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T Cell Requirement for Development of Chronic Ulcerative Dermatitis in E- and P-Selectin-Deficient Mice

S. Bradley Forlow,*† E. James White,* Kennard L. Thomas,*§ Gregory J. Bagby,‖ Patricia L. Foley,‡ and Klaus Ley*‡

C57BL/6 mice deficient in E- and P-selectin (E−/−P−/−) kept under specific pathogen-free barrier conditions have high circulating neutrophil counts and develop hypercellular cervical lymph node and overlying squamous epithelium and superficial colonization by Enterococcus faecium, Staphylococcus xylosus, and Staphylococcus aureus, Escherichia coli, Enterococcus faecium, Staphylococcus xylosus, and Staphylococcus viridans (1, 4). Histopathological analysis of the affected skin showed a loss of the 90% of the mice showing lesions by 6–8 mo (1, 4, 6). Histopathological analysis of the affected skin showed a loss of the overlying squamous epithelium and superficial colonization by bacteria (1). Cultures on infected skin were positive for common bacterial flora, including Staphylococcus aureus, Escherichia coli, Enterococcus faecium, Staphylococcus xylosus, and Staphylococcus viridans (1, 4). Cultures for bacterial and fungal organisms have been consistently negative in the blood, spleen, liver, lymph nodes, and lungs of these mice, suggesting that bacterial colonization of skin lesions is secondary to skin ulceration (1, 4). The causative factors leading to the development of ulcerative dermatitis have not been identified.

Mice lacking E-, P-, and L-selectin (E−/−P−/−L−/−) have been generated by two groups. Robinson et al. (7) reported that the E−/−P−/−L−/− mice (Hynes mutation) developed ulcerative dermatitis although the frequency and age of onset were not evaluated. The E−/−P−/−L−/− mice described by Collins et al. (8) (Baylor mutation) are largely protected from ulcerative dermatitis seen in E−/−P−/− mice. Only 3 of 156 E−/−P−/−L−/− mice showed ulcerative dermatitis by 6 mo of age. However, this same line of E−/−P−/−L−/− mice (Baylor mutation) developed skin lesions at another facility (University of Washington, Seattle, WA), suggesting that lesion development is related to differences in animal husbandry and environment. Mice deficient in E- and P-selectin and ICAM-1 (E−/−P−/−L−/−) show a complete protection from the development of ulcerative dermatitis (9), suggesting a role for ICAM-1 in the development of ulcerative dermatitis that occurs in the absence of E- and P-selectin.

The relative protection from ulcerative dermatitis in E−/−P−/−L−/− and E−/−P−/−/− mice suggests that both L-selectin and ICAM-1 may serve a function critical to lesion development. L-selectin is necessary for lymphocyte homing into peripheral lymph nodes (10) and for lymphocyte recirculation (11). As a result, L-selectin-deficient mice have altered lymphocyte function, impaired T cell responses, and display very small lymph nodes (12). Because trafficking of naive lymphocytes is impaired in E−/−P−/−L−/− mice, the immune response to bacterial Ags present in the normal flora may be reduced. T cell-stimulating activity was also shown to be impaired in mice lacking ICAM-1, suggesting that ICAM-1 is an important costimulatory molecule (13). Based on these findings, we hypothesized that defective lymphocyte function in E−/−P−/−L−/− and E−/−P−/−/− mice protects these mice from lesion development. We inferred that the ulcerative dermatitis seen in E−/−P−/− mice may be lymphocyte-dependent. To test the hypothesis that the development of ulcerative dermatitis in E−/−P−/− mice requires mature T cells, we generated mice deficient in E- and P-selectin and the V(D)J

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recombination activation gene (Rag)−1 (14) (E−/−P−/−/Rag−1−/−) by cross-breeding E−/−P−/− with Rag−1−/− mice. Mice lacking the V(DJ) Rag-1 do not have mature T or B lymphocytes (14). E−/−P−/−Rag−1−/− mice are healthier than E−/−P−/− mice and do not develop ulcerative dermatitis seen in E−/−P−/− mice.

