Plasma Gelsolin Promotes Proliferation of Mesangial Cell in IgA Nephropathy

Lei Zhang, Dan Kong, Hongxue Meng, Changsong Han, Jiang Zhu, Juanjuan Qiao, Yan He, Tianshen Wang, Xiaobo Li, Fengmin Zhang, Xiaoming Jin

Key Words

IgA nephropathy (IgAN) • Plasma gelsolin (pGSN) • Human mesangial cells (HMCs) • Transforming growth factor beta1 (TGF-β1) • Progression and prognosis

Abstract

Background/Aims: Plasma gelsolin (pGSN) is an actin-binding protein that plays a critical role in the pathogenesis of rheumatoid arthritis. However, whether pGSN is involved in other immunological diseases remains unknown. This study focused on the relationship between pGSN and immunoglobulin A (IgA) nephropathy (IgAN).

Methods: Two hundred patients with IgAN, 200 patients each with several other types of nephropathy and healthy controls (HCs) who underwent kidney biopsies between 2000 and 2014 were enrolled in the study. The Oxford classification system was used to predict the risk of disease progression. Serum and renal tissue were used to detect pGSN, and the correlations between pGSN and IgA, galactose-deficient IgA1 (Gd-IgA1), transforming growth factor beta1 (TGF-β1), fibronectin (FN) content, clinical symptoms, and kidney function were analyzed.

Results: We found that the pGSN levels were significantly decreased in sera from IgAN patients compared to sera from patients with other forms of glomerular nephritis and HCs. Furthermore, the serum pGSN levels were negatively correlated with the serum IgA1, FN, and TGF-β1 levels, and positively correlated with the estimated glomerular filtration rate. Conversely, the glomerular pGSN content was significantly elevated in the IgAN patients and was positively correlated with TGF-β1 and FN levels. In renal tissue, the pGSN levels were significantly higher in IgAN patients with M1 and S1 compared to patients with M0 and S0 (p < 0.05). Meanwhile, pGSN promoted human mesangial cell (HMC) proliferation by facilitating cell mitosis in vitro. pGSN also promoted integrin α2β1 expression in HMCs and enhanced the integrin α2β1-pGSN interaction.

Conclusion: Our study suggested that pGSN may play an important role in the development of IgAN by promoting the proliferation of mesangial cells and that serum and glomerular pGSN levels may be new markers for predicting IgAN progression and prognosis.

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Introduction

Immunoglobulin A (IgA) nephropathy (IgAN), the most common form of progressive primary glomerulonephritis (GN), is characterized by the presence of IgA complex deposition in the glomerular mesangium [1-3]. Moreover, most patients develop end-stage renal disease (ESRD) due to renal fibrosis after more than ten years. Renal fibrosis, including glomerular and tubulointerstitial fibrosis, is one of the major pathological changes that are caused by progressive IgAN [4]. Mesangial cells (MCs) play an important role in the fibrotic processes of glomerular diseases by promoting excessive synthesis and secretion of extracellular matrix (ECM). Among the ECM components, transforming growth factor beta1 (TGF-β1) is a pivotal profibrotic cytokine that contributes to renal fibrosis and tubulointerstitial damage [5, 6]. TGF-β1 may specifically promote renal fibrosis through the downstream Smad signaling pathway [7], which can promote the mRNA expression of type I collagen and fibronectin (FN) [8], which are other important constituents of the ECM. Our study examined TGF-β1 and FN levels to evaluate glomerular fibrosis in IgAN.

Gelsolin (GSN) is an intracellular actin-binding protein that affects cell shape, cell motility, phagocytosis and apoptosis [9]. GSN exists in both intracellular (cytoplasmic, or cGSN) and extracellular (secreted/plasma, or pGSN) forms; both of these proteins share a common antigen determinant and partial molecular homology. pGSN circulates in the blood of healthy individuals at a concentration of 200 ± 50 mg/L and has been suggested to be a critical component of the extracellular actin-scavenging system during tissue damage [10]. Osborn [11] found that the pGSN levels in the sera of patients with rheumatoid arthritis were decreased compared to the levels in healthy controls (HCs). Furthermore, GSN-actin complexes have been detected in synovial fluids. Therefore, we hypothesized that pGSN may participate in the initiation and development of immune-related diseases. We previously found that the pGSN levels in the sera of patients with IgAN were decreased compared to those in patients with other types of glomerular nephritis and HCs, and high pGSN levels were detected in the glomeruli of IgAN patients. However, the specific role of pGSN in IgAN has remained unknown. Furthermore, the interactions between pGSN and the proliferation of MCs as well as the progression of IgAN have not previously been investigated.

