CLINICAL STUDY

Altered urinary excretion of aquaporin 2 in IgA nephropathy

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Abstract

Objective: The intrarenal renin–angiotensin system (RAS) activation plays a pivotal role in immunoglobulin A nephropathy (IgAN) pathogenesis, which is still largely undefined. Recently, vasopressin (AVP) has been advocated to contribute to the genesis and progression of chronic kidney diseases (CKD) directly, and indirectly, via RAS activation. Our aim is to explore the intrarenal activity of AVP, its relationship with RAS activity, as well as its modulation by therapies in IgAN.

Design: In this observational study, we measured plasma copeptin, a surrogate marker of AVP, the urine excretion of aquaporin 2 (AQP2), a protein reflecting renal AVP action, and angiotensinogen (AGT), a parameter of renal RAS activation, and their relationship with renal function in 44 IgAN patients at the time of renal biopsy, without any drug therapy, and after 6-month treatment with ACEi or steroid ACEi.

Twenty-one patients with other CKD and 40 healthy subjects were recruited as controls.

Methods: ELISAs were used to measure all variables of interest.

Results: At baseline, IgAN patients showed higher urinary levels of AQP2, compared with controls and patients with other CKD. Urinary AQP2 and AGT levels strongly correlated with the presence of arterial hypertension. Steroids–ACEi caused the decrease of all the variables examined. The fall of urinary AQP2 and AGT following drug treatments was associated with the decrease of daily proteinuria.

Conclusion: Our findings would support the involvement of AVP–AQP2 axis, interacting with the RAS, in the progression of IgAN and candidate AQP2 as a possible novel marker of the disease.

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Introduction

Immunoglobulin A nephropathy (IgAN) is now recognised to be the most common form of glomerulonephritis worldwide. Although the mechanisms underlying the pathogenesis and development of IgA nephropathy remain largely undefined, it is gradually being clarified that intrarenal renin–angiotensin system (RAS) activation plays a pivotal role in the pathogenesis and development of the disease (1, 2). Accordingly, RAS blockade appears to limit proteinuria and reduce glomerular filtration rate decline more effectively than other antihypertensive treatments in IgAN patients (3, 4). Full remission of the disease, however, is seldom achieved, especially when pharmacological intervention is started late, and supports the adoption of more complex strategies than with an isolated pharmacological intervention on the RAS.

In addition to the RAS, an accumulating body of evidence suggests that arginin–vasopressin (AVP) may play a role in the genesis and exacerbation of renal damage and chronic renal insufficiency (5). Particularly, Bankir (6) argued that the activation of AVP specific G protein-coupled V2 renal receptors, by regulating aquaporin 2 (AQP2) expression and subsequent insertion into the luminal membrane of collecting duct principal cells, increases urea recycling from the collecting duct into the loop of Henle; reduces sodium concentration at the macula densa; inhibits tubuloglomerular feedback; stimulates renin release and could in this way result in glomerular hyperfiltration, proteinuria and renal damage (6, 7).

Then, several findings suggest the existence of a crosstalk between AVP and the RAS. Angiotensin II (Ang II) potently induces AQP2 protein expression via AVP V2 and Ang II AT(1) receptors in the mouse renal collecting duct principal cells (8), while a sustained stimulation of AVP receptors induces intrarenal RAS activation (5, 6). Conversely, RAS blockade decreases AQP2 expression (8) and blunts AVP-induced rise in proteinuria in experimental models (7).
Presently, no data are available concerning the activity of AVP in IgAN patients. This study was undertaken to explore the intrarenal activity of AVP, its relationship with RAS activity, as well as its modulation by currently available therapies for patients with IgAN. To this aim, we measured the urinary excretion of AQP2 (uAQP2), a shadow protein reflecting renal AVP action (9), and angiotensinogen (uAGT), a newly proposed parameter of renal RAS activation, at the time of histological diagnosis, and following two different therapeutic approaches, currently representing the most adopted first-line treatments: ACEi alone or steroids in combination with ACEi (3, 4).

