

Combined Mutation of the *GATA2* Gene and *STAT5B* Gene in a Patient with Hypogammaglobulinemia and Autoimmunity

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Antibody deficiency is a type of primary immunodeficiency that often manifests as primary hypogammaglobulinemia, with or without repeated infections. Although primary immunodeficiency appears to be contrary to autoimmunity, they usually occur simultaneously, and the specific pathogenesis remains unknown. We herein describe an adult patient with autoimmune manifestations and recurrent infections. The case was characterized by a sustained decrease in serum immunoglobulin A, accompanied by decreased T lymphocytes, B lymphocytes, monocytes, and platelets in the peripheral blood and the presence of antinuclear and anti-SSA antibodies. Whole-exome sequencing for the patient revealed two spontaneous mutations in *GATA2* (c.1084C>T) and *STAT5B* (c.1924A>C). This case report provides evidence that mutations in the *GATA2* and *STAT5B* genes may be pathogenic in primary immunodeficiency and provides genetic evidence for the possible pathogenesis of primary immunodeficiency with autoimmune symptoms. However, further studies are needed to confirm the causal relationship.

Keywords: antibody deficiency; autoimmunity; gene mutations; primary immunodeficiency; whole-exome sequencing

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Introduction

Antibody deficiency comprises a group of primary immunodeficiency diseases in which the production of antibodies is impaired due to inherent molecular defects of B lymphocytes or a lack of interaction between B lymphocytes and T lymphocytes (Notarangelo et al. 2004). Clinical features include repeated infections of the upper and lower respiratory and gastrointestinal tracts, accompanied by a decrease in the level of one or more serum immunoglobulins, as represented by IgG, IgM, and IgA (Conley et al. 1999). The clinical presentation of antibody deficiency is extremely heterogeneous, involving common clinical manifestations such as recurrent infections or comorbid autoimmune disease. Immunodeficiency and autoimmune diseases were previously considered to be completely opposite conditions, yet there is a high degree of overlap regarding genetic background. In fact, one study reported that approximately 26.2% of primary immunodeficiency patients have autoimmune symptoms (Fischer et al. 2017). In general, autoimmune manifestation in these patients predicts a poor prognosis, though the specific pathogenic mechanism is still unclear. In this study, we carried out whole-exome sequencing for a Chinese Han male patient with hypogammaglobulinemia and autoimmune manifestations to better understand the pathogenesis of antibody deficiency with concomitant autoimmune symptoms.

Case Presentation

The patient was a 29-year-old Chinese Han male who was admitted to the Department of Rheumatology and Immunology, West China Hospital in July 2018 with chief symptoms being repeated cough, fever, and chest tightness. At 2 years of age, the patient experienced sensorineural

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Table 1. Medical history.

Age	Diseases		
2 years	Sensorineural hearing loss of unknown cause		
21 years	Recurrent facial acne		
23 years	Perianal abscess with anal fistula		
26 years	Pulmonary tuberculosis, erythema induratum		
27 years	Bronchiectasis with infection		
29 years	years Pulmonary infection, immunosuppression		
31 years	Pulmonary tuberculosis, idiopathic thrombocytopenic purpura		

Table 2. Infinitute profiles.					
	Age				
	29 years	30 years	31 years		
Immunoglobulins (reference range)					
IgG, mg/dL (8,000-15,500)	8,550	13,500	19,400		
IgA, mg/dL (836-2,900)	436	672	893		
IgM, mg/dL (700-2,200)	1,040	1,050	1,840		
IgE, IU/ml (5-150)	ND	11.30	7.96		
Lymphocytes (reference range)					
CD3+, cells/mL (941-2,226)	636	309	665		
CD3+CD4+, cells/mL (471-1,220)	168	70	136		
CD3+CD8+, cells/mL (303-1,003)	416	215	449		
CD19+, cells/mL (175-332)	ND	4	8		
NK, % (9.26%-23.92%)	ND	2.5	ND		
Hemogram, ×10 ⁹ cells/L (reference range)					
Monocytes (0.1-0.6)	0.01	0	0.02		
Neutrophils (1.8-6.3)	7.41	3.62	2.85		
Platelets (100-300)	92	111	13		
Vaccine titers					
Tetanus toxoid, IU/ml (positive, > 0.1)	Negative	ND	ND		
Tuberculosis antibody	Negative	ND	ND		

Table 2. Immune profiles.