The objective of the present study was to investigate the relationship between pGSN and IgAN progression and to elucidate the mechanisms of pGSN involvement in IgAN progression. Our study demonstrated that pGSN may play an important role in the development of IgAN by promoting the proliferation of MCs and that serum and glomerular pGSN levels may be markers for predicting IgAN progression.

Materials and Methods

Solutions and drugs

Fluorescein isothiocyanate (FITC)-labeled goat anti-human IgA (alpha) was purchased from KPL (Kirkgaard & Perry Laboratories Inc. (KPL), Gaithersburg, Md.). FITC-labeled mouse anti-human IgG1 (Fc), FITC-labeled goat anti-human IgM, mouse monoclonal IgA, mouse anti-FN antibody, mouse anti-human pGSN, and rabbit anti-human GSN were purchased from Abcam Limited (Cambridgeshire, UK). FITC-labeled rabbit anti-human C3c antibody was purchased from Dako (Dako A/S, Glostrup, Denmark). Sambucus nigra agglutinin (SNA) and Vicia villosa lectin (VVL) were purchased from Vector Laboratories (Burlingame) and were used to detect serum α2,6-sialic acid and N-acetylgalactosamine (GalNAc), respectively. Peanut agglutinin (PNA), which was used to detect galactose (Gal), was purchased from Sigma-Aldrich (St. Louis, MO, USA). Mouse anti-human actin, mouse anti-human integrin α2β1 and mouse anti-human cyclin A were purchased from Santa Cruz Biotechnology, Inc. Rabbit anti-human CDK2 antibodies were obtained from Cell Signaling Technology, and FITC-conjugated goat anti-mouse IgG, Alexa Fluor® 594 Goat Anti-Mouse IgG (H+L) Antibody, and peroxidase-conjugated goat anti-mouse IgG were purchased from Invitrogen (Invitrogen, Life Technologies, USA). RNA was isolated using TRIzol reagent Invitrogen, and anti-mouse IgG antibody was purchased from Vector Laboratories (Vector Laboratories, CA). An ECL Plus Detection Kit was
purchased from Amersham Biosciences (Piscataway, NJ, USA). Recombinant human pGSN (rhpGSN) was used as a standard in the enzyme-linked immunosorbent assay (ELISA) and western blotting experiments and was purchased from Cytoskeleton, Inc. (Denver, CO, USA). Finally, the human FN ELISA kit was purchased from Wuhan Boster Bioengineering Company, Limited (China).

Patients and controls

In total, 200 patients with IgAN and 200 patients each with membranous nephropathy (MN), lupus nephritis (LN), mesangial proliferative GN (MsPGN), membrane proliferative GN (MPGN), diffuse endocapillary proliferative GN (DEPGN), focal segmental glomerulosclerosis (FSG), ischemic glomeruli (IG), minimal change glomerulopathy (MCG) and HCs underwent kidney biopsies for diagnosis. The diagnoses were performed in the Department of Pathology of Harbin Medical University from January 2010 to April 2014. Among the IgAN patients, there were 67 males and 113 females, corresponding to a male-to-female ratio of 0.77:1. All of the patients were between 7 and 67 years of age, with an average age of 34.72 ± 9.36 years. Individuals with MN (in situ immune complex deposition-induced GN) or MCG (no immune complex deposition) and HCs were sex and age matched to the IgAN patients. Serum and kidney tissues from the kidney biopsies were collected. The serum was specifically collected from the HCs during outpatient visits at the 2nd Affiliated Hospital of Harbin Medical University. Moreover, the healthy kidney control tissues (30 cases) were collected from anatomical areas that were distant from the surgically resected kidney tumors. All of the patients and HCs provided informed consent prior to the study. Patients with diabetes, Henoch-Schönlein purpura or liver disease were excluded from the study. Serum pGSN levels were measured in the patients with different types of glomerular nephritis, and patients with MN or MCG and HCs were selected as the control groups for the IgAN patients in the detection of all of the studied factors in the serum and renal tissue.

Oxford classification

The renal tissues of the IgAN patients were evaluated using the IgAN Oxford classification (OC) [12]. The presence of the following findings was scored as 1, and the absence was scored as 0. The scoring included a mesangial (M) score of M1 or M0; a segmental glomerulosclerosis (S) score of S1 or S0; an endocapillary hypercellularity (E) score of E1 or E0; and a tubular atrophy/interstitial fibrosis (T) score of T0 (≤ 25%), T1 (26-50%), or T2 (> 50%). The MEST score was calculated as the sum of M + E + S + T.

