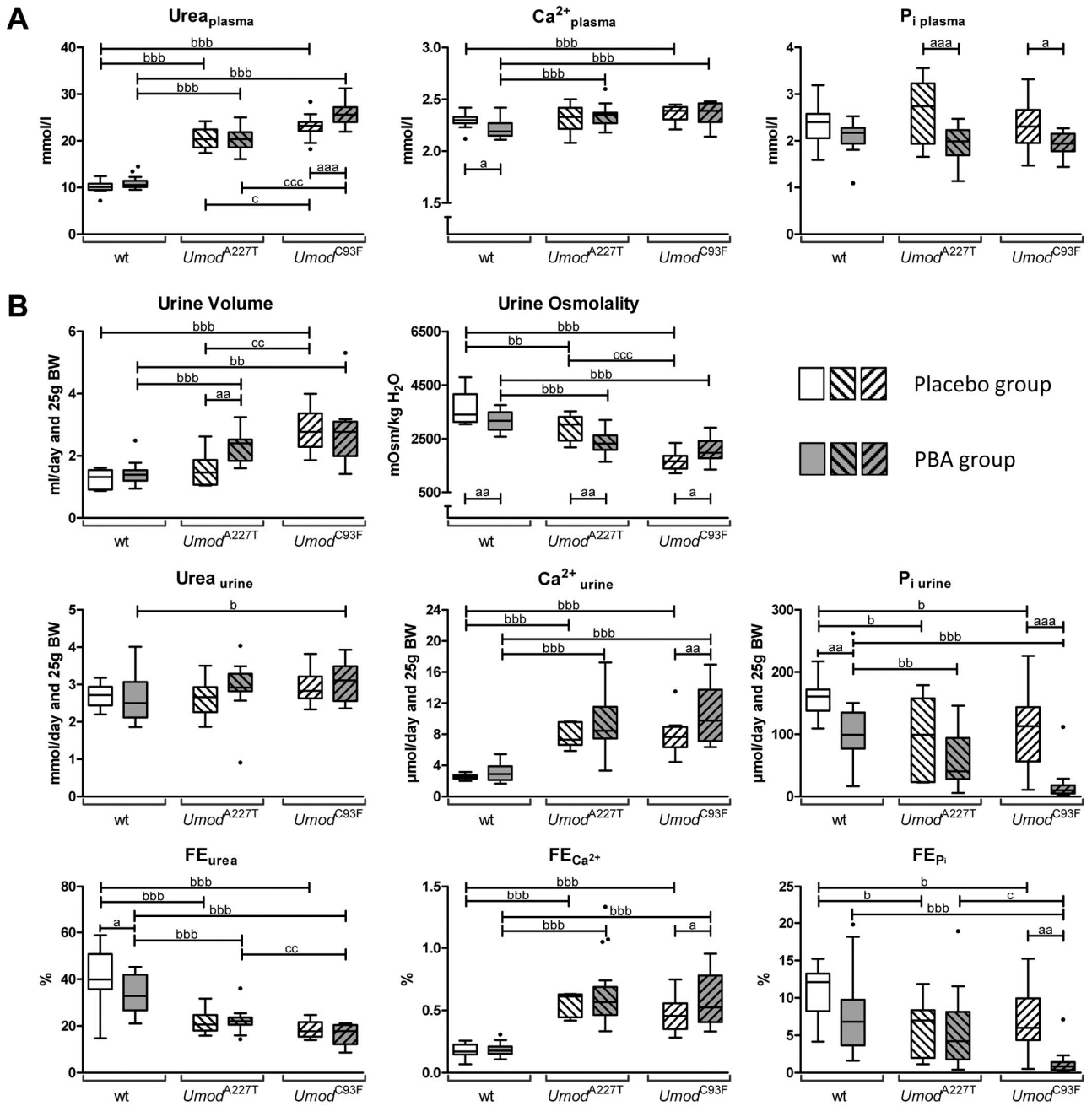


Figure 2