ND, Not Done.

hearing loss of unknown cause; he developed recurrent pulmonary infections after the age of 20 (Table 1). Physical examination revealed thickened breath sound and moist rales in both lung bases. The bronchial symptoms were accompanied by redness and swelling of the skin of both lower limbs, with a flaky, red rash; some of the rash lesions had blistering. Hemogram analysis showed a white blood cell (WBC) count of 7.91 \times 10⁹ /L, with neutrophils accounting for 81.3%. Immunological analysis detected a markedly decreased serum IgA level (436 mg/dL), though levels of IgG and IgM were normal (Table 2). Absolute T and B cell counts were significantly decreased (Table 2). Antinuclear antibody (titer 1:320) and anti-Ro/SSA 52 kDa were positive. A chest CT scan revealed scattered bilateral infective lesions with bilateral pleural thickening, adhesion, and increased mediastinal small lymph nodes, as well as a small amount of pericardial effusion. Sputum culture was positive for Candida albicans. The patient's parents were in good health, with no history of any particular disease.

The patient was treated with antibiotics for bacterial and fungal infections, pulverization inhalation to dispel phlegm, and immunoglobulin (400 mg/kg) during hospitalization. The level of IgA was still low after intravenous immunoglobulin treatment, but IgG and IgM remained normal. Due to repeated infections and persistent low levels of IgA (448 mg/dL), hypogammaglobulinemia (IgA) was initially considered as the clinical possibility. The patient's condition subsequently improved, and he was discharged. After discharge, the patient was regularly followed up monthly in our outpatient clinic. In August 2018 and October 2019, he was hospitalized again for idiopathic thrombocytopenic purpura and pulmonary infection with Mycobacterium tuberculosis. After exclusion of myelodysplastic syndrome by bone marrow aspiration, the patient was discharged after anti-tuberculosis treatment and immunoenhancement treatment. The number of follow-up visits

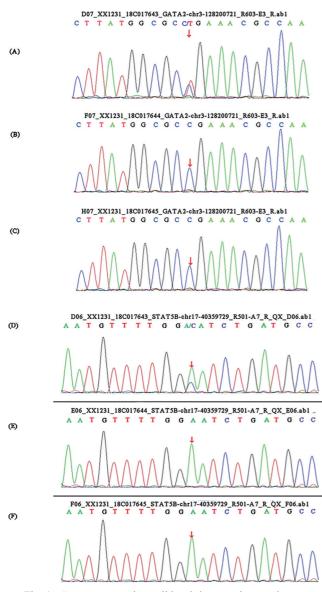


Fig. 1. Sanger sequencing validated the mutations at the two SNP sites.

(A-C) GATA2. (A) c.1084C>T nonsense mutation occurred at chr3-128200721 (GATA2) in the patient (heterozygous mutation) (indicated by an arrow). (B) Father of the patient. (C) mother of the patient.
(D-F) STAT5B. (D) c.1924A>C missense mutation occurred at chr17-40359729 (STAT5B) in the patient (heterozygous mutation) (indicated by an arrow). (E) Father of the patient. (F) Mother of the patient.

was reduced during the novel coronavirus outbreak; his physical condition had recovered very well at the last follow-up visit to the Outpatient Department of our hospital in June 2020.

Whole-exome sequencing was performed for the patient and his parents, revealing two spontaneous heterozygous mutations in *GATA2* (c.1084C>T) and *STAT5B* (c.1924A>C). The SNPs were validated as mutations by Sanger sequencing (Fig. 1).

