Plasma ST6GAL1 regulates IgG sialylation to control IgA nephropathy progression

Youxia Liu^D, Huyan Yu, Sijing Wu, Xia Yang, Congcong Cao, Fanghao Wang, Junya Jia and Tiekun Yan

Abstract

Background: Our previous study revealed that plasma levels of a-2,6-sialyltransferase 1 (ST6GAL1) were increased in patients with IgA nephropathy (IgAN). ST6GAL1 catalyzes terminal sialylation of IgG to shift the antibody effector function to the anti-inflammatory pattern. However, the role of plasma ST6GAL1 in the progression of IgAN and underlying mechanisms are still unknown.

Methods: A total of 180 IgAN patients were included. The kidney outcomes were defined as the eGFR decline or proteinuria remission. Peripheral blood mononuclear cells (PBMCs) were either stimulated with purified sialylated IgG (SA-IgG) or with non-sialylated IgG (NSA-IgG) from IgAN patients to detect the levels of interleukin (IL)-6 and tumor necrosis factor- α (TNF- α) in supernatant.

Results: Compared with the lower ST6GAL1 (reference), the risk of eGFR decline decreased for the higher ST6GAL1 group after adjustment for baseline eGFR, systolic blood pressure (SBP), and proteinuria. The results showed that patients with higher ST6GAL1 levels had a higher rate of proteinuria remission. ST6GAL1, expressed as a continuous variable, was a protective factor for eGFR decline and proteinuria remission. An *in vitro* study showed that the administration of recombinant ST6GAL1 (rST6GAL1) decreased the levels of IL-6 and TNF- α in PBMCs. Furthermore, the administration of rST6GAL1 resulted in the enrichment of SA-IgG in a concentration-dependent manner. In addition, as compared to control, purified SA-IgG-treated PBMCs showed a significant decrease in the expression of IL-6 and TNF- α .

Conclusion: Our study indicated that elevated ST6GAL1 was associated with a slower progression of IgAN, which may play a protective effect by increasing IgG sialylation to inhibit the production of proinflammatory cytokines in PBMCs.

Keywords: IgA nephropathy, IL-6 and TNF-a, kidney outcomes, sialylated IgG, ST6GAL1

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Introduction

Immunoglobulin A nephropathy (IgAN) is the most common glomerulonephritis in the world, with 10–20% of patients progressing to end-stage kidney disease (ESKD) within 10 years after diagnosis.¹ A combination of four processes induces renal injury that culminates in IgAN: Hit 1: production of aberrant glycosylation of IgA1 (Gd-IgA1); Hit 2: synthesis of antibodies (mainly IgG) directed against Gd-IgA1; Hit 3: binding

of the Gd-IgA1 with its antibodies to form circulation immune complexes (CICs); and Hit 4: accumulation of CIC in the glomerular mesangium to initiate renal injury.^{2–4} More and more evidence showed that abnormal release of proinflammatory cytokines, including interleukin-6 (IL-6) and tumor necrosis factor- α (TNF- α) in peripheral blood mononuclear cells (PBMCs) play an important role in the pathogenesis and progression of IgAN.^{5–9} The risk of kidney failure Ther Adv Chronic Dis

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under the current therapy regimen remains high; new interventions are urgently needed to slow the progression of the disease.¹⁰

ST6GAL1 encoding β-galactoside α2,6sialyltransferase mediates the transfer of sialic acid residue with an α -2,6-linkage to a terminal galactose. Genome-wide association studies (GWASs) were performed in the Chinese Han population, which identified the risk allele C at rs7634389 at the ST6GAL1 locus was associated with the susceptibility and slower progression of IgAN.^{11,12} Our previous results showed that administration of recombinant ST6GAL1 (rST6GAL1) inhibited the production of galactose-deficient IgA1 (Gd-IgA1) in cultured PBMCs from IgAN patients.¹³ However, whether increased ST6GAL1 levels are associated with a slower disease progression in IgAN is unknown.

