Review Article

Zinc in Keratinocytes and Langerhans Cells: Relevance to the Epidermal Homeostasis

Youichi Ogawa, Manao Kinoshita, Shinji Shimada, and Tatsuyoshi Kawamura

Department of Dermatology, Faculty of Medicine, University of Yamanashi, Yamanashi 409-3898, Japan

Correspondence should be addressed to Youichi Ogawa; yogawa@yamanashi.ac.jp

Received 24 September 2018; Accepted 10 November 2018; Published 9 December 2018

Guest Editor: Toshiyuki Fukada

Copyright © 2018 Youichi Ogawa et al. This is an open access article distributed under the Creative Commons Attribution License, which permits unrestricted use, distribution, and reproduction in any medium, provided the original work is properly cited.

In the skin, the epidermis is continuously exposed to various kinds of external substances and stimuli. Therefore, epidermal barriers are crucial for providing protection, safeguarding health, and regulating water balance by maintaining skin homeostasis. Disruption of the epidermal barrier allows external substances and stimuli to invade or stimulate the epidermal cells, leading to the elicitation of skin inflammation. The major components of the epidermal barrier are the stratum corneum (SC) and tight junctions (TJs). The presence of zinc in the epidermis promotes epidermal homeostasis; hence, this study reviewed the role of zinc in the formation and function of the SC and TJs. Langerhans cells (LCs) are one of the antigen-presenting cells found in the epidermis. They form TJs with adjacent keratinocytes (KCs), capture external antigens, and induce antigen-specific immune reactions. Thus, the function of zinc in LCs was examined in this review. We also summarized the general knowledge of zinc and zinc transporters in the epidermis with updated findings.

1. Introduction

The epidermis is the outermost layer of the skin and is thus continuously exposed to various kinds of external substances and stimuli that can lead to potential harm. To counteract these risks and maintain homeostasis, the epidermis provides a barrier against the external environment. The importance of preserving epidermal homeostasis is evidenced by a reduction in the development of atopic dermatitis (AD) when moisturizer is applied to the skin during the neonatal period [1, 2]. Reports also suggest that skin barrier disruption leads not only to the development of AD but also to other allergic diseases such as asthma, food allergies, and allergic rhinitis [3, 4].

The epidermis is composed predominantly of keratinocytes (KCs) plus a small number of Langerhans cells (LCs), melanocytes, and epidermal-resident memory T cells, as well as others. The murine epidermis also contains unique dendritic epidermal T cells, a type of γδT cells. The epidermal KCs are categorized into four layers, namely, the stratum basale, stratum spinosum, stratum granulosum (SG), and stratum corneum (SC). A functional epidermal barrier depends on the existence of the SC and tight junctions (TJs) formed in the SG (Figure 1) [5, 6]. LCs can link with KCs to form TJs (Figure 1) [7, 8].

The human body contains 2–3 g of zinc (Zn), with approximately 5% of the total Zn found in the skin [9]. The concentration of Zn is higher in the epidermis than in the dermis and subcutaneous tissue, which may be due to the Zn requirement for active proliferation and differentiation of KCs [10]. Zn facilitates over 1000 enzymatic reactions and is indispensable for over 2000 transcriptional activities [11–13]. Zn finger proteins are involved in various physiological reactions [14–16]. Moreover, approximately 10% of human proteins bind to Zn [17]. Therefore, the dysregulation of epidermal Zn levels due to nutritional deficiency or genetic abnormalities of Zn transporters affects various enzymatic reactions, transcriptional activities, and Zn finger protein functions in the epidermis, leading to the disruption of skin homeostasis [18–20]. This review is aimed at highlighting the association between Zn and epidermal barrier function to understand the importance of Zn in skin immunity. We also summarize the function of Zn and Zn transporters in the epidermis and epidermal cells.
2. Zinc and the Epidermal Barrier

The epidermal barrier function depends largely on the presence of SC and TJs formed in the SG [4, 21]. In this respect, homeostasis of the epidermis depends on normal KC differentiation (keratinization). LCs, a type of tissue-resident macrophages, also form TJs with KCs through linkages with KCs [7]. Most of the skin is covered with stratified epithelia that have both SC and TJs. However, skin appendages such as hair follicles and sweat glands lack SC, so TJs are the sole barrier structure in the skin appendages. The sections below summarized the involvement of Zn in the formation and function of SC and TJs.

2.1. Stratum Corneum. The outer layer of the SC consists of a cornified layer made of flattened and denucleated KCs (or corneocytes) and a SC-specific barrier structure called the corneified envelope that replaces the KC cell membranes. Through the process of terminal differentiation, SG KCs produce two membrane-circumscribed granules, keratohyalin granules and lamellar bodies. The former contains “intracellular” components of the SC such as filaggrin (FLG), loricrin, and keratin filaments, whereas the latter contains “extracellular” components such as lipids, corneodesmosin, and kallikreins. All of these intracellular and extracellular proteins are crucial for the formation and/or function of the epidermal barrier [21].

2.1.1. Zinc and Filaggrin and Its Metabolism. Among the components of keratohyalin granules, FLG has a crucial role in maintaining normal epidermal barrier function. FLG-deficient mice show an impaired SC barrier function and develop spontaneous dermatitis [22, 23]. Studies have shown that loss-of-function mutations in the FLG gene are strongly associated with the development of AD and ichthyosis vulgaris [24, 25]. These mutations were shown to range from 25 to 50% in the Northern European and Asian populations with these ailments [25, 26]. Moreover, FLG gene mutations were demonstrated as the strongest risk factor for AD in the genome-wide association studies (GWAS) [27]. These indicate the critical involvement of FLG in AD pathogenesis mediated by the disruption of the epidermal barrier.

