Multiple Family Members With Delayed Cord Separation and Combined Immunodeficiency With Novel Mutation in *IKBKB*

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**Background:** Inhibitor of kappa kinase 2 (IKK2) deficiency is a recently described combined immunodeficiency. It undermines the nuclear factor-kappa B (NF-κB) activation pathway.

**Methods:** The clinical and immunological data of four patients diagnosed with combined immunodeficiency (CID) from two related Saudi families were collected. Autozygosity mapping of all available members and whole exome sequencing of the index case were performed to define the genetic etiology.

**Results:** The patients had early onset (2–4 months of age) severe infections caused by viruses, bacteria, mycobacteria, and fungi. They all had hypogammaglobulinemia and low absolute lymphocyte count. Their lymphocytes failed to respond to PHA mitogen stimulation. A novel homozygous non-sense mutation in the *IKBKB* gene, c.850C>T (p. Arg284∗) was identified in the index patient and segregated with the disease in the rest of the family. He underwent hematopoietic stem cell transplantation (HSCT) from a fully matched sibling with no conditioning. The other three patients succumbed to their disease. Interestingly, all patients had delayed umbilical cord separation.

**Conclusion:** IKK2 deficiency causes CID with high mortality. Immune reconstitution with HSCT should be considered as early as possible. Delayed umbilical cord separation in CID patients may be a clue to IKK2 deficiency.

**Keywords:** inhibitor of kappa kinase beta/inhibitor of kappa kinase 2, *IKBKB*, combined immunodeficiency, hematopoietic stem cell transplant, delayed separation of the umbilical cord

**INTRODUCTION**

Nuclear factor-kappa B (NF-κB) is a transcription factor that plays a major role in various biological processes including the immune system development, activation, and regulation. This transcription factor complex is held inactive in the cytoplasm by the inhibitor of NF-κB (IκB). The classical activation pathway of NF-κB is triggered by a large number of extracellular stimuli through...
different receptors, which result in the phosphorylation of IκB by IκB kinase (IKK complex; IKKα, IKKβ/IKK2, and IKKγ/NEMO) and the subsequent release of NF-κB complex so it can translocate into the nucleus and drive target gene expression (1).

Different primary immunodeficiency disorders result from genetic defects that involve components linked to the NF-κB activation pathway (2). IKK2 deficiency is a recently described combined immunodeficiency disease that leads to impairment of NF-κB signaling (3). Affected patients develop early severe and recurrent infections caused by bacterial, viral, fungal, and mycobacterial organisms. Unlike patients with IKKα and IKKγ mutations, IKK2-deficient patients do not typically have ectodermal dysplasia. Indeed, out of 22 patients with IKK2 deficiency described in the literature, only one patient has ectodermal dysplasia in the form of conical teeth (3–7).

Herein, we describe the clinical manifestations and the immunologic features of four Saudi patients from two related families with a diagnosis of combined immunodeficiency due to novel homozygous non-sense mutation in IKBKB.

METHODS

The medical records of four Saudi patients from two related families with a diagnosis of combined immunodeficiency were reviewed. The clinical and immunological data were collected. Autozygosity mapping of all available members and whole exome sequencing of the index member were performed as previously described (8). Written consent was obtained from the parents. The study was approved by the Research Advisory Council at King Faisal Specialist Hospital and Research Center (RAC# 2121053).

RESULTS

Our patients (V:1, V:2, V:5, and V:8) belong to two related Saudi families (Figure 1A). They had early onset (2–4 months of age) severe infections caused by viruses, bacteria, mycobacteria, and fungi. The organisms include Klebsiella pneumonuia, Candida albicans, CMV, and BCG (Table 1), which were associated with different clinical manifestations (Table 1). Interestingly, all four patients had delayed umbilical cord separation at 2 months. They all displayed hypogammaglobulinemia. Where data is available, they had low absolute lymphocyte count (420–2,680 cells/mm³), and their lymphocytes failed to respond to PHA mitogen (Table 1). A novel homozygous non-sense mutation in the IKBKB gene, c.850C>T (p. Arg284*) (Figures 1B,C) was identified in patient V:5 within the candidate autozygome, and Sanger sequencing confirmed its segregation with the disease in the remaining living siblings and parents (Figures 1A,B). Three patients (V:1, V:2, and V:8) succumbed to their disease's infectious complications before commencing hematopoietic stem cell transplantation (HSCT).