ST6GAL1 has a modulatory role in the antiinflammatory activity of sialylated IgG (SA-IgG).14 Engineered sialvlation of IgG with ST6GAL1 shifts the antibody effector function to the anti-inflammatory pattern in a mice model of Goodpasture disease, thus effectively attenuating disease severity.¹⁵ Elevated ST6GAL1 inhibits the production of the proinflammatory cytokines in patients with rheumatoid arthritis (RA) by inducing IgG-Fc sialylation.¹⁶ Moreover, several clinical observations indicated that patients with progressive RA have poorly SA-IgG compared with patients with less-severe disease or those in remission.17,18 Whether ST6GAL1 can play antiinflammatory effects by increasing sialylation of IgG in IgAN has not been investigated yet.

In this study, we examined the prognostic utility of plasma levels ST6GAL1 in 180 patients with IgAN. Furthermore, we determined whether the addition of ST6GAL1 played protective effects by an increase in IgG sialylation *in vitro*.

Materials and methods

Sample collection

In this retrospective case–control study, 180 patients with IgAN diagnosed by renal biopsy, and 50 age- and gender-matched healthy participants were included. Of these participants, 10 patients with IgAN and 10 healthy participants were randomly selected to isolate IgG. The

diagnosis of IgAN was based on the presence of IgA deposition in the glomerular mesangium by immunofluorescence and electron-dense material deposition in the mesangium by electronic microscopy. The patients with Henoch-Schönlein purpura and other secondary IgAN were excluded. Plasma from all participants was collected. Written informed consent was obtained from each patient and healthy participant. Clinical informations, including age, gender, blood pressure, 24-h urine protein excretion, serum creatinine, estimated glomerular filtration rate (eGFR), total IgA levels, were collected at the time of renal biopsy. The eGFR was calculated using the Chronic Kidney Disease Epidemiology Collaboration (CKD-EPI) equation.¹⁹ The histological lesions were classified according to Oxford classification-MEST-C (M: mesangial hypercellularity; E: endocapillary hypercellularity; S: segmental glomerulosclerosis; T: tubular atrophy/interstitial fibrosis, and C: crescent).²⁰

Outcomes

The eGFR decline was evaluated as \triangle eGFR, which was calculated as eGFR in the next year minus eGFR at baseline per month.²¹ We defined proteinuria remission as a more than 50% reduction in proteinuria from baseline or proteinuria <300 mg/d.

Isolation of SA-IgG and non-sialylated IgG

The total serum immunoglobulin was separated and purified with a protein G affinity column (Nunc, Rochester, USA). Briefly, the phosphatebuffered saline (PBS)-diluted plasma were added to protein G affinity columns and eluted by washing the affinity columns with 10 mL of buffer B(0.2 M glycine/in chloride acid, pH 2.5). Purified IgG was desalted by PD-10 column. Draw the bulk of IgG into the prepared sambucus nigra lectin (SNA)-agarose column and let the buffer drain by gravity. The outflowing liquid was centrifuged to collected non-sialylated IgG (NSA-IgG). SA-IgG was eluted by adding the eluting solution (VECTOR Labs, USA), followed by washing with PBS. Fractions containing IgG were concentrated by a protein centrifugal filter unit (Millipore, USA). Enrichment of the eluate was verified by Western blotting (WB) with biotinylated SNA, IgG antibody, and horseradish peroxidase (HRP)-conjugated streptavidin.

ELISA assay for SA-IgG in cell supernatant

The level of SA-IgG in cell supernatant was determined using enzyme-linked immunosorbent assay (ELISA) according to the slightly modified procedure (ab195215; Abcam, USA). Assay diluent cell culture supernatants were added into appropriate wells and incubated for 2.5 h at room temperature. After washing with wash solution, the plate was added 100 µL of 1:50 diluted SNA (VECTOR, USA) for 2 h. HRP-Streptavidin solution was incubated for 45 mins. After TMB (3,3', 5,5;-tetramethylbenzidine) solution incubation for 30 mins, the reaction was stopped with a stop solution. The optical densities (ODs) were determined at 450 nm with an EL312 Bio-Kinetics microplate reader (Bio-TekInstruments, Winooski, VT).

