Increased frequencies of memory and activated B cells and follicular helper T cells are positively associated with high levels of activation-induced cytidine deaminase in patients with immunoglobulin A nephropathy

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Received August 20, 2014; Accepted June 9, 2015

DOI: 10.3892/mmr.2015.4071

Abstract. The aim of the present study was to examine the frequencies of different subsets of B and follicular helper T (Tfh) lymphocytes in patients with immunoglobulin A nephropathy (IgAN), and investigate the potential underlying mechanism. A total of 27 patients with IgAN and 10 healthy controls (HC) were recruited for analysis of the frequencies of different subsets of B and Tfh cells. ELISA was used to analyze the concentration of serum interleukin (IL)-21. The transcriptional levels of activation-induced cytidine deaminase (AID) in the B cells were determined using reverse transcription-quantitative polymerase chain reaction, while the translational levels of AID were analyzed using western blotting. The frequencies of circulating memory and activated B cells and Tfh cells were found to be significantly increased in the IgAN groups, compared with those of the HC group, although the number of plasma cells were not significantly different between the two IgAN groups. In addition, the serum levels of IL-21 were found to be higher in the patients with IgAN, and correlated with 24-h proteinuria. IL-21 also enhanced the expression levels of AID in the B cells. The data of the present study revealed that the high levels of memory and activated B cells and Tfh cells were positively associated with the progression of IgAN, and that this may be mediated by the overexpression of AID, which is potentially regulated by IL-21.

Introduction

Immunoglobulin A nephropathy (IgAN) is considered to be the most common type of glomerular disease worldwide (1). IgAN has been confirmed to be an immune complex-mediated glomerulonephritis, defined morphologically by the mesangial deposition of IgA (2,3). Although defects in immune regulation are considered to be important in the pathogenesis of IgAN, the pathogenetic mechanisms remain to be fully elucidated. Previous studies have reported that IgAN is regulated by B lymphocytes (4,5), and increased numbers of T helper (CD4) lymphocytes and reduced numbers of T suppressor (CD8) lymphocytes are associated with the exacerbation of IgAN (6,7). However, overproduction of IgA is most likely the consequence of the involvement of both T and B lymphocytes.

The present study aimed to explore immune status alterations during the progression of IgAN disease. The distributions of different B-cell subsets and Tfh cells were analyzed in patients with IgAN at various disease phases, and the differential contributions of these lymphocytes to IgAN were evaluated. To explain the imbalance of B-cell subsets, the main effector of Tfh cells, IL-21 was also investigated.

IL-21 is not a classic T helper cell 1 or 2 cytokine, but is mainly produced by CD4+ follicular T helper cells, which can be identified by their expression of the chemokine receptor CXCR5. The high levels of IL-21 receptor (IL-21R) expressed by B cells make B cells prime responders to IL-21. B cells faced with IL-21 in the context of antigen-specific BCR stimulation and T cell co-stimulation undergo class switch recombination and differentiate into antibody producing plasma cells (8,9). Activation-induced cytidine deaminase (AID) induces somatic hypermutation, gene conversion, and class-switch recombination of immunoglobulin genes in B cells (10,11). Thus, it is important to clarify the association between IL-21 and AID expression, which may aid understanding of the functions of T helper cells and different B-cell subsets in the process of IgAN disease.
Patients. A total of 27 patients diagnosed with IgAN were recruited for investigation from the Department of Gastroenterology at the Affiliated Hospital of Changchun University of Chinese Medicine (Changchun, China) between 2012 and 2013. The diagnoses of IgAN in the patients were confirmed by biopsy and the presence of proteinuria. The patients with IgAN were divided into two groups, according to the extent of proteinuria: Group A (n=12), proteinuria <4 g/24 h; group B (n=15), proteinuria ≥4 g/24 h. Patients who had received immunosuppressive therapies in the 6 months preceding the investigation were excluded. A total of 10 gender- and age-matched healthy volunteers were selected as the HC group, each of which were recruited from the Department of Physical Examination Center of the Affiliated Hospital of Changchun University of Chinese Medicine. None of the patients or controls had any systemic disorders, viral infections or other autoimmune diseases. All study participants provided written informed consent, and the ethical committee of the Affiliated Hospital of Changchun University of Chinese Medicine approved the experiment. The clinical characteristics of the patients and HC individuals are presented in Table I.

