Case Report

X-Linked Lymphoproliferative Syndrome and Common Variable Immunodeficiency May Not Be Differentiated by SH2D1A and XIAP/BIRC4 Genes Sequence Analysis

Nesrin Gulez, Guzide Aksu, Afig Berdeli, Neslihan Karaca, Sema Tanrıverdi, Necil Kutukculer, and Elif Azarsiz

Department of Pediatric Immunology, Ege University Medical School, 35100 Izmir, Turkey

Correspondence should be addressed to Elif Azarsiz, elif.azarsiz@ege.edu.tr

Received 29 September 2010; Revised 3 January 2011; Accepted 19 February 2011

Academic Editor: Robert A. Eisenberg

Copyright © 2011 Nesrin Gulez et al. This is an open access article distributed under the Creative Commons Attribution License, which permits unrestricted use, distribution, and reproduction in any medium, provided the original work is properly cited.

The X-linked lymphoproliferative syndrome (XLP) is a rare, inherited immunodeficiency characterized by recurrent episodes of hemophagocytic lymphohistiocytosis, hypogammaglobulinemia, and/or lymphomas [1]. It is exceptional among human X-linked immunodeficiencies as critical events occur after EBV infection. However, detailed analyses of the affected individuals revealed that the immune defect was broader than the impaired control of EBV infection [2]. Normally, primary EBV infection can occur without characteristic symptoms, or it can elicit mononucleosis of variable severity, but it regularly subsides. On the contrary, in the XLP patients, mononucleosis can be fatal with explosive activation and proliferation of cellular components of the immune system. The life threatening immunological defect is thus characterized by the defect of protection against the proliferation of EBV-transformed B cells [2–4].

Mutations in the signalling lymphocyte activation molecule- (SLAM-) associated protein SAP are responsible for 60–80% of cases of familial XLP [1, 4–6]. The gene defective in XLP has been identified at Xq25 and has been defined as SH2D1A. Mutation analyses of the gene are currently required for a definitive diagnosis of XLP [6].

Recently, mutations in the X-linked inhibitor of apoptosis (XIAP/BIRC4) gene, in families with XLP but without SAP mutation, have been defined [1, 7]. Zhu et al. evaluated 42 patients having XLP and identified XIAP deficiency at 2.4% of them [8].

XI-linked inhibitor of apoptosis protein (XIAP) has originally been described as an antiapoptotic protein that acts through direct inhibition of caspases. It is ubiquitously expressed in all normal tissues [9–11]. However, XIAP

1. Introduction

The X-linked lymphoproliferative syndrome (XLP) is a rare, inherited immunodeficiency characterized by recurrent episodes of hemophagocytic lymphohistiocytosis (HLH), hypogammaglobulinemia, and/or lymphomas [1]. It is exceptional among human X-linked immunodeficiencies as critical events occur after EBV infection. However, detailed analyses of the affected individuals revealed that the immune defect was broader than the impaired control of EBV infection [2]. Normally, primary EBV infection can occur without characteristic symptoms, or it can elicit mononucleosis of variable severity, but it regularly subsides. On the contrary, in the XLP patients, mononucleosis can be fatal with explosive activation and proliferation of cellular components of the immune system. The life threatening immunological defect is thus characterized by the defect of protection against the proliferation of EBV-transformed B cells [2–4].

Mutations in the signalling lymphocyte activation molecule- (SLAM-) associated protein SAP are responsible for 60–80% of cases of familial XLP [1, 4–6]. The gene defective in XLP has been identified at Xq25 and has been defined as SH2D1A. Mutation analyses of the gene are currently required for a definitive diagnosis of XLP [6].

Recently, mutations in the X-linked inhibitor of apoptosis (XIAP/BIRC4) gene, in families with XLP but without SAP mutation, have been defined [1, 7]. Zhu et al. evaluated 42 patients having XLP and identified XIAP deficiency at 2.4% of them [8].