Patient V:5 underwent HSCT from a fully matched sibling (10/10) at 19 months of age. To save his life and because of the disseminated BCGitis, he received no conditioning and no GVHD prophylaxis. The CD34 dose was $9.2 \times 10^6$/kg. He

![Figure 1](image-url)
had persistent lymphoid engraftment ranging from 7 to 40% (2 weeks to 2.5 years after HSCT) but no myeloid engraftment. He developed acute gut graft-vs.-host disease (GVHD) 2 months after HSCT that responded to cyclosporine A and steroid treatment. Six months later, he developed chronic gut GVHD that was controlled by the same medications. Although his lymphocyte subsets [CD3 2,409/mm$^3$ (79%), CD4 1,211/mm$^3$ (40%), CD8 1,183/mm$^3$ (39%), CD19 296/mm$^3$ (10%), CD16/56 307/mm$^3$ (10%)] and their proliferation response to PHA (relative ratio 127%) showed good recovery, clinically, he did not show good immune reconstitution. He continued to suffer from disseminated BCGitis with bony involvement requiring frequent surgical interventions for fociyosomatic osteomyelitis. More than 4 years after the transplant, he was still on anti-mycobacterial medications (cycloserine, clarithromycin, ethambutol, and levofloxacin), monthly IVIG, and pentamidine prophylaxis. The last chimerism test was done for the patients. Delayed cord separation was not shown good immune reconstitution. The severe disseminated BCGitis and history of GVHD precluded another trial of transplantation.

DISCUSSION

This is the largest cohort of IKK2 deficiency reported in the Saudi population. In general, the clinical features of our patients were not different from previously reported cases. Susceptibility to a wide range of infections caused by opportunistic and non-opportunistic organisms were commonality. The generalized maculopapular rash, hepatosplenomegaly, chorioretinitis, and CNS complications in our patients are probably related to the chronic CMV and disseminated BCG infections. All reported patients succumbed to their disease during infancy unless they had successful HSCT (Table 2). On average, the degree of lymphopenia was more profound in our patients (420–2,680/mm$^3$) compared to others (1,520–11,850/mm$^3$) (7). Interestingly, all our patients had delayed umbilical cord separation [normal separation occurs in 1–2 weeks (8)] Patient V:6, who died in infancy from a febrile illness before diagnosis and the identification of the genetic defect in the family, was also reported to have had delayed separation of the umbilical cord. None of the reported patients with IKK2 deficiency had delayed cord separation except one from Saudi Arabia, but with a different mutation (3–7). This feature may be a clue to IKK2 deficiency in patients with combined immunodeficiency. The mechanism by which IKK2B mutation may predispose to delayed cord separation is not clear. It has been reported that

### TABLE 1 | The clinical and immunological characterization.

| **Cord separation** | **Onset** | **Infections** | **Others** | **Outcome** |
|--------------------|-----------|---------------|------------|-------------|
| V:1 2 months      | 4 months  | Disseminated BCGitis | Extensive Maculopapular skin rash | Died at 6 months |
| V:2 2 months      | 3 months  | Perinatal CMV, Klebsiella pneumoniae sepsis, UTI, and pneumonia | Intraocular calcification | Died at 8 months |
| V:5 3 months      | 3 months  | Oral candidiasis and Disseminated BCGitis | Chorioretinitis | Transplanted |
| V:8 2 months      | 2 months  | Klebsiella pneumoniae sepsis and meningitis | Microcephaly | Died at 3 months |

