Dermatitis due to epiregulin deficiency and a critical role of epiregulin in immune-related responses of keratinocyte and macrophage

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Epidermal growth factor (EGF) family members, including epiregulin (EP), play a fundamental role in epithelial tissues; however, their roles in immune responses and the physiological role of EP remain to be elucidated. The skin has a versatile system of immune surveillance. Biologically active IL-1 α is released to extracellular **space upon damage from keratinocytes and is a major player in skin inflammation. Here, we show that** *EP* **is expressed not only in keratinocytes but also in tissue-resident macrophages, and that** *EP***-deficient (***EP*-**/**-**) mice develop chronic dermatitis. Wound healing in the skin in** *EP*-**/**- **mice was not impaired** *in vivo***, nor was the growth rate of keratinocytes from** *EP*-**/**- **mice different from that of WT mice** *in vitro***. Of interest is that in WT keratinocytes, both IL-1 and the secreted form of EP induced down-regulation of** *IL-18* **mRNA expression, which overexpression in the epidermis was reported to induce skin inflammation in mice, whereas the down**regulation of *IL-18* induced by IL-1 α was impaired in $EP^{-/-}$ kera**tinocytes. Although bone marrow transfer experiments indicated that EP deficiency in non-bone-marrow-derived cells is essential for the development of dermatitis, production of proinflammatory cytokines by** *EP*-**/**- **macrophages in response to Toll-like receptor agonists was much lower, compared with WT macrophages, whose dysfunction in** *EP*-**/**- **macrophages was not compensated by the addition of the secreted form of EP. These findings, taken together, suggested that EP plays a critical role in immuneinflammatoryrelated responses of keratinocytes and macrophages at the barrier from the outside milieu and that the secreted and membranebound forms of EP have distinct functions.**

The system of epidermal growth factor (EGF) superfamily (1)
and EGF receptor (EGFR) family, including EGFR (ErbB1), ErbB2 (HER2), ErbB3 (HER3), and ErbB4 (HER4) (2), play a fundamental role in epithelial tissues. EGF family members, including EGF, transforming growth factor- α , amphiregulin, heparinbinding EGF (HB-EGF), epiregulin (EP), and other members, regulate these receptors by inducing their homo- and/or heterooligomerization (2, 3). EGF family members vary in their ability to activate distinct ErbB heterodimers, and this mechanism may, in part, account for the differences in their bioactivities (4–7).

The membrane-anchored precursor of the EGF family is enzymatically processed externally to release a mature soluble form that acts as autocrine and/or paracrine growth factor $(8, 9)$, whereas some members of the EGF family act in the membrane-anchored form (8, 10). A bioactive transmembrane precursor, pro-HB-EGF, was suggested to induce growth inhibition or apoptosis rather than the proliferative response induced by soluble HB-EGF (10). EP (11–17) acts as an autocrine growth factor in normal human keratinocytes*in vitro* (15); however, its physiological role *in vivo* still remains to be elucidated.

Keratinocytes in the epidermis play critical roles in the cutaneous immune-related responses (18) and contain biologically active IL-1 α (19), which is released to extracellular space upon damage and is a major player in skin inflammation (20). Overexpression of IL-1 α in murine epidermis produces inflammatory skin lesions,

indicating the critical role of IL-1 α in skin inflammation (21). On the other hand, IL-18 is stored as a biologically inactive precursor form in keratinocytes, and overexpression of IL-18 in the mouse skin was reported to induce inflammatory skin lesions without inducing allergen-specific IgE production (22).

Macrophages normally reside in tissues and beneath mucosal surfaces and function at the front line of immune defense against incoming pathogens through ingestion of bacteria by phagocytosis, destruction of bacteria, and recruitment of inflammatory cells to the site of infection by using soluble mediators (23). Toll-like receptors (TLRs) (24, 25), which recognize molecular patterns that are common and shared by many microbial pathogens, are expressed on antigen-presenting cells (APCs), including macrophages and dendritic cells: TLR4 to lipopolysaccharide (LPS), TLR2 to peptidoglycan (PGN), and TLR9 to CpG DNA.