X-linked inhibitor of apoptosis protein (XIAP) has originally been described as an antiapoptotic protein that acts through direct inhibition of caspases. It is ubiquitously expressed in all normal tissues [9–11]. However, XIAP
is also involved in a number of signaling pathways, for example, in nuclear factor kB, transforming growth factor, and bone morphogenetic protein signaling pathways [4, 7, 8]. Polymorphisms in the XIAP/BIRC4 gene may influence XIAP production and activity [12]. In addition, loss of function due to nonsense mutations in the XIAP gene have been observed in patients with XLP [1, 7, 8, 13].

Common variable immunodeficiency (CVID) is the most prevalent symptomatic primary immunodeficiency in humans [13]. Despite the discovery of genetic defects in ICOS, TNFRSF13b/TACI, and CD19, which prove that defects in single genes can be associated with a CVID phenotype, the diagnosis of CVID remains as clinical diagnosis of exclusion in most patients [13]. CVID patients may also have susceptibility to malignancy, especially lymphomas [14, 15]. Several genetically defined primary immunodeficiencies including XLP may mimic CVID.

Based on the previous reports that XLP and CVID may mimic each other, we report a 6-year-old boy, who has the clinical and laboratory signs of both disease. We revealed the difficulty of having an exact diagnosis although molecular analysis for XLP was performed.

2. Case Presentation

A six-year-old male patient, second child of nonconsanguineous healthy parents, was admitted to hospital with complaints of persistent general lymphadenopathy, especially at the cervical region, for the last two years. He had one healthy brother. On admission, he was four years old, and his weight was 15 kg (25th percentile) while his height was 112 cm (50–75th percentile). Clinical findings elsewhere including INFEVERS database. The same finding was confirmed that he had persistent EBV infection accompanied by mild hypogammaglobulinemia, possibility of XLP was searched out.

Initial screening for a mutation in the SH2D1A gene was carried out, and no gene defect was identified. Consequently, XIAP/BIRC4 gene was investigated. Sequence analysis was carried out on genomic DNA extracted from EDTA anti-coagulated venous blood using QiAamp DNA Blood Mini Kit (QiAGEN GmbH, Hilden Germany) according to the manufacturer's instructions. All 7 exons of XIAP/BIRC4 gene were amplified by polymerase chain reaction (PCR) using flanking intronic primers. PCR amplicons were sequenced in both directions with an ABI 3100 automatic DNA sequencer. Before sequencing, the PCR products were purified using Exo-SAP PCR purification Kit (Amersham Life Sciences).

Seven exons of XIAP/BIRC4 gene were searched out, and a heterozygous 1978G > A and 1996T > A nucleotide substitution in region of 3'UTR in exon 7 was detected. Defect in this region of XIAP/BIRC4 gene has not been reported elsewhere including INFEVERS database. The same finding was found in the mother while father and elder brother of the patient were normal for XIAP/BIRC4 sequencing (Table 2). Then, the samples of the patient were sent to a European center experienced in research of XLP patients, and XIAP protein expression was examined by Western Blot method [1]. Expression of XIAP was found to be normal in the lysate of peripheral blood lymphocytes of the patient.

On the third year of his follow-up in out-patient clinic, he developed multiple enlarged intraabdominal lymphadenomegaly and admitted to Pediatric Emergency Department with the diagnosis of invagination of intestines. After operation, in biopsy specimens, EBV-positive non-Hodgkin lymphoma (Burkitt) was observed (stage III and risk group II). Methotrexate, vincristine, cytarabine, etoposide, and intrathecauxi methotrexate, arabinoside-C combined therapies were applied and recovered in four months. Now, he is on regular intravenous immunoglobulin treatment and still in follow-up in pediatric immunology out-patient clinic.

3. Discussion

Diagnosis of XLP is complicated because of its clinical heterogeneity and rare incidence, particularly in males.
Table 1: Laboratory findings of patient.