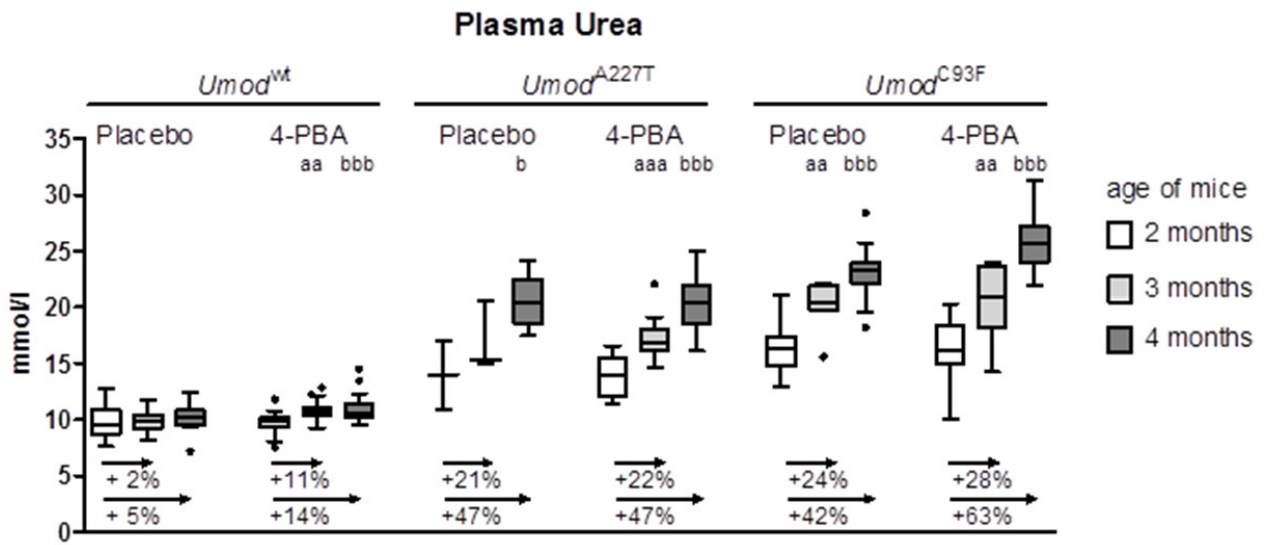


Figure 3

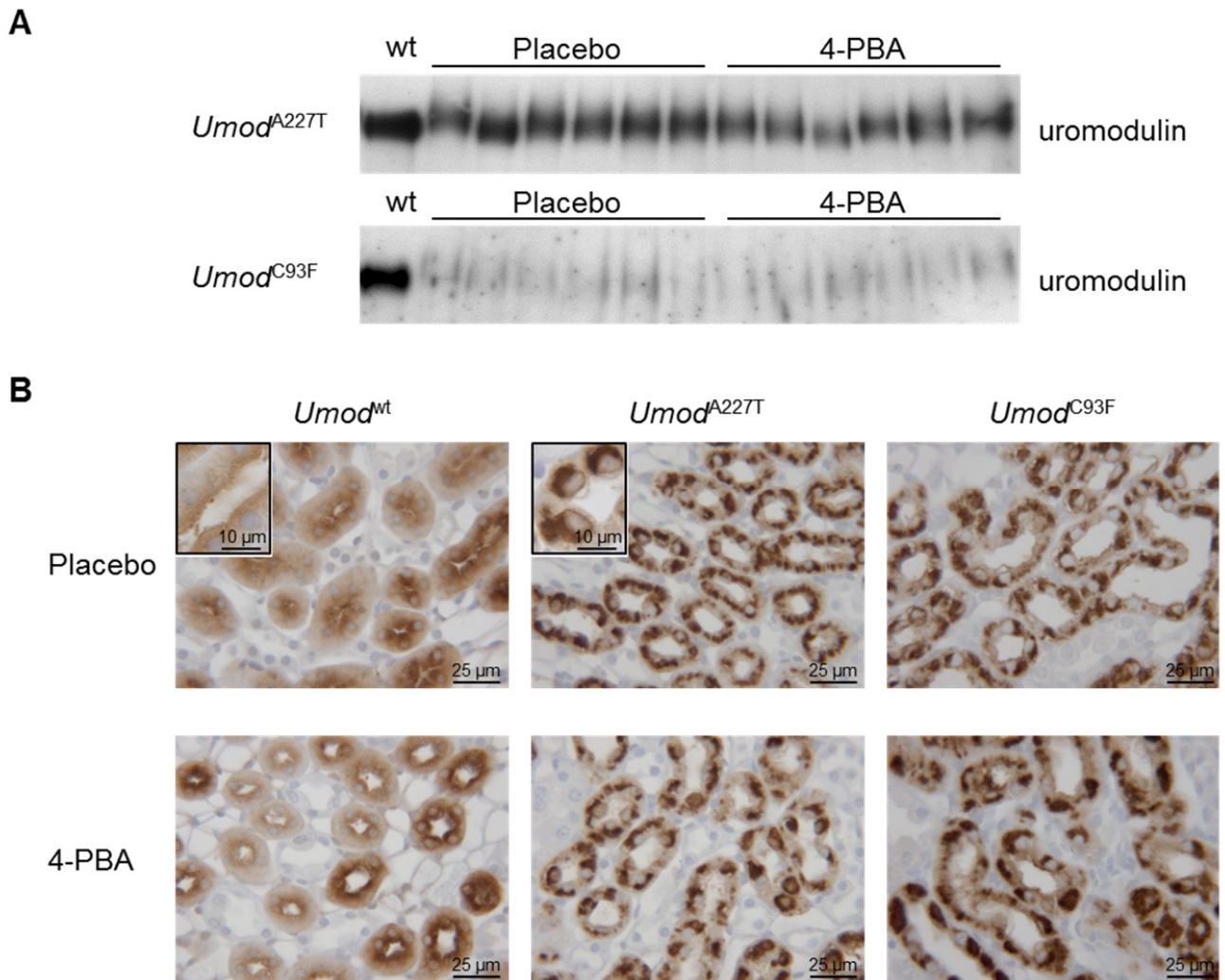