Detection of serum pGSN, IgA, Gal-deficient IgA1 (Gd-IgA1), actin, TGF-β1 and FN by ELISA and western blotting

Gd-IgA1 was detected using a specific lectin-binding ELISA, as previously reported [13]. Briefly, each well of a polystyrene microtiter plate (Corning, NY, USA) was coated with 5 µg/mL anti-human IgA in 100 µL of 0.05 mol/L bicarbonate buffer (pH 9.6) at 4°C overnight. After three washes with 0.01 mol/L phosphate-buffered saline (PBS) containing 0.05% Tween 20 (PBS-T), the plates were incubated with 1% fetal bovine serum (FBS) for 60 min to block their nonspecific reactivity. The plates were then incubated with serum samples from patients with IgAN, MN or MCG and HCs at room temperature for 2 h, followed by washing with PBS-T. Next, PNA, SNA and VVL (1 mg/mL; Sigma) were diluted to 1:200 in a blocking buffer, with serum samples from patients with IgAN, MN or MCG and HCs at room temperature for 2 h, followed by washing with PBS-T. The plates were then incubated with horseradish peroxidase (HRP)-conjugated avidin. After a wash with PBS-T, 3’, 3’, 5’, 5’-tetramethylbenzidine (TMB) peroxidase substrate was added, and the reaction was halted with 2 mol/L H2SO4. The absorbance was then measured with a Bio-Rad 550 microplate reader at a wavelength of 450 nm.

Anti-human pGSN, actin and IgA were diluted to 1:1000 and were used to coat the plates, while the secondary antibody was diluted to 1:2000. The remaining steps followed the above-described methods. Human FN and TGF-β1 ELISA kits were used to measure the FN and TGF-β1 levels, respectively, according to the manufacturer’s instructions. Of the 200 cases of each nephropathy type, 20 were randomly selected for western blotting; we specifically analyzed serum samples from 20 IgAN patients, 20 MCG patients and 20 MN patients and 20 HCs.

The presence of pGSN in the serum was assessed by western blotting. Platelet-poor plasma was first diluted 1:20 in a sample buffer (SB) (10% glycerol, 2% SDS, 62.5 mM Tris-HCl, 0.03% bromophenol blue and 5% β-mercaptoethanol, pH 6.8). rhpGSN (Cytoskeleton, Denver, CO, USA) diluted in 1× SB was then added to form a protein gradient (50 ng, 100 ng, 150 ng and 200 ng) and incubated at 100°C for 10 min.
Afterward, 20 µL of each sample was subjected to 10% SDS-PAGE, and the proteins were transferred to a PVDF membrane. After being blocked in 5% skim milk in PBS-T for 1 h, the membrane was incubated with an anti-pGSN antibody (5 µg/mL in 5% skim milk/PBS-T) for 1 h. After being washed with PBS-T, the membrane was incubated with peroxidase-conjugated goat anti-mouse IgG for 1 h, and reactive bands were detected using the ECL Plus Detection Kit. rhpGSN was used to establish a standard curve, and the pGSN levels were calculated.

**Immunohistochemistry**

Paraffin sections from the 200 cases of IgAN, MN, MCG and HC were pretreated with citrate (pH 6.0) buffer for 3 min at 120°C in a pressure cooker, and endogenous peroxidase was inhibited with 3% H2O2 in PBS for 10 min. Nonspecific reactions in the sections were also blocked with goat serum for 1 h at room temperature. The sections were then incubated with the primary antibody anti-pGSN overnight at 4°C, followed by incubation with the secondary antibody in 1% BSA for 30 min at room temperature. The chromogenic reaction was performed via diaminobenzidine (DAB) staining, and the staining intensity was measured using appropriate software (ACT-1, Nikon).

All frozen sections were blocked with 5% normal goat serum in PBS (pH 7.4). IgA, IgG, IgM and the complement component C3 were detected in human kidney tissues using FITC-labeled anti-human IgA, IgG, IgM and C3 antibodies (1:100), respectively. Additionally, the FN and TGF-β1 levels were measured using an indirect fluorescence detection method. For this purpose, the tissue sections were incubated with human monoclonal anti-FN and anti-TGF-β1 (1:100 dilution for 12 h at 4°C), washed three times in PBS and incubated with FITC-labeled anti-mouse IgG antibody. The total number of glomeruli per slice was counted under 400× magnification in randomly chosen images that were acquired using a fluorescence microscope (Nikon E800), a digital camera (Nikon 1200F) and accompanying software (ACT-1, Nikon).

