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Up-Regulation of the Human Serum and Glucocorticoid-Dependent Kinase 1 in Glomerulonephritis

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Key Words

Glomerulonephritis · Cell volume regulation · SGK1

Abstract

Glomerulonephritis is paralleled by excessive formation of transforming growth factor-beta (TGF- β), which participates in the pathophysiology of the disease. Recently, a novel downstream target of TGF- β has been identified, i.e. the human serum and glucocorticoid-dependent kinase 1 (hSGK1), a serine/threonine kinase participating in the regulation of Na⁺ transport. The present study was performed to elucidate transcriptional regulation of hSGK1 in glomerulonephritis. To this end, in situ hybridization was performed in biopsies from patients with clinical diagnosis of glomerulonephritis. hSGK1 transcript levels were moderately enhanced in 5 out of 9 patients and strongly enhanced in 4 out of 9 patients. Distal nephron epithelial cell hSGK1 transcript levels were low or absent in 7 of the 9 patients but markedly enhanced in 2 of the 9 patients. In conclusion, glomerulonephritis leads to glomerular and in some cases to epithelial up-regulation of hSGK1 transcription.

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Introduction

Ample evidence points to a role of transforming growth factor-beta1 (TGF- β 1) in the pathophysiology of glomerulonephritis [1–8]. Accordingly, TGF- β antagonism reversed matrix accumulation by infusing neutralizing TGF- β antibodies [2], soluble TGF- β receptors [9] and decorin, an inhibitor of TGF- β [10, 11].

Recently, a novel downstream target of TGF- β has been identified, the human serum and glucocorticoiddependent kinase 1 (hSGK1), which is transcriptionally up-regulated by TGF- β in U 937 macrophages [12], HepG2 liver cells [12] and 3T3 fibroblasts [13]. The human hSGK1 has previously been cloned as cell volume sensitive gene [14]. The rat SGK1 was originally cloned from rat mammary tumour cells as serum and glucocorticoid-sensitive kinase [15], but was subsequently shown to be up-regulated by mineralocorticoids [16, 17]. Transcription of the kinase has been demonstrated to be enhanced by excessive extracellular glucose concentrations, an effect blunted by neutralizing anti-TGF- β antibodies [13]. Accordingly, TGF-B at least partially accounts for transcriptional up-regulation of hSGK1 in diabetic nephropathy [13, 18-20].

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Table 1. Clinical details and histopathological grading of the glomerulonephritis patients

| Patient | Age years | Sex | Diagnosis | Crea mg/dl | GFR ml/min | Diabetes mellitus | Hypertension >140/90 mm Hg | Proteinuria g/day | SGK grading (0–4) glomerulus | SGK grading (0–4) DT |
|---------|--------------|-----|------------------------------|---------------|---------------|----------------------|----------------------------------|----------------------|------------------------------------|----------------------------|
| 1 | 62 | f | immune-complex GN | 2.5 | no data | no | yes | no | 3 | 0-1 |
| 2 | 50 | m | perimembranous GN < | <1.1 | 97 | no | yes | 4.9 | 0-1 | 0 |
| 3 | 51 | m | perimembranous GN | 3.2 | no data | no | yes | 8.0 | 0-1 | 0 |
| 4 | 45 | m | mesangio-proliferative GN | 1.3 | 156 | no | yes | 3.2 | 2 | 0-1 |
| 5 | 70 | f | immune-complex GN | 1.8 | no data | no | no | 0.8 | 1 | 0 |
| 6 | 29 | f | mesangio-proliferative GN | 1.4 | no data | no | no | 4.8 | 3 | 0-1 |
| 7 | 30 | m | rapid progressive GN | 9 | no data | no | yes | no data | 3 | 4 |
| 8 | 43 | f | malignant nephrosclerosis | 2.2 | no data | no | yes | mild proteinuria | 4 | 4 |
| 9 | 26 | m | mesangio-proliferative GN | 1.5 | no data | no data | no data | 3.9 | 0-1 | 2 |

Crea = Creatinine; GFR = glomerular filtration rate; GN = glomerulonephritis; DT = distal tubule.

The present study has been performed to test for deranged transcriptional regulation of hSGK1 in glomerulonephritis.

Materials and Methods

Patients

Kidney biopsies from patients with clinical diagnosis of glomerulonephritis were taken for diagnostic reasons. Initially, 13 patients were included in this retrospective study. From those, 4 patients had to be excluded due to initial treatment with steroids or drugs which affect the angiotensin/aldosterone system; the other patients did not receive steroids, immunosuppression or ACE inhibitors at the time of biopsy. From the remaining 9 patients, 3 suffered from mesangioproliferative glomerulonephritis, 1 from rapid progressive glomerulonephritis, 2 from perimembranous glomerulonephritis, 1 from malignant nephrosclerosis and 2 from immune complex glomerulonephritis after streptococci infection. Six out of the 9 patients were hypertensive (RR > 140/90 mm Hg), no patient was diabetic. Normal renal biopsies from 7 patients were obtained from intact parts of kidneys which had been surgically removed because of renal cancer. For full clinical details, see table 1.

