SHORT COMMUNICATION

Darier's Disease: A novel ATP2A2 Missense Mutation at One of the Calcium-binding Residues

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Darier's disease (DD) is a rare autosomal dominantly inherited skin disorder caused by a heterozygous mutation in ATP2A2, which is expressed in both the skin and the brain and encodes for type 2 sarcoendoplasmic reticulum Ca-ATPase (SERCA2) (OMIM 124200). Gain of function due to Ca²⁺ leakage was suggested as the mechanism behind how the mutations cause DD (1). DD sometimes shows neuropsychiatric manifestations. Herein, we report a Japanese patient with DD, who has a novel mutation in one of the Ca²⁺-binding residues (CBR) in the endoplasmic reticulum (ER) membrane. A review of the literature on ATP2A2 mutations in CBR and DD phenotypes suggested differences between the mutations in terms of the accompa-

heterozygous ATP2A2 mutation. **DISCUSSION**

SERCA2 normally pumps Ca2+ from the cytoplasm to the ER. SERCA2 has 2 splice forms: SERCA2a and

ATP2A2 from diverse mammalian species (Fig. S1A¹).

c.2721A>C was not detected in any of the 200 control

alleles (100 control individuals) (data not shown).

Thus, the patient was diagnosed as having DD with the

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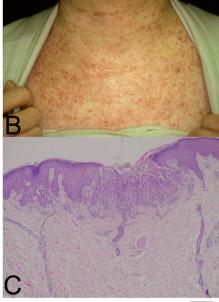
CASE REPORT

nying symptoms.

A 40-year-old woman with no familial history of DD or other serious comorbidities was admitted. Physical examination showed reddish papules on her back and breast (Fig. 1A, B). Histopathology of the papule on the back revealed suprabasal acantholysis with corps ronds and grains (Fig. 1C). Treatment with etretinate effectively resolved the skin lesions.

Following ethical approval, informed written consent was obtained from the patient in compliance with the Declaration of Helsinki guidelines. The coding regions including exon-intron boundaries of ATP2A2 were amplified from genomic DNA by PCR as described elsewhere (2). Direct sequencing of the patient's PCR products revealed the heterozygous missense mutation c.2721A>C (p. Glu907Asp) (Fig. 1D). The glutamine residue at 907, which plays an important role in Ca²⁺ transportation, was confirmed to be highly conserved in





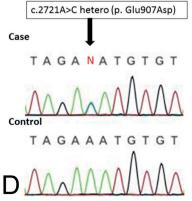


Fig. 1. Clinical feature, histological feature and ATP2A2 sequence data of the patient. The case presented with intensely pruritic papulae on her back (A) and breast (B). The histological features of the case show acantholysis and abnormal keratinisation. They also show a keratotic tier and suprabasal cleft formation. (haematoxylin and eosin staining, original magnification × 100; bar: 1 mm) (C). Sequence data of ATP2A2 in exon 18. Arrow indicates the heterozygous mutation c.2721A>C (p. Glu907Asp) (D).

SERCA2b. SERCA2a is restricted to the slow-twitch skeletal and cardiac muscles, whereas SERCA2b is ubiquitously expressed, and expressed especially highly in keratinocytes. SERCA2b has 11 ER transmembrane domains (TM) and 10 CBR. CBR is necessary for Ca²⁺ transport (Fig. S1B¹, Table SI¹). Each CBR seems to have different levels of importance for Ca²⁺ transport (3).

Three mutations causing amino acid alterations of 2 CBR (p.Asn767 and p.Asn795) have been reported (Table SI¹) (4–7). p.Asn767Ser was found in 4 cases of severe DD accompanied by neuropsychiatric symptoms or haemorrhages in the skin lesions, which suggests severe dysfunction of SERCA2 by the mutation. However, p.Asn767Asp is found in DD without accompanying symptoms. p.Asn795Ser was found in a DD patient whose phenotype was not described in detail. p.Glu907Asp, found in the present case, was the fourth mutation at CBR to be reported in the literature.

In conclusion, the present case and previous cases suggest that even missense ATP2A2 mutations at CBR do not always cause severe DD with accompanying symptoms. This may be because each CBR differs in its importance to Ca^{2+} transport.

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