Moreover, human mesangial cells (HMCs) with or without pGSN treatment (10 mg/L) were harvested, and microscope slides were prepared. After fixation in 4% paraformaldehyde, the cells were blocked with goat serum for 30 min at room temperature. Next, mouse anti-human integrin α2β1 antibody was diluted with 1:50 in PBS and added to the cells at 4°C for 1 h. After washing with PBS, rabbit anti-mouse secondary antibody labeled with HRP was diluted 1:100 and added to the cells for 30 min. Finally, after washing with PBS and mounting with 10% glycerol, the cells were observed under a microscope (OLYMPUS CX31), and images were acquired.

**Isolation of serum IgA1 in patients with IgAN**

IgA1 was isolated from the pooled sera of 20 patients by Jacalin affinity chromatography. The procedures were performed according to the protocol of Wang and Diven [14, 15]. Briefly, the pooled sera were diluted 1:1 with PBS (pH 7.1) and centrifuged at 2000 rpm for 8 min. Then, 3 mL of the supernatant from each sample was loaded onto a Jacalin affinity column. The column was prepared with Jacalin immobilized on cross-linked 4% agarose beads with an IgA1-binding capacity of 2 mg/mL gel. After the pooled serum samples were incubated on the Jacalin affinity column for 30 min, the column was washed with 175 mM Tris-HCl (pH 7.4) until the optical density (OD) at 280 nm was less than 0.10. The IgA1 was then eluted with 0.15 M melibiose (Yuanye Biotechnology Limited Company, Shanghai, China) in 175 mM Tris-HCl in 3.0 mL with PBS and added to the cells at 4°C for 14 h. After washing with PBS, rabbit anti-mouse secondary antibody labeled with HRP was diluted 1:100 and added to the cells at 37°C for 30 min. Finally, after washing with PBS and mounting with 10% glycerol, the cells were observed under a microscope (OLYMPUS CX31), and images were acquired.
Cell culture and cell viability assay

HMCs were purchased from Cobioer Biosciences Co., LTD (China) and cultured at 37°C and 5% CO₂ in RPMI 1640 medium containing FBS (10%), penicillin (100 U/mL), glutamine (2 mmol/L), and streptomycin (100 µg/mL). The proliferation of the HMCs was determined using Cell Counting Kit-8 (CCK-8; Sigma-Aldrich, St. Louis, MO) according to the manufacturer's instructions. HMCs were seeded on 96-well culture plates at 5×10⁵ cells/well and then incubated with 10 mg/L pGSN for 24 h, 48 h, 72 h or 96 h. Next, 10 µL WST-8 was added to each well and incubated for 4 h at 37°C, after which the absorbance was measured on a spectrophotometer/microplate reader (Bio-Rad) at a wavelength of 450 nm. Cells cultured without pGSN were used as a control.

Cell synchronization and cell cycle phase determination

To observe the effect of the cell cycle on growth after pGSN treatment, subconfluent cultures of HMCs were synchronized in the G2/M phase using colchicines. In particular, cells (10⁵) were incubated on 9.6-cm² disks in RPMI 1640 containing 10% FBS in a CO₂ incubator at 37°C for 12 h, after which colchicines (0.1, 0.2, 0.5, 0.8 or 1 µg/mL) were added to the cells for 12 h or 24 h. Adherent cells were collected by treatment with trypsin and then washed with pre-chilled PBS. The cells were then fixed in 2 mL of cold 75% ethanol overnight at 4°C and resuspended in staining buffer (20 µg/mL propidium iodide (PI; Sigma), 0.1% Triton X-100 and 0.2 mg/mL RNase in PBS) for 2 h at 4°C. The PI-stained cells were analyzed using FACS (FACScan; BD), with at least 10⁵ cells counted for each sample. A data analysis was performed using CXP Software (Beckman Coulter, Inc.). The cell cycle phase was also detected in HMCs treated with pGSN (10 mg/L) at 3 h, 6 h, 9 h or 12 h.

Co-immunoprecipitation

HMCs were treated with pGSN alone or with pGSN and aIgA1 or left untreated for 48 h. Proteins were extracted using lysis buffer (10 mM Tris, pH 7.5; 100 mM NaCl; 5 mM EDTA; 1% Triton X-100; and 0.05% SDS with complete protease inhibitor). Protein samples were then immunoprecipitated with 2 µg of rabbit anti-human GSN and mouse anti-human integrin α2β1 antibodies at 4°C overnight, after which the immunoprecipitated proteins were pulled down with Protein A/G Mix Magnetic Beads (protein A-agarose beads; Merck KGaA, Darmstadt, Germany) at 4°C for 4 h. The immunoprecipitates were subsequently boiled in 2× SDS loading buffer and analyzed by western blotting.

