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A Case of Inflammatory Generalized Type of Peeling Skin Syndrome Possibly Caused by a Homozygous Missense Mutation of CDSN

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Key Words

Inflammatory generalized peeling skin syndrome \cdot CDSN \cdot Corneodesmosomes \cdot Homozygous missense mutation

Abstract

A 54-year-old Japanese woman had repetitive superficial skin peeling and ensuing erythematous changes in the sites since infancy. Her parents had a consanguineous marriage, and she was the only individual affected in her family tree. The erythematous changes seemed to worsen in the summer. Histologically, hyperkeratosis and splitting of the epidermis within the stratum corneum was noted, and electron microscopy revealed shedding of corneal cells in the horny layer and normal-looking corneodesmosomes. Gene analysis revealed a homozygous missense mutation at c.1358G>A in *CDSN*. Electron microscopic examination of the length and number of corneodesmosomes revealed statistically significant shortness and sparsity in the affected individual (mean \pm SD 386.2 \pm 149.5 nm) compared with that of an age- and site-matched control (406.6 \pm 182.3 nm). We speculate that this size shrinkage of corneodesmosomes might be the result of a missense mutation of *CDSN* and that this could be one of the factors contributing to the pathological process of skin peeling.

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Introduction

Peeling skin syndrome (PSS) is a rare autosomal recessive cornification disorder that is both clinically and genetically heterogeneous. PSS is characterized by shedding of the stratum corneum upon birth or in early infancy [1] and can be classified into two subtypes: acral PSS (peeling skin confined to the acral surface) and generalized PSS (systemic peeling). A mutation in transglutaminase 5 was identified as giving rise to acral PSS [2, 3]. The generalized type of PSS is further subclassified into non-inflammatory PSS (type A PSS) manifesting year-round peeling of the stratum corneum without inflammatory change, and inflammatory PSS (type B PSS) showing inflammatory changes resembling mild congenital ichthyosiform erythroderma. Type A PSS has been shown to have its genetic basis in the *CHST8* gene encoding a Golgi transmembrane N-acetylgalactosamine-4-O-sulfotransferase (GALNAC4-ST1) in 2012 [4]. Type B PSS is known to be associated with mutations in several loci of corneodesmosin, a specific constituent protein of corneodesmosomes, in all of which the mode of mutation reported so far is exclusively homozygous nonsense, leading either to a truncated corneodesmosin protein or to messenger RNA transcript decay [5–8]. We report a case of type B PSS possibly caused by a homozygous missense mutation of *CDSN*.

Case Report

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The patient was a 54-year-old Japanese woman who had experienced repetitive superficial skin peeling over the entire body and ensuing erythematous color changes at the sites since infancy. She did not have any children and was an offspring of a consanguineous marriage, but neither her four siblings nor anyone in her family tree reported any of the symptoms experienced by the patient. She claimed that the erythematous change worsened in the summer compared to the winter. Shedding of the skin was more prominent on the palms, forearms and intertriginous areas than on the torso (fig. 1). Hair shaft analysis did not show trichorrhexis invaginata.

A skin biopsy specimen was taken from the site of peeling on her right upper arm. Hematoxylin and eosin (H&E) staining revealed basket weave-type hyperkeratosis with half of the stratum corneum in the right upper portion of the image seemingly sloughed off (fig. 2). The figure shows the original stratum corneum with the residual half above it and the area left of the center missing the upper corneal sheet. In the upper dermis, perivascular mononuclear infiltrate and mild dilation of capillaries were observed. Electron microscopy (fig. 3) showed splitting of the corneocyte rows. However, whether this change stemmed from a pathological cause or was an artefact is not clear. Corneodesmosomes were found between the corneocytes, and recognizable abnormalities in the density and morphology of corneodesmosomes could not be detected by viewing the images. The keratin aggregate pattern within the corneocytes and in the cornified envelope appeared normal.

Peripheral blood was collected for genetic analysis with the patient's consent. Biological materials could not be obtained from the patient's family members. As a result, a homozygous missense mutation was revealed at nucleotide position 1358 substituting adenine for guanine (c.1358G>A) in the *CDSN* gene, resulting in an exchange of the amino acid serine for asparagine (p.S453N). This mutation was not detected amongst the 100 healthy Japanese controls.

In order to determine whether there was any structural abnormality in our electron microscopic findings, we measured the length and number of corneodesmosomes per unit length both in our patient and in an age- and site-matched control. The corneodesmosomes

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measured were confined to the second corneocyte layer above the stratum granulosum and carefully chosen from those that were horizontal to the stratum corneum. The results revealed a statistically significant shortening of corneodesmosomes in our case (mean ± SD 386.2 ± 149.5 nm) compared with the healthy control (446.3 ± 185.8 nm), as well as a lower corneodesmosome density in the affected individual (0.860 ± 0.233 μ m⁻¹) than in the control (1.309 ± 0.413) (table 1).

Discussion

The corneodesmosome, a modified form of desmosome found in the stratum corneum and inner root sheath of the hair follicles, differs ultrastructurally from the desmosome and is shown to play an important role in intercellular adhesion between corneocytes in the cornified cell layer [9]. The extracellular part of the corneodesmosome is mainly composed of corneodesmosin, desmoglein 1 and desmocollin 1 [10], which are together incorporated into the normal desquamation process as they lose their adhesive properties from gradual cleavage by enzymes known as kallikrein-related peptidases and cathepsins in the stratum corneum. Secreted from the lamellar granules of keratinocytes between the granular layer and the corneal layer, corneodesmosin, a 52–56-kDa glycoprotein, is incorporated into a structure known as the desmoglea before they are transformed into corneodesmosomes in the process of cornification.