Here, we show the critical role of EP in immune/inflammatoryrelated responses of keratinocytes and macrophages through the establishment and analysis of $EP^{-/-}$ mice.

Methods

Generation of EP-Deficient Mice. We isolated the genomic DNA of the *EP* gene from a 129/Sv mouse genomic library (Stratagene) and constructed the targeting vector by replacing a 6.5-kb *Sal*I-*Xho*I fragment containing exons 2–5 (90% of the coding region) of the *EP* gene, with a pGKneo cassette with opposite transcriptional orientation. The 5' and 3' arm of the targeting construct was composed of 2.0-kb and 11.0-kb genomic DNA, respectively. Diphtheria-toxin A fragment cassette (DT-A) flanked the 3' genomic arm. The targeting vector was linearized with *Sal*I. The mutant embryonic stem cells were microinjected into C57BL/6 blastocysts as described (26), and the resulting male chimeras were mated with C57BL/6 mice. Heterozygous offspring were intercrossed to obtain *EP*-deficient mice.

Histopathology. Tissues were fixed in 4% paraformaldehyde, embedded on paraffin, and stained with hematoxylin/eosin. Mast cells and eosinophils were detected by toluidine blue and Luna staining, respectively.

Quantification of Serum Ig Isotype Concentration. Blood samples were obtained from mouse eye vein. Serum Ig isotype concentration was determined by using an ELISA kit (Bethyl Laboratories, Montgomery, TX).

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Abbreviations: EGF, epidermal growth factor; EGFR, EGF receptor; HB-EGF, heparinbinding EGF; EP, epiregulin; rmEP, recombinant mouse EP; sEP, secreted form of EP; TLR, Toll-like receptor; BMDM, bone marrow-derived macrophage; LPS, lipopolysaccharide; PGN, peptidoglycan; TNF, tumor necrosis factor; APC, antigen-presenting cell.

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Preparation of Mice Macrophages and Keratinocytes. Peritoneal macrophages were isolated as described (27). Bone marrow-derived macrophages were generated as described (28). Primary epidermal keratinocytes were prepared from neonatal foreskins as described (29), then cultured on plates coated with 30 μ g·ml⁻¹ type I collagen $(TOYOBO, Tokyo)$ in Keratinocytes-SFM $(GIBCO/BRL)$ with 50 ng·ml⁻¹ bovine pituitary extract (BPE), 10 ng·ml⁻¹ recombinant murine EGF (Sigma), and 0.03 mM CaCl₂ (9). Keratinocytes (2 \times 105 per well in a 24-well plate) were cultured without BPE and EGF for 24 h, then stimulated with recombinant mouse EP (rmEP) (R & D Systems) or recombinant mouse IL-1 α (PeproTech), and mRNA expressions were determined by RT-PCR.

Quantification of Cytokine Concentration of the Culture Supernatants.

Peritoneal macrophages (5×10^5 per well) or bone marrow-derived macrophages (5×10^5 per well) were cultured in 12-well plates for 24 h, then stimulated with rmEP, LPS, PGN, or CpG [LPS from *Salmonella typhimurium* (Sigma); PGN from *Staphylococcus aureus* (Fluka); CpG oligodeoxynucleotide (5--TCCATGACGTTCCT-GATGCT-3') (Sigma)]. The culture supernatants were subjected to IL-6, IL-12p40, and tumor necrosis factor (TNF)- α by ELISA kit (BioSource International).