Statistical analysis

The data are presented as the means ± standard errors of the mean. The data were analyzed using SPSS 13.0 (SPSS Inc., Chicago, IL, USA). A statistical analysis of the differences between the serum pGSN levels and the other clinical data was performed using the Mann-Whitney U-test. Pearson’s correlation was used to evaluate all correlations. Moreover, two-way ANOVA was used to evaluate the cell proliferation level, and significant differences among the different grades of the Oxford classification were calculated using the unpaired t-test. Significance was defined as \( p < 0.05 \).

Ethics

This study was approved by the Ethics Committee of Harbin Medical University. All of the individuals who were included in the study provided informed consent.

Results

pGSN levels in patients with several types of glomerular nephritis and in HCs

The pGSN levels were significantly decreased in the sera of patients with IgAN compared to those in patients with IG or MCG and in HCs, as determining using an ELISA and western blot assay (**\( p < 0.01 \), Fig. 1A; \( p < 0.05 \), Fig. 1B, 1C, 1D). Moreover, the pGSN levels were significantly decreased in the sera of patients with MN, LN, MPGN, MsPGN, DEPGN or FSG compared to the levels in patients with IG or MCG and in HCs (*\( p < 0.05 \), Fig. 1A).
The serum concentrations of actin, IgA, Gd-IgA1, FN and TGF-β1, as measured using ELISA, in patients with IgAN, MN or MCG and in HCs

The serum IgA, FN, TGF-β1 and Gd-IgA1 levels in patients with IgAN were higher than those in the control groups (**p < 0.01, *p < 0.05 and *p < 0.05, respectively), namely, the MCG and HC groups (Fig. 2A1, 2A2, 2A3 and 2A4, respectively). There was no difference in the actin levels among the groups (data not shown). In the IgAN patients, the serum pGSN levels were negatively correlated with the serum IgA levels (Fig. 2B1, r = -0.2231, *p = 0.0165). In addition, the serum FN and IgA levels were strongly positively correlated in the IgAN patients (Fig. 2B2, r = 0.4994, **p < 0.0001). Finally, the serum pGSN
levels were strongly negatively correlated with TGF-β1 and FN in the IgAN patients (Fig. 2B3, \( r = -0.2300, *p = 0.0134 \); Fig. 2B4, \( r = -0.2141, *p = 0.01 \)). However, there was no correlation between actin and pGSN in this group (\( p = 0.4352 \)).

**pGSN, TGF-β1 and FN expression in renal tissue in IgAN**

pGSN deposition in the tissues from the IgAN and MN patients was significantly higher than that in the tissues from the MCG patients and the HCs (\( *p < 0.05 \), Fig. 3A1). In contrast, there was no difference in pGSN deposition between the MN and IgAN tissues. There were also no differences in actin expression among the IgAN, MN, MCG and HC groups. The IgA fluorescence in the IgAN group was stronger than in the MN, MCG and HC groups (\( **p < 0.01 \), Fig. 3A2). pGSN was mainly deposited in the area of mesangial proliferation of the glomeruli or the epithelial cells of the tubules in IgAN (Fig. 3B1-3B4). The pGSN levels in the renal tissue were not significantly higher in IgAN patients with E1 and T1 compared to those with E0 and T0 (Table 1, Fig. 3B5, \( p > 0.05 \)) but were significantly higher in IgAN patients with M1 and S1 compared to those with M0 and S0 (Table 1, Fig. 3B5, \( *p < 0.05 \)).

TGF-β1 and FN were mainly expressed in the glomeruli and were weakly expressed in the tubules (Fig. 4A-4D). The pGSN levels and both the TGF-β1 and the FN levels in renal tissues were positively correlated (Fig. 4E, 4F, \( r = 0.2183, *p = 0.0191 \) and \( r = 0.1851, *p = 0.0477 \), respectively).
Correlations between the serum and renal tissue pGSN levels and the clinicopathological data of IgAN patients

Serum and kidney pGSN levels were negatively correlated (Fig. 5A, \( r = -0.2733, * p = 0.0031 \)), whereas serum pGSN levels and the estimated glomerular filtration rate (eGFR) were positively correlated (Fig. 5B, Table 2, \( r = 0.3033, * p = 0.0126 \)). However, there were no correlations between the serum BUN and SCr values, CO\(_2\)CP, or the 24-h urine protein level and pGSN levels (Table 2, \( r = -0.2394, p = 0.0510 \); \( r = -0.207, p = 0.0954 \); \( r = -0.05544, p = 0.6559 \); and \( r = -0.05544, p = 0.6559 \)) respectively.

