

## The Bench-to-Bedside Transition

# Paradoxical response to furosemide in uromodulin-associated kidney disease

Laura Labriola<sup>1,\*</sup>, Eric Olinger<sup>2,\*</sup>, Hendrica Belge<sup>2</sup>, Yves Pirson<sup>1</sup>, Karin Dahan<sup>3</sup> and Olivier Devuyst<sup>1,2</sup>

<sup>1</sup>Cliniques Universitaires Saint-Luc, Université Catholique de Louvain, Brussels, Belgium, <sup>2</sup>Institute of Physiology, University of Zurich, Zurich, Switzerland and <sup>3</sup>Institute of Pathology and Genetics, Gosselies, Belgium

Correspondence and offprint requests to: Olivier Devuyst ; E-mail: olivier.devuyst@uzh.ch

\*These authors contributed equally to this study.

### ABSTRACT

Mutations in the *UMOD* gene coding for uromodulin cause autosomal dominant tubulointerstitial kidney disease. Uromodulin is known to regulate transport processes in the thick ascending limb, but it remains unknown whether *UMOD* mutations are associated with functional tubular alterations in the early phase of the disease. The responses to furosemide and to a water deprivation test were compared in a 32-year-old female patient carrying the pathogenic *UMOD* mutation p.C217G and her unaffected 31-year-old sister. A single dose of furosemide induced an intense headache with exaggerated decrease in blood pressure ( $\Delta$ syst: 30 versus 20 mmHg;  $\Delta$ diast: 18 versus 5 mmHg) and body weight ( $\Delta$ 2.6 kg versus  $\Delta$ 0.9 kg over 3 h) in the proband versus unaffected sib. The diuretic response and the fall in urine osmolality were also more important and detected earlier in the affected sib. Water deprivation led to increased plasma osmolality and urine concentration in both siblings; however, the response to desmopressin was attenuated in the affected sib. These data reveal that mutations of uromodulin cause specific transport alterations, including exaggerated response to furosemide and a failure to maximally concentrate urine, in the early phase of the disease.

**Keywords:** autosomal dominant tubulointerstitial kidney disease, furosemide, NKCC2, TAL, Tamm–Horsfall protein

### INTRODUCTION

Uromodulin (Tamm–Horsfall protein) is exclusively produced in the epithelial cells lining the thick ascending limb (TAL) of Henle's loop and is the most abundant protein in

the normal urine. Evidence obtained from *in vitro* studies and knock-out mouse studies have shown that uromodulin protects against urinary tract infection and kidney stones, and regulates transport systems operating in the TAL [1]. The latter include the sodium–potassium–chloride cotransporter NKCC2 and the potassium channel ROMK, which operate in parallel and are essential to reabsorb 25% of the filtered NaCl and to mediate the urinary concentrating ability [2].

Mutations in the *UMOD* gene encoding uromodulin have been shown to cause 'medullary cystic kidney disease type 2' (MCKD2; MIM 603 860), 'familial juvenile hyperuricaemic nephropathy' (FJHN; MIM 162 000) and 'glomerulocystic kidney disease' (GCKD; MIM 609 886) [3–5] which are collectively referred to as uromodulin-associated kidney diseases (UAKD). UAKD are autosomal dominant disorders characterized by hyperuricaemia and gout early in life, alteration of urinary concentrating ability and tubulointerstitial fibrosis with occasional cysts at the cortico-medullary junction [1]. UAKD invariably lead to chronic renal failure during the third to seventh decade of life [6]. There is no specific therapy to slow renal disease progression, and most patients end up with end-stage renal disease.

The fact that hyperuricaemia is the most consistent feature observed in patients harbouring *UMOD* mutations has led to suggestions that the disease causes a dysfunction of the TAL, with NaCl loss and secondary reabsorption of uric acid in the proximal tubule [6, 7]. This hypothesis has been supported by functional defects evidenced in the first transgenic mouse model of disease [8]. However, the existence of a specific dysfunction of the TAL as an early consequence of defective uromodulin processing has not been evaluated in patients harbouring a pathogenic *UMOD* mutation.

