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A novel homozygous *RAG1* mutation is associated with severe combined immunodeficiency and neurological presentations

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Abstract

Introduction and objectives: Severe combined immunodeficiency (SCID) is a subset of primary immunodeficiency diseases caused by a hereditary deficiency of the adaptive immune system. Mutation in recombination activating gene (*RAG*) is known as the underlying genetic cause of SCID. *RAG* protein plays a pivotal role in V(D)J recombination which is the main process to assemble lymphocyte antigen receptors during T- and B-cell development. The patients are characterized by recurrent infections, failure to thrive, chronic diarrhea, and fever, in early infancy. Herein, we present a case of SCID with rare neurological manifestations affected by a mutation in *RAG1*. **Patients and methods:** The patient was a 15-month-old infant born to a consanguineous family. She was presented with neurological abnormalities including facial nerve palsy, seizure,

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and decreased consciousness. Next-generation sequencing (NGS)-based primary immunodeficiency disease (PID)-gene panel screen and Sanger sequencing were performed to identify the genetic mutation.

Results: We found a novel homozygous missense mutation in *RAG1*, c.1210C>T,p.Arg404Trp, which was predicted to be deleterious (combined annotation dependent depletion, CADD score of 27.4). Both parents were heterozygous carriers for this mutation. According to her laboratory data, both T cell and B cell numbers were decreased and the patient was diagnosed as *RAG1*- SCID.

Conclusions: SCID is a pediatric emergency with a variety of manifestations in infants. Therefore, accurate diagnosis importantly in the case of rare manifestations must be considered in these patients. Our findings point toward the importance of genetic assessment for early diagnosis and timely treatment of this disorder.

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Introduction

Severe combined immunodeficiency (SCID), is a hereditary group of disorders with the highest rate of mortality between the inborn errors of immunity (IEI).¹ The incidence of the disease is one in 40,000 to 75,000 newborns.² Both cellular and humoral immune systems can be involved in SCID. The patients suffer from either a very low number of T-cells or decreased function of these cells (CD3+, CD4+, and CD8+) with or without a reduction in the number of B lymphocytes (CD19+, CD20+) and/or even natural killer cells (CD16+, CD56+).³ Hence, SCID is classified into four sub-groups with regard to the lack or presence of T, B, and natural killer cells (NK) including T-B-NK-, T-B-NK+, T-B+NK+, and T-B+NK-.⁴

The primary clinical manifestations frequently present between 4 and 7 months of age including chronic diarrhea, insufficient growth, and recurrent infections.⁵ The neurodevelopmental defects are not common and progressive in SCID patients, nevertheless, seizure and microcephaly have been observed in some cases.⁶ Moreover, infants who suffer from SCID are more vulnerable to be infected with community-acquired infections that lead to end-organ dysfunction. Such infections might affect multiple organs like the lungs and liver which makes SCID a pediatrics' emergency.⁷

Genetic mutations in SCID patients are divided into X-linked and autosomal recessive forms.⁸ To date, multiple known mutations have been reported which are associated with different types of SCID including recombination activating gene 1 (*RAG1*), *RAG2*, interleukin 2 receptor gamma (*IL2RG*), adenosine deaminase (*ADA*), purine nucleoside phosphorylase (*PNP*), etc. Except for *IL2RG* mutation with X-linked inheritance, other mutations are autosomal recessive.^{9,10}

Mutations in *RAG1* and *RAG2* are the most common causative gene in countries where a high rate of consanguinity is known.¹¹⁻¹³ *RAG1* encodes proteins that are involved in DNA breakage and repair during rearrangement of the genes encoding the variable (V), diversity (D), and joining (J) segments. This process leads to provide a variety of antigen recognition sites for T and B cell receptors.^{14,15}

The phenotypic spectrum of primary immunodeficiency (PID) associated with *RAG1* mutations differs depending on the nature of the defect, and its impact on V(D)J

recombination. Amorphic or null mutations of both alleles of *RAG1* result in complete *RAG1* deficiency leading to a severe combined immunodeficiency phenotype which is characterized by a lack of T and B lymphocytes.^{16,17} The patients with this phenotype exhibit recurrent infections and failure to thrive (FTT) in early infancy.^{18,20} Hypomorphic *RAG1* mutations result in very low levels of recombination activity characterized by residual development of T and B lymphocytes. Different phenotypes associated with hypomorphic mutations are including Omenn syndrome characterized by hepatosplenomegaly, lymphadenopathy, eosinophilia, elevated IgE levels in serum, and erythroderma^{21,23}; atypical Omenn syndrome characterized by skin inflammation without T-cell expansion, more than 3% B-cells, a low frequency of CD31 cells,²⁴ and atypical SCID with $\gamma\delta$ T-cell expansion characterized by autoimmunity, severe cytomegalovirus infection, and partially functioning B-cells with a limited ability for antibody production.^{25,27}

