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NIJMEGEN BREAKAGE SYNDROME COMPLICATED WITH PRIMARY CUTANEOUS TUBERCULOSIS

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Abstract: Nijmegen breakage syndrome (NBS) is a rare autosomal recessive chromosomal instability syndrome characterized by severe immunodeficiency, growth retardation, microcephaly, a distinct facial appearance, and a high predisposition to lymphoid malignancy. We report a 7-year-old white girl with NBS associated with cutaneous tuberculosis. The patient presented with multiple red-brown, centrally scarring plaques on the leg and had neither pulmonary nor systemic manifestation of tuberculosis. Polymerase chain reaction testing using *Mycobacterium* genus- and *Mycobacterium tuberculosis* species-specific primers confirmed the clinical diagnosis of cutaneous tuberculosis. This is the first report describing the simultaneous presentation of NBS and cutaneous tuberculosis.

Key Words: Nijmegen breakage syndrome, cutaneous tuberculosis

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Nijmegen breakage syndrome (NBS) is a rare, autosomal recessive DNA-repair disorder characterized by severe immunodeficiency, progressive microcephaly, characteristic face,

growth retardation, spontaneous chromosomal instability, and a peculiar predisposition to cancer development.¹ The syndrome is caused by mutations in the *NBS1* gene located on chromosome 8q21, which encodes for nibrin, a DNA double strand break repair protein.^{2,3} The immune deficiency in affected individuals is profound and highly variable, with a tendency to progress with time.⁴ In particular, defective antibody response to common pathogens and low concentrations of serum immunoglobulin isotypes and IgG subclasses are common.⁴ Cellular immunity is also impaired and manifested in reduced T-lymphocyte proliferation and lymphopenia. The number of CD3⁺ and CD4⁺ T cells are usually decreased, whereas CD8⁺ cell depletion is rarely seen. Infectious complications mostly affect the respiratory tract and are caused by community-acquired rather than opportunistic pathogens.

Cutaneous tuberculosis, a very rare manifestation of mycobacterial infections has recently been reported from areas with a high incidence of HIV infection and multidrug-resistant pulmonary tuberculosis. Lupus vulgaris is a characteristic clinical form of cutaneous tuberculosis which typically occurs through hematogenous, lymphatic, or contiguous spread from underlying tuberculosis, also after primary inoculation.^{5,6} It may also be acquired exogenously following secondary inoculation, which is a rare complication of BCG vaccination.⁷ We present here a unique association of NBS with lupus vulgaris in a 7-year-old girl with neither pulmonary nor systemic manifestations of tuberculosis.

METHODS

Measurement of serum isotypes and lymphocyte surface markers were performed by routine immunochemical and flow cytometry assays. EDTA blood from the patients and family members was obtained and genomic DNA was isolated by using GenElute Blood Genomic DNA kit (Sigma-Aldrich GmbH, Budapest). Polymerase chain reaction was performed by using 1 primer pair to selectively amplify products of the hot spot region of *NBS1* gene in exon 6. Primer sequences are available on request. Amplified segments were purified using a MICROCON YM-100 Centrifugal Filter Devices (Millipore Co, Bedford, MA). Mutational analysis was performed using the BigDye Terminator Cycle sequencing kit (Applied Biosystems, Foster City, CA) and an ABI PRISM 3130 Genetic Analyzer (Applied Biosystems, Foster City, CA).

DNA was isolated from sections of the formalin-fixed paraffin-embedded skin biopsy samples. *Mycobacterium* DNA was detected by polymerase chain reaction, using *Mycobacterium* genus- and *Mycobacterium tuberculosis* species-specific primers.⁸ To demonstrate the presence of suitable DNA, a 268 bp segment of the human β -globin gene was also amplified.

All studies were approved by the institutional review board of the Medical and Health Science Center of the University of Debrecen. Informed consent was obtained from the parents.

CASE REPORT

This 7-year-old girl was admitted to our Department for immunologic investigation because of microcephaly, mild psychomotor retardation, and chronic multiple leg scars. She was the fourth child of a nonconsanguineous Hungarian family. The pregnancy was uncomplicated and the child was delivered at term with a birth weight of 3050 g; length, 48 cm; and head circumference, 31.5 cm (<third percentile). She had received BCG vaccination on the third day of life without complication and with no history of revaccination later. The patient received medical attention because of microcephaly, failure to thrive, and dysmorphic facial features (Fig., A and B, Supplemental Digital Content 1, <http://links.lww.com/INF/A607>). At 5 years of age, brain magnetic resonance imaging revealed occipital hypoplasia with dilatation of the occipital horns of ventricle. She was

hospitalized for *Campylobacter* enterocolitis at the age of 5, and for bacterial pneumonia at the age of 7. At age 6, she presented with painless, slowly, progressively expanding, reddish, discoid plaques on both legs (Fig., E and F, Supplemental Digital Content 1, <http://links.lww.com/INF/A607>).