Isolation of RNA and RT-PCR. Epidermis was isolated from the ear. T, B, CD11b⁺, and CD11c⁺ cells were isolated by magnetic cell sorting (MACS) (Miltenyi Biotec, Auburn, CA). Total RNA from the tissues and sorted cells were isolated by ISOGEN (Nippon Gene, Toyama, Japan), and total RNA from primary epidermal keratinocytes and bone marrow-derived macrophage (BMDM) were isolated by using the RNeasy Mini kit (Qiagen, Valencia, CA). First-strand cDNA were synthesized from 1 μ g of total RNA by Omniscript Reverse Transcriptase (Qiagen), and PCR was done by using LAtaq polymerase (Takara Shuzo, Kyoto) and gene-specific primers: for *EP*, 5'-TTGGGTCTTGACGCTGCTTTGT-3' and 5'-TGAGGTCACTCTCATATTC-3'; for *IL-18*, 5'-ATGGCTGC-CATGTCAGAAGACTCT-3' and 5'-ACTCCATCTTGTTGT-GTCCTGGAAC-3'; for *TLR2*, 5'-TCTCCTGTTGATCTT-GCTCGTAGG-3' and 5'-TTACCCAAAACACTTCCT-GCTGGC-3'; for *TLR4* 5'-TCAGTCTTCTAACTTCCCTC-CTGC-3' and 5'-AGCTGTCCAATAGGGAAGCTTTCT-3'; for *NOD1*, 5'-CAGATAACTGATATCGGAGCCAGG-3' and 5'-TTTGGCCTCCTCGGGCTTAATCAA-3'; for *NOD2*, 5'-AAGCAGAACTTCTTGTCCCTGAGG-3' and 5'-TCACAA-CAAGAGTCTGGCGTCCCT-3'; for *MD2*, 5'-ATGTT-GCCATTTATTCTCTTTTCGACG-3' and 5'-ATTGACAT-CACGGCGGTGAATGATG-3'; GAPDH primers were from Clontech. The optimal cycle number for each gene was determined empirically under nonsaturating conditions.

Measurement of T Cell Responses and Growth Rate of Keratinocytes.

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 $CD4⁺$ T cells were isolated from spleens of mice immunized with 100 μ g of ovalbumin (OVA) emulsified with complete Freund's adjuvant after 14 days. CD4⁺ T cells (4×10^5 cells) were incubated with irradiated peritoneal macrophages (5×10^5 cells) in 96-well plates at the indicated concentration of OVA in RPMI medium 1640 containing 10% FCS for 72 h. They were pulsed with 1 μ Ci $(1 \text{ Ci} = 37 \text{ GBq})$ [³H]thymidine for the final 12 h, and radioactivity was measured by liquid scintillation counting (30). For proliferation assay, primary epidermal keratinocytes $(4 \times 10^4 \text{ cells})$ were plated on a type I collagen-precoated 96-well plate and cultured in the presence of 10 ng·ml⁻¹ EGF, then pulsed with 0.5 μ Ci of [³H]thymidine for the final 8 h, and radioactivity was measured by using liquid scintillation counting (29).

Phagocytosis Assay. Mouse peritoneal macrophages $(5 \times 10^6 \text{ ml}^{-1})$ were incubated for 60 min at 37 $^{\circ}$ C in the presence of 0.75 μ M fluorescent beads (Polysciences). These cells were washed with ice-cold PBS twice and fixed with 2.5% formaldehyde in PBS for 20 min on ice. Phagocytosis was measured by the percentage of phagocytic cells per 100 cells by fluorescent microscopy, as described (31).

Cell Staining and Flow Cytometry. Single-cell suspensions were incubated at 2×10^5 cells per 100 μ l on ice in staining buffer (PBS containing 2.5% FCS and 0.01% NaN₃) with the mAbs for 15 min (28). The mAbs used were anti-I- A^b , anti-CD80, anti-CD86, anti-CD40, and anti-CD11b (Pharmingen). Flow cytometry analysis was performed on a FACScan cytometer (Becton Dickinson).

Bone Marrow Transplantation. Bone marrow transplantation experiments were done as described (32). In brief, bone marrow cells were isolated from $EP^{-/-}$ and WT mice, and recipient mice were irradiated (11 Gy) 12 h before injection (i.v.) of 2×10^7 donor bone marrow cells.