To investigate whether a pathogenic *UMOD* mutation is associated with functional alterations in the TAL in the early stage of the disease, we compared the response to a furosemide test and to a water deprivation test in a proband carrying a pathogenic mutation of *UMOD* and her unaffected sister.

## MATERIALS AND METHODS

The proband, a 32-year-old female harbouring a missense (p.C217G) mutation in *UMOD* (III,2), and her 31-year-old sister tested negative for the mutation (III,4) belong to a four-generation family. The pathogenic nature of the p.C217G mutation of *UMOD* was evidenced by the typical course of UAKD observed in the other sister of the proband (III,3) who reached end-stage renal disease at age 37 years (Figure 1A). The proband and her unaffected sister had an estimated glomerular filtration rate (eGFR) according to the CKD-EPI equation from the Chronic Kidney Disease - Epidemiology Collaboration  $\geq 60$  mL/min/1.73 m<sup>2</sup> at time of testing, and none of them was diabetic. They were not taking any drugs, including diuretics, allopurinol or antihypertensive medications. The protocol was approved by the Ethical Committee of the UCL Medical School (Brussels), and informed consent was provided.

A 24-h urine collection was obtained in both sisters immediately before the furosemide test. On the day of the furosemide test, a baseline clinical and biological evaluation was performed at 9:30 AM, 4 h before giving a single oral dose of 80 mg furosemide. Baseline values for the furosemide test (Figure 1B) correspond to a 3-h urine collection preceding immediately the administration of furosemide. After furosemide administration, urine samples were collected hourly during 3 h and analysed for Na<sup>+</sup> and K<sup>+</sup>, creatinine, urea and uric acid concentrations, osmolality and pH. Clinical and biological evaluations were repeated at the end of the test (time +3 h; Table 1). The following day (i.e. after 18 h of normal hydration), both individuals underwent an 8-h water deprivation test, including a 30-min intravenous infusion of desmopressin acetate (Minirin®, 0.3 µg/kg; Ferring AG, Baar, Switzerland) after 6 h of water deprivation. Desmopressin (or deamino-8D-arginine vasopressin, dDAVP) is a peptidic analogue of endogenous antidiuretic hormone (ADH) with a higher affinity for V2 receptors and negligible vasopressive potential. A clinical evaluation was performed hourly and plasma and urine samples were taken every 2 h during the water deprivation and 1-h and 2-h after initiation of desmopressin perfusion. Dietary intake was identical in the two sisters before the furosemide test and throughout the water deprivation test.

## RESULTS

The baseline osmolar balance in the two sisters immediately before the functional tests was similar, thus allowing direct inter-individual comparison of the responses to furosemide and to water restriction (values for affected sib *versus* control sib: 24-h Na<sup>+</sup> excretion: 228 *versus* 307 mmol; 24-h urea excretion:

29.1 *versus* 23.1 g; osmolar excretion rate: 779.2 *versus* 787.5 µosm/min; plasma renin activity: 1.1 *versus* 0.8 ng/mL/h; plasma aldosterone: 0.31 *versus* 0.30 nM, respectively. Plasma ADH: <0.2 pg/mL in both sibs).

The furosemide test was poorly tolerated by the proband, who reported intense headache, a severe drop in blood pressure ( $\Delta 30/\Delta 18$  mmHg *versus*  $\Delta 20/\Delta 5$  mmHg in the unaffected sister) and a weight loss of 2.6 kg (*versus* weight loss of 0.9 kg in the unaffected sister) in 3 h (Table 1). After furosemide administration, both siblings showed the expected increase in diuresis with concomitant fall in urine osmolality. However, the changes were more important and were detected earlier in the proband, who also showed a more pronounced and earlier increase of fractional excretion of Na<sup>+</sup> than the control sibling (Figure 1B).