Subjects and methods

Patients

Between January 2005 and December 2009, we selected 80 consecutive IgAN patients biopsied at the Section of Nephrology, University of Foggia. Entry criteria were: age > 18 years, grades I–IV lesions (Lee’s classification (10)), absence of nephrotic syndrome, concomitant systemic disease and urological abnormalities and/or infection and plasma creatinine levels of ≤ 2.0 mg/dl. Thirty-six patients, taking RAS blockers at the time of renal biopsy, were excluded, thus 44 IgAN patients were included for this study (Table 1). All patients received ACEi, titrated at the maximally tolerated dose (5–10 mg ramipril). Nineteen subjects with moderate to severe daily proteinuria (1.0–8.4 g/die) and grades III and IV histological lesions were assigned to steroids and received 0.5 g methylprednisolone i.v. for three consecutive days at the beginning of the steroid course and again 2 and 4 months later; they were also given oral prednisone at a daily dose of 0.5 mg/kg for 6 months. The remaining 25 patients received ACEi only. Hypertension was defined as blood pressure (BP) of more than 140/90 mmHg in two repeated measurements in a standing position, or the need for antihypertensive agents. All patients in both groups were targeted to achieve BP values ≤ 130/80 mmHg by ACEi, and calcium channel blockers if needed. All patients were studied before and after the designed therapy. None of the patients took

Table 1 Demographic and laboratory features of healthy controls, IgAN patients (before and after 6-month treatment), and patients with chronic kidney diseases other than IgAN. Data are expressed as mean ± s.d.

<table>
<thead>
<tr>
<th></th>
<th>Healthy controls (n=40)</th>
<th>CKD (n=21)</th>
<th>All (n=44)</th>
<th>ACEi (n=25)</th>
<th>Steroids + ACEi (n=19)</th>
</tr>
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<tbody>
<tr>
<td></td>
<td></td>
<td>Baseline</td>
<td>6 months</td>
<td>Baseline</td>
<td>6 months</td>
</tr>
<tr>
<td></td>
<td></td>
<td>Baseline</td>
<td></td>
<td>Baseline</td>
<td></td>
</tr>
<tr>
<td>Age (years)</td>
<td>38 ± 14</td>
<td>37 ± 18</td>
<td>40 ± 14</td>
<td>39 ± 13</td>
<td>45 ± 14</td>
</tr>
<tr>
<td>Serum creatinine (mg/dl)</td>
<td>0.8 ± 0.2</td>
<td>0.90 ± 0.32</td>
<td>1.03 ± 0.36</td>
<td>0.90 ± 0.27</td>
<td>0.91 ± 0.21</td>
</tr>
<tr>
<td>Proteinuria (g/day)</td>
<td>2.49 ± 2.93</td>
<td>1.59 ± 1.62</td>
<td>1.07 ± 0.91</td>
<td>1.07 ± 0.59</td>
<td>2.39 ± 2.15†</td>
</tr>
<tr>
<td>IgAN Histology grading</td>
<td>NA</td>
<td>1</td>
<td>–</td>
<td>–</td>
<td></td>
</tr>
<tr>
<td>G1</td>
<td></td>
<td>1</td>
<td>–</td>
<td>–</td>
<td></td>
</tr>
<tr>
<td>G2</td>
<td></td>
<td>9</td>
<td>–</td>
<td>–</td>
<td></td>
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<tr>
<td>G3</td>
<td></td>
<td>12</td>
<td>–</td>
<td>10</td>
<td></td>
</tr>
<tr>
<td>G4</td>
<td></td>
<td>3</td>
<td>–</td>
<td>9</td>
<td></td>
</tr>
<tr>
<td>MDRD GFR (ml/min per 1.73 m²)</td>
<td>102.0 ± 38.2</td>
<td>78.7 ± 25.9</td>
<td>88.9 ± 22.9</td>
<td>94.0 ± 22.2</td>
<td>66.1 ± 24.4²</td>
</tr>
<tr>
<td>Hypertension (yes/not)</td>
<td>–</td>
<td>4/17</td>
<td>21/23</td>
<td>8/17</td>
<td>13/6</td>
</tr>
</tbody>
</table>
| *P<0.05, †P<0.01, ‡P<0.005: steroids + ACEi versus ACEi alone (Mann–Whitney U test) and §P=0.05 versus CKD.

Figure 1 (A) Urinary levels of aquaporin 2 (AQP2), (B) plasma levels of copeptin and (C) urinary levels of angiotensinogen (AGT) in healthy individuals (CTRL), and in patients with IgAN measured at the time of renal biopsy, in the absence of any drug therapy. The boxes extend from the first quartile to the third quartile and the whiskers extend to the 10th and 90th percentiles of the data. Individual values beyond the 10th and 90th percentiles are also shown. The central horizontal line within each box represents the 50th percentile (median) (Mann–Whitney U test).
diuretics throughout the study period. We further enrolled 21 patients with chronic kidney disease (CKD) other than IgAN (seven with membranous glomerular nephropathy, eight with focal segmental glomerular sclerosis and six with benign nephroangiosclerosis) (Table 1). All patients were investigated at the time of histological diagnosis. Forty healthy controls without any personal or family history of renal disease and arterial hypertension were enrolled as well. The study was approved by the local ethics committee, and a written informed consent was obtained from all participants. All investigations were conducted according to the principles expressed in the Declaration of Helsinki.