Results

EP Expression in Epidermis and Tissue-Resident APCs. Because human *EP* mRNA is expressed predominantly in peripheral blood lymphocytes (PBL) (12), mouse *EP* expression in immune-related tissues was examined by RT-PCR. Mouse *EP* expression was detected not only in the epidermis (15) but also in peritoneal macrophages, and in macrophages and $CD11c⁺$ dendritic cells in Peyer's patches (Fig. 1*A*). On the other hand, *EP* was rarely detected in the immune system, including bone marrow, thymus, lymph nodes, and spleen (Fig. 1*A*). Furthermore, mouse *EP* expression in PBL was much lower, compared with that in tissueresident APCs (Fig. 1*A*). These results suggested that EP may be involved in the functions of keratinocytes in the epidermis and tissue-resident APCs.

EP Deficiency Results in Dermatitis in Specific Pathogen-Free (SPF) Conditions. To elucidate the function of EP *in vivo*, *EP*-deficient mice were generated, in which 90% of the coding region was deleted (Fig. 1 *B* and *C*). $EP^{-/-}$ mice were born from heterozygous matings and seemed to be indistinguishable from WT mice. However, $EP^{-/-}$ mice had chronic dermatitis at age 5 months at the earliest, even under SPF conditions. The rate of dermatitis observed in $EP^{-/-}$ mice was \approx 20%, 50%, and 75% at the age of 9 months, 12 months, and 15 months, respectively, in the mixed genetic background of 129SV/J and C57BL6/J (Fig. 1D), suggesting that a genetic background and/or environmental factors affect the development of dermatitis in $EP^{-/-}$ mice.

Pathology of the Skin Legions in EP^{-/-} Mice. Most of the skin lesions started as focal skin alterations in the ears or faces and gradually extended to the necks (Fig. 2*A*). The pathological findings of the skin legions in $EP^{-/-}$ mice were typical chronic dermatitis characterized by thickness of epithelial layers and fibrosis with infiltration of inflammatory cells (Fig. 2*B*), including mast cells (Fig. 2*C*) and eosinophils (Fig. 2*D*).

Serum Ig and Cytokine Concentrations in EP-**/**- **Mice.** Serum Ig isotype concentrations, including IgG1, IgG2a, IgG2b, IgG3, IgA, IgM, and IgE, in $EP^{-/-}$ mice were not different from WT mice (Fig. 1*E*). Furthermore, the serum concentrations of cytokines, including IL-2, IL-4, IL-5, IL-6, IL-12, IL-13, IL-18, TNF- α , and IFN- γ did not differ between WT and $EP^{-/-}$ mice without dermatitis (data not shown). All these results suggested that dermatitis in $EP^{-/-}$ mice is not IgE-dependent and systemic immunological disturbance might not exist in $EP^{-/-}$ mice.

Wound Healing Was Not Impaired in EP-**/**- **Mice.** Although EP was reported to be involved in tissue repairing and growth of keratinocytes, wound healing in $EP^{-/-}$ mice was not impaired *in vivo* (Fig. 3*A*), nor was the growth rate of keratinocytes from $EP^{-/-}$ mice different from that of WT mice *in vitro* (Fig. 3*B*). These findings

Fig. 1. *EP* expression in mouse immune-related tissues and generation of *EP*-deficient mice. (*A*) *EP* expression level was determined by using RT-PCR with *GAPDH* as control. PBL, peripheral blood lymphocytes. (*B*) Schematic representations of targeting construct, WT *EP* genomic DNA, and homologous recombinant (HR) allele. The numbers denote exons (filled boxes). External probe (ext) used in Southern blot to detect HR events is indicated by a dashed box. The germ-line and recombinant *Kpn*I restriction fragments are denoted by lines with arrowheads. Restriction sites: K, *Kpn*I; R, *Eco*RI; S, *Sal*I; X, *Xho*I; neo, pGK-neo; D, diphtheria toxin A fragment. (*C*) Southern blot of *Kpn*I-digested genomic DNA with the external probe. (*D*) The rate of chronic dermatitis observed in *EP*/ mice. (*E*) Serum Ig isotype concentration. Serum Ig isotype concentration was determined by ELISA. WT mice (open circles; *n* = 22) and *EP*^{-/-} mice without dermatitis (filled circles; $n = 20$) at the age of 4 months. Mean values are indicated by short horizontal lines.

taken, together, suggested that EP will not play a crucial role in the growth of keratinocytes *in vivo*, which will be possibly due to the functional redundancy of the EGF family.

