ORIGINAL ARTICLE

Clinical, Laboratory, and Molecular Findings for 63 Patients With Severe Combined Immunodeficiency: A Decade's Experience

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Abstract

Introduction: Severe combined immunodeficiency (SCID) is a life-threatening pediatric disease. We report on the clinical evaluation, immunological assessment, molecular analysis, and outcomes of SCID patients in a tertiary referral center in Iran.

Methods: From January 2006 to December 2015, we performed a prospective cohort study in which initial screening and advanced immunological tests were carried out on patients suspected of having SCID. Genetic analysis was also performed to confirm the diagnosis.

Results: A total of 63 patients were diagnosed with SCID (43 male [68.3%]). The median age at onset and diagnosis and diagnostic delay were 40 and 110 days respectively. A total of 49 patients (77.8%) had a history of BCG vaccination, and of these, one-third experienced BCG-associated complications. The most common clinical manifestations were pneumonia, recurrent oral candidiasis, chronic diarrhea, and failure to thrive. Of the thirteen patients who underwent hematopoietic stem cell transplantation, 8 survived and 5 died before they could receive the transplant. Most patients (34.9%) were classified as having T B NK+ SCID and had a mutation in the RAG2 or RAG1 gene.

Conclusion: Autosomal recessive SCID is the most common type in Iranian patients. Providing high-quality training to physicians and patients’ families to reduce the diagnostic delay should be prioritized. It is also important to raise awareness of live vaccination and to expand stem cell donor registries to speed up the transplantation process.

Key words: Severe combined immunodeficiency. Newborn screening. BCG complications. Stem cell transplantation.

Resumen

Introducción: La inmunodeficiencia combinada severa (SCID) es una grave enfermedad pediátrica que puede comprometer la vida del paciente. El artículo recoge la evaluación clínica e inmunológica, el análisis molecular y la supervivencia de los pacientes con SCID atendidos en un hospital de referencia de Irán.

Métodos: Desde enero de 2006 a diciembre de 2015, se realizó un estudio prospectivo en los pacientes con SCID en el que se realizó un screening inicial junto a diferentes análisis inmunológicos. Se realizó un análisis genético para confirmar el diagnóstico.

Resultados: Sesenta y tres pacientes fueron diagnosticados de SCID, cuarenta y tres (63,8%) de los mismos eran varones. La mediana de la edad de inicio de la enfermedad, diagnóstico y retraso en su diagnóstico, fueron de 40, 110 y 60 días respectivamente. Cuarenta y nueve pacientes (77,8%) recibieron vacunación con BCG y un tercio de los mismos presentó complicaciones como consecuencia de la misma. Las manifestaciones clínicas más frecuentes de estos pacientes fueron: neumonía, candidiasis oral recidivante, diarrea crónica y retraso en el crecimiento. Ocho de los trece pacientes que recibieron trasplante de progenitores hematopoyéticos, lograron sobrevivir. Los restantes pacientes fallecieron antes de poder recibir dicho trasplante. El 34,9% de los pacientes tuvieron TBNK+ SCID y la mayoría de los pacientes eran portadores de mutaciones en los genes RAG2 o RAG1.

Conclusión: La variante autosómica recesiva de la SCID es la forma más común en los pacientes iraníes. Se debe considerar prioritario concienciar para evitar la vacunación con gérmenes vivos y expandir los registros de donantes de células madre para agilizar el trasplante de estos pacientes.

Palabras clave: Inmunodeficiencia combinada severa. Screening del recién nacido. Complicaciones BCG. Transplante de células madre.

Introduction

Severe combined immunodeficiency (SCID) is a prototype of primary immunodeficiency disease (PID). It is a heterogeneous, life-threatening syndrome comprising a group of rare and monogenic diseases [1].

The prevalence of SCID worldwide is estimated to be 1 in 50,000 to 100,000 live births [2]. However, the actual number of PIDs has been shown to be higher than previously reported, especially in countries such as Iran, which have a high rate of consanguinity [3], and in other countries, where a large number of early deaths of unknown cause have been reported [4]. It is difficult to determine the exact number of cases of SCID in Iran before conducting newborn screening for SCID, although it seems to be greater than that we initially thought [5].