ELISA assay for ST6GAL1, IL-6, and TNF- α

The ST6GAL1 levels from plasma and IL-6 and TNF- α levels from cell supernatants were measured using commercial ELISA kits (R&D Systems, USA) according to the manufacturer's specifications. At last, the absorbance was detected at 450 nm with an EL312 Bio-Kinetics microplate reader (Bio-TekInstruments, Winooski, VT).

PBMCs culture and treatment

About 5-mL peripheral blood was taken into ethylenediaminetetraacetic acid (EDTA)-anticoagulated tubes. PBMCs were separated by density-gradient centrifugation on Ficoll (TBD, China) and resuspended in PBS. In the in vitro experiment, PBMCs were seeded into 24-well plates. The cells were stimulated with lipopolysaccharide (LPS) (Sigma-Aldrich, USA) and incubated with 0, 500, 1000 ng/mL human recombinant ST6GAL1 (rST6GAL1, R&D Systems, USA) and CMP-sialic acid (Sigma, USA) at a final concentration of 1.0 mM for 24 h to detect SA-IgG levels. PBMCs were incubated with purified SA-IgG (0.5 mg/mL) and NSA-IgG (0.5 mg/mL) for 24 h to detect the levels of IL-6 and TNF- α in cell supernatants.

Statistical analysis

For continuous variables, data with a normal distribution was expressed as the mean \pm SD and compared by an unpaired *t*-test. For nonnormally distributed variables, data were expressed as the median (first quartile and third quartile) and analyzed by the Mann-Whitney U-test. Categorical variables were summarized as proportions and were compared by a chi-square test. Correlation analysis was performed by Pearson's correlation test. Linear regression analyses were used to evaluate levels of ST6GAL1 on the rate of eGFR decline. Kaplan-Meier analysis was used to compare proteinuria remission with different levels of ST6GAL1. We estimated the regression coefficient and the corresponding 95% confidence interval (CI) between ST6GAL1 and eGFR decline rates with unadjusted and multivariable adjusted linear regression. The odds ratio (OR) and the 95% CI between ST6GAL1 and proteinuria remission were examined in logistic models. Power calculations were used to determine the number of participants required with 0.90 power at an alpha level of 0.05. A two-tailed p value < 0.05 was considered statistically significant. Statistical analysis was performed using SPSS, version 16.0 software.

Results

Baseline clinical and pathological data

The demography characteristics of the study population, clinical findings, and pathological lesions at the time of renal biopsy were listed in Supplement Table 1. There were 91 males (51%) with a mean age of 40 ± 13 years in patients with IgAN and were 22 males (44%) with a mean age of 38 ± 10 years in healthy controls. The average systolic blood pressure (SBP) was 132 ± 17 mmHg and diastolic blood pressure (DBP) was 82 ± 11 mmHg. The mean eGFR was 93.5 ± 44.29 mL/min per 1.73 m², and the proteinuria level was 1763.35 ± 1721.48 mg/d. After a mean follow-up of 17 months, 85 (47%) participants reached the proteinuria remission.