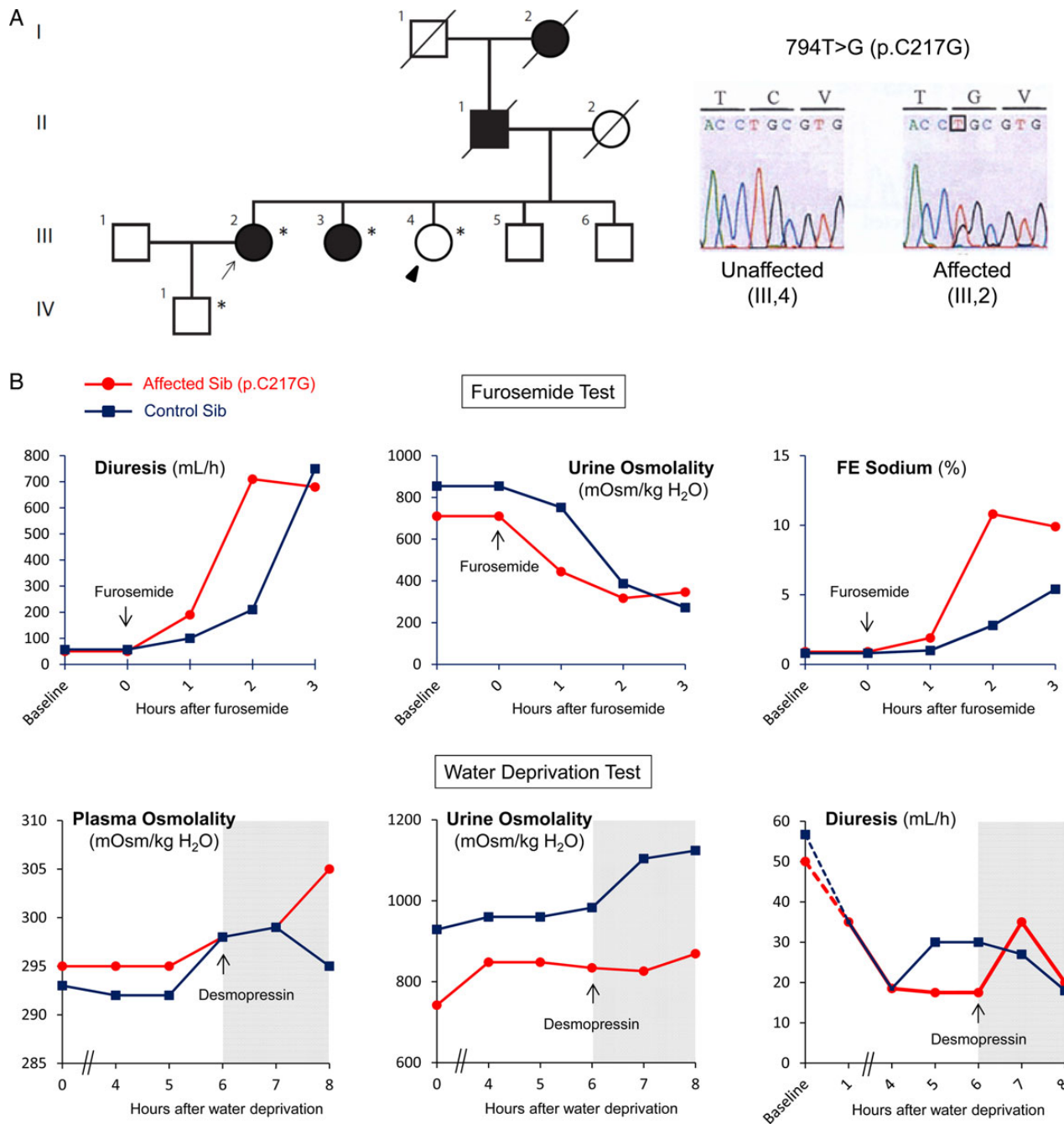
The water deprivation test was well tolerated by both participants. Blood pressure and weight remained constant during all the procedures. As expected, water deprivation resulted in a progressive increase in plasma osmolality with an increased urine osmolality and a reduced urine flow in both subjects (Figure 1B). The response to desmopressin was normal in the control sibling, with an increase in urine osmolality (from 983 to 1124 mOsm/kg H<sub>2</sub>O, 2 h after infusion), a decrease in urine flow (from 30 to 18 mL/h) and a decrease in plasma osmolality (from 298 to 295 mOsm/kg H<sub>2</sub>O). In contrast, the affected sib showed a blunted increase in urine osmolality (from 834 to 869 mOsm/kg H<sub>2</sub>O), an increase in urine flow (from 17.5 to 20 mL/h) and a paradoxical increase in plasma osmolality (from 298 to 305 mOsm/kg H<sub>2</sub>O) (Figure 1B).