Therefore, the mutation of this gene could affect the immune system due to the impaired development of lymphocytes.²⁸ Timely genetic assessment and implementation of appropriate therapies could improve symptoms and decrease morbidity in these patients. Herein, we for the first time describe an Iranian case of SCID with neurological manifestations affected by a novel homozygous mutation in *RAG1* which resulted in TlowB-NK+ type of SCID.

Patients and Methods

Case presentation

A 15-month-old female was admitted to Namazi Hospital of Shiraz University of Medical Sciences with decreased vital signs, cyanosis, seizure, poor feeding, and right facial nerve palsy. Her parents were first cousins and there was a history of fatality in her parents' family because of the several cancers as well as congenital palsy in childhood (Figure 1a). The patient has no siblings and her mother had one miscarriage due to unknown causes. The patient's maternal grandmother has experienced repetitive abortions of unknown etiology. Her uncle passed away in infancy due to unknown causes. Besides, two grandchildren of the patient's maternal aunt passed away in childhood because

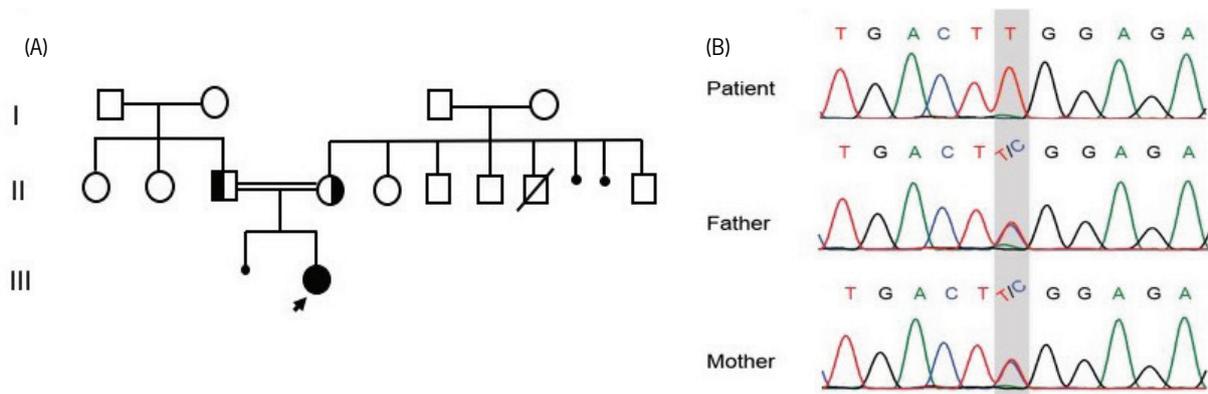


Figure 1 (A) Pedigree of the patient. The patient was homozygous for c.1210C>T,p.Arg404Trp mutation and is shown by a solid black symbol. (B) Sanger sequencing of the patient and her parents. A homozygous and heterozygous missense mutation in *RAG1*, c.1210C>T,p.Arg404Trp, was shown in the patient and both parents, respectively.

of blood cancer and congenital palsy. Also, the patient's paternal aunt and cousin passed away due to an unknown type of cancer.

She had a history of hospitalization at the age of 6 months (April 2018) due to the diagnosis of BCGitis with axillary and supraclavicular lymphadenopathies along with purulent secretion. Her blood sample was positive for acid-fast bacilli (AFB) and tuberculosis (TB) culture. Additionally, she had pustular lesions around her anus leading to necrosis. She received antibiotics, antifungal, and antiviral drugs (Figure 2).

At the time of second admission (Dec 2018), general examinations showed normal chest, HEENT evaluation (head, ears, eyes, nose, and throat), abdominal, and extremity exam. Abdomen and pelvic sonography appeared normal. The patient suffered from lethargic mental status during the first days of administration. Also, the patient experienced a partial seizure with loss of consciousness afterward. She was intubated and admitted to the neurology ward to receive antiepileptic medicine including Topiramate, Levetiracetam, Phenytoin, Phenobarbital, and Clobazam for status epilepticus. Besides, she received broad-spectrum antibiotics and antifungal drugs (Figure 2).