FLG is produced in the SG as proflaggrin (FLG polymer) and is stored in keratohyalin granules. At the transition to the SC, the polymer is processed to the monomer by proteases such as Prss8 and SasPase [28, 29] and then binds to keratin and forms the fundamental structure of the corneocytes. At the outermost layer of the SC, FLG is citrullinated by peptidylarginine deiminase and then dissociated from keratin filaments [30]. These dissociated FLG are degraded to free amino acids, including glutamine, arginine, and histidine. The FLG-derived histidine-rich proteins are converted into urocanic acid (UCA) and pyrrolidine carboxylic acid (PCA) by proteases. UCA absorbs ultraviolet, maintains the acidic pH, and suppresses excess LC activation [31, 32]. PCA is a source of natural moisturizing factors. As a result, FLG is indispensable to the framework of SC and its metabolites are important for maintaining the epidermal barrier function (Figure 2).

Zn is involved in the regulation of FLG expression as well as its metabolism. It facilitates FLG production by increasing the activity of Prss8 [33]. Alternatively, Zn can also suppress FLG metabolism by decreasing PAD activity [34]. Moreover, Zn is required for histidine conversion to UCA [35] (Figure 2). Propionibacterium acnes (P. acnes) induce excess KC proliferation and FLG expression in the epidermis through the induction of IGF-1, which activates the IGF-1 receptor (IGF-1R) on KCs. This causes follicular plugging that is frequently observed in patients with acne vulgaris. In these cases, Zn helps to maintain homeostasis by directly suppressing the induction of IGF-1 and IGF-1R and the overexpression of FLG [36].

OVO-like proteins (OVOls) are transcribed from ubiquitously conserved genes encoding a C2H2 zinc finger transcription factor in mammals [37]. Mutations in the Ovol1 gene, as well as the FLG gene, were demonstrated as a risk factor for AD in the GWAS [38–40]. Consistent with this finding, Ovol1-knockout mice show rapid disruption of the epidermal barrier [41]. Ovol1 regulates transcription...
Zn can also suppress FLG metabolism by decreasing PAD activity.

Alternatively, and pyrrolidine carboxylic acid (PCA) by proteases. Zn facilitates histidine-rich proteins are converted into urocanic acid (UCA) including glutamine, arginine, and histidine. The FLG-derived fligmatic proteins are involved in the regulation of FLG expression [43, 44]. Given this, it is evident that Zn facilitates histidine transcription via Sp1, a C2H2-type Zn finger motif [42]. Additionally, HAT activity of p300 is negatively regulated by histone deacetylases (HDACs), which are Zn-dependent hydrolases [42]. Thus, OVOL1 expression and activity are regulated by Zn and Zn-related epigenetic enzymes. Aryl hydrocarbon receptor (AHR) is a ubiquitous ligand-activated transcription factor that is activated by both endogenous and exogenous ligands. AHR activation induces nuclear translocation of OVOL1, leading to the upregulation of FLG expression [43, 44]. Given this, it is evident that Zn and various Zn finger proteins are involved in the regulation of FLG expression.

### 2.1.2. Zinc and Cornified Envelope, Intercellular Lipid Lamellae, and Corneodesmosome

The cornified envelope (CE) is formed of keratins enclosed within an insoluble protein shell just beneath the corneocyte cell membrane [45]. It provides a solid physical barrier and consists of a 10 nm thick layer of highly cross-linked insoluble proteins, such as involucrin, envoplakin, periplakin, loricrin, and small proline-rich protein. A 5 nm thick layer of intercellular lipid lamellae composed of ceramides, free fatty acids, and cholesterol is covalently bound to these proteins. Adhesion between corneocytes is assumed to occur by the desmosome apparatus (corneodesmosome), composed of desmosomal cadherin, armadillo proteins, and plakins. The CE, intercellular lipid lamellae, and corneodesmosome are essential for effective physical and water barrier functions in the epidermis. The role of Zn in the formation and function of these proteins has not yet been elucidated.

#### 2.1.3. Zinc and Corneocyte Desquamation

Shedding of corneocytes at the outermost layer of the SC is called desquamation and is an important step for maintaining epidermal homeostasis and preventing hyperkeratosis. This step is primarily assumed by a proteolytic cleavage of kallikrein (KLK-) related peptidases. Out of 15 KLK proteins, human KCs express nearly all of them [46]. Among these KLKs, corneocyte desquamation is mainly conducted by KLK5, KLK7, and KLK14 [47]. The proteolytic activity is dependent on pH in the SC and is regulated by a cocktail of protease inhibitors, including lymphoepithelial Kazal-type 5 serine protease inhibitor (LEKTI) encoded by the serine protease inhibitor Kazal-type 5 (SPINK5) [48]. Importantly, the proteolytic activity of KLK5, KLK7, and KLK14 is impaired in the presence of Zn [49–51]. Furthermore, the expression of several KLK-related peptidases, including KLK5 and KLK7, is negatively regulated by transcriptional factor specificity protein 1 (Sp1), a C2H2-type Zn finger protein [52].

Several reports demonstrated the association between AD and Zn deficiency in humans and mice [53–55]. Consistent with this finding, the skin pH is increased in patients with AD [56], possibly due to the reduced UCA production caused by Zn deficiency (see Section 2.1.1). Additionally, Sp1 expression is decreased in the epidermis of patients with AD [52], and, as a consequence, KLK activity is often enhanced. Since KLKs process prointerleukin- (IL-) 1α and IL-1β that are abundantly stored in the corneocytes to active forms, this enhanced KLK activity leads to skin inflammation [57]. Additionally, KLK5 and KLK14 activate proteinase-activated receptor- (PAR-) 2, which is a G-protein-coupled receptor expressed on KCs, leading to the elicitation of itch common with AD [58]. Collectively, Zn suppresses excess inflammation and itching by inhibiting excess KLK activity. This action helps to maintain the epidermal barrier by suppressing IL-1-mediated FLG downregulation [57].