<table>
<thead>
<tr>
<th></th>
<th>Patient</th>
<th>Normal values for his age</th>
</tr>
</thead>
<tbody>
<tr>
<td>IgG (mg/dL)</td>
<td>694</td>
<td>986.2 ± 209.6</td>
</tr>
<tr>
<td>IgM (mg/dL)</td>
<td>84.7</td>
<td>105.8 ± 40.8</td>
</tr>
<tr>
<td>IgA (mg/dL)</td>
<td>39</td>
<td>91.9 ± 37.4</td>
</tr>
<tr>
<td>WBC/mm³</td>
<td>14600 (26% pnl, 70% lym, 4% mono)</td>
<td>4000–10400</td>
</tr>
<tr>
<td>Absolute lymphocyte counts/mm³</td>
<td>10220</td>
<td>1500–5200</td>
</tr>
<tr>
<td>CD3⁺ (%-absolute count)</td>
<td>51–5212</td>
<td>55–79/1900–3600</td>
</tr>
<tr>
<td>CD19⁺ (%-absolute count)</td>
<td>37–3781</td>
<td>11–31/300–1200</td>
</tr>
<tr>
<td>CD3⁺CD4⁺ (%-absolute count)</td>
<td>17–1737</td>
<td>26–49/600–2000</td>
</tr>
<tr>
<td>CD3⁺CD8⁺ (%-absolute count)</td>
<td>27–2759</td>
<td>9–35/300–1300</td>
</tr>
<tr>
<td>CD19⁺CD40⁺ (%-absolute count)</td>
<td>37–3781</td>
<td>Expressed on all B lymphocytes</td>
</tr>
<tr>
<td>CD40L⁺ on activated T cells (%)</td>
<td>75</td>
<td>50 (24 hours after stimulation)</td>
</tr>
<tr>
<td>CD3–CD56⁺CD16⁺ (NK) (%-absolute count)</td>
<td>8–817</td>
<td>4–26/90–900</td>
</tr>
<tr>
<td>CD3–CD56⁺CD16⁺ (NKT) (%-absolute count)</td>
<td>1.8</td>
<td>&gt;4</td>
</tr>
<tr>
<td>Anti-nuclear antikor</td>
<td>Negative</td>
<td>Negative</td>
</tr>
<tr>
<td>Direct Coombs</td>
<td>Negative</td>
<td>Negative</td>
</tr>
<tr>
<td>AntiEBV VCA (IgM)</td>
<td>Negative</td>
<td>Negative</td>
</tr>
<tr>
<td>AntiEBV VCA (IgG)</td>
<td>Positive</td>
<td>Negative</td>
</tr>
<tr>
<td>AntiEBV EBNA (IgG)</td>
<td>Positive</td>
<td>Negative</td>
</tr>
<tr>
<td>EBV anti-EA</td>
<td>Negative</td>
<td>Negative</td>
</tr>
<tr>
<td>EBV DNA</td>
<td>Positive (lymph node biopsy)</td>
<td>Negative</td>
</tr>
</tbody>
</table>

Table 2: Molecular analysis of SH2D1A and XIAP/BIRC4 genes in patient and family members.

<table>
<thead>
<tr>
<th></th>
<th>Patient</th>
<th>Father</th>
<th>Mother</th>
<th>Brother</th>
</tr>
</thead>
<tbody>
<tr>
<td>SH2D1A</td>
<td>N</td>
<td>N</td>
<td>N</td>
<td>N</td>
</tr>
<tr>
<td>XIAP/BIRC4</td>
<td>Heterozygous 1978 G &gt; A and 1996 T &gt; A nucleotide substitution in 3′ UTR region of exon 7</td>
<td>Heterozygous 1978 G &gt; A and 1996 T &gt; A nucleotide substitution in 3′ UTR region of exon 7</td>
<td>N</td>
<td></td>
</tr>
</tbody>
</table>

manifesting with a phenotype consistent with XLP but without a family history. Lymphadenopathy, splenic and/or liver enlargement, fever, dysgammaglobulinemia, anemia, and thrombocytopenia are also common features of various immunodeficiencies, including CVID.