Plasma ST6GAL1 levels were correlated with the severity of IgAN

The mean ST6GAL1 level in plasma in participants with IgAN at biopsy was 6914 ± 2624 pg/mL, significantly higher than that of healthy controls $(4321 \pm 827 \text{ pg/mL}, p < 0.001$, Supplement Figure 1). The patients were then divided into two equal subgroups according to the median level of plasma ST6GAL1 levels (low ST6GAL1 < 6412 pg/mL and high ST6GAL1 > 6412 pg/mL) to explore the association of plasma ST6GAL1 with clinical pathological lesions in patients with IgAN (Table 1). We found that patients with higher ST6GAL1 levels had higher levels of triglyceride, serum creatinine, uric acid, serum IgA, serum C3, serum C4 and proteinuria, and lower levels of eGFR and IgG compared with those patients with lower SA-IgG levels. As for the Oxford classification, patients with higher ST6GAL1 levels tend to have a higher proportion of glomerulosclerosis.

Plasma ST6GAL1 levels were negatively correlated with kidney disease progression

Mean rates of eGFR decline over 1 year in the higher and lower ST6GAL1 level groups, measured by the average of \triangle eGFR, were 0.25 and -0.76 mL/min/1.73 m², respectively. Patients with lower levels of ST6GAL1 had a more rapid decline in the eGFR (p=0.03, Figure 1). Multiple linear regression analyses suggested a significant correlation of eGFR decline rate with ST6GAL1 after adjusting for baseline eGFR, SBP, and proteinuria (regression coefficient: 2.98×10^{-4} ; 95% CI: 4.45×10^{-4} to 1.51×10^{-4}). Compared with the lower ST6GAL1 (reference), the risk of eGFR decline decreased for the higher ST6GAL1 group (regression coefficient: 1.27; 95% CI: 0.48-2.05, Table 2).

The plasma ST6GAL1 level was separately tested for association with the proteinuria remission. Thirty-five patients (38%) got proteinuria remission in the lower ST6GAL1 group, and 50 patients (56%) reached remission in the higher ST6GAL1 group. Kaplan–Meier analysis showed that the proteinuria remission rate in the high ST6GAL1 group was significantly better than that in the low group (p=0.004, Figure 2).

The logistic regression analysis showed that compared with the low ST6GAL1 (reference), the rate of proteinuria remission increased for the high ST6GAL1 level after adjustment for baseline eGFR, SBP, and proteinuria (OR: 2.09; 95% CI: 1.01–4.31, Table 3).

Recombinant ST6GAL1 downregulated the secretion of IL-6 and TNF- α in PBMCs

To address the possible influence of rST6GAL1 on the production of proinflammatory cytokines, IL-6, and TNF- α levels were analyzed in supernatant from PBMCs. As depicted in Figure 3, rST6GAL1 at a concentration of 1000 ng/mL significantly decreased the production of IL-6 (p=0.015, Figure 3(a)) and TNF- α (p=0.017, Figure 3(b)) in PBMCs.

Recombinant ST6GAL1 upregulated SA-IgG expression in PBMCs

We detected the SA-IgG levels in culture supernatant of PBMCs under rST6GAL1 challenge. Our results indicated that in the presence of the corresponding CMP-sialic acid, ST6GAL1 resulted in enrichment of SA-IgG (p=0.025, Figure 4).

SA-IgG downregulated IL-6 and TNF- α secretion in PBMCs

As depicted in Figure 5, the releases of IL-6 and TNF- α were significantly decreased in PBMCs cultured with SA-IgG from patients with IgAN. The findings implied that SA-IgG protected against inflammatory response in PBMCs.

Discussion

Consistent with our previous findings,13 the results of this study reconfirmed that increased plasma levels of ST6GAL1 were associated with the severity of IgAN. Most importantly, it demonstrated that high levels of ST6GAL1 predicted a slower eGFR decline rate and a higher proteinuria remission rate in IgAN. Furthermore, the data provided evidence that ST6GAL1 increased sialylation of IgG-Fc to counteract the actions of the proinflammatory cytokines IL-6 and TNF- α in IgAN. Preliminary observations indicated that ST6GAL1 played protective effects in IgAN by an increase in IgG sialylation. It lends support to the notion that ST6GAL1 can be used as a promising addition to improve existing strategies for the treatment of IgAN.