In situ Hybridization

Kidney biopsy tissue was fixed in 4% paraformaldehyde/0.1 M sodium phosphate buffer (pH 7.2) for 4 h and embedded in paraffin. Four-micrometer tissue sections were dewaxed and hybridized basically as described [21–23]. Hybridization mixture contained either the ³⁵S-labeled RNA antisense or sense control hSGK1 probe [13] (500 ng/ml) in 10 mM Tris HCl, pH 7.4/50% (vol/vol) deionized formamide/600 mM NaCl/1 mM EDTA/0.02% polyvinylpyrrolidone/ 0.02% Ficoll/0.05% bovine serum albumin/10% dextrane sulphate/ 10 mM dithiothreitol/denatured sonicated salmon sperm DNA at 200 µg/ml/rabbit liver tRNA at 100 µg/ml [12]. Hybridization with RNA probes proceeded at 42 °C for 18 h. The slides were then

washed as described [21–23] followed by 1 h at 55 °C in 2 × standard saline citrate. Non-hybridized single-stranded RNA probes were digested by RNAse A (20 μ g/ml) in 10 mM Tris HCl, pH 8.0/0.5 M NaCl for 30 min at 37 °C. Tissue slide preparations were autoradiographed [23] and stained with haematoxylin/eosin.

Statistical Analysis

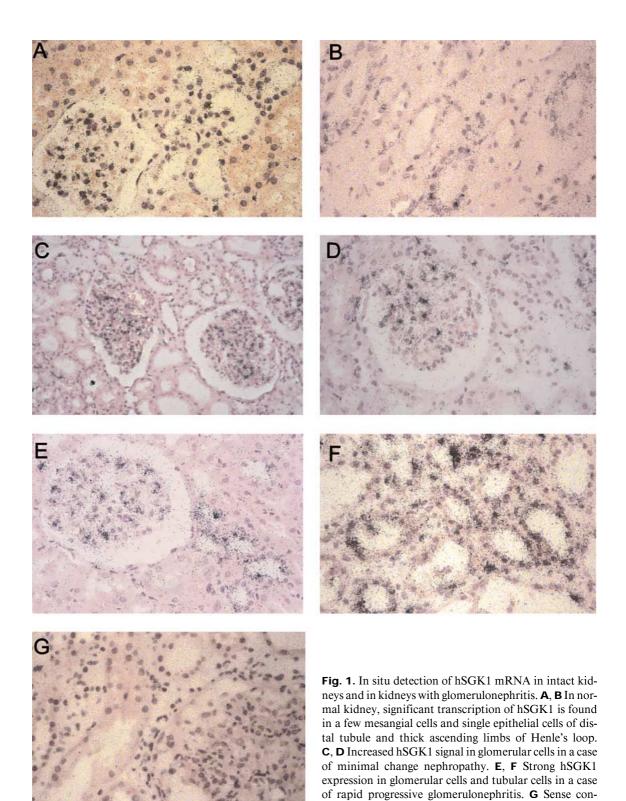
Data are expressed as arithmetic means \pm SEM. Statistical analysis has been made by the Student t test or ANOVA, where applicable. A p value of <0.05 was considered statistically significant.

Results

As shown in figure 1, expression of hSGK1 mRNA in intact kidneys was restricted to some mesangial cells of the glomeruli and a few epithelial cells in distal convoluted tubules and thick ascending limbs of Henle's loop. In 4 out of the 9 glomerulonephritic kidneys, glomerular hSGK1 transcription was markedly enhanced, in 5 out of the 9 patients, hSGK1 expression remained low or was only moderately enhanced. In 2 of the 9 patients, markedly enhanced transcript levels were observed in epithelial cells from distal tubule and thick ascending limb. Both patients were hypertensive. In addition, high levels of hSGK1 transcripts were observed in interstitial cells, most likely representing macrophages and fibroblasts. Unspecific labelling of kidney cells was excluded by hybridization of tissue sections with the ³⁵S-labeled sense control RNA hSGK1 probe.

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trol.

Discussion

The present study demonstrates up-regulation of hSGK1 transcription in kidneys from glomerulonephritic patients. The up-regulation may be expected, as glomerular injury is paralleled by up-regulation of TGF- β [4], which is a strong stimulator of hSGK1 transcription [12, 13] and is thought to damage renal tissue by stimulation of matrix deposition [24]. This does, of course, not rule out the contribution of further cytokines and hormones to the up-regulation of hSGK1 in glomerulonephritic kidneys. The transcription of hSGK1 has proved to be highly sensitive to a wide variety of stimuli [25].

In 2 patients, hSGK1 was not only up-regulated in affected glomeruli but as well in epithelial cells, particularly in distal tubule and thick ascending limb. As hSGK1 has been shown to be a strong stimulator of the renal epithelial Na⁺ channel ENaC [16, 13, 17, 26, 27] and the thick ascending limb Na⁺, K⁺, 2Cl⁻ cotransport [13], the up-regulation of hSGK1 in renal epithelial cells could lead to renal Na⁺ retention and thus favour the development of hypertension. However, in the epithelial cells of the remaining 7 patients, the hSGK1 levels remained low

even though 4 of those patients developed hypertension. Possibly, hSGK1 is up-regulated in confined nephron segments such as medullary collecting duct outside the available tissue specimen. Nevertheless, widespread epithelial up-regulation of hSGK1 does not appear to be a prerequisite for the development of hypertension in glomerulonephritis.

In conclusion, glomerulonephritis is paralleled by upregulation of hSGK1 in affected glomeruli and in a subset of patients in renal epithelial cells of thick ascending limb and distal tubule.

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