The study protocol was approved by the Ethics

Committee of West China Hospital, Sichuan University. All participants provided their written informed consent for the study.

Discussion

Although the coexistence of immunodeficiency (a hypoimmune state) and autoimmunity (a hyperimmune state) seems paradoxical, it is common for them to coexist clinically. However, primary immunodeficiency with autoimmune symptoms tends to be more difficult to treat and has a worse prognosis. In the current report, the patient harbored the GATA2 mutation c.1084C>T, which causes a cytosine to thymine change at nucleotide position 1,084, resulting in an amino acid change to a stop codon (p. R362X). Based on bioinformatic software SIFT, PolyPhen 2 and REVEL, this nonsense mutation has a predicted unknown effect on GATA2 protein function. Nevertheless, according to ACMG (Chinese Board of Genetic Counseling, http://acmg.cbgc.org.cn) guidelines, the mutation is pathogenic and reportedly related to myelodysplastic syndrome (Chen et al. 2019). The zinc finger transcription factor GATA2 affects cell fate in multiple tissue types by recognizing the common sequence of GATA in target gene promoters. In addition to its role in the development of hematopoietic stem cells, GATA2 has been shown to be an important regulator of the adult stem-cell pool. In addition, a lower dose of the GATA2 gene negatively affected proliferation and survival of primitive cells, though the differentiation or self-renewal ability of the remaining stem cells did not change (Rodrigues et al. 2005). Moreover, GATA2 mutations are associated with monocyte depletion, dendritic cell and B cell defects, myeloproliferation, immunodeficiency, and problems of the vascular and lymphatic systems (Hsu et al. 2011). Therefore, immunodeficiency caused by GATA2 dysfunction or insufficiency may result in infections, including mycobacterial, viral, and fungal infections. In one case report, a patient diagnosed with common variable immunodeficiency early due to hypogammaglobulinemia and defective antibody response was found to carry a heterozygous GATA2 mutation, with significantly decreased monocyte and lymphocyte counts during adolescence (Chou et al. 2014). Our case was mainly characterized by repeated infections, which is consistent with previous reports. In addition, Hoshino et al. (2019) found that GATA2 deficiency can lead to sensorineural hearing loss in mice, which may explain the hearing impairment in our patient.

In addition, we detected a missense mutation in one copy of *STAT5B* in the present patient, causing nucleotide 1,924 to change from adenine to cytosine, which results in amino acid 642 changing from asparagine to histidine (p.N642H). This *STAT5B* mutation (p.N642H) has been found in over 90 leukemia and lymphoma patients (Pham et al. 2018). According to ACMG guidelines, the mutation is preliminarily considered likely pathogenic. A previous study reported that patients with mutations in *STAT5B* often

develop infections, autoimmune manifestations, eczemalike manifestations, and growth hormone insensitivity syndrome (Kofoed et al. 2003), but manifestations such as eczema and growth deficiency were not observed in our patient. Although study had proposed that STAT5B is related to development of primary immunodeficiency (Caldirola et al. 2018), we do not have enough evidence on the pathogenicity of STAT5B and more work is worth exploring in the future.

In conclusion, our study enriches our understanding of the pathogenic genes of antibody deficiency disorders, suggesting that coinheritance of mutations in immunoregulatory genes may affect the pathogenesis of primary immunodeficiency with autoimmunity. Further research with a large patient cohort is needed to clarify the interactive pathogenic roles of *GATA2* and *STAT5B*, especially when autoimmunity is concurrent.

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Author Contributions

Z.L. was responsible for the conception and design of the study, collection and interpretation of the clinical data, and for drafting of the manuscript. P.Q. and Y.Z. were responsible for suggestions and data acquisition and for revising the manuscript. Y.L. and T.M. participated in the clinical evaluation of the case and in critically revising the manuscript. All authors read and approved the final manuscript.

Conflict of Interest

The authors declare no conflict of interest.

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