Pathology examination of the end-stage kidney biopsy of patient III,3 (Figure 1C) revealed marked tubulointerstitial lesions, with thickening of the tubular basement membrane and abnormal processing and accumulation of uromodulin in the tubular cells. Of note, there was no detectable immunostaining for NKCC2 in the biopsy.

## DISCUSSION

In this study, we performed a sib-pair functional testing to evaluate whether pathogenic mutations of *UMOD* are associated with dysfunction of the TAL in the early phase of UAKD. Our data reveal that the proband carrying an *UMOD* mutation shows a clinically and biologically exaggerated response to furosemide and a failure to maximally concentrate urine after desmopressin administration. To the best of our knowledge, this is the first report of abnormal tubular functional testing in a subject harbouring a pathogenic *UMOD* mutation.

Several lines of evidence suggest that uromodulin plays an important role in regulating NKCC2 and ROMK in the TAL [9–11]. These two transport processes mediate the furosemide-sensitive NaCl reabsorption and the generation of the osmotic gradient that drives the urine concentrating ability. Patients with *UMOD* mutations are characterized by an early defect in urine concentration [3–5], which could be paralleled with a discrete NaCl-losing phenotype explaining hyperuricaemia [6, 7]. Studies of mice harbouring a pathogenic mutation of uromodulin revealed that the urinary concentrating



**FIGURE 1:** Pedigree of the family, *UMOD* mutation, response to furosemide and water deprivation and end-stage kidney biopsy. **A**, Pedigree of the family and *UMOD* mutation. Individuals with a history of renal disease are depicted by black symbols. Circles denote females, squares males. The proband carrying the *UMOD* mutation (III,2) and her unaffected sister (III,4) are marked by an arrow and an arrowhead, respectively. Individuals tested for mutation are tagged by an asterisk (\*). Patient III,3 showed a typical course of uromodulin-associated kidney disease, with hyperuricaemia at age 21 years and end-stage renal disease at age 37 years. Sequence analysis of the *UMOD* gene revealed a thymine to guanine transition at nucleotide 794 resulting in the substitution of a well-conserved cysteine by a glycine at position 217 (p.C217G). Encoded amino acid sequence is indicated above the DNA sequence. Mutated nucleotide is boxed. The mutation has been previously reported by Dahan *et al.* [4]. **B**, Biological parameters at baseline and evaluation after furosemide administration and water deprivation with desmopressin perfusion. The patient harbouring the *UMOD* mutation is indicated by red dots and her unaffected sister by blue squares. Baseline values in the furosemide test have been obtained from a 3-h urine collection immediately preceding furosemide administration. This same baseline diuresis has been plotted in the water deprivation test curve. Urine and plasma osmolality values at time 0 of the water deprivation test were obtained from urine and blood samples collected before the test. **C**, End-stage kidney biopsy from patient III,3 harbouring the p.C217G *UMOD* mutation. Haematoxylin–eosin staining (panel a) reveals interstitial fibrosis and tubular atrophy with thickening of the tubular basement membrane (inset, arrows). Immunostaining (panel b) reveals diffuse intracellular accumulation of uromodulin in a subset of cells lining enlarged tubules (arrow) as well as cells displaying the normal apical staining for uromodulin (arrowhead). There is no detectable immunostaining for NKCC2 (panel c) in the kidney of patient III,3 compared with normal staining pattern (inset). Original magnifications: a,  $\times 350$ ; b,  $\times 200$ , c,  $\times 700$ .