SCID patients generally present severe and repeated infections by opportunistic microorganisms (eg, Pneumocystis jiroveci, Candida albicans, and cytomegalovirus), skin rashes, persistent diarrhea, and failure to thrive within the first year of life. SCID patients often need immediate pediatric emergency care [1,6].

According to their fundamental defects, SCID patients are classified into 4 major groups: (1) imperfect function of pre–T-cell antigen receptor complex (eg, CD3ε/δ/ζ and CD45); (2) impaired signaling function based on common γ-chain–dependent cytokine receptors, such as defects in the interleukin 2 receptor γ-chain, interleukin 7 receptor α-chain, and Janus kinase 3; (3) V(D)J recombination defects (eg, Artemis and RAG1/ RAG2 deficiency); and (4) early lymphocyte death mediated by purine metabolic dysfunction (ADA deficiency) [7].

Given the scarcity of publications on Iranian newborns with SCID, the aim of this study was to present the clinical features of SCID patients, available laboratory findings, and results of genetic analyses.

Methods

Data Collection

The study sample comprised 63 SCID patients referred to the Immunology, Asthma and Allergy Research Institute (IAARI) of Tehran University of Medical Sciences, Tehran,
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Iran from various health centers across the country. We conducted a prospective cohort study over a 10-year period (January 2006 to December 2015).

A questionnaire was completed for each patient to record demographic information, history of complicated infections in the past and associated symptoms, family history of immunodeficiency, disseminated infection following BCG vaccination, and patients’ outcomes. This study was approved by the Ethics Committee of the IAARI.

Patients were screened early, and an advanced immunological work-up was performed. This included a complete blood count, serum immunoglobulin levels (IgG, IgA, IgM, and IgE) measured by nephelometry, lymphocyte count and immunophenotype analysis using flow cytometry (CD3, CD4, CD8, CD19, and CD16/56), lymphocyte proliferation response to phytohemagglutinin (PHA), nitro-blue tetrazolium (NBT) test, and the total hemolytic complement (CH50) assay.

Based on the guidelines of the Pan-American Group for Immunodeficiency and the European Society for Immunodeficiencies [8], the clinical diagnostic criteria included at least 1 invasive bacterial, viral, or fungal/opportunistic infection and/or persistent diarrhea, recurrent pneumonia, persistent thrush, as well as failure to thrive and/or affected family members. Patients with symptoms during the first year of life in whom HIV infection was ruled out were enrolled in the study. Participating patients had low or zero counts of CD3, CD4, or CD8 T cells (except for atypical SCID). Based on normal lymphocyte subsets in healthy children [9] and lymphocyte immunophenotyping, patients were classified into 5 groups, as follows: T+B+NK− SCID, T+B+NK+, T−B+NK− SCID, T−B+NK+, and atypical T− SCID.

Genetic Analysis

Molecular evaluations were performed for specific genes known to be involved in SCID (including IL2R, IL7RA, ADA, RAG1, and RAG2). Briefly, in patients with T+B+NK−, T−B+NK+, and T−B+NK+ phenotypes, a genetic study was performed, respectively, for RAG1/RAG2, ADA, and IL7RA. A genetic study of RAG1/RAG2 was also performed for 3 atypical patients with the T−B+NK+ phenotype. Males with the T+B+NK− phenotype were evaluated to assess mutations in IL2R. Genomic DNA was extracted from peripheral blood leukocytes. The exons and exon-intron boundaries were amplified by PCR runs with a final volume of 25 µL containing 200 ng of genomic DNA, 10 pmol of each primer, 100 µM of dNTP, various concentrations of MgCl2, 1x Taq DNA polymerase buffer, and 2 U Taq DNA polymerase (Cingene). Initial denaturation of genomic DNA was performed for 5 minutes at 95°C, followed by 35 cycles of amplification as follows: 35 seconds at 95°C, 35 seconds at 54-64°C, and 30 seconds at 72°C. After evaluation on 1.5% agarose gel, PCR products were sequenced.

The definitive diagnosis was made based on data from the clinical findings, immunological assessments, and molecular analysis.