Recently, several genetic studies explored the association between ST6GAL1 and IgAN pathogenesis and progression. A GWAS follow-up study including 613 patients with IgAN revealed that the patients carrying the risk allele rs7634389-C at the *ST6GAL1* locus had a decreased risk for the progression of IgAN.¹² Fu *et al.*²² reported that rs7634389-C allele was associated with increased susceptibility to IgAN, higher risks of hyperuricaemia, and segmental glomerulosclerosis. Interestingly, they found patients carrying the risk

Characters	Mean ± SD or <i>n</i> (%)		
	Low ST6GAL1 (<i>n</i> < 6412 pg/mL) High ST6GAL1 (<i>n</i> > 6412 pg/mL		_
Gender (M/F)	44 (49)/46 (51)	47 (52)/43 (48)	0.65
Age (mean \pm SD, year)	40.01 ± 13.11	39.84 ± 13.63	0.93
SBP (mmHg)	130.43±17.81	134.84 ± 17.4	0.09
DBP (mmHg)	80.59 ± 11.69	83.47 ± 11.72	0.1
BMI (kg/m²)	24.14±4.21	25.28 ± 3.35	0.06
Hemoglobin (g/L)	131.58±18.13	133.08 ± 20.13	0.6
Serum albumin (g/L)	37.04 ± 5.06	37.03 ± 5.03	0.98
Triglyceride (mmol/L)	1.69±1	2.78 ± 2.66	0.001
Serum creatinine (µmol/L)	87.47 ± 47.81	100.96 ± 47.78	0.06
eGFR (mL/min/1.73 m²)	102.18±51.62	84.81 ± 33.6	0.008
Uric acid (µmol/L)	357.21 ± 97.52	401.99 ± 107.09	0.004
Serum IgA (mg/dL)	313.22±121.19	573.74 ± 418.8	0.001
Serum IgG (mg/dL)	1051.3±261.4	889.42 ± 420.69	0.003
Serum IgM (mg/dL)	119.39±64.23	113.31 ± 54.85	0.513
Serum IgE (mg/dL)	66.4±167.19	158.64 ± 495.06	0.11
Serum C3 (mg/dL)	86.63±15.6	94.47 ± 17.86	0.003
Serum C4 (mg/dL)	21.95±5.95	25.15 ± 8.48	0.005
Proteinuria (mg/d)	1409.83±1373.14	2116.87 ± 1955	0.006
Urine RBC (/HP)	35.93±61.24	80.58 ± 257.26	0.11
Oxford classification			
M score (M0/M1)	11 (12)/79 (88)	10 (11)/80 (89)	0.82
E score (E0/E1)	53 (59)/ 37 (41)	52 (58)/38 (42)	0.89
S score (S0/S1)	44 (49)/46 (51)	28 (31)/62 (69)	0.02
T score (T0/T1/T2)	35 (39)/50 (56)/5 (5)	27 (30)/50 (55)/13 (15)	0.15
C score (C0/C1/C2)	23 (26)/54 (60)/13 (14)	32 (36)/50 (55)/8 (9)	0.25

 Table 1. The baseline data for IgAN patients with lower and higher ST6GAL1 levels.

BMI, body mass index; DBP, diastolic blood pressure; eGFR, estimated glomerular filtration rate; IgAN, immunoglobulin A nephropathy; RBC, red blood cell; SBP, systolic blood pressure; SD, standard deviation.



Figure 1. The correlation between ST6GAL1 and eGFR decline rate.

Characteristic	Mean ± SD (pg/mL)	Univariate analysis		Multivariate analy	Multivariate analysis ^a	
		β (95% CI)	р	β (95% CI)	р	
ST6GAL1	6914±2624	3.45×10^{-4} (1.92 × 10 ⁻⁴ to 4.98 × 10 ⁻⁴)	<0.001	2.98×10^{-4} (1.51 × 10 ⁻⁴ to 4.45 × 10 ⁻⁴)	<0.001	
ST6GAL1 dichotomy						
Lower level	5009 ± 951	1 (reference)		1 (reference)		
Higher level	8365 ± 2079	1.56 (0.75–2.37)	< 0.001	1.27 (0.48–2.05)	0.002	

Table 2. Univariate and multivariate linear regression analysis of ST6GAL1 influencing eGFR decline rate.