Fig. 2. Pathology of the skin legions in *EP*-deficient mice. (*A*) Chronic dermatitis around the face of *EP^{-/-}* mice (8 months of age). (*B-D*) Chronic dermatitis in skin in *EP^{-/-}* mice. Sections of skin were subjected to staining with hematoxylin/eosin (*B*), toluidine blue staining for mast cells (*C*), and Luna staining for eosinophils (*D*).

Induction of EP mRNA Expression and Down-Regulation of IL-18 mRNA Expression in Keratinocytes by IL-1 α **.** To elucidate the molecular mechanisms by which *EP* deficiency leads to dermatitis, we next analyzed the inflammatory responses of primary keratinocytes stimulated with IL-1 α (21). IL-1 α induced *EP* mRNA expression at 1 h after stimulation, and it was recovered to the basal level at 8 h after stimulation (Fig. 3C). Of note is that IL-1 α evidently reduced the expression of *IL-18* mRNA at 4 h after stimulation in WTderived keratinocytes and the reduced *IL-18* expression was still observed at 24 h after stimulation, whereas the decreased *IL-18* expression in $EP^{-/-}$ keratinocytes was not prominent, compared with the finding in WT keratinocytes (Fig. 3*C*). These results, together, suggested that IL-1 α -induced down-regulation of *IL-18* expression is impaired in keratinocytes of $EP^{-/-}$ mice.

Down-Regulation of IL-18 mRNA Expression Induced by the Secreted Form of EP. To elucidate whether the secreted form of EP (sEP) affects the down-regulation of *IL-18* expression, keratinocytes were stimulated with rmEP. rmEP reduced *IL-18* expression in WT- and $EP^{-/-}$ -derived keratinocytes at 8 h after stimulation, and there was no obvious difference between WT and $EP^{-/-}$ (Fig. 3D). The expressions were still reduced at 24 h after stimulation in both keratinocytes, although their expressions were gradually increased at 24 h (Fig. 3*D*). These results, together, suggested that IL-1 α induced down-regulation of *IL-18* will be, in part, dependent on sEP. Furthermore, preexposure of keratinocytes with an EGFR kinase blocker (PD168393) (33) prevented rmEP-induced downregulation of *IL-18* expression, whereas a platelet-derived growth factor receptor (PDGFR) blocker (AG1296) as a control did not do so (Fig. 3*E*). These findings, together, suggested that sEP downregulates the *IL-18* expression through EGFR in keratinocytes and that loss of EP leads to the disturbance of tight regulation of *IL-18* expression in keratinocytes, which will, in part, cause the development of dermatitis in $EP^{-/-}$ mice.

Fig. 3. Disturbance of IL-1 α -induced down-regulation of *IL-18* mRNA expression in *EP*^{-/-} mice-derived keratinocytes. (A) Wound healing in WT and *EP*^{-/-} mice. A full-thickness skin excision (6 mm in diameter) was made on the back of WT (open circles; *n* = 10) and *EP^{-/-}* (filled circles; *n* = 10) mice. Values indicate the mean diameter \pm SD. (*B*) Proliferation of primary epidermal keratinocytes from WT (open bars) and *EP^{-/-}* (filled bars) mice. Keratinocytes (4 \times 10⁴ per well in a 96-well plate) were cultured, and cell proliferation at the indicated times was determined by [3H]thymidine incorporation. Data shown in triplicate are the mean \pm SD. (C and *D*) Down-regulation of *IL-18* mRNA expression by IL-1 α (C) and EP (*D*) in primary keratinocytes. Keratinocytes (2 × 10⁵ per well in a 24-well plate) were stimulated with 50 ng·ml⁻¹ recombinant mouse IL-1 a (C) and 100 ng·ml⁻¹ rmEP (D) for the indicated time, and the expressions were determined by using RT-PCR. Experiments were repeated five times with similar results. (*E*) Down-regulation of *IL-18* expression by EP depends on EGFR. Keratinocytes (2 105 per well in a 24-well plate) were preincubated with 2 μM of PD168393 (Calbiochem), 2 μM of AG1296 (Calbiochem), or vehicle for 30 min, followed by incubation with 100 ng·m I^{-1} rmEP in culture medium for 12 h. The expressions were determined by using RT-PCR.