The result of the electroencephalogram showed Status Epilepticus due to refractory complete partial seizure. The Brain CT scan was normal. There were no signs of increased

intracranial pressure and no abnormalities in the cerebello-pontine angle areas on both sides and the calvarium.

The echocardiography result was normal and no pulmonary hypertension was observed. The result for brain MRI disclosed evidence of T1 increase intensity in both basal ganglia that is suggestive for early calcification or anoxia damage due to the postictal phase of the seizure. Otherwise, there was no sign of mass or hemorrhage or ischemic infarction within the brain parenchyma. No hydrocephalus or shift of midline structures was seen. 7th-8th nerve root complexes appeared normal. The pituitary gland was unremarkable. No extra-axial mass or hematoma or fluid collection was seen. We did not receive any shreds of evidence for T2-weighted images. Besides, the primitive electrocardiogram was abnormal with low voltage evidence.

Laboratory investigations including complete blood count (CBC), immunoglobulin levels, flow-cytometry for immunophenotyping, Dihydrorhodamine test (DHR), cerebrospinal fluid (CSF) analysis including polymerase chain reaction (PCR) for herpes simplex virus (HSV), and CSF culture were performed. Besides, next-generation sequencing (NGS)-based PID-gene panel screen, which targets >500 genes in parallel, and Sanger sequencing were requested to investigate gene mutations.

The patient had an uncontrolled seizure and had to be intubated. She fell into a coma and vegetative state

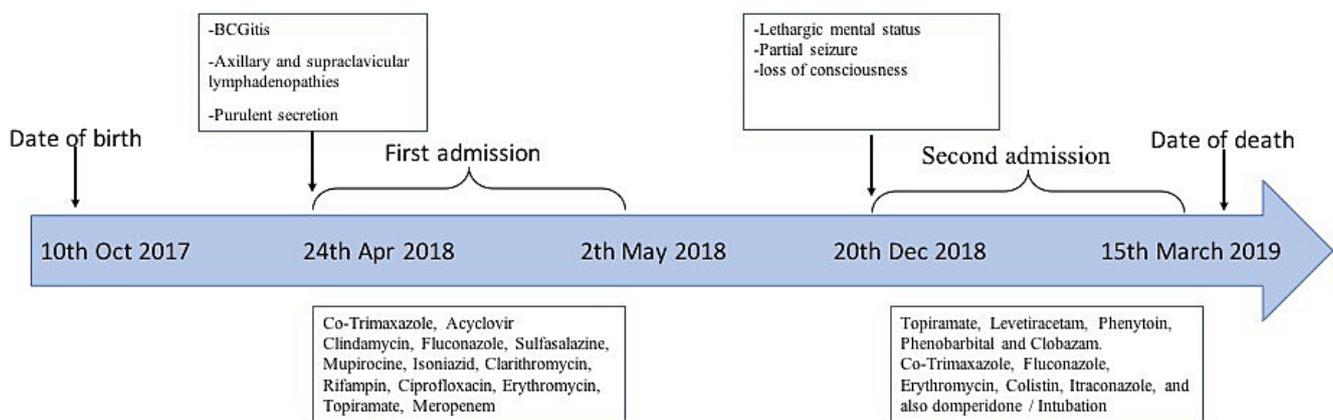


Figure 2 Timeline graph depicting the patient's clinical course and treatments.

after administering anesthesia for a tracheostomy. Then, she was under the mechanical ventilator in the neurology ward. Unfortunately, 3 months after hospitalization she passed away due to persistent seizures and cardiopulmonary arrest.

Results

The laboratory results are summarized in Tables 1 and 2. Immunophenotyping using flow-cytometry showed decreased levels of both T- and B-cell markers including CD3, CD4, CD8, CD19, and CD20. Moreover, the markers for NK cells including CD16 and CD56, CD14 for monocytes, CD11b, and interferon-gamma receptor were normal. The DHR test was also within the normal range (220, normal >50). Hypogammaglobulinemia and lymphopenia were detected in the evaluation of immunoglobulin levels and CBC results, respectively. No CNS infection was documented. PCR HSV and CSF culture were negative. CSF findings are shown in Table 2.