C

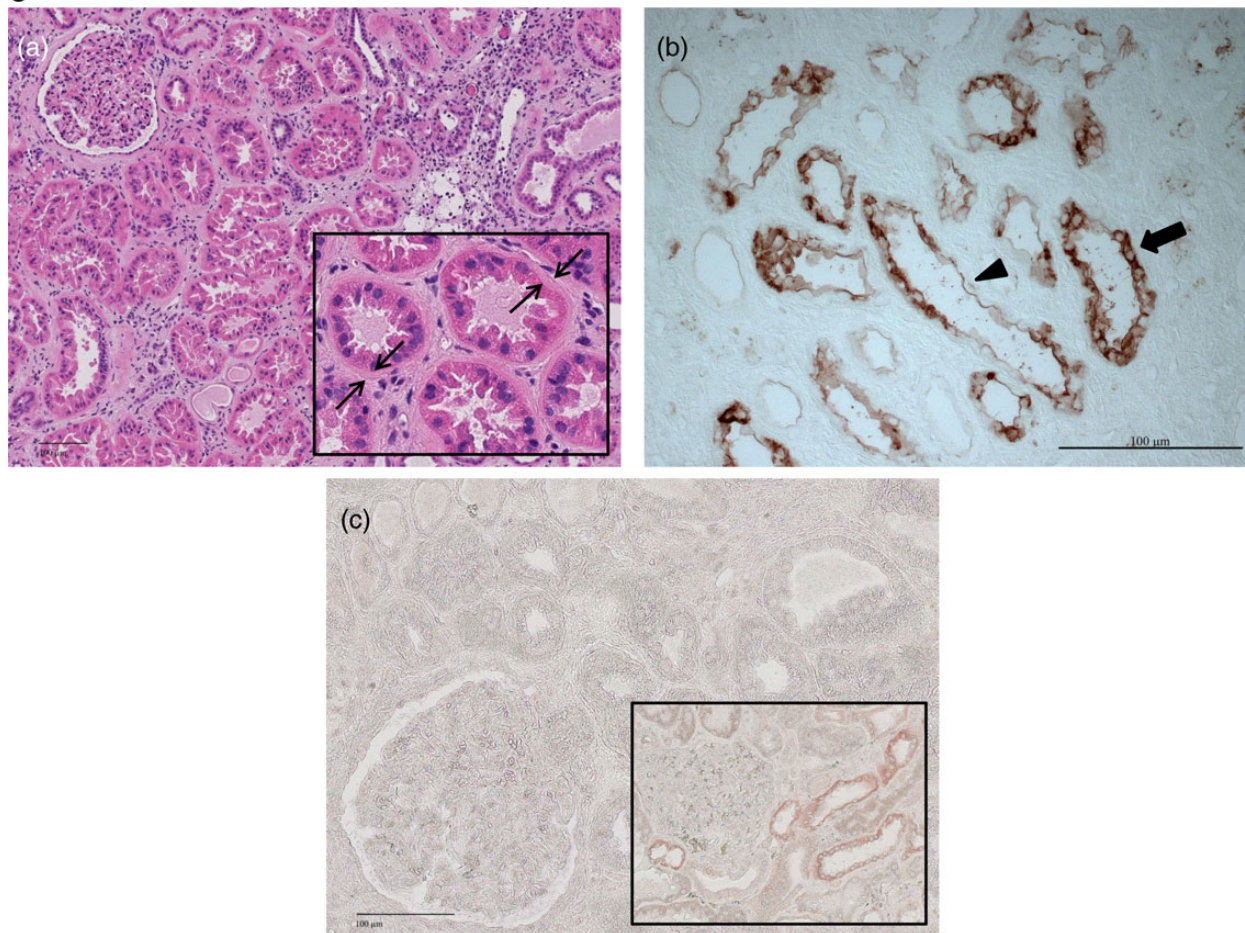


FIGURE 1: *Continued*

defect precedes renal failure and is related to a specific defect in the TAL [8]. The latter includes defective expression of NKCC2 and other markers, secondary to the accumulation of mutant uromodulin in the endoplasmic reticulum [8]. We confirm such lesions in the end-stage kidney biopsy of patient III,3, showing extensive tubulointerstitial damage with abnormal processing of uromodulin and loss of NKCC2 immunoreactivity.

The demonstration of an exaggerated response to furosemide in the proband, despite the lower eGFR than her control sister, is of particular interest. Since uromodulin expression levels regulate NKCC2 activity [10, 11], it is conceivable that the reduced trafficking of native uromodulin in UAKD might lead to a decreased amount of active NKCC2 on the apical cell membrane. This effect would be amplified by the fact that disease-causing *UMOD* mutations also lead to decreased surface expression of ROMK [9]. The paradoxically increased response to furosemide, as observed here, might therefore indicate that there is still sufficient NaCl uptake capacity (e.g. due to hyperactivated residual NKCC2) to maintain a steady-state in the early stage of disease. Administration of furosemide may disrupt this equilibrium, by inhibiting active NKCC2 transporters with no immediately available compensation in other tubule segments. The normal serum uric acid levels

(Table 1) and the slightly higher uric acid fractional excretion in the proband (5.9 versus 5.1% in the healthy sib calculated from the baseline 3-h urine collection) may indeed suggest a maintained sodium balance by residual TAL function without proximal compensatory adaptation.