Statistical Analysis

Questionnaire and laboratory data were analyzed using SPSS, version 16 (SPSS Inc). The data were presented using descriptive statistics (mean [SD], median [IQR]). The normality of quantitative variables was determined using Kolmogorov-Smirnov test. The independent-sample t test (or Mann-Whitney test for variables with a nonnormal distribution) was applied to compare the means of 2 quantitative variables. Statistical significance was set at $P< .05$.

Results

During the study, 63 patients with SCID from 61 families were diagnosed and followed up. Most patients with a typical feature of SCID (n=52, 82.5%) had CD3− T-cell counts <500/µL. Furthermore, a CD3− T-cell count <300/µL was found to be positive in 77.8% of patients. Table 1 shows the laboratory findings and cell counts for the different patient phenotypes. Lymphocytopenia (<3000 cells/µL) was present in 82.5% of patients. The median lymphocyte count was 1380 (602-2304) for all patients, including atypical SCID patients. The results of the other immunological assessments of phagocytic function (NBT) and complement activity (CH50) were normal in all patients. According to typical immunologic classifications based on the immunophenotyping findings, 22 patients (34.9%) were classified as T+B+NK− SCID, 15 (23.8%) as T+B+NK+, 10 (15.9%) as T−B+NK−, 9 (14.3%) as T+B−NK− (X-linked SCID), and 7 (11.1%) as atypical SCID (T− SCID).

Forty-three patients (68.3%) were male. Consanguinity was found in 87.3% (n=55) of patients’ families. A positive family history of SCID was recorded in 34 patients (54%), and most families had only 1 child with SCID. The mean age at onset and diagnosis was 52 (48) and 113 (90), and the median age was 40 (15-60) and 110 (45-150) days, respectively. Interestingly, mean and median time to diagnosis decreased to 97 (85) and 60 (35-150) days in the second half of the study.

Except for 2 patients who were diagnosed before the onset of symptoms based on the death of a sibling, the median diagnostic delay for the remaining 61 patients was 60 (15.5-90) days. Similarly, the median diagnostic delay decreased to 45 (13.7-60) days in the second half of the study. In addition, the difference in diagnostic delay between patients with a positive family history of SCID (45 [5.5-101]) and patients with a negative family history of SCID (60 [33.5-90]) was only 15 days.

In total, 49 patients had received a BCG vaccination (77.8%), and subsequently localized complications (adenitis) were observed in 22 (44.9%), of whom 14 patients had disseminated infection (BCGosis). Among patients with a positive family history of PID, about 65% had received BCG vaccines. Since the complications were not seen in all vaccinated patients, the median absolute numbers of T, B, and NK cells at the time of diagnosis in patients with complications of the BCG vaccine (except for atypical patients) were evaluated. The respective median absolute counts of CD3, CD19, and CD16/56 cells in SCID patients with BCG complications were 329 (124-748), 73 (7.5-711), and 69 (14-139), whereas in SCID patients without localized or disseminated BCG complications, the counts were 23 (9-114), 8 (3-426), and 117 (31-336), respectively. However, no significant differences were found between these 2 groups of patients.

In the present study, the most common clinical manifestations were pneumonia (n=49, 77.8%), recurrent oral candidiasis (n=29, 46%), chronic diarrhea (n=15, 23.8%), failure to thrive (n=13, 20.6%), and oral ulcer (n=10, 15.4%).

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Table 1. Clinical Characteristics and Laboratory Findings