#### 2.2. Zinc and Tight Junctions

TJs that are formed in the SG seal the intercellular spaces between SG KCs, thereby regulating the movement of water and inorganic ions via the paracellular pathway [59]. The major components of TJs are the transmembrane claudin proteins. Among the claudins expressed in the epidermis, claudin-1 is most critical for the formation and function of TJs, as confirmed in claudin-1 knockout mice, which showed severe dehydration and death soon after birth without modulating SC formation [60]. Providing further support to the importance of claudin-1, the expression is decreased in the epidermis of patients with AD [61]. Claudin-1 acts as a paracellular tightness barrier, maintaining SC hydration and barrier function. This is consistent with the findings that Zn deficiency is associated with increased SC hydration and barrier function in humans and mice [53–55].

**Figure 2:** Filaggrin and its metabolism. FLG is produced in the SG as profilaggrin (FLG polymer) and is stored in keratohyalin granules. At the transition to the SC, the polymer is processed to the monomer by proteases such as Prss8 and SASPase and then binds to keratin and forms the fundamental structure of the corneocytes. At the outermost layer of the SC, FLG is citrullinated by peptidylarginine deiminase and then dissociated from keratin filaments. These dissociated FLG are degraded to free amino acids, including glutamine, arginine, and histidine. The FLG-derived histidine-rich proteins are converted into urocanic acid (UCA) and pyrrolidine carboxylic acid (PCA) by proteases. Zn facilitates FLG production by increasing the activity of Prss8. Alternatively, Zn can also suppress FLG metabolism by decreasing PAD activity. Moreover, Zn is required for histidine conversion to UCA.
3. Zinc and Langerhans Cells

LCs are antigen-presenting cells that occupy approximately 3% of the epidermis [70]. They were previously considered a subtype of dendritic cells (DCs) because of their ability to capture antigens (Ag), migrate to the draining lymph nodes (dLNs), and then present the Ag to T cells and initiate the immune response. However, recent evidence revealed that LCs originate from macrophage lineage of fetal liver progenitors and not from DC lineage [71–74]. Therefore, LCs are currently considered tissue-resident macrophages with the ability to migrate to the dLNs.

3.1. LCs and Tight Junctions. LCs express claudin-1 and form TJs with adjacent SG KCs [75]. As such, LCs are an important component of TJs to create an effective epidermal barrier. In steady state, LC dendrites lie beneath the TJs. Upon activation, LCs elongate their dendrites to penetrate the KC TJs by forming tricellulin-dependent TJs between KC-LC-KC. The elongated dendrites are able to reach beneath the SC and take up Ag from the extra-SC environment without destroying barrier integrity (Figure 1) [7]. These LCs then induce Th2-type, but not Th1-type, humoral immune responses [8]. Furthermore, LCs take up KC-derived auto-Ag and present it on their MHC class II. These LCs expand polyclonal Ag-specific regulatory T cells and keep peripheral tolerance against auto-Ag [76]. Therefore, LCs assume the dual role of T component and surveillance agent of foreign and auto-Ag.

3.2. LCs and Zinc Deficiency. The association between Zn and LCs was revealed by analysis of skin specimen from patients with acrodermatitis enteropathica (AE; OMIM 201100) [77]. AE is an autosomal recessive disease caused by mutations in the SLC39A4 gene that encodes ZRT/IRT-like protein 4 (ZIP4) [78]. ZIP4 is abundantly expressed in the apical side of the intestinal epithelium, thereby working as the primary gate absorbing Zn into the enterocytes. This absorbed Zn is subsequently transported to the bloodstream by Zn transporter 1 (ZnT1) [79, 80]. Therefore, ZIP4 dysfunction due to mutations results in decreased serum Zn levels. Interestingly, epidermal LCs were absent in AE skin lesions [77] (Figure 3). However, LCs were restored in the epidermis after patients were given Zn supplementation. These phenomena suggest that LCs disappeared from the epidermis when patients are deficient in Zn, but the effect can be easily reversed.

The association between LC loss and the development of characteristic AE skin lesion was investigated using dietary Zn-deficient (ZD) and Zn-adequate (ZA) mice [77]. Consistent with observations in patients with AE, epidermal LCs also disappeared in ZD mice (Figure 4).

“Allergic” contact dermatitis was significantly impaired in ZD mice compared with ZA mice. On the other hand, “irritant” contact dermatitis (ICD) was significantly enhanced in ZD mice compared to ZA mice. ICD is mediated by adenosine triphosphate (ATP), which is secreted by KCs in response to environmental irritants through lytic and nonlytic mechanisms [81–84]. As expected, ATP production from croton oil-applied skin was significantly increased in ZD mice compared with ZA mice. Additionally, ATP production from Pam212 KCs (murine immortalized KCs) was significantly increased by incubation with TPEN. These data suggest that KCs in ZD mice produce more ATP than KCs in ZA mice [77].

ATP is a potent inflammation inducer. Therefore, the epidermis is equipped with a mechanism to prevent ATP-mediated inflammation in the steady state. CD39 (ectonucleoside triphosphate diphosphohydrolase-1; ENTPD-1) plays a central role in ATP hydrolysis [83]. In the epidermis, CD39 is predominantly expressed in LCs, but not in KCs [83, 85, 86]. Thus, ATP is not hydrolyzed in the epidermis.
of patients with AE and ZD mice, thereby eliciting ATP-mediated ICD in skin. In line with this concept, characteristic AE skin lesions develop in anogenital and periorificial areas and distal portions of the extremities, where frequent contact with external irritants including feces, urine, saliva, food, shoes, or excessive sweating is expected.

Taken together, the nature of AE skin lesions is ICD mediated by (1) increased ATP production by ZD KCs and (2) disabling of ATP hydrolysis due to the loss of CD39-expressing LCs.