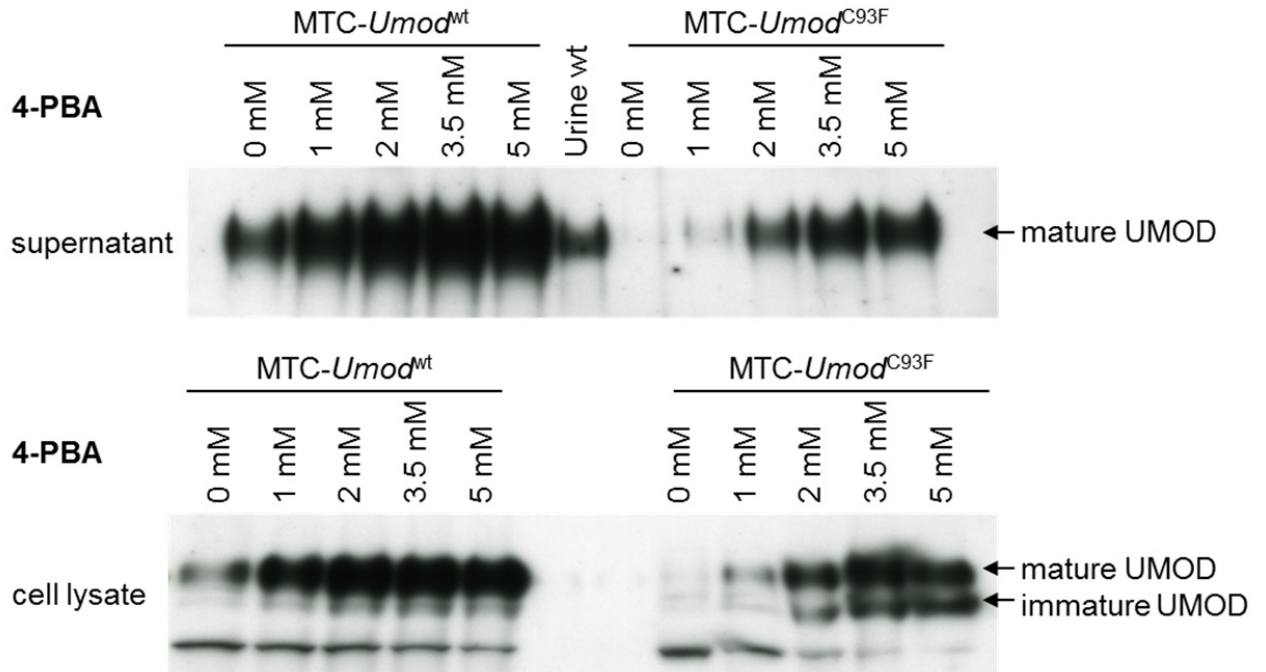