Sample collection

Blood and first-void urine were collected early in the morning, after an overnight fast, all subjects being on their usual diet, with free access to fluids.

Urine samples were first tested for standard parameters using Multistix reactive stripes (Bayer Diagnostics), then centrifuged and filtered, after the addition of protease inhibitors cocktail (P8340, Sigma–Aldrich).

Blood samples were collected in BD Vacutainer Rapid Serum tubes and centrifuged.

All processed samples were stored as aliquots at −80 °C until use.

Table 2 Relationship between systemic hypertension and urinary levels of AQP2 and AGT in IgAN patients, analysed by logistic regression.

<table>
<thead>
<tr>
<th></th>
<th>Simple logistic analysis</th>
<th>Multiple logistic analysis</th>
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<tbody>
<tr>
<td></td>
<td>OR 95% CI</td>
<td>R²</td>
</tr>
<tr>
<td>AQP2 (100 fmol/mg Ucr)</td>
<td>1.22 1.02–1.45</td>
<td>0.0014</td>
</tr>
<tr>
<td>AGT (ng/ml Ucr)</td>
<td>1.16 0.93–1.45</td>
<td>0.0078</td>
</tr>
</tbody>
</table>

*likelihood ratio; †comparing AQP2 and AGT CI levels.

Results

Baseline evaluation

At the time of histological diagnosis, IgAN patients had significantly higher levels of uAQP2, compared with healthy controls (Fig. 1) as well as to patients with CKD other than IgAN (1964.62 ± 275.44 vs 720.96 ± 114.43 fmol/mg UrCr; P=0.0004). Vesicular trafficking and long-term changes in the renal abundance of AQP2 water channels are mainly modulated by AVP. Therefore, we next measured copeptin, a marker of endogenous AVP (12), and found that IgAN patients

Statistical analysis

All data are presented as proportions of the patient population, mean ± s.d. or median and range, as appropriate. Differences between quantitative non-parametric variables were tested by the Mann–Whitney U test and Wilcoxon test, as appropriate. The correlation between non-parametric variables was determined by Spearman’s rank correlation test. Adjustment was made for relevant covariates that had been found significant (P<0.05) in univariate analysis. Logistic regression was used to determine factors significantly related to the presence of arterial hypertension and to deterioration of kidney function over time (i.e. change of proteinuria following drug therapies). Variables with a significance level <0.05 at simple logistic analysis were next fit in a multivariate model. The risk is expressed as odds ratio (OR) ± 95% confidence interval (CI). The Statview software package, SAS (5.0 version; Cary, NC, USA) was used for all analyses.

Urinary AGT, AQP2, bradykinin and plasma copeptin assays

Copeptin plasma levels, and bradykinin (BK) and AGT urinary levels were measured using commercially available ELISA kits (BK and copeptin: Uscn Life Science, Inc., Wuhan, China; AGT: Immuno-Biological Laboratories Co., Gunma, Japan). Urinary AQP2 was measured by ELISA as described previously (11). Urine concentration of BK, AGT and AQP2 was expressed as protein-to-urinary creatinine (UrCr, in mg) ratio.

Figure 2 (A) Urinary levels of aquaporin 2 (AQP2) in healthy individuals (CTRL), in normotensive CKD patients, and in IgAN patients, distinguished in normotensives (NT) and hypertensives (HT). (B) Urinary levels of angiotensinogen (AGT) in healthy individuals and in hypertensive and normotensive IgAN patients. All variables were measured at the time of renal biopsy, in the absence of any drug therapy. Box-and-whisker plots, as in Fig. 1 (ANOVA test).
had a concomitant increase in the plasma levels of the glycopeptide (Fig. 1).

Then, we found significantly higher levels of uAGT, confirming the RAS activation in IgAN patients (Fig. 1).

Daily proteinuria higher than 1 g was associated with higher levels of uAQP2 (2739.0 ± 553.5 vs 1579.6 ± 162.5 fmol/mg UrCr; \( P = 0.02 \)) and uAGT (19.6 ± 5.9 vs 6.4 ± 2.7 ng/mg UrCr; \( P = 0.004 \)).