As described, CD39 (ENTPD-1) potently hydrolyzes ATP. However, there are three groups of molecules that hydrolyze ATP, including ENTPDs, ectonucleotide pyrophosphatase/phosphodiesterases (ENPPs), and alkaline phosphatase (ALP) [87, 88]. The latter two molecules are Zn-dependent molecules. Among ENTPDs and ENPPs, ENTPD-1 (CD39), -2, -3, and -8 and ENPP-1, -2, and -3 aid in ATP hydrolysis. The expression of these molecules in LCs and KCs was not previously understood, with the exception of CD39. Thus, we determined that LCs strongly

\[ \text{Zn deficiency + skin irritants} \rightarrow \begin{align*} &\text{(1) Loss of epidermal LCs (CD39+)} \\
&\text{(2) Impaired ENPP activity} \\
&\text{Increased ATP release by KCs} \end{align*} \]

\[ \text{Impaired ATP hydrolysis} \]

\[ \text{(i) ATP ↑ in skin} \]

\[ \text{Skin inflammation (neutrophils, etc.)} \rightarrow \text{Aberrant chemokine production by KCs (CXCL1, CXCL2, etc.)} \]

\[ \text{(ii) Severe and prolonged ICD} \]

\[ \text{Acrodermatitis enteropathica} \]

\[ \text{(Histologically)} \]

**Figure 3:** Loss of epidermal LCs in patients with AE. Immunohistochemical staining for langerin (red) in normal skin (a) and the erythematous lesions in atopic dermatitis (b), psoriasis vulgaris (c), or three AE (d–f) patients. Original magnification: 200x.

**Figure 4:** Model for the etiology of AE. (i) Skin irritants increase ATP release in the skin in Zn deficiency, due to increased ATP release by KCs, impaired ATP hydrolysis by LCs, and impaired ENPP activity. (ii) Increased ATP release induces severe and prolonged ICD via aberrant chemokine production by KCs and neutrophil-mediated skin inflammation. Histologically, cutaneous lesions in AE and ICD lesions in ZD mice demonstrate common histological features, such as subcorneal vacuolization and epidermal pallor.
expressed CD39 and weakly expressed ENTPD-2 and ENPP-1, -2, and -3. Normal human epidermal KCs weakly expressed CD39, ENTPD-2 and -3, and ENPP-1 and -2. Neither LCs nor KCs expressed ENTPD-8 or ALP. Therefore, although LCs strongly express CD39, other ATP-hydrolyzing molecules are weakly expressed in both LCs and KCs. KCs occupy approximately 97% of the epidermis, whereas LCs occupy approximately 3%. Therefore, we determined the degree of contribution of LCs and KCs to ATP hydrolysis. ATP hydrolysis was impaired by approximately 80% in LC-depleted epidermal suspension compared with sham-sorted epidermal suspension. This suggests that LCs assume approximately 80% of epidermal ATP hydrolysis, whereas KCs assume the remaining 20% [89]. Recently, it was demonstrated that Zn deficiency impairs the activity of ENPP-1 and ENPP-3, as well as ALP and CD73 [90]. As described, KCs weakly express ENPP-1 and ENPP-2. This explains one underlying mechanism by which ZD KCs increase ATP production.

3.3. LCs and Zinc Finger Proteins. In the steady state, LCs form the firm connections with adjacent KCs by claudin-1 and E-cadherin. Additionally, EpCAM (epithelial cell adhesion molecule) in mice and its human homolog, tumor-associated calcium signal transducer 2, also participate in this cell adhesion [91]. Claudin-1 and EpCAM colocalize in LCs [92]. Upon LC activation, LCs downregulate these adhesion molecules and then migrate to the dLNs. LCs upregulate ZEB-1 and ZEB-2 (see Section 2.2) during maturation, subsequently downregulating E-cadherin expression in LCs [93, 94]. Additionally, downregulation of EpCAM in LCs impairs claudin-1 expression in LCs [92]. This suggests that Zn deficiency leads to LC retention in the epidermis. Nevertheless, the presence of LCs is reduced with Zn deficiency [77]. For LC development and survival, LC-derived, but not KC-derived, autocrine transforming growth factor beta (TGF-β) is crucial [95–98]. This latent secreted protein is processed and activated by αvβ6 and αvβ8 integrins on KCs and then is recognized by LCs [99–101]. In patients with AE and ZD mice, the epidermal expression of TGF-β is strongly impaired, and thus, LCs are reduced [77]. However, the association between Zn deficiency and impaired epidermal TGF-β expression is not fully understood. Collectively, Zn deficiency in the epidermis results in (1) disruption of epidermal barrier function (see Section 2), (2) LC disappearance due to impaired epidermal TGF-β expression, (3) impaired ATP hydrolysis due to the reduced number of CD39-expressing LCs and impaired ENPP activity, and (4) elicitation of ATP-mediated skin inflammation. Because of its importance, we are currently investigating the impact of LC loss on TJ development and function using AE skin specimens and ZD mice.

4. Zinc and Zinc Transporters in the Epidermis

The subject of Zn transporters in the epidermis has been widely reviewed by us and others [18–20]. Thus, here, we briefly summarized what is known and added recent findings. Zn is distributed to a higher degree in the epidermis than in the dermis. Within the epidermis, Zn is distributed primarily in the stratum spinosum [102]. LCs are present in the stratum spinosum, where Zn is found, and research suggests that LCs definitively require Zn for their survival and function [77]. The fundamental functions of Zn in the KCs are proliferation and anti-inflammation. KCs treated with nontoxic concentration of Zn increased their proliferation and survival [103]. Conversely, the chelation of intracellular Zn by TPEN facilitates KC apoptosis by activating caspase-3 and DNA fragmentation [104]. Zn has been reported to suppress the production of tumor necrosis factor-α, inducible nitric oxide synthase, and subsequent nitric oxide in KCs [65, 105, 106]. Zn also suppresses the expression of toll-like receptor 2 on KCs [107].