Loss of EP in Non-Bone-Marrow-Derived Cells Is Essential for the Development of Dermatitis. To elucidate that*EP* expression in which cells, keratinocytes, or bone marrow-derived cells including macrophages is essential for the proper homeostasis of the skin region, bone marrow transfer experiments were done. The chimera mice, in which WT-derived bone marrow cells were transplanted into $EP^{-/-}$ mice, developed dermatitis, but not vice versa (Table 1), suggesting that loss of EP in non-bone-marrow-derived cells will be essential for the development of dermatitis and that EP expression in keratinocytes may be critical for the homeostasis of the skin.

Antigen Presentation and Phagocytosis of Peritoneal Macrophages. As *EP* was strongly expressed in tissue-resident macrophages (Fig. 1*A*), we focused on the functions of peritoneal macrophages. First, we examined the potential of antigen presentation of macrophages through T cell responses to the antigen by using immunized mice with ovalbumin. There was no significant difference in proliferation of T cells from the immunized WT and $EP^{-/-}$ mice, when stimulated by WT- or $EP^{-/-}$ -derived peritoneal macrophages with ovalbumin (Fig. 4*A*), indicating that antigen presentation of macrophages and the T cell responses were not impaired in $EP^{-/-}$ mice. Furthermore, the potential of phagocytosis was not impaired in macrophages from $EP^{-/-}$ mice (Fig. 4*B*).

The number of mice with dermatitis was determined 6 months after transplantation.

*The number of mice with dermatitis/the number of recipients.

Low Responsiveness to TLR Agonists in Proinflammatory Cytokine Production by EP^{-/-} Peritoneal Macrophages. To elucidate a possibility that EP is involved in the innate immunity, we examined TLR signaling in macrophages. LPS induced surface expressions of CD80, CD86, and CD40, by peritoneal macrophages from WT and $EP^{-/-}$ mice, and differences were not observed between WT and $EP^{-/-}$ (data not shown). However, IL-6 and TNF- α productions were remarkably reduced in macrophages from $EP^{-/-}$ mice in response to LPS and PGN, compared with findings in WT mice (Fig. 4 *C* and *D*).

Membrane-Anchored EP, but Not Secreted EP, May Play Critical Roles in LPS- and PGN-Induced Proinflammatory Cytokine Production by Macrophages. To elucidate molecular mechanisms underlying the low responsiveness to TLR agonists, we analyzed BMDM. The three TLR agonists, including LPS, PGN, and CpG, induced the MHC class II molecule I-A^b and costimulatory molecules, including CD80, CD86, and CD40, by WT- and $EP^{-/-}$ -derived BMDM, and there was no difference in the surface expressions of these molecules between WT- and $EP^{-/-}$ -derived BMDM (Fig. 4*E* and data not shown). Obvious differences of the responses were observed in the IL-6, IL-12, and TNF- α secretion by BMDM between WT and $EP^{-/-}$ mice, when stimulated with LPS and PGN, but not CpG (Fig. 4 *F*–*H*). Expression levels of *TLR-2*, *TLR-4*, *nucleotide-binding oligomerization domain 1* (*NOD1*), *NOD2* (34), and *MD-2* (28) in *EP*^{-/-} mice–derived BMDM were not different from findings in WT mice (Fig. 4*K*). These results, together, suggested that EP affects the TLR4- and TLR2-mediated signaling in macrophages.