PBMC culture and stimulation. Peripheral blood mononuclear cells (PBMCs) were isolated from venous blood samples of all patients using standard Ficoll-Paque Plus (GE Healthcare Life Sciences, Pittsburgh, PA, USA) density-gradient centrifugation (1,500 x g for 10 min at room temperature). Subsequent to washing in phosphate-buffered saline (PBS), the PBMCs were diluted at 4x10^6/ml in Dulbecco’s modified Eagle's medium (GE Healthcare Life Sciences, Logan, UT, USA), containing 10% fetal bovine serum ( Gibco Life Technologies, Carlsbad, CA, USA) and penicillin-streptomycin (100 U/ml) solution (GE Healthcare Life Sciences) and were cultured in 24-well plates (Corning Incorporated, Corning, NY, USA). The isolated PBMCs were cultured with CpGB (3 µg/ml; R&D Systems, Inc., Minneapolis, MN, USA) and 10 ng/ml recombinant interleukin (IL)-2 (R&D Systems, Inc.) for 72 h to detect B cell sub-populations. In order to stimulate T cells, 50 ng/ml phorbol myristate acetate, 1.0 mg/ml ionomycin and 5.0 mg/ml lipopolysaccharide (all from Sigma-Aldrich, St. Louis, MO, USA) were added to each well and they were incubated for an additional 4 h at 37˚C.

Flow cytometric analysis. To analyze the distribution of different B cells subsets and Th cells, the PBMCs were stimulated in vitro, as described above, and then harvested, washed with ice-cold PBS, and stained with fluorescein-labeled monoclonal antibodies (1:500 dilution) against various B cell markers [anti-CD19 PerCP (cat. no. 340421), anti-CD27 APC (cat. no. 561786), anti-CD86cy5.5 (cat. no. 561129), anti-CD138 fluorescein isothiocyanate (cat. no. 561703)] and Th cells [anti-CXCR5 PerCP (cat. no. 562781) and anti-CD4 APC (cat. no. 340443)]. All antibodies were from BD Pharmingen (San Diego, CA, USA). The control PBMCs were cultured in medium alone. Following a 1 h incubation with the primary antibodies at room temperature, the cells were washed with PBS, and at least 20,000 events were recorded. The data was obtained using a FACSCalibur analytical instrument (BD Biosciences) and four-color analysis was performed using FlowJo software, version 7.6 (FlowJo, LLC, Ashland, OR, USA).

ELISA. The serum level of IL-21 was quantified in the IgAN patients and HC individuals using a Human IL-21 ELISA kit, according to the manufacturers’ instructions (Roche Diagnostics, Ltd, Burgess Hill, UK). The plate was read at 450 nm using an Infinite M200pro plate reader (Tecan, Männedorf, Switzerland) and the sensitivity of the ELISA kits used in the experiment was 19 pg/ml. All samples were analyzed in duplicate using the average optical density values to calculate the concentrations.