Figure 4

A Stable transfected MTC cells



B Primary kidney cells of a homozygous *Umod*^{A227T} mutant mouse

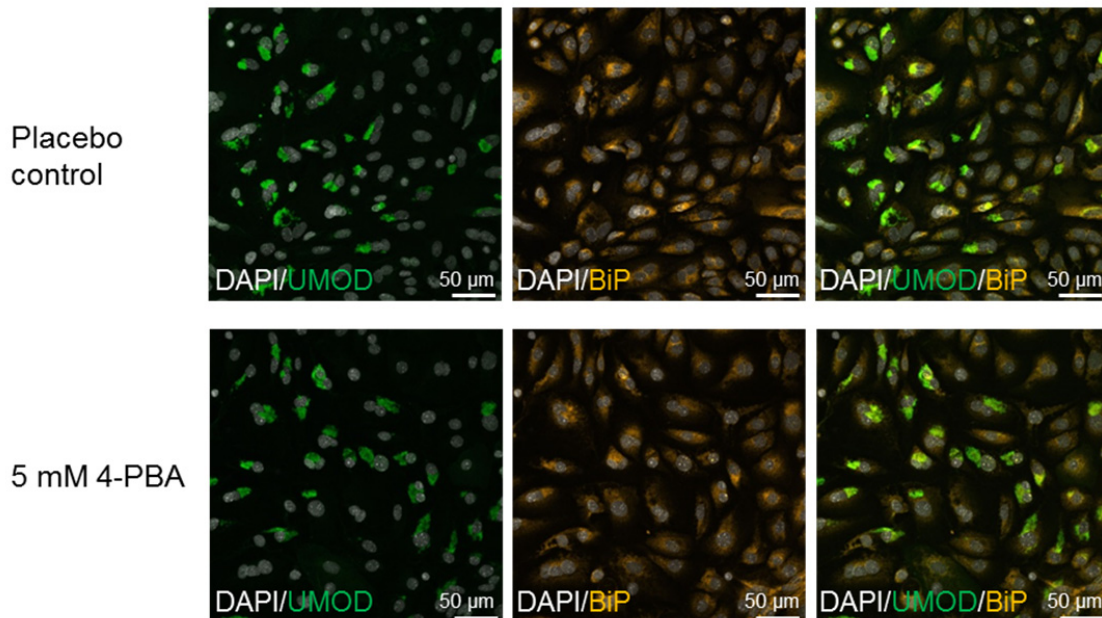


Figure 5

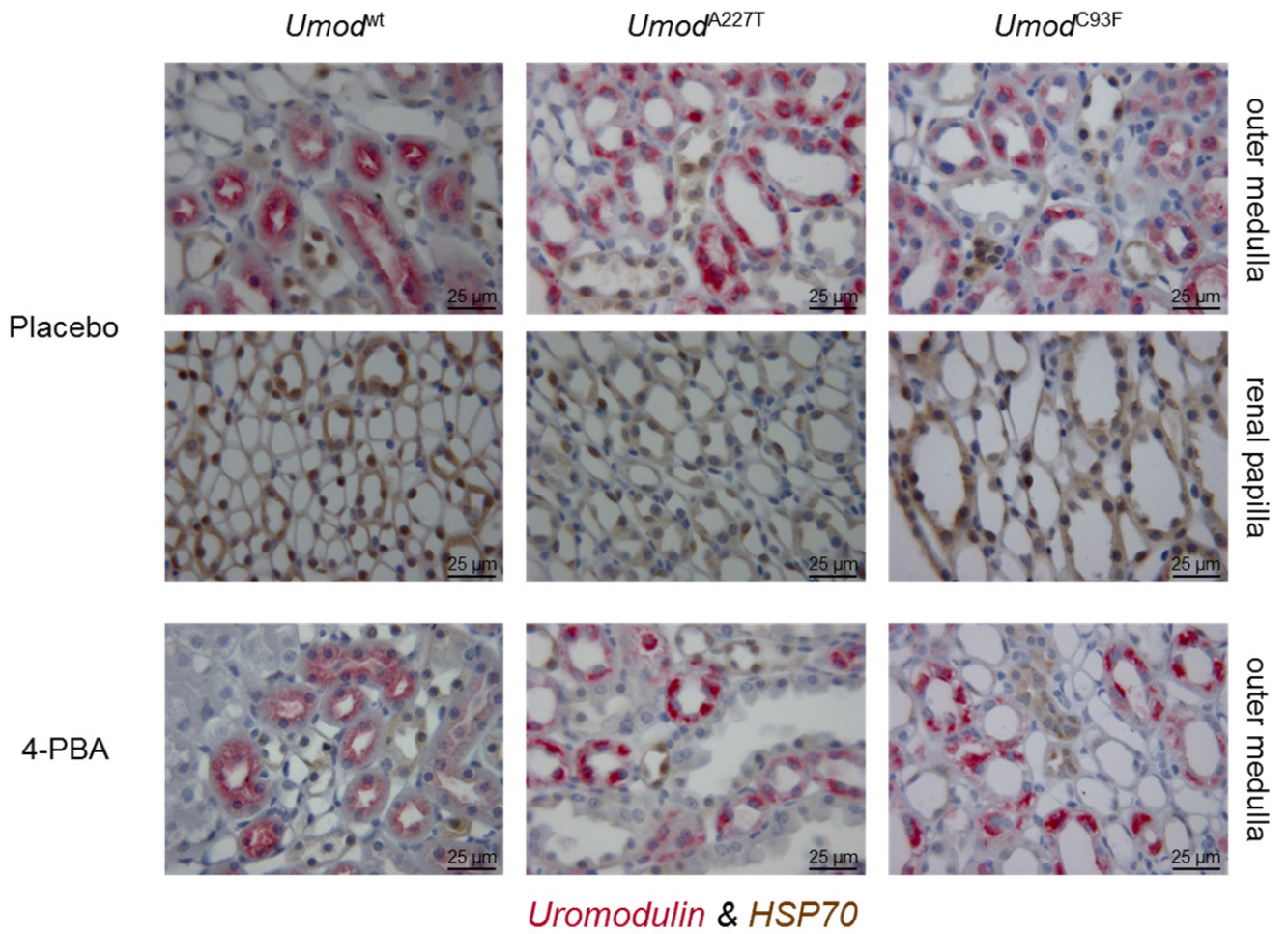
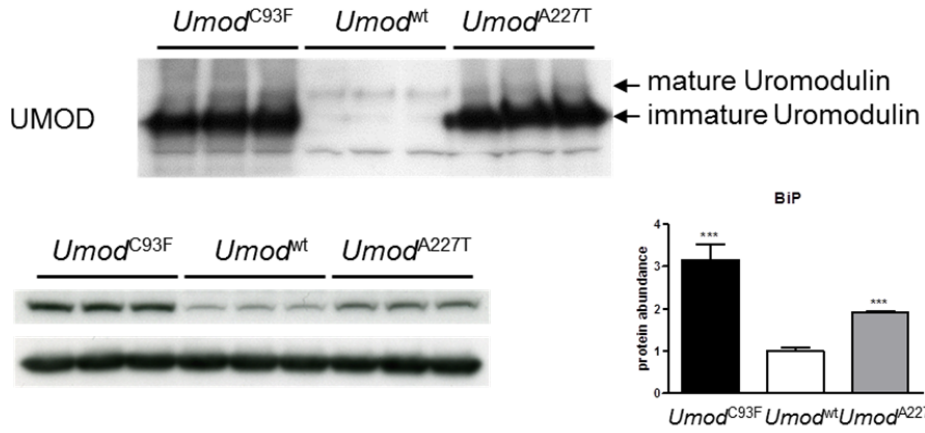