Table 1. Relationship between pGSN in renal tissue and pathologic variables of IgAN

<table>
<thead>
<tr>
<th>P value</th>
<th>MEST</th>
<th>pGSN (gray scale)</th>
</tr>
</thead>
<tbody>
<tr>
<td>M</td>
<td>*p&lt;0.05</td>
<td>M0 (n=61) 38.54±11.956</td>
</tr>
<tr>
<td></td>
<td></td>
<td>M1 (n=139) 47.90±10.103</td>
</tr>
<tr>
<td>E</td>
<td>NS</td>
<td>E0 (n=99) 38.75±12.248</td>
</tr>
<tr>
<td></td>
<td></td>
<td>E1 (n=101) 42.20±11.946</td>
</tr>
<tr>
<td>S</td>
<td>*p&lt;0.05</td>
<td>S0 (n=144) 37.68±12.128</td>
</tr>
<tr>
<td></td>
<td></td>
<td>S1 (n=56) 47.80±8.889</td>
</tr>
<tr>
<td>T</td>
<td>NS</td>
<td>T0 (n=162) 39.46±11.823</td>
</tr>
<tr>
<td></td>
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<td>T1 (n=38) 40.97±12.363</td>
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</table>
Correlations between pGSN and HMC proliferation

We chose two different time points (24 and 48 h) at which to add 10 mg/L pGSN to stimulate MC proliferation. pGSN promoted the proliferation of HMCs in vitro, especially at 48 h (Fig. 6A, two-way ANOVA, *p < 0.05). Cell cycle phases were also detected in the MCs that were treated with pGSN (10 mg/L) at 3 h, 6 h, 9 h or 12 h. After pGSN treatment for 6 h, the percentage of cells in the S phase of the cell cycle increased, while the percentage

Table 2. Relationships between the serum pGSN levels and clinicopathological data of IgAN patients. (a) Reference value for hypertension: >140/90 mmHg. (b) Reference value for BUN: 2.9-8.2 mmol/L. (c) Reference value for Scr: 45-104 mmol/L. (d) Normal range of CO₂ CP: 22-30 mmol/L. (e) 24-h urinary protein: 0.15 g. (f) eGFR (mL/min/1.73 m²)

<table>
<thead>
<tr>
<th>Clinicopathological data</th>
<th>n=200 (%)</th>
<th>Correlations between the serum pGSN level and IgAN clinicopathological variables</th>
</tr>
</thead>
<tbody>
<tr>
<td>Symptom before seeing a doctor</td>
<td>Cases (%)</td>
<td>Mann-Whitney U test</td>
</tr>
<tr>
<td>Edema</td>
<td>152 (76)</td>
<td>p=0.013*</td>
</tr>
<tr>
<td>Flu and fever</td>
<td>99 (49.5)</td>
<td>p=0.636</td>
</tr>
<tr>
<td>Pharyngalgia and antidiocncus</td>
<td>85 (42.5)</td>
<td>p=0.177</td>
</tr>
<tr>
<td>Gross hematuria</td>
<td>61 (30.5)</td>
<td>p=0.663</td>
</tr>
<tr>
<td>Edema</td>
<td>31 (15.5)</td>
<td>p=0.899</td>
</tr>
<tr>
<td>Kidney function change</td>
<td></td>
<td></td>
</tr>
<tr>
<td>1) BUN</td>
<td>r=0.2394, p=0.0510</td>
<td></td>
</tr>
<tr>
<td>2) Scr</td>
<td>r=0.207, p=0.0954</td>
<td></td>
</tr>
<tr>
<td>3) CO₂ CP</td>
<td>r=0.0554, p=0.6559</td>
<td></td>
</tr>
<tr>
<td>4) 24-h urine protein level</td>
<td>r=0.1392, p=0.1379</td>
<td></td>
</tr>
<tr>
<td>5) eGFR</td>
<td>r=0.3033, p=0.0126</td>
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</tbody>
</table>

