A Plakophilin-1 Gene Mutation in an Egyptian Family with Ectodermal Dysplasia-Skin Fragility Syndrome

Ebtesam M. Abdalla\textsuperscript{a} Cristina Has\textsuperscript{b}

\textsuperscript{a}Department of Human Genetics, Medical Research Institute, Alexandria University, Alexandria, Egypt; \textsuperscript{b}Department of Dermatology, University Medical Center Freiburg, Freiburg, Germany

Abstract

Ectodermal dysplasia-skin fragility syndrome (ED-SFS) is a rare genodermatosis caused by mutations in the \textit{PKP1} gene, encoding the desmosomal plaque protein plakophilin-1. Since its initial description in 1997, few individuals with this disorder have been reported to date. Here, we present the first Egyptian cases of ED-SFS, carrying a novel homozygous mutation in the \textit{PKP1} gene. Direct sequencing of the amplified DNA from the affected cases disclosed a G-to-T transversion at nucleotide position c.203–1 within intron 1 of \textit{PKP1} (c.203–1G>T). To the best of our knowledge, this mutation has not been previously described in the databases.

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www.karger.com/msy

E-Mail: karger@karger.com

E-Mail: drebtesamabdalla@yahoo.com

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Clinical Report

The 5- and 4-year-old female probands are the offspring of healthy first-cousin Egyptian parents and are the only affected members of the entire family (fig. 1A). Both patients had an average physical growth and normal intellectual developmental milestones, although they showed some motor delay because of the skin condition. The disease proceeded similarly in the 2 sibs; they were both born at full term with normal skin and scalp hair. Three days after birth, the condition started with extensive blistering all over the body, more severe on the limbs, while milder on the face and trunk. After rupture of the blisters, the underlying skin healed without scarring. At that time, the condition was thought to be a form of epidermolysis bullosa. Few months later, nail dystrophy started with recurrent falling off of the nails. Concurrently, the normal original hair was lost and replaced by scanty abnormal hair. At the age of ~1 year, the skin blistering improved dramatically, although the skin was still fragile and scalded very easily on minimal trauma. When the sibs were 2 years old, hyperkeratosis of the palms and soles became evident. Dentition started at the age of 1 year; the teeth were initially normal and then became dysplastic after 2 years. Moreover, the patients suffered from frequent chest infections that necessitated hospitalization. On examination, the patients displayed sparse friable scalp hair, curly eyelashes, cheilitis with perioral fissuring, dysplastic teeth, yellowish thickened nails with subungual hyperkeratosis, and palmoplantar hyperkeratosis, which was very severe in the soles, with painful fissuring. We also noted numerous erosions and crusts over the face and limbs (fig. 1B, C).

Since the constellation of clinical findings displayed by both patients was typical of ED-SFS, we obtained peripheral blood samples from the affected girls and their parents and performed DNA extraction. All coding exons and flanking intronic regions of the PKP1 gene (GenBank NM_001005337.2) were amplified by PCR. Direct sequencing of the amplified DNA from the affected cases disclosed a G-to-T transversion at nucleotide position c.203–1 within intron 1 of PKP1 (c.203–1G>T). The patients were shown to carry the mutation in a homozygous state, while the parents were heterozygous carriers. To the best of our knowledge, this variant has not been described before in the Human Gene Mutation Database, and it is not referenced in the Single Nucleotide Database (dbSNP 141, http://www.ncbi.nlm.nih.gov/projects/SNP/), in 1000 Genomes (http://www.1000genomes.org/) or on the Exome Variant Server (http://evs.gs.washington.edu/EVS).

Discussion

In 1997, McGrath et al. described the first case of a human genetic disease resulting from inherited mutations in a structural component of desmosomes. The clinical features included skin erosions and peeling with histological evidence of widening of intercellular spaces between keratinocytes, extending from the suprabasal to the granular layer [McGrath et al., 1997]. However, there were also early changes in hair and nails, suggesting that PKP1 had an additional role in the development of certain
ectodermal structures. Although all cases of ED-SFS reported so far showed similar clinical abnormalities, subtle differences between their phenotypic manifestations were noted [Hamada et al., 2002; Steijlen et al., 2004]. Hypohidrosis and cheilitis were constant features, while the extent of the alopecia and the degree of disability associated with the skin fissures varied between patients. This clinical diversity probably reflects the impact of the individual *PKP1* mutations on the levels of functional plakophilin-1 protein. However, the paucity of cases with this syndrome makes it difficult to establish accurate genotype-phenotype correlations.

The mutation c.203–1G>A was detected in a case of ED-SFS, and it was shown to generate 2 aberrant transcripts, predicted to lead to premature termination codons and absence of plakophilin-1 expression [McGrath et al., 1999]. Subsequently, the same mutation was recurrently reported as disease-causing in other patients from different origins [Sprecher et al., 2004]. In this report, the patients, who belonged to a consanguineous family of Arab Christian origin, were found to carry a G>T transversion at the same nucleotide position. Thus, the nucleotide position c.203–1 is likely to represent a mutational hot spot. Interestingly, splice site mutations are overrepresented in this gene. The various outcomes of leaky splice site anomalies, allowing residual protein expression, may correlate with improvement of the clinical phenotype [Wessagovit and McGrath, 2005]. For that reason, we intend to follow the progress of our patients with this rare genodermatosis. Given the clinical similarities between the present cases and the previously described male and female patients from different races [Zheng et al., 2005; Ersoy-Evans et al., 2006; Adhe et al., 2011; Boyce et al., 2012], it seems that neither sex nor ethnic origin significantly influence the phenotypic expression of mutations in the *PKP1* gene.

In summary, ED-SFS is a rare inherited skin disorder. We describe the first Egyptian patients suffering from ED-SFS, caused by a splice site mutation within the *PKP1* gene. This result provides further evidence for the previously established role of *PKP1* mutations in the pathophysiology of ED-SFS and expands the database of *PKP1* mutations.

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**References**