Alternatively, the exaggerated response to furosemide in the proband with early UAKD could unmask a shift of steady-state NaCl reabsorption inside the TAL. Based on functional and anatomical differences, the TAL segment could indeed be subdivided into a cortical (cTAL) and a medullary (mTAL) portion. The cTAL, which lies in an environment isosmotic to the plasma, is responsible for the majority of NaCl reabsorption and is the true ‘diluting segment’ of the nephron [12, 13]. In the rat kidney, two different cell types coexist in the TAL: cells with a smooth surface and a dense subapical NKCC2-containing vesicle pool, which predominate in the mTAL, and cells with a rough surface and a much less abundant vesicle system, which largely dominate in the cTAL [14]. Although human *UMOD* transcript levels have been found to be quite similar in mTAL and cTAL [15], human uromodulin protein levels were 4-fold greater in the TAL-enriched outer medulla than in the cortex by immunoblotting [16]. In healthy individuals, steady-state NaCl reabsorption is probably achieved by the early part of cTAL, leaving a ‘physiological reserve’ in the

**Table 1. Clinical and biological response to furosemide in the proband and the unaffected sibling**

	Proband (III,2)		Unaffected sib (III,4)	
	Baseline	Furosemide + 3 h	Baseline	Furosemide + 3 h
Weight (kg)	76.1	73.5	81.1	80.2
Blood pressure (mmHg)	140/90	110/72	135/80	115/75
eGFR (CKD-EPI; mL/min/1.73 m <sup>2</sup> )	60	60	99	99
Plasma urea (mg/dL)	45	43	25	23
Plasma sodium (mmol/L)	141	138	139	138
Plasma potassium (mmol/L)	4.4	3.7	3.8	3.8
Plasma uric acid (mg/dL)	6.3	6.3	6.4	6.4
Plasma osmolality (mOsm/L)	298	298	287	288
Plasma bicarbonate (mmol/L)	27.5	29.5	22.5	27.5