<table>
<thead>
<tr>
<th>Type</th>
<th>(T–B+NK–)</th>
<th>(T–B+NK+)</th>
<th>(T–B–NK–)</th>
<th>(T–B–NK+)</th>
<th>Atypical</th>
<th>Total</th>
</tr>
</thead>
<tbody>
<tr>
<td>n</td>
<td>9</td>
<td>10</td>
<td>15</td>
<td>22</td>
<td>7</td>
<td>63</td>
</tr>
<tr>
<td>Gender (M/F)</td>
<td>7/2</td>
<td>6/4</td>
<td>11/4</td>
<td>13/9</td>
<td>6/1</td>
<td>43/20</td>
</tr>
<tr>
<td>Age at presentation, d</td>
<td>30 (0-150)</td>
<td>65 (0-150)</td>
<td>45 (8-150)</td>
<td>48 (0-180)</td>
<td>14 (0-60)</td>
<td>40 (0-180)</td>
</tr>
<tr>
<td>Diagnostic delay, d</td>
<td>45 (7-270)</td>
<td>97 (10-300)</td>
<td>90 (27-360)</td>
<td>120 (0-390)</td>
<td>60 (30-240)</td>
<td>110 (0-390)</td>
</tr>
<tr>
<td>Granulocytes/µL</td>
<td>4209 (2080-11,218)</td>
<td>4355 (2180-10,841)</td>
<td>3720 (410-23,610)</td>
<td>3350 (250-17,453)</td>
<td>3810 (6894-11,110)</td>
<td>3800 (250-30,736)</td>
</tr>
<tr>
<td>Lymphocytes/µL</td>
<td>1910 (960-3880)</td>
<td>1625 (40-5590)</td>
<td>360 (90-2108)</td>
<td>1330 (300-2584)</td>
<td>5600 (3136-25,520)</td>
<td>1380 (40-25520)</td>
</tr>
<tr>
<td>Eosinophils/µL</td>
<td>295 (0-760)</td>
<td>20 (100-950)</td>
<td>120 (0-1210)</td>
<td>395 (0-6970)</td>
<td>71 (0-11440)</td>
<td>235 (0-11,440)</td>
</tr>
<tr>
<td>CD3, cells/µL</td>
<td>23 (3-512)</td>
<td>57 (3-492)</td>
<td>26 (1-553)</td>
<td>43 (2-874)</td>
<td>2999 (1255-23,478)</td>
<td>42 (1-23,478)</td>
</tr>
<tr>
<td>CD4, cells/µL</td>
<td>6 (3-307)</td>
<td>31 (0-711)</td>
<td>16 (3-387)</td>
<td>35 (0-784)</td>
<td>2576 (68-5869)</td>
<td>31 (0-5869)</td>
</tr>
<tr>
<td>CD8, cells/µL</td>
<td>31 (4-153)</td>
<td>55 (0-1422)</td>
<td>7 (0-518)</td>
<td>77 (0-784)</td>
<td>2507 (60-16,588)</td>
<td>32 (0-16588)</td>
</tr>
<tr>
<td>CD19, cells/µL</td>
<td>1382 (365-2754)</td>
<td>1057 (487-14081)</td>
<td>8 (2-100)</td>
<td>7 (0-187)</td>
<td>44 (5-2848)</td>
<td>44 (0-14 081)</td>
</tr>
<tr>
<td>CD56, cells/µL</td>
<td>117 (9-156)</td>
<td>342 (183-2800)</td>
<td>33 (6-120)</td>
<td>722 (169-3181)</td>
<td>1020 (64-16,588)</td>
<td>321 (6-3379)</td>
</tr>
<tr>
<td>CD16+</td>
<td>72 (0-760)</td>
<td>57 (3-492)</td>
<td>26 (1-553)</td>
<td>43 (2-874)</td>
<td>2999 (1255-23,478)</td>
<td>42 (1-23,478)</td>
</tr>
<tr>
<td>IgA, mg/dL</td>
<td>10 (0-70)</td>
<td>4 (0-26)</td>
<td>4 (0-66)</td>
<td>10 (1-68)</td>
<td>7 (0-16)</td>
<td>7 (0-70)</td>
</tr>
<tr>
<td>IgM, mg/dL</td>
<td>26.5 (0-56)</td>
<td>42 (2-284)</td>
<td>33 (6-120)</td>
<td>42 (2-284)</td>
<td>33 (6-120)</td>
<td>42 (2-284)</td>
</tr>
<tr>
<td>IgG, mg/dL</td>
<td>79.5 (30-164)</td>
<td>358 (20-2041)</td>
<td>79.5 (30-164)</td>
<td>358 (20-2041)</td>
<td>1020 (64-16,588)</td>
<td>321 (6-3379)</td>
</tr>
<tr>
<td>IgE, IU/mL</td>
<td>1 (0-1608)</td>
<td>1 (0-1608)</td>
<td>1 (0-1608)</td>
<td>1 (0-1608)</td>
<td>1 (0-1608)</td>
<td>1 (0-1608)</td>
</tr>
</tbody>
</table>

Quantitative variables are shown as median (min-max).