The rigorous Zn^2+ regulation is conducted by Zn transporters (ZnTs and ZIPS) and metallothioneins (MTs) [108, 109]. So far, 10 ZnTs, 14 ZIPS, and 4 MTs were identified in humans [110, 111]. ZnTs and ZIPS mediate Zn efflux and uptake, respectively. MTs are ubiquitously expressed throughout various types of cells and are predominantly distributed in the cytoplasm and to a lesser extent in the nuclei and lysosomes. MTs can bind to metal ion including Zn via a unique cysteine-rich amino acid sequence and essentially control the availability of Zn [112]. Among 24 Zn transporters (10 ZnTs and 14 ZIPS), the function of only four Zn transporters (ZnT1, ZIP2, ZIP4, and ZIP10) in the epidermis or KCs has been elucidated so far. ZnT1 in KCs is involved in the development of epidermodyplasia verruciformis (EV; OMIM 226400), which is a rare autosomal-recessive skin disease that can lead to nonmelanoma skin cancers resulting from selective susceptibility to oncogenic human papillomaviruses (HPVs) [113]. In KCs, EVER1 and EVER2 proteins form a complex with ZnT1 primarily in the ER and to a lesser extent in the nuclear membrane and Golgi apparatus. This complex maintains Zn homeostasis, which inhibits activator protein-1 (AP-1) activation that promotes HPV replication. Patients with EV have mutations in either the EVER1 or EVER2 genes [114, 115]. The complex of ZnT1 and mutated EVERs increases free Zn transport in the nucleus and subsequently enhances AP-1 activity, leading to aberrant replication of EV-related oncogenic HPVs and thereby developing skin cancers.

ZIP2 is expressed on the differentiating KCs of humans and mice. Since ZIP2 knockdown in KCs decreases intracellular Zn, suppresses KC differentiation, and downregulates involucrin expression, Zn taken up by KC ZIP2 is required for proper KC differentiation and CE formation (also see Section 2.1.2) [102]. ZIP4 is expressed on the undifferentiating KCs of humans. Since ZIP4 knockdown in KCs decreases intracellular Zn, suppresses KC differentiation, downregulates the expression of FLG and involucrin, and impairs the activity of p63 that is a critical regulator of epidermal formation, Zn taken up by KC ZIP4 is required for proper KC differentiation and SC formation (also see Section 2.1) [116]. Murine ZIP10 is expressed on the epidermal progenitor cells of the outer root sheath of hair follicles. Therefore, ZIP10 depletion in keratin 14-expressing cells leads to a thin epidermis and a hypoplasia of hair follicles via downregulation of
The epidermal regulator p63 also controls the activity of ZNF750, a C2H2-type Zn finger protein. ZNF750 strongly regulates terminal epidermal differentiation. Thus, ZNF750 knockdown downregulates the expression of epidermal barrier-related proteins including FLG, loricrin, and SPINK5 [118]. In human skin, MT-1 and MT-2 are expressed in the actively proliferating KCs, such as the hair matrix, outer hair roots, and stratum basale [119]. Knockdown of both MT-1 and MT-2 in mice impairs KC proliferation [120]. MT expressions are upregulated in the hyperplastic KCs of inflamed skin lesions and skin cancers [121]. Collectively, Zn supports KC proliferation and survival. ZIP2 and ZIP4 are required for proper KC differentiation and subsequent epidermal barrier formation. ZIP10 is required for the successful epidermal formation. MTs are involved in KC proliferation.

5. Conclusions and Perspectives

Epidermal barrier homeostasis is the first line of defense for preventing the initiation of atopic march. SC and TJs are responsible for epidermal barrier function. Zn and Zn finger proteins regulate SC formation as well as its metabolism, while the contribution of Zn to TJ function is less understood. However, when Zn is deficient, LCs are not present, and this important component of TJs is lacking. Therefore, Zn deficiency might lead to epidermal barrier dysfunction. Disruption of the epidermal barrier induces a Th2-type immune response by producing KC-derived Th2-promoting cytokines and T cell-derived Th2 cytokines. Meanwhile, these Th2-related cytokines impair the structure and function of the SC and TJs. In this way, epidermal barrier disruptions trigger a negative spiral of inflammation. Zn homeostasis in cells, including KCs, is maintained by ZnTs, ZIPS, and MTs. MT-mediated Zn in KCs facilitates its proliferation. On the other hand, the function of ZnTs and ZIPS in KC biology is less understood, because only the function of ZnT1, ZIP2, ZIP4, and ZIP10 has been elucidated from among 10 ZnTs and 14 ZIPS. The epidermis is composed of KCs, LCs, melanocytes, epidermal-resident memory T cells (TRMs), and others. The murine epidermis, but not the human epidermis, also contains dendritic epidermal T cells (DETCs), a type of γδ T cells. The role of Zn and Zn transporters in LCs, melanocytes, TRMs, and DETCs has not been analyzed to date. Additional research must be conducted to thoroughly understand the role of Zn and Zn transporters in maintaining healthy skin.

Conflicts of Interest

The authors declare no competing financial interest.

References

[1] K. Horimukai, K. Morita, M. Narita et al., “Application of moisturizer to neonates prevents development of atopic dermatitis,” The Journal of Allergy and Clinical Immunology, vol. 134, no. 4, pp. 824–830.e6, 2014.

[2] E. L. Simpson, J. R. Chalmers, J. M. Hanifin et al., “Emollient enhancement of the skin barrier from birth offers effective atopic dermatitis prevention,” The Journal of Allergy and Clinical Immunology, vol. 134, no. 4, pp. 818–823, 2014.

[3] J. M. Spergel and A. S. Paller, “Atopic dermatitis and the atopic march,” The Journal of Allergy and Clinical Immunology, vol. 112, no. 6, pp. S118–S127, 2003.

[4] G. Egawa and K. Kabashima, “Multifactorial skin barrier deficiency and atopic dermatitis: essential topics to prevent the atopic march,” The Journal of Allergy and Clinical Immunology, vol. 138, no. 2, pp. 350–358.e1, 2016, e351.

[5] M. J. Cork, S. G. Danby, Y. Vasilopoulos et al., “Epidermal barrier dysfunction in atopic dermatitis,” The Journal of Investigative Dermatology, vol. 129, no. 8, pp. 1892–1908, 2009.

[6] A. De Benedetto, A. Kubo, and L. A. Beck, “Skin barrier disruption: a requirement for allergen sensitization?” The Journal of Investigative Dermatology, vol. 132, no. 3, pp. 949–963, 2012.