The parameters were similarly recorded 4 h before (Baseline) and 3 h after furosemide administration (Furosemide + 3 h).

late part of the cTAL [12]. In the proband with early UAKD, a reduced function of NKCC2 in mTAL could be compensated by a larger implication of the cTAL, possibly spared from the transport alterations associated with UAKD. Thus, a shift of NaCl reabsorption to downstream/late part of the cTAL may have used up the ‘physiological reserve’ in the affected sib, which could then contribute to the differential response to furosemide. The fact that, in contrast to the control sib, the proband was unable to lower its urinary sodium concentration below plasma sodium (data not shown) might further indicate that the diluting function of the TAL was completely abolished by furosemide. Taken together, these data indicate that the response to furosemide in UAKD patients is diphasic, with an exaggerated response in early disease and an anticipated loss of response as predicted by the loss of NKCC2 in end-stage kidney. Loop diuretics in these patients should thus be managed with great caution.

At baseline, the proband showed a trend for increased plasma osmolality contrasting with lower urine osmolality, compatible with a slight alteration of water homeostasis. The observation of a preserved response to water deprivation followed by a blunted response to desmopressin may again indicate a reduced ‘physiological reserve’ of the urinary concentrating ability in early UAKD. The normal response of the proband to water deprivation is distinct from the severe nephrogenic diabetes insipidus observed in the transgenic mutant mouse model [8]. The difference is probably due to the fact that the tubular and interstitial lesions observed in the patient are less severe than in the transgenic mice analysed at a late stage of the disease. Additionally, the water restriction test challenges mostly the inner medulla with its ADH-sensitive urea transporters in the terminal collecting duct, where uromodulin expression is absent [13]. Possible explanations for the blunted response to desmopressin include (i) a failure to increase or maintain interstitial medullary hyperosmolality and/or (ii) a decreased response to desmopressin. A less active osmolar gradient generator, namely NKCC2 transporter, or a leaky TAL allowing water influx into the interstitium or NaCl backleak into the lumen could underlie the former hypothesis. Considering the interactions of uromodulin with NKCC2 and the physical properties (including water impermeability) of uromodulin polymers formed under specific ionic conditions [1], an altered interstitial osmolality could be very well

explained by a reduced excretion of uromodulin. In fact, the *Umod* KO mice showed no increase in pNKCC2 after stimulation with dDAVP, suggesting that uromodulin is important for the TAL sensitivity to vasopressin [10].

The main limitations of this study include the small number of siblings involved and the lack of histopathology data in the early stage of disease. However, we feel that these observations give insights into the biology of uromodulin and the pathophysiology of UAKD. We investigated a patient with pre-clinical disease, with presumably limited interstitial inflammation/fibrosis (normal urinary sediment), and an unaffected sibling of the same gender and similar age. The deleterious effect of the mutation was evidenced by the end-stage kidney biopsy of a third sibling.

In conclusion, our study suggests that, in the early phase of UAKD, the lack of functional uromodulin leads to a discrete dysfunction of the TAL, with maintenance of a precarious clinical equilibrium that can be disturbed by specific testing. As the disease progresses, the aggravation of the tubulointerstitial lesions leads to overt tubular dysfunction and compensatory mechanisms, culminating with chronic kidney disease.

## ACKNOWLEDGEMENTS

We acknowledge Mrs Y. Cnops, Prof. J-P. Cosyns and Mrs N. Van Oost for their help and the reviewers for their helpful suggestions and comments.

## FUNDING

These studies were supported by the European Community’s 7th Framework Programme (FP7/2007–2013) under grant agreement no 246539 and 608847 (IKPP Marie Curie) and grant no 305608 (EURenOmics); Action de Recherche Concertée (ARC10/15-029, Communauté Française de Belgique); the FNRS and FRSM; Inter-University Attraction Pole (IUAP, Belgium Federal Government); supported by the Fonds National de la Recherche, Luxembourg (6903109); the NCCR Kidney.CH program (Swiss National Science Foundation); the Gebert Rūf Stiftung (Project GRS-038/12); and the Swiss National Science Foundation 310030–146490.

