

CASE REPORT

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# A novel mutation in *ITGB4* gene in a newborn with epidermolysis bullosa, pyloric atresia, and aplasia cutis congenita

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## Abstract

**Background:** Epidermolysis bullosa with pyloric atresia (EB-PA), also known as Carmi syndrome, is an uncommon, autosomal recessive genodermatosis that typically affects the skin and gastrointestinal tract. EB-PA is caused by homozygous or compound heterozygous mutations in the integrin alpha 6 (*ITGA6*) gene on chromosome 2q31.1 or in the integrin beta 4 (*ITGB4*) gene on 17q25.1.

**Case presentation:** A male premature infant was born with aplasia cutis, atresia of the pylorus, and bilateral hydronephrosis. His clinical and imaging findings were compatible with EB-PA. A novel, small deletion of the last two bases in exon 6 and the first two nucleotides of intron 6 (c.565\_566+2del) in *ITGB4* gene was identified.

**Conclusion:** EB-PA-aplasia cutis congenita is known to be a non-treatable condition with a poor prognosis as the reported case. The novel mutation reported in this patient may lead to the lethal form of this disease. Identification of underlying genetic abnormality is critical to give genetic counseling.

**Keywords:** Epidermolysis bullosa, Carmi syndrome, Integrin alpha 6, Integrin beta 4

## Background

Epidermolysis bullosa (EB) is a clinically and genetically heterogeneous group of inherited skin diseases characterized by blisters followed by skin and mucosal erosions. EB has subgroups based on the ultrastructural level. There are four subgroups: EB simplex, junctional epidermolysis bullosa (JEB), dystrophic EB, and Kindler syndrome [1–3].

JEB with pyloric atresia (JEB-PA, OMIM: 226730) is a rare form of EB with multisystem involvement. Cutaneous manifestations, congenital pyloric atresia, and ureteral and renal anomalies such as renal collecting system defects, multicystic/dysplastic kidney, hydronephrosis, and absent bladder are the characteristic features of the disorder [4]. Cutaneous manifestations of

JEB-PA include severe mucocutaneous blisters, extreme skin fragility, atrophic scarring, and milia (small white spots). JEB-PA is caused by mutations in the *ITGA6* or *ITGB4* genes coding for subunit alpha 6 ( $\alpha 6$ ) or beta 4 ( $\beta 4$ ) of integrin [5, 6]. Additional features shared by JEB-PA include aplasia cutis congenita (ACC) that means congenital localized absence of the skin affecting the extremities, head, nail dystrophy, scarring alopecia, enamel hypoplasia, contractures, and dilated cardiomyopathy [7–9].

*ITGA6*, mapped to 2q31.1, is one of the genes known to cause this phenotype. *ITGA6* consists of 28 exons and encodes the integrin  $\alpha 6$  which is a member of integrin alpha chain family. On the other hand, *ITGB4* is located at 17q25.1 and encodes integrin  $\beta 4$  which is a component  $\alpha 6\beta 4$  integrin with integrin  $\alpha 6$ . Most of the patients with JEB-PA bear pathogenic variants in *ITGB4* gene, while *ITGA6* mutations are less common. Integrins are heterodimeric transmembrane receptors that play a

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critical role in cell surface adhesion and signaling. The  $\alpha6\beta4$  integrin is a hemidesmosomal protein, and its expression is altered by the mutations in *ITGA6* and *ITGB4* genes that cause structural defects in hemidesmosome [10–12].

Here, we describe a clinical case of a newborn with JEB-PA and ACC due to a novel mutation in *ITGB4* gene.