Furthermore, the addition of rmEP to LPS stimulation did not show differences in cytokine production, compared with a single LPS stimulation (Fig. 4*I*), suggesting that sEP does not affect these responses, although LPS and PGN did induce rapid *EP* expression in macrophages (Fig. 4*L*). Preexposure of BMDM with high concentration of PD168393 before LPS stimulation

Fig. 4. TLR agonist-induced proinflammatory cytokine production in macrophages. (*A*) Shown are proliferative responses of T cells to the immunized antigen presented by macrophages. T cell proliferation to the irradiated peritoneal macrophages with the indicated concentration of ovalbumin was evaluated by [³H]thymidine incorporation. M ϕ , macrophage. (*B*) Activity of phagocytosis of macrophages. Phagocytosis of fluorescent beads by macrophages was determined by using fluorescent microscopy. Solid bars, *EP^{-/-}* mice; open bars, WT mice). The values are the means \pm SD of three independent experiments. (C and *D*) Production of IL-6 (C) and TNF- α (D) by the peritoneal macrophages stimulated with LPS (100 ng·ml⁻¹), PGN (10 μ g·ml⁻¹), and EP (100 ng·ml⁻¹). Cytokine concentration in the supernatant after 24 h was examined by using ELISA. Experiments were repeated five times in triplicate with similar results. Data shown in triplicate are the mean \pm SD. Shown are WT (open bars) and $EP^{-/-}$ (filled bars). (E) Surface expression of CD40 on the BMDM stimulated with LPS (100 ng·ml⁻¹), PGN (1 µg·ml⁻¹), and CpG (1 µg·ml⁻¹). Expressions were determined at 24 h after stimulation by FACS. (*F–H*) Production of IL-6 (*F*), TNF- α (G), and IL-12p40 (*H*) by the BMDM stimulated with LPS, PGN, and CpG, respectively. (*I*) Production of IL-6 by the BMDM stimulated by rmEP (100 ng·ml⁻¹) in addition to LPS stimulation. (*J*) Production of IL-6 by the BMDM stimulated with LPS in the presence of EGFR blocker. BMDM (5 × 10⁵ per well in 12-well plates) were preincubated with 2 μ M or 20 μ M of PD168393 for 30 min, followed by stimulation with LPS (100 ng·ml $^{-1}$). Cytokine concentration in the supernatant after 24 h was examined by using ELISA. Experiments were repeated five times in triplicate with similar results. Data shown in triplicate are the mean \pm SD. Shown are WT (open bars) and *EP*/ (filled bars). (*K*) Expression of *Toll-like receptor 2* (*TLR2*), *TLR4*, *nucleotide-binding oligomerization domain 1* (*NOD1*), *NOD2*, and *MD-2* in BMDM from WT and *EP^{-/-}* mice. The expressions were determined by RT-PCR. (*L*) Induction of *EP* expression by LPS and PGN in peritoneal macrophages. Peritoneal macrophages were stimulated with 100 ng·ml⁻¹ LPS and 1 μ g·ml⁻¹ PGN for the indicated time, and the expressions were determined by RT-PCR.

seemed to reduce the total amount of IL-6 production (35); however, the obvious difference in IL-6 secretion between \overline{WT} and $EP^{-/-}$ BMDM still existed (Fig. 4*J*), indicating that the difference is not dependent on EGFR-mediated signaling. All these results suggested that the membrane-anchored EP will play critical roles in the cytokine production by macrophages and/or in macrophage differentiation (17, 36).

Discussion

Here, we demonstrate that *EP* deficiency results in chronic dermatitis in mice and that EP is physiologically a critical molecule for tight control of *IL-18* mRNA expression of keratinocytes and for proper production of proinflammatory cytokines by macrophages in response to LPS and PGN.