## CONFLICT OF INTEREST STATEMENT

None declared.

(See related article by Zacchia and Capasso. The importance of uromodulin as regulator of salt reabsorption along the thick ascending limb. *Nephrol Dial Transplant* 2015; 30: 158–160.)

## REFERENCES

1. Rampoldi L, Scolari F, Amoroso A *et al*. The rediscovery of uromodulin (Tamm-Horsfall protein): from tubulointerstitial nephropathy to chronic kidney disease. *Kidney Int* 2011; 80: 338–347
2. Devuyst O. Salt wasting and blood pressure. *Nat Genet* 2008; 40: 495–496
3. Hart TC, Gorry MC, Hart PS *et al*. Mutations of the *UMOD* gene are responsible for medullary cystic kidney disease 2. *J Med Genet* 2002; 39: 882–892
4. Dahan K, Devuyst O, Smaers M *et al*. A cluster of mutations in the *UMOD* gene causes Familial Juvenile Hyperuricemic Nephropathy with abnormal expression of uromodulin. *J Am Soc Nephrol* 2003; 14: 2883–2893
5. Rampoldi L, Caridi G, Santon D *et al*. Allelism of MCKD, FJHN and GCKD caused by impairment of uromodulin export dynamics. *Hum Mol Genet* 2003; 12: 3369–3384
6. Bollée G, Dahan K, Flamant M *et al*. Phenotype and outcome in hereditary tubulointerstitial nephritis secondary to *UMOD* mutations. *Clin J Am Soc Nephrol* 2011; 6: 2429–2438
7. Scolari F, Caridi G, Rampoldi L *et al*. Uromodulin storage diseases: clinical aspects and mechanisms. *Am J Kidney Dis* 2004; 44: 987–999
8. Bernascone I, Janas S, Ikehata M *et al*. A transgenic mouse model for uromodulin-associated kidney diseases shows specific tubulo-interstitial damage, urinary concentrating defect and renal failure. *Hum Mol Genet* 2010; 19: 2998–3010
9. Renigunta A, Renigunta V, Saritas T *et al*. Tamm-Horsfall glycoprotein interacts with renal outer medullary potassium channel ROMK2 and regulates its function. *J Biol Chem* 2010; 286: 2224–2235
10. Mutig K, Kahl T, Saritas T *et al*. Activation of the bumetanide-sensitive Na<sup>+</sup>, K<sup>+</sup>, 2Cl<sup>-</sup>-cotransporter (NKCC2) is facilitated by Tamm-Horsfall protein in a chloride-sensitive manner. *J Biol Chem* 2011; 286: 30200–30210
11. Trudu M, Janas S, Lanzani C *et al*. Common noncoding *UMOD* gene variants induce salt-sensitive hypertension and kidney damage by increasing uromodulin expression. *Nat Med* 2013; 19: 1655–1660
12. Burg M. Thick ascending limb of Henle's loop. *Kidney Int* 1982; 22: 454–464
13. Bankir L, De Rouffignac C. Urinary concentrating ability: insights from comparative anatomy. *Am J Physiol* 1985; 249: 643–666
14. Nielsen S, Maunsbach AB, Ecelbarger CA *et al*. Ultrastructural localization of Na-K-2Cl cotransporter in thick ascending limb and macula densa of rat kidney. *Am J Physiol* 1998; 275: 885–893
15. Chabardès-Garonne D, Mejéan A, Aude JC *et al*. A panoramic view of gene expression in the human kidney. *Proc Natl Acad Sci USA* 2003; 100: 13710–13715
16. Serafini-Cessi F, Malagolini N, Cavallone D. Tamm-Horsfall glycoprotein: biology and clinical relevance. *Am J Kidney Dis* 2003; 42: 658–676

Received for publication: 4.9.2014; Accepted in revised form: 22.11.2014