Determination of the effect of IL-21 on levels of AID. To detect the expression of AID, the B cells were collected from the PBMCs of the HC group, were stained with anti-CD19 PerCP and anti-CD3 PE antibodies, and were sorted using a FACS Aria flow cytometer (BD Biosciences). The purified B cells were identified as CD3-CD19+ cells. The isolated B cells were cultured with recombinant human IL-21 (20 ng/ml; Peprotech, Inc., Rocky Hill, NJ, USA), CpGB (3 µg/ml) and recombinant IL-2 (10 ng/ml) for 72 h, and were then lysed on ice for 30 min in radioimmunoprecipitation assay buffer (Beyotime Institute of Biotechnology, Haimen, China), containing 50 mM Tris-HCl (pH 7.4), 150 mM NaCl, 0.1% SDS, 1% deoxycholate, 1% Triton X-100, 1 mM EDTA, 5 mM NaF, 1 mM sodium vanadate and protease inhibitor cocktail. Following quantification of the proteins with a Bicinchoninic Acid Protein Assay kit (Beyotime Institute of Biochemistry), the total proteins (70 µg) were then subjected to 10% SDS-PAGE (Beyotime Institute of Biotechnology), and transferred onto a nitrocellulose membrane (Pall Corporation, Port Washington, NY, USA). The membrane was incubated with primary antibodies targeting AID and GAPDH (1:2,000 dilution; cat. nos. sc-25620 and sc-25778; Santa Cruz Biotechnology, Inc., Dallas, TX, USA). The animal-matched horseradish peroxide-conjugated secondary antibody was also purchased from Santa Cruz Biotechnology, Inc. (1:2,000 dilution; cat. no. sc-2370). The total RNA of these cultured cells was extracted using TRIzol reagent (Invitrogen Life Technologies, Carlsbad, CA, USA). Total RNA (1 µg) was used for cDNA synthesis using the RevertAid First Strand cDNA Synthesis kit (Thermo Fisher Scientific, Inc., Waltham, MA, USA). PCR was performed using Hot Start Taq DNA Polymerase (Takara Bio Inc., Otsu, Japan), and PCR cycling conditions were as follows: 94°C for 3 min, followed by 35 cycles at 94°C for 30 sec, 58°C for 30 sec and 72°C for 45 sec, and a final extension step at 72°C for 10 min. Differences in the expression levels of cDNA were normalized to GAPDH. The primer [Generay Biotech (Shanghai) Co., Ltd., Shanghai, China] sequences were as follows: AID, forward 5'-CAATAAGAACGGCTGCCC-3' and reverse 5'-TTGCGGTTCCTCACAGAAGTAG-3'; and GAPDH, forward 5'-CACCACTGGGAGCCAT-3' and reverse 5'-ACAGCCTGGTAGCAACG-3'.

Statistical analysis. All data are expressed as the mean ± standard deviation in the text and figures. Statistical
analyses were conducted using GraphPad Prism 5.0 software (GraphPad Software, Inc., La Jolla, CA, USA). Significant differences were calculated using the two-tailed, unpaired Student's t-test, with a 95% confidence interval. P<0.05 (two-tailed) was considered to indicate a statistically significant difference.
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Results and Discussion

Increased frequencies of different B cell subsets in IgAN. IgAN is characterized by circulating immune complexes, composed of galactose‑deficient IgA1 and a glycan‑specific IgG antibody (12,13), which is secreted by plasma cells. To determine the immune status of B cells in IgAN, 27 patients with IgAN were recruited in the present study and were divided into two groups: Group A, proteinuria/24 h <4 g and group B, proteinuria/24 h ≥4 g. No statistically significant differences were observed in the level of glomerular filtration rate or the distribution of age and gender between the IgAN groups and the HC group (Table I).

Marked increases in the percentages of CD19+CD138+ plasma cells, CD19+CD27+ memory B cells and CD19+CD86+ activated B cells (14-16) were observed in the IgAN groups, compared with the HC group. Although no significant difference in the percentage of plasma B cells was observed between the two IgAN groups, higher levels of memory and activated B cells were observed in the IgAN group B, compared with the group A (Fig. 1). These data indicated that the frequencies of memory and activated B cells were associated with the progression of IgAN.

Frequencies of circulating Tfh cells are higher in IgAN. Tfh cells are a subset of T cells that localize in the GC and specialize in assisting B cell growth, differentiation and class switching (17). Tfh cells are characterized by increased expression levels of molecules, including CXCR5, PD‑1, ICOS, CD40L and IL‑21, and reduced expression of CCR7.