NGS and Sanger sequencing results of our patient showed a novel homozygous missense mutation in *RAG1* (NM_000448.3:c.1210C>T, p.Arg404Trp) that was predicted to be deleterious (CADD score of 27.4).²⁹ Both parents were heterozygous carriers for this mutation (Figure 1b).

Discussion

RAG1 and *RAG2* are known as the most common underlying genetic cause of SCID. These genes are restricted to lymphocytes and encode the proteins involved in gene rearrangement and site-specific V(D)J recombination of the lymphocyte receptor. Thereby, the mutations in *RAG* can result in a lack or reduced levels of lymphocytes or producing immature lymphocytes with impaired function. T negative (-) B negative (-) NK positive (+) phenotype is the frequent type of SCID caused by *RAG* mutation and counts for about 20 percent of all SCID cases.¹⁰

Table 1 CBC results of the patient.

	Patient			Normal range
	Jan. 2019	Feb. 2019	March. 2019	
WBC (mm ³)	6000	10,900	5600	3100–21,600
Neut	3660	6649	3752	4940–7410
Lymph	600	654	280	2470–4940
Eos	240	327	224	123–494
RBC (*10 ¹² /L)	3.85	3.53	3.4	3.12–7.3
Hb (g/dL)	10.9	10.6	9.5	11–17.3
HCT (%)	31.4	29.5	29.1	35.4–56.5
MCV (fL)	81.6	83.6	85.4	90.4–128
MCH (pg)	28.3	30	28.2	34.41–2.36
Plat (*10 ⁹ /L)	93	401	268	152–472

WBC: white blood cells; Neu: neutrophils; Lymph: lymphocyte; Eos: eosinophils; RBC: red blood cells; Hb: hemoglobin; HCT: hematocrit; MCV: mean cell volume; MCH: mean corpuscular hemoglobin; Plat: platelet.

Table 2 Summary of laboratory evaluation.

Blood	Result		
Culture	<i>Pseudomonas aeruginosa</i>		
CSF*	Result		
RBC	Negative		
WBC	Negative		
Sugar (mg/dL)	61		
Protein (mg/dL)	18		
Culture	Negative		
HSV (PCR)	Negative		
Immunoglobulins	Result	Normal range	
IgG (mg/dL)	13.8	200–1000	
IgM (mg/dL)	1.66	25–150	
IgA (mg/dL)	3.47	4–90	
IgE (IU/mL)	80	<200	
Blood Flow cytometry analysis: lymphocytes subsets /mm ³ (%)	Apr, 2018	July, 2018	Normal range
CD3	795 (38)	334 (16)	1852–2853 (50–77)
CD4	544 (26)	188 (9)	1222–2149 (33–58)
CD8	146 (7)	83 (4)	482–963 (13–26)
CD19	42 (2)	20 (1)	482–1297 (13–35)
CD20	21 (1)	20 (1)	482–1297 (13–35)
CD16	1255 (60)	1737 (83)	74–482 (2–13)
CD56	1255 (60)	1737 (83)	74–482 (2–13)
CD14	272 (13)	N/A	111–222 (3–6)

*CSF: cerebrospinal fluid; Ig: Immunoglobulin; N/A: not available.

Here, we identified a novel mutation in the *RAG1* gene (c.1210C>T, p.Arg404Trp) for which the child was homozygous and both parents were found to be heterozygous. Our patient was presented with FTT and rare neurological manifestations including nerve palsy, loss of consciousness, and persistent seizures since the age of 15 months. The patient developed recurrent infections and BCGitis in early infancy that could dictate early investigation regarding the possibility of primary immunodeficiency.

A heterozygous mutation in this position (*RAG1*, p.Arg404Trp) has been previously reported in a patient with T-B-NK+ Omenn syndrome. Corneo *et al.*, reported a 2.5 months old boy with early onset of diffuse erythroderma, eosinophilia, diarrhea, and FTT. His laboratory findings have been shown a decreased level of lymphocytes and CD3, lack of CD19, and normal NK cells.³⁰ Our patient had lymphadenopathy and suspicious lesions around her anus and Omenn syndrome could be a probable consideration for this case. Since she had neither hepatosplenomegaly

nor eosinophilia in the peripheral smear, the possibility of Omenn syndrome was ruled out.