*Normal range for age: at 3 months: IgG (1.76–5.81 g/L), IgM (0.24–0.89 g/L), IgA (0.046–0.46 g/L), IgE (0.18–3.76 IU/ml). At 6 months: IgG (2.15–7.04 g/L), IgM (0.35–1.02 g/L), IgA (0.081–0.69 g/L), IgE (0.44–16.3 IU/ml). Absolute lymphocyte count (ALC): 4,054–7,048 cells/mm$^3$. CD3: 2,409/mm$^3$ (79%), CD4: 1,211/mm$^3$ (40%), CD8: 1,183/mm$^3$ (39%), CD19: 296/mm$^3$ (10%), CD16/56: 307/mm$^3$ (10%) and their proliferation response to PHA (relative ratio 127%) showed good recovery, clinically, he did not show good immune reconstitution. The severe disseminated BCGitis and history of GVHD precluded another trial of transplantation.**
| Patient (references) | Onset | Clinical manifestation | HSCT | Age at 1st/2nd HSCT | Donor | HLA match | Conditioning GVHD prophylaxis | Outcome/cause of death | Age at death |
|----------------------|-------|-----------------------|------|---------------------|-------|-----------|-----------------------------|------------------------|-------------|
| P1 (7)               | Unknown | Disseminated BCGitis | No   | NA                  | NA    | NA        | NA                          | Died/infection        | <4 months   |
| P2 (7)               | 3 months | Bacteremia, otitis media, pyelonephritis, pneumonia, tinea capitis, and oropharyngeal and perineal candidiasis. Causative organisms: *Klebsiella pneumoniae*, *E. coli* and polymicrobial gram negatives | No   | NA                  | NA    | NA        | NA                          | Died/infection        | 14 months   |
| P3 (7)               | 2 weeks | Recurrent bacteremia, recurrent pneumonia, oral thrush, and disseminated HSV. Causative organisms are *E. coli* | Yes  | 19 months           | Father | 6/6       | MAC                         | Died/not reported     | 7 years     |
| P4 (7)               | 2.5 months | Disseminated BCGitis, disseminated CMV, disseminated adenovirus, bacteremia. Causative organisms: Streptococcus pneumoniae, *Stenotrophomonas maltophilia* | Yes  | 4 months            | Half-sibling | 10/10 | None              | Died/not reported     | 4 months     |
| P5 (7)               | 1 month | Disseminated BCGitis, disseminated CMV, candidal meningitis, oral thrush, varicella zoster infection, and bacteremia. Causative organisms: *E. coli*, *Morganella morgani*, *Staphylococcus aureus* | No   | NA                  | NA    | NA        | NA                          | Died/infection        | 5 months     |
| P6 (7)               | 3.5 months | Disseminated adenovirus, disseminated CMV, oral thrush, and bacteremia. Causative organisms: *E. coli*, *Pseudomonas aeruginosa* | Yes  | 4.5 months/not recorded | Mother/mother | 10/10 | None/MAC          | Died/not reported     | 8 months     |
| P7 (7)               | 2 months | *Listeria monocytogenes* bacteremia and meningitis and oral thrush | No   | NA                  | NA    | NA        | NA                          | Died/infection        | 2 months     |
| P8 (7)               | 1 month | Oral thrush, pneumonia and bacteremia. Causative organism: *E. coli* | Yes  | 5 months            | Unrelated | 6/6    | RiC                         | Died/not reported     | 2 years      |
| P9 (7)               | 1 month | Oral and perineal candidiasis, RSV pneumonia, bacteremia and meningitis. Causative organisms: *Staphylococcus aureus*, *Enterococcus faecalis*, *Enterococcus faecium*, and *Leuconostoc lactis* | No   | NA                  | NA    | NA        | NA                          | Died/infection        | 15 months    |
| P10 (7)              | 3 months | Oral thrush, bacteremia, and pneumonia. Causative Organisms: Parainfluenza, *Mycoplasma pneumoniae*, *Staphylococcus hominis*, and *Enterococcus faecium* | Yes  | 7 months            | Brother | 10/10 | MAC                         | Alive 7 years         | NA          |
| P11 (7)              | 2 months | Oral thrush, scabies, pneumonia, and meningitis. Causative organisms: rhinovirus and *Klebsiella pneumoniae* | No   | NA                  | NA    | NA        | NA                          | Died/infection        | 2.5 months   |