Similarly, IgAN patients with eGFR lower than 60 ml/min (\( n = 10 \)) showed significantly higher levels of both AQP2 (2397.1 ± 178.9 vs 1634.1 ± 179.7 fmol/mg UrCr; \( P = 0.005 \)) and uAGT (36.2 ± 8.5 vs 5.7 ± 1.7 ng/mg UrCr; \( P = 0.001 \)).

At the start of the study, in the absence of any therapy, we were unable to find a direct correlation between uAQP2 and uAGT levels.

None of the biomarkers measured resulted to correlate with age or sex of the patients studied.

Hypertension

IgAN patients presenting with arterial hypertension at the time of histological diagnosis displayed the highest levels of uAQP2 and uAGT (Fig. 2). Simple logistic analysis showed that both the variables examined were highly correlated with the presence of systemic hypertension. The OR associated with a 100 fmol/mg UrCr increase in mean uAQP2 or 1 ng/mg UrCr increase in mean uAGT yielded a 22 or 15% increase in risk for arterial hypertension respectively (Table 2). When uAQP2 and uAGT were fitted in a multiple logistic regression model, both variables maintained their relationship with the presence of hypertension (Table 2).

End-of-study evaluation

Six-month treatment did not significantly modify plasma osmolarity, fractional sodium excretion and urine osmolality of IgAN patients (Table 3), which would rule out major modifications of renal salt and water handling in a cohort of subjects on a free diet and with free access to fluids. Of note, the change in urine osmolality following drug treatment correlated with the change of uAQP2 (\( P = 0.007 \)). Corticosteroids caused a significant decrease of both the variables examined (Fig. 3). At variance, 6-month treatment with ACEi failed to significantly modify uAGT or uAQP2 (Fig. 3). RAS blockade by ACEi is acknowledged to increase BK concentration (13), and the kinin would directly impair cell surface expression of AQP2 water channel (14). Therefore, we measured uBK at both the start and the end of the study and found that ACEi caused a significant increase of the kinin, but only in the absence of steroids (Fig. 3). This seemingly negates a relevant interfering effect of BK on the urine excretion of AQP2 in the IgAN patients studied.

Finally, in the whole group of patients, the change of uAGT following pharmacological treatments significantly correlated with the change of uAQP2 (\( r = 0.502; \ P = 0.0013 \)).

Proteinuria

Drug therapy caused > 50% decrease of daily proteinuria in 45% patients treated with ACEi alone and in 58% patients treated with steroids + ACEi. The estimated glomerular filtration rate was stable during the whole observation period (Table 1).

Table 3 Effects of drug therapy (ACEi and ACEi+steroids) on renal salt and water handling in the cohort of IgAN patients. Data are presented as median (range).

<table>
<thead>
<tr>
<th></th>
<th>ACEi Baseline</th>
<th>ACEi 6 months</th>
<th>ACEi+steroids Baseline</th>
<th>ACEi+steroids 6 months</th>
</tr>
</thead>
<tbody>
<tr>
<td>Plasma sodium (mEq/l)</td>
<td>139 (133–143)</td>
<td>140 (134–143)</td>
<td>138.5 (134–143)</td>
<td>139 (136–147)</td>
</tr>
<tr>
<td>Plasma osmolarity (mOsm/l)</td>
<td>291.4 (278.8–305.8)</td>
<td>291.1 (282.9–300.3)</td>
<td>291.5 (280.4–301.3)</td>
<td>297.4* (285.7–314.6)</td>
</tr>
<tr>
<td>Urine osmolality (mOsm/kg per mg urine creatinine)</td>
<td>4.0 (2.9–7.6)</td>
<td>5.1 (1.3–8.4)</td>
<td>4.0 (1.9–5.7)</td>
<td>4.3 (1.3–6.3)</td>
</tr>
<tr>
<td>FENa (%)</td>
<td>1.2 (0.5–3.0)</td>
<td>1.1 (0.4–1.9)</td>
<td>1.6 (0.4–5.2)</td>
<td>1.7 (0.7–3.2)</td>
</tr>
</tbody>
</table>

*\( P < 0.05 \) versus baseline.

\[\text{Figure 3} \text{ Changes in urinary AQP2 (A), AGT (B) and bradykinin (C) excretion following 6-month treatment with ACEi alone or steroids + ACEi. Box-and-whisker plots, as in Fig. 1 (Wilcoxon test).}\]
protein AQP2. Furthermore, patients with a greater deterioration of kidney function (i.e., eGFR lower than 60 ml/min and daily proteinuria higher than 1 g) displayed a higher urine excretion of both AGT and AQP2.