Figure 2. Fluorescence‑activated cell sorting analysis of the frequency of Tfh cells. Peripheral blood mononuclear cells (5x10⁵) were stained with anti‑CXCR5 PerCP and anti‑CD4 APC for detection of the Tfh cells. (A) Representative results for the frequency of each group of Tfh cells, measured using flow cytometry. (B) Percentage of Tfh cells in the immunoglobulin A nephropathy (group A and B) and HC groups (*P<0.05). Each data point represents an individual subject, and horizontal lines represent the median. Tfh, follicular helper T; HC, healthy control; SSC, side scatter; FSC, forward scatter; group A, 24‑h proteinuria <4 g; group B, 24‑h proteinuria ≥4 g.
The CD4+CXCR5+ T cells have been previously identified as Tfh cells (18-20), and the present study demonstrated that the frequency of CD4+CXCR5+ Tfh cells was significantly higher in the two IgAN groups, compared with that in the HC group. In addition, the percentage of Tfh cells was higher in the IgAN group B than group A (Fig. 2). These data indicated that high levels of Tfh cells may accelerate IgAN exacerbation by promoting the proliferation and function of B cells.

Serum IL-21 levels are increased in patients with IgAN. As a key effector of Tfh cells, IL-21 is involved in the process of promoting the growth, differentiation and class switching of B cells (21,22). Due to the high frequency of Tfh cells observed in the patients with IgAN, the concentration of IL-21 in the serum was also determined in the above-mentioned groups. In addition, the percentage of Tfh cells was higher in the IgAN group B than group A (Fig. 2). These data indicated that high levels of Tfh cells may accelerate IgAN exacerbation by promoting the proliferation and function of B cells.

The CD4+CXCR5+ T cells have been previously identified as Tfh cells (18-20), and the present study demonstrated that the frequency of CD4+CXCR5+ Tfh cells was significantly higher in the two IgAN groups, compared with that in the HC group. In addition, the percentage of Tfh cells was higher in the IgAN group B than group A (Fig. 2). These data indicated that high levels of Tfh cells may accelerate IgAN exacerbation by promoting the proliferation and function of B cells.

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Effect of IL-21 on AID in IgAN. AID belongs to the APOBEC family of cytidine deaminases and is capable of deaminating dCs into dUs in vitro on ssDNA substrates and ssDNA generated by the formation of RNA-DNA hybrids (10,23). The key function of AID is to induce somatic hypermutation (SHM), Ig class-switch recombination (CSR) and gene conversion (11,24,25). IgA class switching occurs in IgAN (26,27), therefore, the present study investigated whether the high levels of IL-21 were associated with the expression of AID. Following culture of the B cells in vitro, a significantly increasing level of AID was observed in the IL-21-treated group (Fig. 4). This suggested that the excessive production of IgA may be mediated by the high levels of IL-21 secreted by Tfh cells.

IgAN is characterized by circulating immune complexes, which are composed of galactose-deficient IgA1 and a glycan-specific IgG antibody. In addition, antigen-antibody immune complexes against hepatitis B (HB) surface, HB core or HB e antigens, together with complement components, have been demonstrated to be deposited in renal tissue (28,29). B cells are considered to be the key regulators of IgA production in mucosal tissues, and CD138, CD86 and CD27 are reliable markers, which are expressed by different subsets of B cells (30,31). In the present study, CD19+CD138+ cells were defined as plasma cells, CD19+CD27+ cells as memory B cells and CD19+CD86+ cells as activated B cells, and the frequencies of these subsets of B cells were analyzed in patients with IgAN. Significant increases in the percentage of these three subsets of B lymphocytes were observed in the two IgAN groups, compared with the HC group. Furthermore, the frequencies of memory and activated B cells in the advanced IgAN group (group B) were markedly higher than that of the less advanced IgAN group (group A). Notably, although the IgA produced by plasma cells mediated immune complex deposition in the glomerular mesangial area, resulting in IgAN, the numbers of plasma cells were not significantly different between the IgAN groups. These data indicated that high levels of memory and activated B cells resulted in the progression of IgAN, however, but that the IgA immune complex deposition was not solely associated with excessive IgA secretion by the plasma cells. Hernández et al (32), demon-
strated that levels of plasma von Willebrand Factor, a specific marker for endothelial cell injury, are abnormal in patients with IgAN, who exhibit elevated levels or defective molecules. Elevated levels of sFlt-1, a receptor for vascular endothelial growth factor, may lead to widespread endothelial dysfunction and also contributes to the progression of IgAN (33). Thus, the IgA immune complex deposition is associated with immune cells and abnormal endothelial function.