### Case presentation

A male preterm infant was born at 31 weeks' gestation by vaginal delivery as the first child to consanguineous parents with a birth weight of 1620 g. On prenatal ultrasonography, gastric dilatation and unilateral hydronephrosis were described. He received positive pressure ventilation at the delivery room and transferred to neonatal intensive care unit after stabilization. On physical examination, the skin was absent on the face, anterior side of the left forearm, both legs, and scrotum. He had multiple bullous lesions on the right arm, along with dystrophic nails. The radiograph of the abdomen showed gastric bubble resembling that of pyloric stenosis (Figs. 1, 2, and 3). Bilateral hydronephrosis and left ureterocele were detected by abdominal ultrasonography. Echocardiography and cranial ultrasonography were normal. The patient was intubated due to respiratory failure at the 2nd day of his life. Skin biopsy was performed and confirmed the diagnosis of EB. During follow-up, intravenous hydration was increased and parenteral nutrition and intravenous albumin were administered. He could



**Fig. 2** Dystrophic nails of the patient

not achieve the required stability for surgery and died on the 4th day on his life.

Written informed consent was obtained from the parent of the patient.



**Fig. 1** Aplasia cutis from the left arm, legs, and scrotum of the patient



**Fig. 3** Radiograph of the abdomen showing gastric dilatation due to pyloric atresia

### Genetic studies

Genomic DNA was isolated from 200  $\mu$ L of peripheral blood sample of the patient and his parents using Magna Pure LC DNA Isolation Kit-Large Volume and Magna Pure LC instrument (Roche Applied Science, Mannheim, Germany). All exons and the flanking regions of *ITGA6* and *ITGB4* were amplified by polymerase chain reaction using the 100 ng genomic DNA (the 260/280 ratios of DNA samples were  $\sim$  1.8) as a template for each PCR reaction. Conditions and primers for generating PCR products spanning all exons of the coding regions and flanking intronic sequences of the genes have been previously described elsewhere [6, 13]. After that, bidirectional Sanger sequencing analysis was performed using BigDye Terminator v3.1 Cycle Sequencing Kit on an ABI PRISM 3130 Genetic Analyzer (Thermo Fisher Scientific, Waltham, MA, USA).

A novel homozygous c.565\_566+2del mutation was found in *ITGB4* gene of the patient. His parents were found to be heterozygous for the same mutation (Fig. 4). This homozygous deletion in *ITGB4* disrupts the intron 6 donor splice site and expected to result in aberrant splicing and loss of function for the encoded  $\beta$ 4 integrin subunit. This variant was not listed in the 1000 Genomes (<http://browser.1000genomes.org>) or in the ExAC database (<http://exac.broadinstitute.org>).

### Discussion

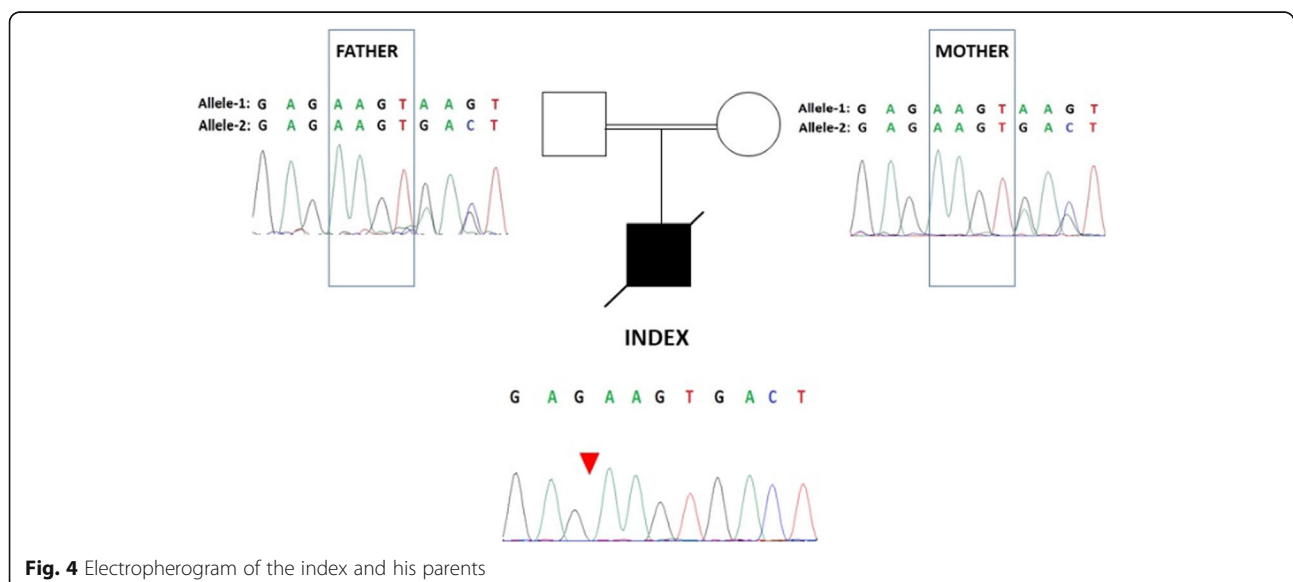
Here, we describe a case with JEB-PA and ACC emerging from a novel splice site mutation in *ITGB4* gene. EB is an inherited disease with an estimated frequency at 1 in 300,000, whereas pyloric atresia (PA) has an incidence of 1 in 100,000 live births. Familial PA with EB was first described in 1968, and pathophysiology was