Figure 6

A



B

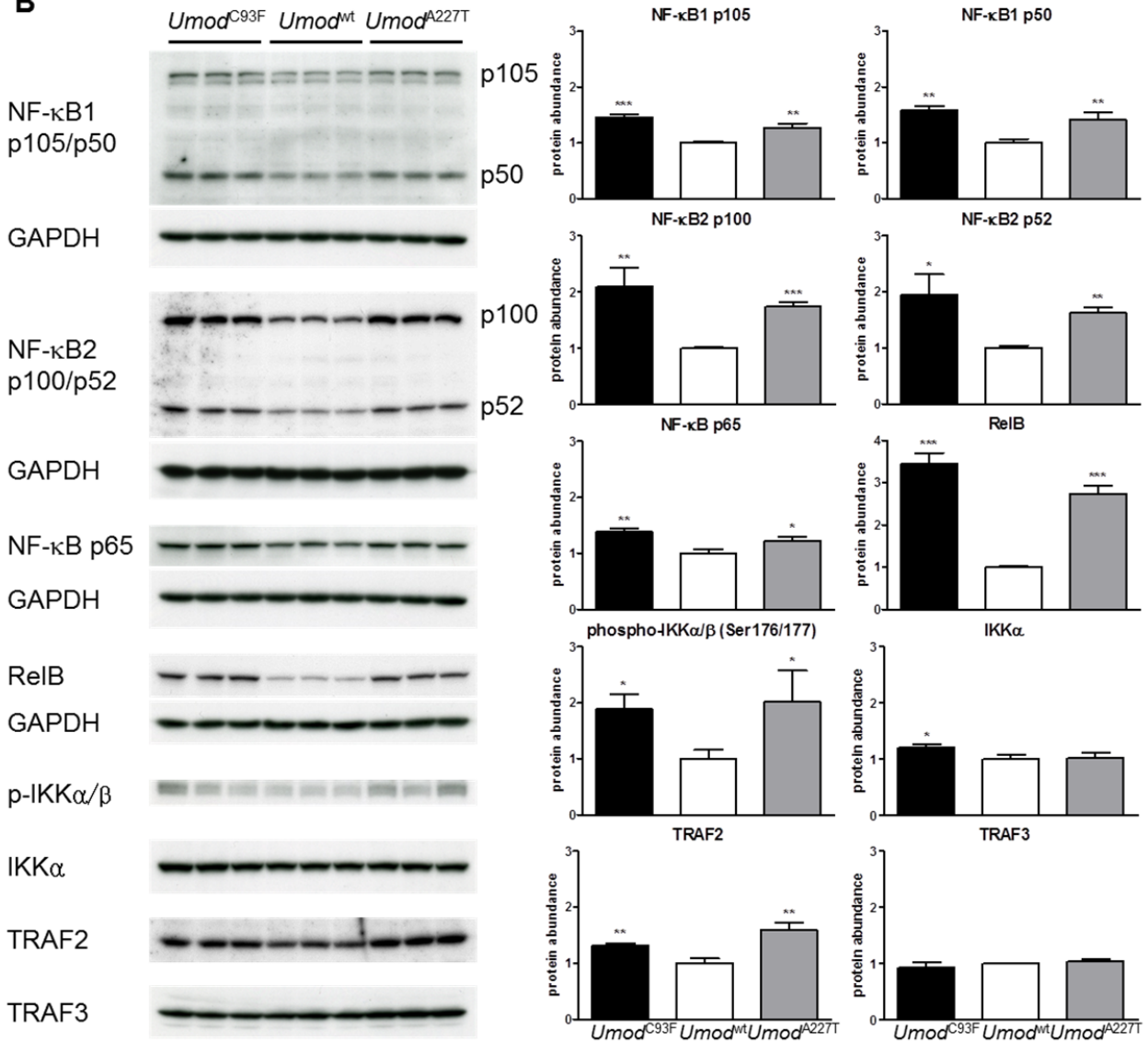
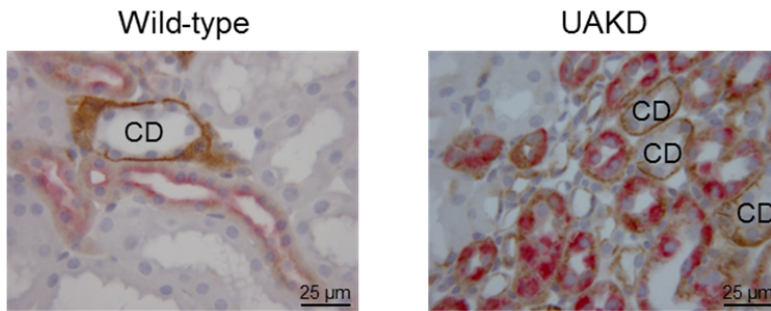


Figure 7

A



Uromodulin & phospho-IKK α/β

B