[7] A. Kubo, K. Nagao, M. Yokouchi et al., “External antigen uptake by Langerhans cells with reorganization of epidermal tight junction barriers,” The Journal of Experimental Medicine, vol. 206, no. 13, pp. 2937–2946, 2009.

[8] T. Ouchi, A. Kubo, M. Yokouchi et al., “Langerhans cell antigen capture through tight junctions confers preemptive immunity in experimental staphylococcal scalded skin syndrome,” The Journal of Experimental Medicine, vol. 208, no. 13, pp. 2607–2613, 2011.

[9] M. J. Jackson, “Physiology of zinc: general aspects,” in Zinc in Human Biology, C. F. Mills, Ed., pp. 1–14, Springer, London, 1989.

[10] G. Michaelsson, K. Ljunghall, and B. G. Danielson, “Zinc in epidermis and dermis in healthy subjects,” Acta Dermato-Venerologica, vol. 60, no. 4, pp. 295–299, 1980.

[11] C. Andreini, I. Bertini, and G. Cavallaro, “Minimal functional sites allow a classification of zinc sites in proteins,” PLoS One, vol. 6, no. 10, article e26325, 2011.

[12] C. Andreini and I. Bertini, “A bioinformatics view of zinc enzymes,” Journal of Inorganic Biochemistry, vol. 111, pp. 150–156, 2012.

[13] J. M. Berg and Y. Shi, “The galvanization of biology: a growing appreciation for the roles of zinc,” Science, vol. 271, no. 5252, pp. 1081–1085, 1996.

[14] T. Fukada, S. Yamasaki, K. Nishida, M. Murakami, and T. Hirano, “Zinc homeostasis and signaling in health and diseases: zinc signaling,” Journal of Biological Inorganic Chemistry, vol. 16, no. 7, pp. 1123–1134, 2011.

[15] A. Klug, “The discovery of zinc fingers and their development for practical applications in gene regulation and genome manipulation,” Quarterly Reviews of Biophysics, vol. 43, no. 1, pp. 1–21, 2010.

[16] J. H. Laitly, B. M. Lee, and P. E. Wright, “Zinc finger proteins: new insights into structural and functional diversity,” Current Opinion in Structural Biology, vol. 11, no. 1, pp. 39–46, 2001.

[17] C. Andreini, L. Banci, I. Bertini, and A. Rosato, “Counting the zinc-proteins encoded in the human genome,” Journal of Proteome Research, vol. 5, no. 1, pp. 196–201, 2006.

[18] Y. Ogawa, T. Kavamura, and S. Shimada, “Zinc and skin biology,” Archives of Biochemistry and Biophysics, vol. 611, pp. 113–119, 2016.
...
Molecular Biology of the Cell, vol. 18, no. 9, pp. 3607–3619, 2007.

M. Debela, P. Goettig, V. Magdolen, R. Huber, N. M. Schechter, and W. Bode, “Structural basis of the zinc inhibition of human tissue kallikrein 5,” Journal of Molecular Biology, vol. 373, no. 4, pp. 1017–1031, 2007.

M. Debela, P. Hess, V. Magdolen et al., “Chymotryptic specificity determinants in the 1.0 A structure of the zinc-inhibited human tissue kallikrein 7,” Proceedings of the National Academy of Sciences of the United States of America, vol. 104, no. 41, pp. 16086–16091, 2007.

C. A. Borgoño, I. P. Michael, J. L. V. Shaw et al., “Expression and functional characterization of the cancer-related serine protease, human tissue kallikrein 14,” Journal of Biological Chemistry, vol. 282, no. 4, pp. 2405–2422, 2007.

L. Bin, B. E. Kim, C. F. Hall, S. M. Leach, and D. Y. M. Leung, “Inhibition of transcription factor specificity protein 1 alters the gene expression profile of keratinocytes leading to upregulation of kallikrein-related peptidases and thymic stromal lymphopoietin,” The Journal of Investigative Dermatology, vol. 131, no. 11, pp. 2213–2222, 2011.

H. Takahashi, M. Nakazawa, K. Takahashi et al., “Effects of zinc deficient diet on development of atopic dermatitis-like eruptions in DS-Nh mice,” Journal of Dermatological Science, vol. 50, no. 1, pp. 31–39, 2008.

T. J. David, F. E. Wells, T. C. Sharpe, and A. C. C. Gibbs, “Low serum zinc in children with atopic eczema,” The British Journal of Dermatology, vol. 111, no. 5, pp. 597–601, 1984.

J. E. Kim, S. R. Yoo, M. G. Jeong, J. Y. Ko, and Y. S. Ro, “Hair zinc levels and the efficacy of oral zinc supplementation in patients with atopic dermatitis,” Acta Dermato-Venereologica, vol. 94, no. 5, pp. 558–562, 2014.

F. Rippke, V. Schreiner, T. Doering, and H. I. Maibach, “Stratum corneum pH in atopic dermatitis: impact on skin barrier function and colonization with Staphylococcus aureus,” American Journal of Clinical Dermatology, vol. 5, no. 4, pp. 217–223, 2004.

S. Kezic, G. M. O’Regan, R. Lutter et al., “Filaggrin loss-of-function mutations are associated with enhanced expression of IL-1 cytokines in the stratum corneum of patients with atopic dermatitis and in a murine model of filaggrin deficiency,” Journal of Allergy and Clinical Immunology, vol. 129, no. 4, pp. 1031–1039.e1, 2012.

K. Stefansson, M. Brattsand, D. Roosterman et al., “Activation of proteinase-activated receptor-2 by human kallikrein-related peptidases,” The Journal of Investigative Dermatology, vol. 128, no. 1, pp. 18–25, 2008.

N. Kirschner, P. Houdek, M. Fromm, I. Moll, and J. M. Brandner, “Tight junctions form a barrier in human epidermis,” European Journal of Cell Biology, vol. 89, no. 11, pp. 839–842, 2010.

M. Furuse, M. Hata, K. Furuse et al., “Claudin-based tight junctions are crucial for the mammalian epidermal barrier: a lesson from claudin-1-deficient mice,” The Journal of Cell Biology, vol. 156, no. 6, pp. 1099–1111, 2002.