first described by Carmi, so the disease has been called as ‘‘Carmi syndrome.’’ Gastrointestinal, urinary, pulmonary, and eye involvement are also reported to be associated with EB-PA. ACC has been previously described with EB-PA in few reports. Literature shows that JEB-PA-ACC is the most severe spectrum of the disease and suggested that ACC is the poorest prognostic factor of the combination [7–9, 11, 14].

Mutations in *ITGA6* and *ITGB4* cause an altered expression of  $\alpha$ 6 $\beta$ 4 integrin resulting in a structural defect of hemidesmosome which has an important role in stabilizing the association of the dermis with the epidermis. The defective hemidesmosomes cause blisters and erosions on the skin and gastrointestinal and urogenital abnormalities [11, 15]. Mutations in *ITGA6* are less frequent than in *ITGB4* [16]. There are more than 80 mutations that cause EB-PA and some of them are lethal. In lethal cases, the expression of  $\alpha$ 6 $\beta$ 4 integrin is absent [17]. We identified a novel, small deletion of the last two bases in exon 6 and first two nucleotides of intron 6 (c.565\_566+2del) in *ITGB4* gene in our patient with Carmi syndrome and ACC. This mutation causes splicing defect and premature stop codon which is responsible for the clinical features of our patient.

### Conclusions

EB-PA-ACC is known to be a non-treatable condition with a poor prognosis. There are few patients reported to be survived [16]. Although our patient was a premature baby and intestinal surgery could not be performed, we thought that the novel mutation reported in this patient may lead to a lethal form of the disease. Identification of underlying genetic abnormality is critical to give genetic counseling.



**Abbreviations**

ACC: Aplasia cutis congenita; EB: Epidermolysis bullosa; EB-PA: Epidermolysis bullosa with pyloric atresia; JEB: Junctional epidermolysis bullosa; JEB-PA: Junctional epidermolysis bullosa with pyloric atresia; JEB-PA-ACC: Junctional epidermolysis bullosa with pyloric atresia and aplasia cutis congenita

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**Authors' contributions**

GT and EO contributed to the conception. CDD, AG, EO, NYK, McGJ, and LL contributed to the acquisition and analysis. NYK, OE, McGJ, and LL contributed to the interpretation of the data. GT, CDD, and EO drafted the work. NYK, OE, BA, and SA critically revised the manuscript. GT, CDD, AG, EO, NYK, OE, BA, SA, McGJ, and LL approved the final manuscript.

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**Consent for publication**

Written parental informed consent was obtained for the publication of this clinical report and images.

**Competing interests**

The authors declare that they have no competing interests.

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