Regarding the effect of Tfh cells on B cell differentiation and activation, the frequency of Tfh cells was analyzed in PBMCs in the present study. CD4+CXCR5+ T cells were identified as Tfh cells, and the frequency of circulating Tfh cells was higher in the advanced IgAN group than in the less advanced IgAN group, which was similar to the results observed in the memory and activated B cells. Although Tfh cells activate the proliferation and differentiation of B cells, the selective stimulation targeting different subsets of B cells requires further investigation. Tfh cells are classified into three subsets: Tfh1, Tfh2 and Tfh17, (CXCR3+CCR6+, CXCR3 CCR6 and CXCR3 CCR6+ cells, respectively) (20). In addition, Tfh2 and Tfh17 cells have been reported to assist in the activation of B cells via the production of IL-21, resulting in the secretion of various isotypes, including IgM, IgA and IgG, and IgE for Tfh2 cells (20). Thus, the present study hypothesized that different B cell subsets are stimulated by different Tfh cell subsets, warranting investigation of the mechanism underlying the regulation of Tfh cells.

As a key effector of activated Tfh cells, IL-21 induces B cell proliferation, mediates the differentiation of activated B cells into plasma and promotes IgM, IgG and IgA production (34,35). Thus, the concentration of serum IL-21 was investigated, which revealed that increased levels of IL-21 were positively correlated with the extent of 24 h proteinuria. Thus, the activation of B cells was suggested to be predominantly mediated by high levels of IL-21 secreted by Tfh cells, and IL-21 may act as a biomarker of IgAN progression.

The association between high levels of IL-21 and the expression of AID was subsequently investigated to clarify the function of IL-21. By stimulating B cells using recombinant human IL-21, a significant upregulation of AID was observed. This indicated that IL-21 was positively associated with the expression of AID in B cells. As a cytidine deaminase, the predominant function of AID is inducing SHM and Ig CSR in GC or GC-like states at extracellular loci (36). In the GC, B cells differentiate into either plasma cells, which secrete antibodies, or memory cells, enabling long-term memory of antigens (37). IgA CSR is important during the process of IgA production. B cell activation factor has been reported to induce the expression of germline AID and IgA class switching in a CD40-independent manner (38). Therefore, a high level of AID is a key mediator for excessive IgA production, resulting in the development of IgAN. The activity of AID is regulated at the transcriptional level by HoxC4 and nuclear factor-κB, and at the translational level by AID phosphorylation and ubiquitination (39-42). The results of the present study confirmed that IL-21 enhanced the expression of AID. Therefore, it was hypothesized that high levels of IL-21 secreted by Tfh cells may promote the differentiation of B cells and induce IgA CSR in patients with IgAN.

The data of the present study demonstrated that the progression of IgAN was closely associated with high levels of memory B cells, activated B cells and Tfh cells, and that the potential mechanism was predominantly associated with the selective effect of Tfh cells on different B cell subsets. The significantly increased levels of IL-21 upregulated the expression of AID in the B cells, which mediated IgA class switching during the differentiation of activated B cells into plasma B cells. However, further investigation is required to fully elucidate the selective mechanism underlying the stimulation of different types of Ig.

Acknowledgements

The authors would like to thank Dr Munan Sun (Jilin Province People’s Hospital) for the collection of clinical samples and supporting information.

References