Local RAS hyperactivity has long been demonstrated in patients with IgAN (2, 15). Recently, uAGT has been proposed as a reliable marker of activated intrarenal RAS in IgAN: increases in uAGT levels were correlated with augmented intrarenal AGT gene expression and Ang II levels (16). Then, uAGT concentration is deemed to reflect the extension of renal damage in patients with CKD, including IgA nephropathy (17, 18).

For the first time, we demonstrated increased copeptin plasma levels, along with enhanced levels of uAQP2, in untreated IgAN patients, suggesting an increase of the intrarenal action of the neuropeptide in the disease. This increase seems to be specific for IgAN, and is seemingly modest in other forms of CKD. No studies have previously explored the excretion of uAQP2 in human glomerulonephritis. We assume that the enhanced uAQP2 excretion measured in IgAN patients could likely be the end result of a complex series of driving signals on the kidney expression and trafficking of the water channel protein, comprising the increased levels of AVP, the hyperactivity of renal RAS and, on the other side, the enhanced renal concentration of BK, as inferred by the increase of uBK recorded at the start of the study. It is worth noting that the increase of uBK mostly reflects the increase in the local concentration of the kinin, since BK serum levels of IgAN patients

Discussion

In this study, we report that at the time of histological diagnosis IgAN patients displayed an early and strong activation of the renal RAS, which was associated with a rise in the renal excretion of the water channel

Figure 4 (A) Correlation between the change (Δ) of urinary AQP2 and the change (Δ) of 24 h proteinuria (g/die) in IgAN patients after 6-month treatment with either ACEi or steroids + ACEi (Spearman’s rank correlation test). (B) Change of AQP2 urine excretion in patients, distinguished in responders (>50% decrease of proteinuria) and non-responders (<50% decrease of proteinuria) (Mann–Whitney U test). (C) Change of urinary AQP2 excretion according to the type of treatment and the modification of daily proteinuria (ANOVA test). Box-and-whisker plots, as in Fig. 1.

When patients in both treatment groups were distinguished according to the change of proteinuria at the end of the 6-month period, those with a fall of proteinuria >50% displayed also a significant decrease of uAQP2 and uAGT: the higher the decrease of uAQP2 or uAGT, the higher the decrease in protein excretion (Figs 4 and 5). Univariate logistic analysis confirmed that the modifications of uAQP2 and uAGT were independent predictors of the change of proteinuria, regardless of the type of therapy (Table 4, left panel). When fitted in a multiple logistic regression model, both the independent variables maintained their relationship with the change of proteinuria (Table 4, right panel).

Finally, we tried to ascertain whether the baseline levels of the biomarkers investigated would predict treatment response. Baseline uAQP2 levels, but not serum copeptin or uAGT, were significantly associated with the change of proteinuria recorded at the end of the study ($\rho = -0.489; P = 0.004$). Accordingly, patients with higher levels of uAQP2 at baseline showed higher decrements in the urine excretion of the water channel protein following drug therapy ($\rho = -0.745; P < 0.0001$).

Figure 5 (A) Correlation between the change (Δ) of urinary angiotensinogen (AGT)) and the change (Δ) of 24 h proteinuria (g/die) in IgAN patients after 6-month treatment with either ACEi or steroids + ACEi (Spearman’s rank correlation test). (B) Change of AGT urine excretion in patients, distinguished in responders (>50% decrease of proteinuria) and non-responders (<50% decrease of proteinuria) (Mann–Whitney U test). (C) Change of urinary AGT excretion according to the type of treatment and the modification of daily proteinuria (ANOVA test). Box-and-whisker plots, as in Fig. 1.
Therapies resulted to correlate with the change of mechanisms, our findings suggest that 6-month steroids potently inhibit tissue inflammation. Moreover, AGT transcription in IgAN AGT II-induced stimulation of AT1 receptors (26), and in this way may dampen Ang II. However, glucocorticoids would also down-regulate Ang II, been shown to activate human tubular epithelial cells, glucocorticoids have independent mechanisms. Then, the down-modulation of uAGT by steroids is rather puzzling. In cultured tubular cells (27), AGT transcription in IgAN involves Ang II AT1 receptors (26), and in this way may dampen Ang II-induced stimulation of AGT expression by proximal tubular cells (27). Moreover, AGT transcription in IgAN correlates with tubulointerstitial inflammation (1), and steroids potently inhibit tissue inflammation. Whatever the mechanisms, our findings suggest that 6-month treatment with corticosteroids depresses intrarenal RAS activity more markedly than ACEi alone in IgAN.