A. De Benedetto, N. M. Raefas, L. Y. McGirt et al., “Tight junction defects in patients with atopic dermatitis,” Journal of Allergy and Clinical Immunology, vol. 127, no. 3, pp. 773–786.e7, 2011.

Y. Miyoshi, S. Tanabe, and T. Suzuki, “Cellular zinc is required for intestinal epithelial barrier maintenance via the regulation of claudin-3 and occludin expression,” American Journal of Physiology. Gastrointestinal and Liver Physiology, vol. 311, no. 1, pp. G105–G116, 2016.

M. N. Tatari, B. De Craene, B. Soen et al., “ZEB2-transgene expression in the epidermis compromises the integrity of the epidermal barrier through the repression of different tight junction proteins,” Cellular and Molecular Life Sciences, vol. 71, no. 18, pp. 3599–3609, 2014.

C. Becker-Pauly, M. Höwel, T. Walker et al., “The alpha and beta subunits of the metalloprotease meprin are expressed in separate layers of human epidermis, revealing different functions in keratinocyte proliferation and differentiation,” The Journal of Investigative Dermatology, vol. 127, no. 5, pp. 1115–1125, 2007.

I. Wessels, M. Maywald, and L. Rink, “Zinc as a gatekeeper of immune function,” Nutrients, vol. 9, no. 12, 2017.

W. Amano, S. Nakajima, H. Kunugi et al., “The Janus kinase inhibitor JTE-052 improves skin barrier function through suppressing signal transducer and activator of transcription 3 signaling,” The Journal of Allergy and Clinical Immunology, vol. 136, no. 3, pp. 667–677.e7, 2015.

J. Seltmann, L. M. Roesner, F. W. von Hesler, M. Wittmann, and T. Werfel, “IL-33 impacts on the skin barrier by downregulating the expression of filaggrin,” Journal of Allergy and Clinical Immunology, vol. 135, no. 6, pp. 1659–1661.e4, 2015.

W. I. Ryu, H. Lee, H. C. Bae et al., “IL-33 down-regulates CLDN1 expression through the ERK/STAT3 pathway in keratinocytes,” Journal of Dermatological Science, vol. 90, no. 3, pp. 313–322, 2018.

C. Cornelissen, Y. Marquardt, K. Czaja et al., “IL-31 regulates differentiation and filaggrin expression in human organotypic skin models,” Journal of Allergy and Clinical Immunology, vol. 129, no. 2, pp. 426–433.e8, 2012.

M. Merad, F. Ginhoux, and M. Collins, “Origin, homeostasis and function of Langerhans cells and other langerin-expressing dendritic cells,” Nature Reviews Immunology, vol. 8, no. 12, pp. 935–947, 2008.

L. Chorro, A. Sarde, M. Li et al., “Langerhans cell (LC) proliferation mediates neonatal development, homeostasis, and inflammation-associated expansion of the epidermal LC network,” The Journal of Experimental Medicine, vol. 206, no. 13, pp. 3089–3100, 2009.

G. Hoeffel, Y. Wang, M. Greter et al., “Adult Langerhans cells derive predominantly from embryonic fetal liver monocytes with a minor contribution of yolk sac-derived macrophages,” The Journal of Experimental Medicine, vol. 209, no. 6, pp. 1167–1181, 2012.

C. Schulz, E. G. Perdiguerro, L. Chorro et al., “A lineage of myeloid cells independent of Myb and hematopoietic stem cells,” Science, vol. 336, no. 6077, pp. 86–90, 2012.

L. C. Davies, S. J. Jenkins, J. E. Allen, and P. R. Taylor, “Tissue-resident macrophages,” Nature Immunology, vol. 14, no. 10, pp. 986–995, 2013.

S. C. Zimmerli and C. Hauser, “Langerhans cells and lymph node dendritic cells express the tight junction component claudin-1,” The Journal of Investigative Dermatology, vol. 127, no. 10, pp. 2381–2390, 2007.

D. Y. Kitashima, T. Kobayashi, T. Woodring et al., “Langerhans cells prevent autoimmunity via expansion of keratinocyte antigen-specific regulatory T cells,” eBioMedicine, vol. 27, pp. 293–303, 2018.
to an epidermally applied protein antigen,” The Journal of Investigative Dermatology, vol. 136, no. 8, pp. 1627–1635, 2016.

[93] S. Konrad, N. Yasin, D. Hashwanter et al., “Langerhans cell maturation is accompanied by induction of N-cadherin and the transcriptional regulators of epithelial-mesenchymal transition ZEB1/2,” European Journal of Immunology, vol. 44, no. 2, pp. 553–560, 2014.

[94] J. Comijn, G. Berx, P. Varmassen et al., “The two-handed E box binding zinc finger protein SIP1 downregulates E-cadherin and induces invasion,” Molecular Cell, vol. 7, no. 6, pp. 1267–1278, 2001.

[95] T. A. Borkowski, J. J. Letterio, A. G. Farr, and M. C. Udey, “A role for endogenous transforming growth factor beta 1 in Langerhans cell biology: the skin of transforming growth factor beta 1 null mice is devoid of epidermal Langerhans cells,” The Journal of Experimental Medicine, vol. 184, no. 6, pp. 2417–2422, 1996.

[96] D. H. Kaplan, M. O. Li, M. C. Jenison, W. D. Shlomchik, R. A. Flavell, and M. J. Shlomchik, “Autocrine/paracrine TGFβ1 is required for the development of epidermal Langerhans cells,” The Journal of Experimental Medicine, vol. 204, no. 11, pp. 2545–2552, 2007.

[97] J. M. Kel, M. J. H. Girard-Madoux, B. Reizis, and B. E. Clausen, “TGF-beta is required to maintain the pool of immature Langerhans cells in the epidermis,” Journal of Immunology, vol. 185, no. 6, pp. 3248–3255, 2010.