Regarding the other types of SCID, X-linked SCID and Janus kinase 3 (*Jak3*) deficiency were ruled out. These disorders are determined by an elevated number of B-cells while our patients had reduced levels of B-cells. Besides, in adenosine deaminase (*ADA*)-deficient SCID, bone abnormalities presentations are more frequent, which were absent in this case.³⁰

Neurological presentations and facial nerve palsy occurred in our case are uncommon presentations of SCID patients. Neurological presentations in the patients with *RAG1* mutation except for a limited number have not yet been reported. Dhingra *et al.*,³¹ described a 15-month-old boy with *RAG1*-SCID who had progressive encephalopathy and partial seizures. His immunological evaluation showed hypogammaglobulinemia, T- and B-cells lymphopenia, and normal NK cells. His genetic investigations revealed a homozygous mutation in *RAG1* (c:2881T>C; p:l794T). In accordance with this, the immunophenotyping results and immunoglobulin levels of our patient showed a decreased level of T- and B-cells, normal NK cells, and hypogammaglobulinemia, and the patient was suspected to have TlowB-NK+ SCID.

In addition, some of the neurological abnormalities in SCID have been described in *ADA* deficiency, DNA ligase IV deficiency, and Cernunnos deficiency. Neurological manifestations in *ADA*-deficient SCID are including mental retardation, seizure, behavioral abnormalities, motor delay, and nystagmus that might be related to toxic accumulation of adenosine in the brain.^{32,34} Microcephaly and developmental delay were recognized in DNA ligase IV deficiency and Cernunnos deficiency patients.^{32,33,35}

There is not enough evidence to explain the relation between *RAG* deficiency and neurological presentation. It has been shown that the *RAG1* gene is expressed in the central nervous system, especially in the brain of vertebrates and the transcription of *RAG1* initiates as soon as the neural stages of development. Hence, *RAG1* protein could play an important role in the central nervous system development.³⁶ Although *RAG1* expression has been found in the central nervous system of the mouse, no apparent neuroanatomical or behavioral abnormalities have been detected in the *RAG1*-deficient mice.³⁷ We believe further investigation about the expression of *RAG1* in neurological tissues might be helpful.

On the other hand, we cannot rule out the possible role of autoreactive antibodies or T-cells in SCID patients. SCID could be associated with autoimmune disorders, usually in the Omenn's syndrome phenotype. It has been shown that central tolerance that is responsible for the elimination of autoreactive T-cell clones, has a defect in SCID. Besides, peripheral tolerance is reduced in SCID due to several factors such as the expansion of T-cell clones in charge of the consequence of the lymphopenia as well as T regulatory (FOXP3+) cell numbers reduction, which all allowing autoreactive T-cells to proliferate and infiltrate various organs in the body of patients with SCID. It is thus of no surprise when both central and peripheral tolerance is impaired that autoimmunity can be observed in SCID.³⁸

Although these findings indicate the autoimmunity and autoreactive T-cells might appear in SCID as in our case, the role of autoimmunity in CNS is not yet proven.

On the other hand, since CNS infections have been reported in some SCID patients, the possibility of CNS infection in these patients should not be ignored. Waruiru *et al.*,³⁹ described eight patients of SCID including patients with *RAG* deficiency and T-B- NK+ SCID with CNS viral infections confirmed by PCR. The CSF analysis in our case was completely normal.

We present this *RAG1*-deficient patient with rare neurological presentations to hallmark the significant role of early genetic analysis. Timely genetic assessment in patients with the same clinical presentations could be important in early diagnosis and implement appropriate therapy to manage the severe outcomes.

Hematopoietic stem cell transplantation (HSCT) can be an efficient and permanent cure for children with *RAG1*-SCID.^{40,41} Although HSCT is considered a life-saving treatment, it could be limited due to the obstacles such as high rates of graft versus host disease (GVHD) and transplant-related mortality.⁴² Gene therapy is shown to be successful in treating patients with X-linked SCID and *ADA*-SCID.⁴³⁻⁴⁸ As there are some hopeful results obtained from the pre-clinical phase, this approach might receive the attention of researchers for treating the patients with *RAG1*-SCID in future studies.^{49,50}

Authors' contributions

All authors have been contributed significantly to the work, have read the manuscript, attest to the validity, and agree to its submission.

Conflicts of interest

The authors have no conflicts of interest to declare.

Ethical disclosure

After describing the novelty of the genomic mutation causing the disease for the patient's family, they orally consented to the authors to use the patient's medical records for publication.

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