A series of reports has established that type B PSS is caused by deleterious mutations in the *CDSN* gene encoding corneodesmosin. A comparison of all reported loci of mutation, their histological findings and serum IgE levels (table 2) revealed that the homozygous missense mutation in the present case lay very far downstream from the previously known mutations, which were all homozygous nonsense. Most of the type B PSS cases presented remarkably high serum IgE compared with the present case, suggesting that the relative mildness of symptoms and the relatively lower serum IgE level may correlate with less barrier function deficiency in our patient.

The biopsy specimen in all of the reported cases demonstrated compact hyperkeratosis and detachment of the corneal sheet immediately above the stratum granulosum as shown by H&E staining (table 2). In contrast, our case showed neither stuck hyperkeratosis nor subcorneal cleavage, and the image was rather characterized by basket weave-type hyperkeratosis and possible shedding of the corneal sheet amid the stratum corneum. This histopathological aberrance from previously reported cases might be explained by the presence of solid corneodesmosomes detected by electron microscopy in our patient. Although it is difficult to determine to what extent our missense mutation affected the adhesive property of the corneodesmosomes, it is possible that they may have remained partly functional, though not sufficiently so to execute the normal desquamation process.

Measurements of the corneodesmosomes by electron microscopy revealed shrinkage and sparsity in our case (table 1), but the structure itself seemed intact, as evidenced by the homogeneously electron-dense structure. Only one report of type B PSS, in which a homozygous nonsense mutation was identified at the upstream locus of *CDSN*, examined the electron microscopic view of corneodesmosomes and revealed hardly visible corneodesmosomes immediately above the stratum granulosum [5]. This might explain how the desmoglea does not phenotypically form and become visible without non-truncated corneodesmosin, and furthermore how the downstream locus with homozygous missense mutation in our case might affect morphologically normal-looking corneodesmosome formation.

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We have reported a case of type B PSS possibly caused by a homozygous missense mutation of CDSN. Further cumulative, analytic studies of CDSN mutations will undoubtedly elucidate the genotype-phenotype correlation as well as the detailed function of corneodesmosin.

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	Our case	Healthy control	F value
Length of corneodesmosomes, nm (mean ± SD)	386.2±149.5	446.3±185.8	0.015 (<0.05)
Density of corneodesmosomes, µm-1 (mean ± SD)	0.860±0.233	1.309±0.413	0.038 (<0.05)

Table 1. Comparison of the length and number of corneodesmosomes per unit length between our case and an age- and site-matched control

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 Table 2. Comparison between previously reported cases and our case in terms of mutation loci, mode of mRNA translational errors, histopathological features and serum IgE level

Reference	Genetic mutations (affected amino acids)	Mode of mRNA translational errors	H&E staining	Serum IgE, IU/ml
0ji et al. [5], 2010	c.175A>T (p.K59X)	homozygous nonsense	compact hyperkeratosis; detachment directly above the striatum granulosum	>2,000
Telem et al. [7], 2012	c.164_167dup GCCT (p.T57PfsX6)	homozygous duplication \rightarrow stop codon	compact hyperkeratosis; subcorneal separation	unknown
Israeli et al. [6], 2011	c.746delG (p.G249VfsX40)	homozygous single nucleotide deletion → stop codon	compact hyperkeratosis; intracorneal and subcorneal separation	30,375
Mazereeuw et al. [8], 2011	c.754delC (p.P252LfsX37)	homozygous single nucleotide deletion → stop codon	compact hyperkeratosis; subcorneal separation	23,000
Our case	c.1358G>A (p.S453N)	homozygous missense	basket weave-type hyperkeratosis; intracorneal separation	369

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Fig. 1. Peeling of the skin was noted on both palms, thighs and buttocks. Erythematous color changes at the site of peeling were observed. Shedding of the skin was observed more prominently on the palms, forearms and intertriginous areas than on the torso.

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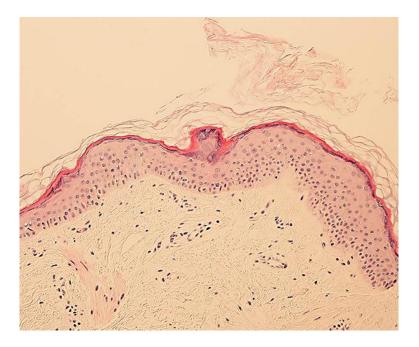


Fig. 2. Basket weave-type hyperkeratosis with half of the stratum corneum in the right upper portion of the image seemingly sloughed off. The figure shows the original stratum corneum with the residual half above it and the area left of the center missing the upper corneal sheet.

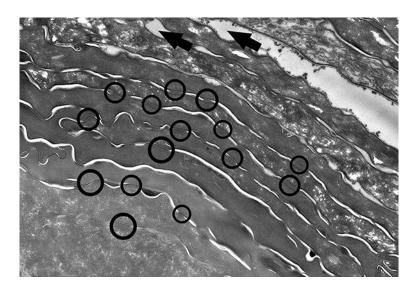


Fig. 3. Electron microscopy showed split formation (arrows) between the corneocyte rows. However, whether this change stemmed from a pathological cause or was an artefact is not clear. As shown by the circles, corneodesmosomes were found between the corneocytes, and abnormalities in the density and morphology of the corneodesmosomes were not obvious.

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