CI, confidence interval; eGFR, estimated glomerular filtration rate; SD, standard deviation. ^aAdjusted for eGFR, proteinuria, and systolic pressure.

allele rs7634389-C allele had a lower risk for disease progression.²² The authors explained that this discrepancy might be caused by the complex interactions between genetic, epigenetic, and environmental factors during the progression of IgAN.²² These findings indicated that genetic variation in *ST6GAL1* participated in the development and progression of IgAN. However, the plasma levels of ST6GAL1 were not explored in these studies. In this study, we expanded on the findings of previous studies by showing that the elevated levels of ST6GAL1 were associated with disease severity in patients with IgAN with larger sample sizes. More importantly, compared with patients with lower ST6GAL1, those with higher levels of ST6GAL1 had a lower eGFR decline rate after adjusting for confounding factors. The results stressed the importance of ST6GAL1 in delaying renal function decline in IgAN. There is growing evidence that suggests proteinuria



reduction can be used as a reasonably surrogate endpoint for a treatment's effect on progression to end-stage renal disease (ESRD) in patients with IgAN.²³ We showed a higher proteinuria remission rate in IgAN patients with higher plasma levels of ST6GAL1 than those in patients with lower levels of ST6GAL1. Together, the data suggest that high levels of baseline ST6GAL1 have a prognostic utility in IgAN patients with better outcomes.

The underlying protective mechanism of ST6GAL1 should be discussed. Previous research showed that lots of proinflammatory cytokines, IL-6 and TNF- α , were triggered in the development of IgAN and associated with the pathogenesis, aggravation, and progression of IgAN.^{6,9,24} The present data confirmed that the administration

Figure 2. Kaplan–Meier analysis of proteinuria remission in IgA nephropathy (IgAN) patients with the median plasma of ST6GAL1.

Table 3. Univariate and multivariate logistic regression analysis of ST6GAL1 at baseline influencing proteinuria remission.

Characteristic	Univariate analysis	variate analysis		Multivariate analysisª	
	OR (95% CI)	p	OR (95% CI)	p	
ST6GAL1	1.0001 (0.999–1.0004)	0.11	1.0001 (0.9999–1.0002)	0.1	
ST6GAL1 dichotomy					
Lower level	1 (reference)		1 (reference)		
Higher level	2.05 (1.03–4.06)	0.04	2.09 (1.01–4.31)	0.04	

CI, confidence interval; OR, odds ratio.

^aAdjusted for eGFR, proteinuria, and systolic pressure.







Figure 4. SA-IgG production after rST6GAL1 stimulation in supernatant of PBMCs.

of recombinant ST6GAL1 decreased IL-6 and TNF- α levels in PBMCs and advanced the idea that the ability of sialyltransferase ST6GAL1 to SA-IgG in vitro. Furthermore, we explored whether the anti-inflammatory effect of ST6GAL1 in IgAN was mediated by increasing sialylation of IgG. It showed that increased SA-IgG induced by ST6GAL1 decreased inflammation activation of PBMC with the downregulatory synthesis of IL-6 and TNF- α . The galactosylation is a prerequisite for sialylation of ST6GAL1 at the Asn297 site of IgG-Fc. In contrast to these findings in sialylation, we did not detect a difference in the expression of B4GALT1, the enzyme-mediating attachment of galactose to the IgG-Fc glycan, between IgAN patients and healthy control. The galactosylated IgG-Fc were then sialylated with 2,6 sialyltransferases The use of artificially sialylated IgG-Fc has been confirmed to increase the efficacy of anti-inflammatory treatment.^{25,26} The sialvlation of IgG reduced the pathogenicity of autoantibodies and diminished the arthritogenic activity in the collagen-induced arthritis (CIA) model.27 Immune complexes with antigenspecific SA-IgG inhibited the production of IL-6 in autoimmune mouse models of lupus nephritis and RA.28 Troelsen et al.29 showed that the reduction of IgG glycosylation in RA was correlated with IL-6 and C-reactive protein. These data suggested that upregulation of IgG sialylation modulating by ST6GAL1 was a potent approach to attenuate harmful autoantibodymediated inflammation in autoimmune disease. In this study, we also observed consistent