With SCID and Omenn syndrome belonged to the T–B–NK– group and that the XL-SCID group had the lowest number of patients (14%). These findings are in contrast with reports from Europe and America [10], possibly owing to genetic variations between populations and a higher rate of consanguinity in Iran.

SCID is similar to another common PID in Iran, namely, chronic granulomatosis disease, which is more frequently autosomal recessive than X-linked [11]. The dominant phenotype of SCID varies according to the country. For example, in China, 25 of 44 SCID patients had X-linked SCID [12], and in Serbia and Montenegro, 12 of 21 patients with SCID and Omenn syndrome belonged to the T–B–NK– group; 11 had mutations in RAG1 or RAG2 and that the XL-SCID group had the lowest number of patients (14%). These findings are in contrast with reports from Europe and America [10], possibly owing to genetic variations between populations and a higher rate of consanguinity in Iran.

**Discussion**

To our knowledge, this is the first comprehensive national report from Iran to show the clinical manifestations and laboratory and molecular findings for a large sample of patients with SCID. We found that most patients (34.9%) were positive for NK cells but negative for both T and B cells (T–B–NK–). The genetic analysis performed in 53% of patients revealed that most alterations were in RAG1 and RAG2 and that the XL-SCID group had the lowest number of patients (14%). These findings are in contrast with reports from Europe and America [10], possibly owing to genetic variations between populations and a higher rate of consanguinity in Iran.

Prophylaxis (antituberculosis, antibacterial, antiviral, and antifungal) was administered in 14 (22.2%), 36 (57.1%), 13 (20.6%), and 22 (34.9%) patients, respectively. Intravenous immunoglobulin was administered to 43 patients (68.3%). During the study period, appropriate donors were found for 21 patients, and 13 underwent hematopoietic stem cell transplantation (HSCT). However, HSCT was able to rescue only 8 patients (12.7%), who are still alive. The remaining 5 patients died of graft-versus-host disease or sepsis following transplantation. The mean age at the time of transplantation was about 8.3 months (range, 3-13 months). The mean survival time after transplantation until the end of this study was 17.7 months (range, 2-63 months). Overall, 50 of 63 patients (87.3%) died because suitable donors were not found.

The lymphocyte proliferation response to PHA was assessed in 11 patients. A low response was found in 10 cases. In spite of a mutation found in the RAG2 gene in the remaining patient, the response was normal (>50%).

Although mutation analysis was performed for 33 patients (53%), defects due to mutations were detected in only 17. Fourteen different mutations, including 5 novel mutations, were identified. As shown in Table 2, most alterations were detected in RAG2 and RAG1.
Serbia (2 months), and the Netherlands (2 months) [12 15 17], and better than Brazil (4.2 months) [16].

According to Roifman et al [18], a CD3+ T-cell count <500/μL is the most specific and sensitive parameter for separating patients with SCID from those with combined immunodeficiency. The authors also explained that a significant decrease in the response to the PHA test was found in all patients with CD3+ T cells <500/μL, although they hypothesized that the CD3 count might be normal in patients with profound T-cell dysfunction. Similarly, most of our patients were considered to have typical SCID in terms of the relevant feature of presenting CD3+ T-cell counts <500/μL. Nevertheless, new diagnostic criteria for SCID proposing <300/μL as the cut-off for SCID [18] have opened the debate on setting the most accurate value. Therefore, we also evaluated patients with CD3+ <300/μL, although we found no significant differences compared with patients with CD3+ <500/μL (77.8% vs 82.5%).

Most patients in our study had typical clinical manifestations and laboratory findings, including lymphopenia and low Ig levels except for maternal IgG; however, some had a high absolute lymphocyte count and normal Ig levels. The fact that these patients had an atypical presentation of lymphocyte subsets and normal serum Ig suggests that they comprise a subgroup of SCID patients with normal B lymphocytes who have the potential to produce measurable amounts of serum IgE and eosinophils [2,7]. In atypical cases, different clinical and immunological features were described, even within one family [19].