The relationships between the pGSN concentration and major clinical symptoms (e.g., upper respiratory tract infection, sore throat, hypertension, edema and gross hematuria) were also analyzed (Table 2). There was a significant difference in the serum pGSN levels between patients with and without edema (Mann-Whitney U-test, *p < 0.05). The total number of patients in each group (n = 200) was used for statistical analysis.
Fig. 5. Analysis of the correlations between the serum pGSN levels and both the pGSN levels in the glomerulus and the eGFR. (A) Analysis of the correlation between the serum pGSN levels and the pGSN levels in renal tissue ($r = -0.2733$, *$p = 0.0031$). pGSN was specifically deposited in the glomerulus. (B) Analysis of the correlation with eGFR ($r = 0.3033$, *$p = 0.0126$). The correlations between the results were evaluated using Pearson’s correlation. $p < 0.05$ was considered significant. The total number of patients in each group ($n = 200$) was used for statistical analysis.

Fig. 6. pGSN promoted the proliferation of HMCs by facilitating cell mitosis. pGSN promoted the proliferation of HMCs (*$p < 0.05$, two-way ANOVA). The HMCs were incubated with 10 mg/L pGSN for 24 h, 48 h, 72 h or 96 h. (B) Effects of pGSN on the expression of cell cycle regulatory proteins. MCs were treated with pGSN (10 mg/L) at 3 h, 6 h, 9 h or 12 h. Western blot analysis showed the levels of CDK2 and cyclin A in MCs with or without pGSN treatment. (C-F) pGSN promoted the mitosis of HMCs. Cell cycle phases were detected in MCs that were treated with pGSN (10 mg/L) at 3 h, 6 h, 9 h or 12 h. (C) Detection of the cell cycle in HMCs treated with or without pGSN. (D) Dynamic change in the percentage of cells in the G1 phase of the cell cycle in HMCs treated with or without pGSN at different time points. (E) Dynamic change in the percentage of cells in the S phase of the cell cycle in HMCs treated with or without pGSN at different time points. (F) Dynamic change in the percentage of cells in the G2/M phase of the cell cycle in HMCs treated with or without pGSN at different time points.
in the G2/M phase decreased, which suggested that pGSN could promote the mitosis of HMCs (Fig. 6C, 6D-F). Furthermore, western blotting revealed that pGSN treatment resulted in a significant increase in the expression of CDK2 and cyclin A at different time points (3 h, 6 h, 9 h and 12 h) and that pGSN also promoted the mitosis of HMCs in association with regulation of CDK2 and cyclin A expression (Fig. 6B, t-test, \( * p < 0.05, ** p < 0.01 \)).

**Correlations between pGSN and integrin \( \alpha 2 \beta 1 \) expression in HMCs**

Integrins \( \alpha 2 \beta 1 \) were obviously up-regulated after pGSN treatment (Fig. 7A), and the integrin \( \alpha 2 \beta 1 \)-GSN interaction was enhanced after pGSN treatment or co-treatment with pGSN and aIgA1 (Fig. 7B).

**Discussion**

IgAN is the most common glomerular disease in the world. The pathological characteristics of IgAN include IgA immune complex deposition in the mesangium, glomerular MC proliferation, expansion of mesangial regions, massive ECM accumulation and renal fibrosis. MCs play important roles in renal physiological functions, including by regulating intraglomerular capillary flow, phagocytosing and clearing foreign bodies, secreting cytokines [16], and generating ECM. However, excessive MC proliferation can lead to glomerular fibrosis. A lack of effective biomarkers limits the capacity to evaluate MC proliferation, glomerular fibrosis stages and IgAN progression, especially during the period before clinical symptoms become evident. Renal biopsy remains the gold standard for the assessment of the pathological degree in IgAN patients; nonetheless, this procedure is invasive and is associated with certain risks and sampling variabilities. In addition, the diagnostic accuracy of renal biopsy depends on the size and position of the biopsy specimen, and biopsy diagnoses made by different pathologists may be discordant. The present study represents the first proposal of potentially using pGSN as a biomarker for MC proliferation in IgAN patients. Specifically, our findings suggest the potential utility of pGSN as a prognostic indicator in IgAN.

GSN is an important actin-binding protein and a regulator of cellular cytoskeleton dynamics. In particular, pGSN is secreted into extracellular fluids. Serum pGSN levels decrease markedly in a variety of clinical conditions, such as lung injury [17], hepatitis B-associated liver cirrhosis [18], sepsis [19], major trauma [20], brain injury [21, 22], certain cancers [23, 24], Alzheimer’s disease [25], rheumatoid arthritis [11] and multiple sclerosis [26]. Furthermore, certain studies have demonstrated that GSN is up-regulated in patients with idiopathic pulmonary fibrosis (IPF) or fibrotic nonspecific interstitial pneumonia [27, 28]. Another study found that the serum pGSN levels decreased in patients with hepatitis B virus (HBV)-related cirrhosis and concluded that pGSN is a new candidate fibrosis marker in this condition [18]. However, the function of pGSN in IgAN has remained unclear.