Clinical and laboratory findings of our patient were similar to previous XLP reports [1, 7, 13]. All patients, in these series had splenic enlargement. In Salzer et al. series [13], lymph node enlargement was reported in 55% of patients and hypogammaglobulinemia was reported in all of patients as in our case. Rigaud et al. [1] reported hypogammaglobulinemia in only three patients out of 12. They reported HLH in 11 of 12 patients, but in our case and also in other two series, HLH was not present [1, 7, 13].

There was a proven EBV infection in our patient. The presence of EBV infection was 75% in Rigaud et al. [1] and 22% in Salzer et al. [13] reports, respectively. Although colitis was the prominent symptom in Rigaud et al. [1] and Salzer et al. [13] series, it was absent in our and Doshi et al. cases. Rigaud et al. [1] reported the mean age of the patients as six years. Salzer et al. examined sequence analysis of BIRC4/XIAP genes in male patients who were previously diagnosed as CVID; therefore, the mean age of these patients reported in this study was 32 years [13].

NKT cells are a unique population of T cells that express an invariant T cell receptor which recognizes glycosphingolipid antigens presented by the CD1d molecule. These cells are known to be absent in patients with XLP due to SAP gene defect [4]. XIAP-deficient patients also show no T, B, or NK cell lymphopenia, but very low numbers of NKT cell [1, 4]. Rigaud et al. [1] thought that NKT cells might be particularly sensitive to apoptosis, and XIAP might be required for their survival and/or development [1]. In contrast, Marsh et al. [17] concluded that invariant NKT cells (iNKT) (defined as CD3 lymphocytes bearing an invariant TcR Vα14 Vβ11) were not decreased in the majority of XIAP-deficient patients. In our patient, the percentages of total T and NK cells are normal. B cells and CD8⁺ cells were slightly elevated, probably because of recurrent and chronic infections. NKT cell percentage (defined by the coexpression of CD3 and CD16/CD56) was found to be low by flow cytometric analyses compared to age-matched reference values of healthy children [18–21].

The clinical and some laboratory signs were good evidences to diagnose XLP. However, SH2D1A gene was normal, and the finding in 3′UTR region of seventh exon of XIAP/BIRC4 gene was not thought to be disease causing, because of reported public databases. In the previous reports,
Salzer et al. [13] pointed out that these 3’UTR nucleotide changes are polymorphisms. In addition, the mother who had the same amino acid changes was very healthy. Normal expression of XIAP protein confirmed our suggestions. Furthermore, it is very unlikely that XIAP is involved in the pathology of this patient as no association with lymphoma has been reported yet [22]. The clinical and laboratory resemblance and the findings of no gene mutation and normal XIAP protein expression led us to think that there may be another responsible gene for XLP.

Le Guerr et al. [23] described two CVID cases who developed B cell lymphomas, one related to EBV infection, 5 and 12 years after CVID had been diagnosed. Polizzotto et al. [24] reported a case of Burkitt lymphoma in the setting of CVID. Because of the occurrence of lymphomas during the course of CVID, the other diagnosis for our patient is still CVID. This patient also fulfills the criteria for CVID [25]. He will be followed up and managed as CVID until he presents new signs or until the identification of a new gene.

In conclusion, the differential diagnosis is not always easy between XLP and CVID patients. Molecular analysis for well-known mutated genes of XLP may not solve the problem and the patients have to be carefully long-term monitored and treated for life-threatening complications.

Acknowledgment

The authors thank Dr. Slyvain Latour (Institut National de la Sante et de la Recherche Medicale (INSERM) Unite 768, Lab du Developpement Normal et Pathologique du Systeme Immunitaire, Hopital Necker-Enfants Malades, Paris, France) for molecular analysing of SH2D1A and XIAP/BIRC4 genes and XIAP protein expression.

References


Submit your manuscripts at
http://www.hindawi.com