Pulmonary diseases were the most relevant complication (80%) in the patients we report. The other affected organs were the reticuloendothelial system, the gastrointestinal tract, skin, bones and joints, and the ear, nose, and throat. However, autoimmune diseases were not found.

We observed BCG-associated complications in only one-third of the patients, possibly because of factors such as genetic differences, the immune repertoire, vaccine ingredients, and lower T-cell counts [20] (the last factor was not confirmed in our study and requires further investigation).

Both BCG and other live attenuated vaccines are contraindicated in infants with SCID [20]. To avoid inoculating the patients with a positive family history (65% of patients in our study), awareness of the dangers of vaccination needs to be increased both in physicians and in patients’ families.

A recent approach to the early diagnosis of SCID involves the implementation of a newborn screening program (T-cell receptor excision circles assay) to hasten treatment and prevent the use of live vaccines in immunocompromised children [21-23].

The main limitations of the present study were the lack of laboratory facilities for performing genetic studies and the fact that the lymphocyte proliferation test was not performed in all participants. Nevertheless, our findings in this long-term study with a large sample may be of considerable value.

**Conclusion**

We found that autosomal recessive SCID was the most frequent inherited type of SCID in Iranian patients. However, diagnostic delay and the lack of suitable donors are the main

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**Table 2. Genetic Analysis**

<table>
<thead>
<tr>
<th>Immunophenotype</th>
<th>No. of Patients Who Underwent Genetic Evaluation</th>
<th>Genetic Tests, No.</th>
<th>No. of Cases With a Positive Result</th>
<th>Positive Mutations</th>
<th>Results Reported/Not Reported</th>
</tr>
</thead>
<tbody>
<tr>
<td>T–B–NK+ (22)</td>
<td>14</td>
<td>RAG1, RAG2</td>
<td>3 (15.8)</td>
<td>c.2687G&gt;A, p.W896X</td>
<td>Not reported</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td></td>
<td>c.835Cdel, p.A280fsX9</td>
<td>Not reported</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td></td>
<td>c.749_750Cdel, p.T250fsX17</td>
<td>Reported</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td></td>
<td>c.685G&gt;A, p.R229Q</td>
<td>Reported</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td></td>
<td>c.1432T&gt;C</td>
<td>Reported</td>
</tr>
<tr>
<td>Atypical (7)</td>
<td>3</td>
<td>RAG1, RAG2, ADA</td>
<td>1 (5.8)</td>
<td>c.2192C&gt;T, p.T731I</td>
<td>Reported</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td></td>
<td>c.685G&gt;A, p.R229W</td>
<td>Reported</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td></td>
<td>c.415G&gt;A, p.G139S</td>
<td>Not reported</td>
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<tr>
<td></td>
<td></td>
<td></td>
<td></td>
<td>c.1432T&gt;C, p.C478R</td>
<td>Not reported</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td></td>
<td>c.685C&gt;T, p.R229W</td>
<td>Reported</td>
</tr>
<tr>
<td>T–B–NK– (15)</td>
<td>7</td>
<td>ADA, IL-2, IL-7</td>
<td>2 (11.8)</td>
<td>c.2192C&gt;T, p.T731I</td>
<td>Reported</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td></td>
<td>c.956_960AAGAG, p.G319X</td>
<td>Reported</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td></td>
<td>c.704G&gt;A, p.R235Q</td>
<td>Reported</td>
</tr>
<tr>
<td>T–B+NK– (9)</td>
<td>4</td>
<td>IL-2, IL-7</td>
<td>3 (15.8)</td>
<td>c.31Gdel</td>
<td>Not reported</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td></td>
<td>c.685C&gt;T, p.R229W</td>
<td>Reported</td>
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<td></td>
<td></td>
<td>c.670C&gt;T, p.S225R</td>
<td>Reported</td>
</tr>
<tr>
<td>T–B+NK+ (10)</td>
<td>5</td>
<td>IL-7</td>
<td>1 (5.8)</td>
<td>c.31Gdel</td>
<td>Not reported</td>
</tr>
</tbody>
</table>

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causes of the high mortality observed among SCID patients. Increasing physicians’ awareness of the onset of serious infections and irreversible outcomes of BCG vaccination could improve the management of these patients.

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Conflicts of Interest

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References


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