On admission, her height was 97 cm; weight, 15 kg; and head circumference, 48 cm, all below the third percentile. She had a “bird-like” face characterized by a prominent midface and large ears, with a receding forehead and mandible, long philtrum, and micrognathia (Fig., A and B, Supplemental Digital Content 1, <http://links.lww.com/INF/A607>). She was fully cooperative and communicated adequately despite of striking retardation of physical growth and severe microcephaly. Café au lait spots and hypopigmented areas were noticed all over her body. Erythematous, indurated, scaly plaques on the right leg with areas of scarring was observed (Fig., E and F, Supplemental Digital Content 1, <http://links.lww.com/INF/A607>). The lesions were not painful on palpation.

RESULTS

Histopathologic examination revealed lupus vulgaris with well-developed noncaseating epithelioid granulomas, with inchoate central necrosis and some Langhans-type giant cells (Fig., C and D, Supplemental Digital Content 1, <http://links.lww.com/INF/A607>). No mycobacteria were identified in tissue by the Ziehl-Neelsen stain.

Molecular analysis of DNA extracted from paraffin-embedded skin specimen was positive for the *M. tuberculosis* IS6110 insertion sequence. Chest radiograph and abdominal sonography were normal. There was no history of tuberculosis and exposure to patients with tuberculosis. Screening for active tuberculosis in the family was negative.

Laboratory tests showed decreased concentrations of IgG (0.51 g/L) and IgA (undetectable) and normal IgM (1.14 g/L). Flow cytometry analysis of peripheral blood mononuclear cells disclosed decreased number of CD19⁺ cells (4%; 0.06 G/L) and CD4⁺ cells (24%; 0.25 G/L), and decreased CD4⁺/CD8⁺ ratio (0.55). The diagnosis of NBS was confirmed by mutation analysis of *NBS1* which revealed a c.657del5bp deletion mutation in exon 6 in both alleles, predicting a p.K219fsX234 change in the nibrin protein. Her 16-year-old sister was also diagnosed with NBS. Treatment was commenced with 400 mg/kg intravenous immunoglobulin infusions every 3 weeks. In addition, the patient was daily given rifampicin (150 mg), isoniazid (150 mg), and ethambutol (200 mg). A favorable outcome of the skin disease was observed, suggesting the beneficial effect of antimycobacterial therapy.

DISCUSSION

Cutaneous manifestations of tuberculosis are seen in less than 0.5% of cases.⁹ On the basis of data available from the ESID Registry, association of NBS with lupus vulgaris has not been reported before. Lupus vulgaris is a progressive and common form of cutaneous tuberculosis and usually results from dissemination from pulmonary tuberculosis. It follows a chronic course and without treatment may slowly extend with time. Lupus vulgaris is generally under-diagnosed because of its rarity and diverse clinical manifestations, and lack of awareness of the disease. Disseminated mycobacterium infection in a patient with NBS was reported by Resnick et al.¹⁰ This patient died at the age of 10 and had no skin manifestations.

NBS patients have defective humoral and cellular immunity which predispose them to infections. The impaired cellular immunity may play a role in the predisposition to infections by intracellular pathogens including mycobacteria. However, association of this pri-

mary immunodeficiency with tuberculosis is surprisingly rare. The case presented in this report is unique, because cutaneous tuberculosis is typically seen in patients with pulmonary tuberculosis. Cutaneous tuberculosis is extremely rare in patients having neither pulmonary nor systemic manifestation of tuberculosis. Our case is also notable because she developed lupus vulgaris, the well-organized form of the pathologic lesions seen in cutaneous tuberculosis. The clinical and pathologic lesions seen in cutaneous tuberculosis vary from scrofuloderma to lupus vulgaris. In scrofuloderma occurring mostly in patients with immunodeficiencies, histopathological examinations usually show a large number of mycobacteria in the lesions with a small to moderate number of lymphocytes in the granuloma. In contrast, lupus vulgaris is often seen in immunocompetent patients. In lupus vulgaris, lesional tuberculous bacilli are usually absent or scant, and cultures are often negative. Finally, our child had a favorable response to combined antituberculous therapy despite severe immunodeficiency.

Inborn errors of immunity including interferon γ receptor α and β chain deficiencies, signal transducer and activator of transcription 1 deficiency, interleukin (IL)-12 p40 deficiency, IL-12 receptor β 1 chain deficiency, and nuclear factor- κ B essential modulator deficiency have been proposed to predispose patients to infections with various mycobacterial species.^{11–14} Intriguingly, IL-12 β 1 receptor deficiency has been shown to associate mostly with *M. tuberculosis* infections but not with invasive BCG or environmental mycobacterial disease. The occurrence of *M. tuberculosis* infection in our patient should prompt research to define the expression and function of IL-12 and IL-12 receptor in NBS.

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