Of note, the change in uAQP2 following drug therapies resulted to correlate with the change of uAGT, which would support a functional relationship between the two systems investigated, as suggested by in vitro models (8) and further supported by multivariate analysis (Table 3).

We found that patients with a higher decrease in proteinuria (>50%), and therefore with a diminished risk of progression (28), exhibited a definitely higher drop of uAQP2 and uAGT (Figs 4B and 5B), regardless of the type of therapy done (Figs 4C and 5C). Then, baseline levels of uAQP2 were strong predictors of patients’ response to the therapy: the higher its levels at histological diagnosis, the higher its decrease following pharmacological treatments, and the higher the decrease of daily proteinuria.

Urinary AGT has been reported to significantly correlate with urinary protein in patients with CKD, including IgAN (17, 18). Consequently, its decrease is expected to associate with a decrease in daily proteinuria. The association between the change in proteinuria and the change in AQP2 excretion is striking. Bardoux et al. (7) showed that stimulation of AVP V2 receptors induces a rise in urinary albumin excretion in humans, which is only partially prevented by ACEi in rats. In a large population study, plasma copeptin levels were lower than in controls (61.38 ± 7.5 vs 114.19 ± 28.57 pg/ml; P = 0.04). Recent evidence suggests that AVP may play a role in the genesis and exacerbation of renal damage (5). The urinary excretion of AQP2 is taken to closely parallel changes in intrarenal AVP action. Thus, the increase of uAQP2, described here, would support a contributory role of AVP in the progression of renal damage of IgAN, and its levels of excretion are seemingly associated with the degree of deterioration of renal function.

We then found that the presence of arterial hypertension was associated with higher levels of uAGT and uAQP2 in IgAN patients, in the absence of any treatment (Fig. 2). As for uAGT, our findings confirm previous studies (19), and extend the correlation between uAGT and BP also to IgAN patients. Then, we demonstrate for the first time that uAQP2 associates with the presence of systemic hypertension in humans, similar to animal models (20).

Following 6-month treatment with ACEi alone, uAGT and uAQP2, on average, failed to show any significant modification (Fig. 3). At variance, the addition of steroids caused a significant decrease in the concentration of both the urine variables examined (Fig. 3A and B). As for the diminished AQP2 excretion, glucocorticoids exert a powerful inhibitory effect on hypothalamic regulatory pathways of AVP secretion (21, 22). Moreover, long-term aldosterone stimulation has been reported to decrease apical expression of AQP2 (23, 24). In other words, steroids can decrease AQP2 expression through both AVP-dependent and AVP-independent mechanisms. Then, the down-modulation of uAGT by steroids is rather puzzling. In cultured human tubular epithelial cells, glucocorticoids have been shown to activate AGT gene transcription (25). However, glucocorticoids would also down-regulate Ang II AT1 receptors (26), and in this way may dampen Ang II-induced stimulation of AGT expression by proximal tubular cells (27). Moreover, AGT transcription in IgAN correlates with tubulointerstitial inflammation (1), and steroids potently inhibit tissue inflammation. Whatever the mechanisms, our findings suggest that 6-month treatment with corticosteroids depresses intrarenal RAS activity more markedly than ACEi alone in IgAN.

Of note, the change in uAQP2 following drug therapies resulted to correlate with the change of uAGT, which would support a functional relationship between the two systems investigated, as suggested by in vitro models (8) and further supported by multivariate analysis (Table 3).
exerted by ACEi, and interfere with AVP-dependent induction of renal damage.

Collectively, the above findings seem to support a role of AVP–AQP2 axis, possibly interacting with the RAS, in the progression of IgAN and suggest AQP2 as a possible novel marker of IgAN and of disease progression.

On the other hand, we are aware of some limitations of this study preventing firm conclusions. First, we examined a relatively limited number of patients over a short time period, which precluded the use of hard end points, such as doubling of serum creatinine, and forced to the adoption of a surrogate marker of disease progression, namely daily proteinuria. Then, we failed to measure serum copeptin during and/or at the end of drug therapy. Finally, a conclusive and direct demonstration of the role of AVP–AQP2 axis on the progression of IgAN would require the administration of a selective AVP V2-receptor antagonist, such as tolvaptan, for relatively long periods.

Declaration of interest
The authors declare that there is no conflict of interest that could be perceived as prejudicing the impartiality of the research reported.

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