Similarly, in our study, the serum pGSN levels were significantly reduced in the IgAN patients compared to the patients with other types of nephropathy and HCs. We also found
that the pGSN levels were significantly higher in the kidneys of the IgAN group than in the kidneys of the MCG and HC groups. Furthermore, we found a negative correlation between the serum and kidney pGSN levels. Finally, we observed that pGSN promotes the proliferation of HMCs by facilitating cell mitosis in association with regulation of CDK2 and cyclin A expression, which in turn causes accumulation of the cytokines (such as TGF-β) secreted by HMCs and thus promotes the evolution of IgAN. These data indicated that serum pGSN levels correlate with MC proliferation and indirectly reflect glomerular fibrosis in IgAN.

In our study, the serum IgA and Gd-IgA1 levels in the IgAN patients were higher than those in the other groups. Moreover, the serum pGSN levels were negatively correlated with IgA in the IgAN patients, indicating that pGSN participates in the pathogenesis of IgAN. The above results suggest that IgA may interact with pGSN through an unknown mechanism. In addition to the transferrin receptor, integrins α1β1 and α2β1 are new candidate IgA1 receptors in human MCs [29], and GSN plays an important regulatory role in the integrin α2β1-mediated initial steps of collagen fibril attachment and engulfment in fibroblasts [30, 31]. To observe the receptors on HMCs treated with pGSN in the current study, integrins alpha1, 2/ beta1, 2 were detected by western blotting. Integrins α2β1 were obviously up-regulated on the MCs after pGSN treatment, confirming the preferential interaction between GSN and integrins α2β1. We deduced that IgA1 and/or pGSN may combine with the integrin receptors to promote HMC proliferation and ECM production. We also added IgA1 from IgAN patients to MCs to mimic the environment in IgAN, and the interaction between GSN and integrins α2β1 was remarkably strengthened after co-treatment with pGSN and IgA1. This result further demonstrated that pGSN is easily deposited in the renal tissue in IgAN.

TGF-β1 is a key factor in the pathogenesis of IgAN, and its levels correlate with the progress of IgAN [7]. The pathogenesis of TGF-β1/Smad-mediated glomerular fibrosis has specifically been characterized [32]. The TGF-β1/Smad signaling pathway adjusts the transcription and expression of its target genes, such as FN. FN is an important component of the ECM that exists as a soluble glycoprotein in the blood and other body fluids. The insoluble form of FN constitutes a major ECM protein in the course of glomerular fibrosis and sclerosis. In particular, the up-regulation of FN indicates excessive ECM deposition, eventually leading to glomerular sclerosis and renal fibrosis. In our study, the TGF-β1 and FN levels in the sera and glomeruli were significantly higher than what would be associated with the degree of glomerular fibrosis. Moreover, the serum levels of TGF-β1 and FN and the serum pGSN levels were negatively correlated, whereas the levels of these factors in the glomeruli were positively correlated. The pGSN levels in the glomeruli of the IgAN patients reflected the proliferation of the MCs that induce glomerular fibrosis.

In IgAN patients, the serum pGSN levels were significantly reduced and were positively correlated with the eGFR. Repeated attacks of gross or microscopic hematuria are common symptoms of IgAN; however, edema (76.84%) is another indicator of IgAN that is often ignored, particularly in the early stages. In the present study, the pGSN levels in patients with edema were significantly decreased compared to the levels in patients without edema. A low pGSN level in the serum indicates serious renal injury, which could easily induce edema. The correlation between serum pGSN levels and several critical clinical conditions suggests the potential utility of pGSN as a prognostic marker.

In summary, our study revealed that the serum pGSN concentration reflected the progress and prognosis of IgAN; a low serum pGSN level specifically indicated a severe degree of proliferation of MCs and a worse prognosis. pGSN promotes the proliferation of MCs, thus contributing to IgAN progression, and the integrin α2β1 receptors may be the main binding sites for pGSN on MCs. Monitoring the pGSN levels in IgAN patients could thus help to predict prognosis and glomerular fibrosis and could be used to assess the progression of IgAN.

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Disclosure Statement

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References

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