evidence for the anti-inflammation effect of SA-IgG. We speculate ST6GAL1 is actively upregulated in an effort to protect the body from damage. Plasma ST6GAL1 is increased in patients with an initial onset of IgAN and the more serious the disease, the higher the ST6GAL1 level. Furthermore, elevated ST6GAL1 catalyzes terminal sialylation of IgG to inhibit the production of inflammatory cytokines in PBMCs to delay the progression of IgAN. It indicated the therapeutic application of sialylation of IgG in IgAN.

Patients with IgAN are characterized by a highly variable clinical course ranging from a totally benign incidental condition to rapidly progressive renal failure. Many risk factors including high proteinuria, hypertension, low GFR, and high MEST-C score have been confirmed to predict the adverse kidney outcomes in IgAN.¹ However, the current therapy mainly focused on proteinuria reduction and antihypertension. No effective therapies are available. Disease-specific therapies which slow the progression of disease are urgently needed. Many pieces of clinical evidence suggested a pathogenic role of autoantibodies (mainly IgG) in IgAN. The serum levels of IgG autoantibodies at the time of biopsy were significantly associated with the clinical progression of IgAN.^{30,31} The modulation of sialylation patterns of IgAN-associated IgG autoantibodies with recombinant ST6GAL1 administration may be a possible novel potential candidate for therapeutic intervention of IgAN.

Limitations

There are several limitations in this study. First, the small sample size and the relatively short follow-up with fewer endpoints reduced the study power. On further expansion, the sample size and extension of the follow-up period would provide more accurate effect estimates of the associations between the levels of ST6GAL1 and disease severity and progression of IgAN. Second, we did not stimulate cells at different concentrations of SA-IgG because of the small amount of extraction. Third, the detailed mechanistic links of how cell surface sialylation affects cell behavior remain to be elucidated. Additional longitudinal studies of more diverse patient cohorts are needed to validate the protective role of ST6GAL1 in IgAN.



Figure 5. The expression of IL-6 (a) and TNF- α (b) after SA-IgG and NSA-IgG stimulation in supernatant of PBMCs.

Conclusion

Our study indicated that elevated ST6GAL1 was associated with a slower progression of IgAN. It played a protective effect by increasing IgG sialylation to inhibit the production of proinflammatory cytokines in PBMCs. This was the first study to highlight the critical role of ST6GAL1 in delaying the progression of IgAN, which provided a potential therapeutic target for the treatment of IgAN.

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Author contributions

Y.X.L. and J.Y.J. conceived the study, Y.X.L., J.Y.J., and T.K.Y. participated in its design and coordination. H.Y.Y., C.C.C., and F.H.W. collected clinical data. H.Y.Y., X.Y., and S.J.W. performed the experiment. Y.X.L. contributed to the writing of the manuscript. All authors read and approved the final manuscript for submission.

Ethical approval

All subjects provided written informed consent. The study protocol was approved by the Institutional Ethical Committee of Tianjin Medical University General Hospital (approval id: ZYY-IRB-SOP-016(F)-002-04).

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Conflict of interest statement

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Availability of data and materials

Raw data used during this study are available from the corresponding author on reasonable request for non-commercial use.

Supplemental material

Supplemental material for this article is available online.

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