[98] A. Bobr, B. Z. Igyarto, K. M. Haley, M. O. Li, R. A. Flavell, and D. H. Kaplan, “Autocrine/paracrine TGF-β1 inhibits Langerhans cell migration,” Proceedings of the National Academy of Sciences of the United States of America, vol. 109, no. 26, pp. 10492–10497, 2012.

[99] Z. Yang, Z. Mu, B. Dabovic et al., “Absence of integrin-mediated TGFβ1 activation in vivo recapitulates the phenotype of TGFβ1-null mice,” The Journal of Cell Biology, vol. 176, no. 6, pp. 787–793, 2007.

[100] P. Aluwihare, Z. Mu, Z. Zhao et al., “Mice that lack activity of avy6- and avy8-integrins reproduce the abnormalities of Tgfb1- and Tgfb3-null mice,” Journal of Cell Science, vol. 122, no. 2, pp. 227–232, 2008.

[101] J. Mohammed, L. K. Beura, A. Bobr et al., “Stromal cells control the epithelial residence of DCs and memory T cells by regulated activation of TGF-β,” Nature Immunology, vol. 17, no. 4, pp. 414–421, 2016.

[102] Y. Inoue, S. Hasegawa, S. Ban et al., “ZIP2 protein, a zinc transporter, is associated with keratinocyte differentiation,” The Journal of Biological Chemistry, vol. 289, no. 31, pp. 21451–21462, 2014.

[103] E. Emri, E. Miko, P. Bai et al., “Effects of non-toxic zinc exposure on human epidermal keratinocytes,” Metallomics, vol. 7, no. 3, pp. 499–507, 2015.

[104] D. Wilson, G. Varigos, and M. L. Ackland, “Apoptosis may underlie the pathology of zinc-deficient skin,” Immunology and Cell Biology, vol. 84, no. 1, pp. 28–37, 2006.

[105] A. Gueniche, J. Via, G. Lizard, M. Charveron, and D. Schmitt, “Protective effect of zinc on keratinocyte activation markers induced by interferon or nickel,” Acta Dermato-Venereologica, vol. 75, no. 1, pp. 19–23, 1995.

[106] J. Yamaoka, T. Kume, A. Akaike, and Y. Miyachi, “Suppressive effect of zinc ion on iNOS expression induced by interferon-gamma or tumor necrosis factor-alpha in murine
keratinocytes,” *Journal of Dermatological Science*, vol. 23, no. 1, pp. 27–35, 2000.

[107] V. Jarrousse, N. Castex-Rizzi, A. Khammari, M. Charveron, and B. Dréno, “Zinc salts inhibit in vitro Toll-like receptor 2 surface expression by keratinocytes,” *European Journal of Dermatology*, vol. 17, no. 6, pp. 492–496, 2007.

[108] T. Kambe, T. Tsuji, A. Hashimoto, and N. Itsumura, “The physiological, biochemical, and molecular roles of zinc transporters in zinc homeostasis and metabolism,” *Physiological Reviews*, vol. 95, no. 3, pp. 749–784, 2015.

[109] M. A. Hediger, M. F. Romero, J. B. Peng, A. Rolfs, H. Takanaga, and E. A. Bruford, “The ABCs of solute carriers: physiological, pathological and therapeutic implications of human membrane transport proteins,” *Pflügers Archiv European Journal of Physiology*, vol. 447, no. 5, pp. 465–468, 2004.

[110] R. D. Palmiter and L. Huang, “Efflux and compartmentalization of zinc by members of the SLC30 family of solute carriers,” *Pflügers Archiv European Journal of Physiology*, vol. 447, no. 5, pp. 744–751, 2004.

[111] D. J. Eide, “The SLC39 family of metal ion transporters,” *Pflügers Archiv European Journal of Physiology*, vol. 447, no. 5, pp. 796–800, 2004.

[112] H. Tapiero and K. D. Tew, “Trace elements in human physiology and pathology: zinc and metallothioneins,” *Biomedicine & Pharmacotherapy*, vol. 57, no. 9, pp. 399–411, 2003.

[113] M. Lazarczyk, C. Pons, J. A. Mendoza, P. Cassonnet, Y. Jacob, and M. Favre, “Regulation of cellular zinc balance as a potential mechanism of EVER-mediated protection against pathogenesis by cutaneous oncogenic human papillomaviruses,” *The Journal of Experimental Medicine*, vol. 205, no. 1, pp. 35–42, 2008.

[114] N. Ramoz, M. Favre, G. Orth et al., “Evidence for a nonallelic heterogeneity of epidermodysplasia verruciformis with two susceptibility loci mapped to chromosome regions 2p21-p24 and 17q25,” *The Journal of Investigative Dermatology*, vol. 114, no. 6, pp. 1148–1153, 2000.

[115] N. Ramoz, L. A. Rueda, B. Bouadjar, L. S. Montoya, G. Orth, and M. Favre, “Mutations in two adjacent novel genes are associated with epidermodysplasia verruciformis,” *Nature Genetics*, vol. 32, no. 4, pp. 579–581, 2002.

[116] B. H. Bin, J. Bhin, N. H. Kim et al., “An acrodermatitis enteropathica-associated Zn transporter, ZIP4, regulates human epidermal homeostasis,” *The Journal of Investigative Dermatology*, vol. 137, no. 4, pp. 874–883, 2017.

[117] B. H. Bin, J. Bhin, M. Takaishi et al., “Requirement of zinc transporter ZIP10 for epidermal development: implication of the ZIP10-p63 axis in epithelial homeostasis,” *Proceedings of the National Academy of Sciences of the United States of America*, vol. 114, no. 46, pp. 12243–12248, 2017.

[118] G. L. Sen, L. D. Boxer, D. E. Webster et al., “ZNF750 is a p63 target gene that induces KLF4 to drive terminal epidermal differentiation,” *Developmental Cell*, vol. 22, no. 3, pp. 669–677, 2012.

[119] J. J. van den Oord and M. De Ley, “Distribution of metallothionein in normal and pathological human skin,” *Archives of Dermatological Research*, vol. 286, no. 1, pp. 62–68, 1994.