(Continued)
| Patient (references) | Onset | Clinical manifestation | HSCT | Age at 1st/2nd HSCT | Donor | HLA match | Conditioning | GVHD prophylaxis | Outcome/cause of death | Age at death |
|----------------------|-------|------------------------|------|---------------------|-------|-----------|-------------|------------------|----------------------|-------------|
| P12 (7)              | 3 months | Oral thrush, candidal UTI, bacteremia and brain abscesses, and pneumonia. Causative organisms: Serratia marcescens, E. coli, Klebsiella pneumoniae, RSV, coronavirus, Hemophagocytic lymphohistiocytosis | Yes | 8 months | Unrelated | 5/6 | MAC | Tacrolimus/ methotrexate | Alive 6 years | NA |
| P13 (7)              | 4 months | Sepsis | No | NA | NA | NA | NA | NA | Died/infection | 2.5 months |
| P14 (7)              | 2.5 months | Disseminated BCGitis, candidemia, and bacteremia. Causative organisms: E. coli, Stenotrophomonas maltophilia, Enterobacter cloacae | No | NA | NA | NA | NA | NA | Died/infection | 3.5 months |
| P15 (7)              | Newborn (newborn screening) | None | Yes | 2 months | Unrelated | 5/6 | MAC | Tacrolimus/ methotrexate | Died | 11 months |
| P16 (7)              | Newborn (family history) | None | Yes | 1 month | Unrelated | 5/6 | MAC | Tacrolimus/ methotrexate | Alive 6 months | NA |
| P17 (5)              | 5 months | Pneumocystis jiroveci pneumonia, disseminated BCGitis | No | NA | NA | NA | NA | NA | Died/BCGitis and cardiac arrest | 14 months |
| P18 (4)              | 2 months | Omphalitis, delayed separation of the umbilical cord, Salmonella sepsis, disseminated BCGitis, recurrent infections. Causative organisms: Acinetobacter, Enterobacter, Stenotrophomonas, and Achromobacter species, rotavirus, and Candida. Conical teeth. | No | NA | NA | NA | NA | NA | Died/massive bleeding after central line insertion | 25 months |
| P19 (6)              | 5 months | Candida, BCGitis | Yes | Not recorded | Father identical | MAC | Not recorded | Alive |
| P20 (6)              | 11 months | Candida, BCGitis, rotavirus | Yes | Not recorded | Cord blood | Not recorded | MAC | Not recorded | Died/infection | 2 months post HSCT |
| P21 (6)              | 7 months | Candida | Yes | Not recorded | Father | T-cell depleted haploidentical | MAC | Not recorded | Died/infection | 1 year post HSCT |
| P22 (6)              | 6 months | Candida, Klebsiella, OMV | Yes | Not recorded | Mother | T-cell depleted haploidentical | MAC | Not recorded | Died/infection | 13 months post HSCT |

MAC, myeloablative conditioning; RIC, reduced intensity conditioning; NA, not available.
Deficiency and adhesion was illustrated in cases of leukocyte adhesion process in healthy babies, and a defect in neutrophil chemotaxis was prominent during the cord separation in patients with IKKB2 deficiency. On the other hand, a gain-of-function (GOF) mutation in IKKB2 leads to a relatively mild form of combined immunodeficiency, ectodermal dysplasia, and immune dysregulation, where affected patients may live to their fourth decade. Although both diseases result in hypogammaglobulinemia and lymphopenia, patients with IKKB2 GOF mutation have lower number of naïve T lymphocytes with overactivated memory cells (14).

In our index patient, his unstable clinical condition and the disseminated BCGitis dictated the decision of non-conditioned HSCT. The fact that he had a sustained lymphoid engraftment without conditioning is likely due to the severe T-cell dysfunction in this type of immunodeficiency. We do not have a clear explanation why our index case did not clear the disseminated BCGitis inspite of the recovered in vitro lymphoproliferative response to PHA. One possibility is that the donor cells are anergic to BCG, but we could not test for that. So far, 13 out of 27 patients with IKKB2 deficiency including our cohort had undergone HSCT. Only five were alive at the time of reporting, while those who did not receive HSCT succumbed to their disease in their infancy or early childhood. All survivors received myeloablative conditioning except our patient, while the remaining eight patients received non-myeloablative conditioning (n = 5), reduced intensity conditioning (n = 1), or no conditioning (n = 2) (14).

Although it is difficult to draw a conclusion from such limited number of reported patients, it seems that no conditioning or reduced intensity conditioning are not sufficient to cure the disease, and myeloablative protocol is needed to establish a good immune reconstitution.

DATA AVAILABILITY STATEMENT

All datasets generated for this study are included in the article/supplementary material.

ETHICS STATEMENT

The studies involving human participants were reviewed and approved by The Research Advisory Council (RAC) at King Faisal Specialist Hospital and Research Center (RAC# 2121053). Written informed consent to participate in this study was provided by the participants' legal guardian/next of kin.

AUTHOR CONTRIBUTIONS

ZA acquired data and wrote the manuscript. FA and HS designed and conducted the genetic study. HA-M contributed to writing the transplantation and post transplantation part of the manuscript. MA, NA, and AAls substantially contributed to the acquisition and analysis of the clinical data for the work. AAls acquired data, supervised, critically revised and edited the work for intellectual content. All authors provided approval for publication of the content and agree to be accountable for all aspects of the work in ensuring that questions related to the accuracy or integrity of any part of the work are appropriately investigated and resolved.

FUNDING

The authors would like to thank the Deanship of Scientific Research for funding the study through the research Group Project No. RGP-190.

ACKNOWLEDGMENTS

An abstract of this manuscript was presented in the European Society of Immunodeficiency (ESID) 2017 conference. They kindly granted permission to publish this manuscript.

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**Conflict of Interest:** The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

The reviewer SP declared a past co-authorship with the author HA-M to the handling editor.

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