Wound healing in the skin in $EP^{-/-}$ mice and the growth of in $EP^{-/-}$ keratinocytes were not impaired (Fig. 3A and *B*), suggesting that dermatitis in $EP^{-/-}$ mice will be caused rather by the dysregulation of immune-related responses of the skin, but not due to the dysfunction of the proliferative activities of keratinocytes. Skin is the place to be damaged frequently, and indigenous bacteria exist in the epidermis, the barrier from the outside milieu. It is intriguing that a member of the EGF family is critically involved in immune regulation of keratinocytes and macrophages. Elucidation of the functions of the EGF family on APCs and epithelial cells in immune regulation might lead to a better understanding of immune defense at the primary interface between the body and the environment and might provide insight into the pathogenesis of inflammatory epithelial disorders.

IL-1 α rapidly induced *EP* expression in keratinocytes (Fig. 3*C*) and the skin lesions in $EP^{-/-}$ mice were mostly restricted to faces and ears, suggesting that IL-1 α release by repeated groominginduced injury may accumulate IL-18 in the skin of $EP^{-/-}$ mice, resulting in the initiation of development of dermatitis in $EP^{-/-}$ mice (22). IL-1 α alone did not induce the active form of IL-18 by primary keratinocytes (data not shown), suggesting that some particular signals, in addition to IL-1 α , may be necessary for the production of the active form of IL-18 by keratinocytes. On the other hand, dysregulation of *IL-18* expression in the skin legions in $EP^{-/-}$ mice *in vivo* was not constantly observed (data not shown), suggesting that dysregulation of IL-1 α -induced *IL-18* downregulation may not be involved in the promotion of dermatitis. Because IL-1 α -deficient mice do not develop dermatitis (37), analysis of $EP^{-/-}/IL$ -1 $\alpha^{-/-}$ doubly mutant mice will show the relation between IL-1 α and EP for the development of dermatitis in $EP^{-/-}$ mice.

Bone marrow transfer experiments supported the critical role of EP in keratinocytes for homeostasis of the skin; however, there are possibilities that EP by keratinocytes might affect the function and/or differentiation of tissue-resident macrophages and that loss of EP in macrophages might affect the progression of dermatitis.

Although *EP* expression in other epithelial tissues was not extensively examined in this study, there is a possibility that EP

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might be involved in immune/inflammatory responses in other epithelial tissues, including intestine and lung. Dendritic cells (DCs) initiate and regulate immune responses, linking innate and adaptive immunity (38). EP expression in Langerhans cells, the DCs in epidermis (39, 40), was not determined in this study, whereas EP expression in DCs in Peyer's patches was strongly observed. There might be a possibility that EP plays a crucial role not only in DCs in Peyer's patches but also in Langerhans cells.

The sEP is involved in the IL-1 α -induced down-regulation of *IL-18* expression (Fig. 3*D*) and the mechanism by which sEP regulates *IL-18* expression depends on the EGFR (Fig. 3*E*). On the other hand, addition of sEP or EGFR blocker did not affect the proinflammatory cytokine production by macrophages (Fig. 4 *I* and *J*), suggesting that the membrane-anchored EP may be involved in the proinflammatory cytokine production by macrophages stimulated with LPS and PGN through non-EGFR-mediated signaling. A bioactive membrane precursor, pro-HB-EGF, functions as the diphtheria toxin receptor in primates (41) and distinctively signals growth inhibition or apoptosis rather than the proliferative response induced by soluble HB-EGF (10). Like HB-EGF, sEP and membrane-anchored EP will have distinct function and there might be a possibility that membrane-anchored EP transduces signals into the own cells as receptor (8).

In conclusion, we have demonstrated that $EP^{-/-}$ mice are susceptible for the development of dermatitis even in specific pathogen-free conditions and that sEP is critical for the proper regulation of *IL-18* in keratinocytes and membrane-anchored EP will be essential for the proper proinflammatory cytokine production in response to LPS and PGN by tissue resident macrophages. Further elucidation of EP function may shed light on epithelial inflammatory disorders.

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