Materials and Methods

Mice

Mice deficient in both E- and P-selectin and Rag-1 were generated by cross-breeding E−/−P−/− C57BL/6 N6 (1) and Rag−1−/− C57BL/6 N10 (The Jackson Laboratory, Bar Harbor, ME) mutants. F1 hybrids were back-crossed to E−/−P−/− to generate E−/−P−/−Rag−1−/− and E−/−P−/−Rag−1−/− mice. F1 hybrids were back-crossed to Rag−1−/− mice to generate E+/−P−/−Rag−1−/− and E+/−P−/−Rag−1−/−. Breeding was established from these offsprings to generate all necessary control groups, including E−/−P−/−Rag−1−/−, E−/−P−/−Rag−1−/−, E+/−P−/−Rag−1−/−, E+/−P−/−Rag−1−/−, and E−/−P−/−Rag−1−/− mice. Because there is no evidence for haplo-insufficiency in any of the selectin mutants, E+/−P−/− mice were pooled with E−/−P−/− mice. Wild-type refers to E−/−P−/−Rag−1−/− and E+/−P−/−Rag−1−/− mice; Rag−1−/− refers to E+/−P−/−Rag−1−/− and E+/−P−/−Rag−1−/− mice; E−/−P−/−Rag−1−/− mice; and E−/−P−/−Rag−1−/− mice. Wild-type refers to E−/−P−/−Rag−1−/− mice. Because there is no evidence for haplo-insufficiency in any of the selectin mutants, E+/−P−/− mice were pooled with E−/−P−/− mice. Wild-type refers to E−/−P−/−Rag−1−/− and E+/−P−/−Rag−1−/− mice; Rag−1−/− refers to E+/−P−/−Rag−1−/− and E+/−P−/−Rag−1−/− mice; E−/−P−/−Rag−1−/− mice; and E−/−P−/−Rag−1−/− mice.

Polymerase chain reaction

Genomic DNA was analyzed using PCR for the wild-type and knockout E- and P-selectin alleles. The P-selectin PCR assay used a forward primer from exon 3 of murine P-selectin (5′-TCA CGG GTG TTC TGC TGT AGG AGG-3′) and a reverse primer from exon 3 (5′-GGG GCC GAG TTA CTC TTG ATGC-3′), yielding a 260-bp wild-type fragment. The P-selectin knockout allele was detected using a forward primer from exon 2 (5′-GGT GCC GTC CGA AAA GGT T-3′) and a reverse primer from the inserted neomycin resistance gene (5′-ACC CGT GAT ATT CCT GGA GAG C-3′). This reaction yielded a 1300-bp fragment. PCR conditions for the detection of the wild-type alleles were 94°C for 2 min, 55°C for 30 s, and 72°C for 30 s for 25 cycles. PCR conditions for the detection of the P-selectin knockout alleles were 94°C for 4 min, 35 cycles of 94°C for 1 min, 61°C for 1 min, and 72°C for 1 min, followed by 72°C for 7 min.

The E-selectin PCR assay used a forward primer from exon 5 of murine E-selectin (5′-TTG GCT GAT GTA AAA GGG GCT ACC-3′) and a reverse primer from exon 6 (5′-CAT GAT GCC GTC TCG TTA TTA CCT G-3′), yielding a 1000-bp wild-type fragment. The E-selectin knockout allele was detected by a forward primer from hypoxanthine phosphoribosyltransferase gene (5′-ACT ATC AGT TCC CTT TGG GCC-3′) and a reverse primer from exon 3 (5′-CAT GAT GCC GTC TCG TTA TTA CCT G-3′), which yields a 650-bp knockout fragment. PCR conditions for the detection of the E-selectin wild-type alleles were 94°C for 4 min, 35 cycles of 94°C for 1 min, 57°C for 1 min, 72°C for 1 min, and 72°C for 7 min. PCR conditions for the detection of the E-selectin knockout alleles were 94°C for 4 min, 35 cycles of 94°C for 1 min, 59°C for 1 min, and 72°C for 1 min, followed by 72°C for 7 min.

Serum G-CSF measurements

Serum G-CSF protein concentrations were determined using a specific enzyme-linked immunoassay using Ab pairs purchased from R&D Systems (Minneapolis, MN). For this purpose, 96-well plates (NuncImmunoPlates Maxisorb; Nunc, Neptune, NJ) were coated with 2 µg/ml of capture Ab and incubated overnight at 4°C. The plates were then washed five times with wash buffer (0.05% Tween 20 in PBS) and blocked with 200 µl 2% BSA in wash buffer for 2 h at room temperature. G-CSF standards and samples were diluted in wash buffer containing 2% FCS. Both standards (31.25-1000 pg/ml) and samples (50 µl) were added to wells and the plates were incubated for 1 h at 37°C. After washing, 50 µl of biotinylated anti-G-CSF (0.1 µg/ml in dilution buffer) was added and the plates were incubated for 1 h at 37°C. Wells were then washed and incubated for 1 h after adding 100 µl of 0.1 µg/ml peroxidase-conjugated streptavidin (Jackson Immunoresearch Laboratories, West Grove, PA) in dilution buffer. After washing the plate, 100 µl of tetramethylbenzidine (Sigma-Aldrich, St. Louis, MO) was added as substrate and color was allowed to develop for 30 min in the dark. After stopping the reaction with 50 µl of 3 M H2SO4, ODs were determined at 450 nm. G-CSF concentrations were calculated from the standard curve using log-log linear regression.

Flow cytometry

Peripheral blood was collected from the tail vein. Expression of CD4 and CD8 on peripheral lymphocytes was determined by direct immunofluorescence to identify mice lacking the V(DJ) Rag-1. Lymphocytes were identified and gated by forward and side scatter properties. CD8+ cells were identified by FITC-labeled mAb 53-6.7 (BD Pharmingen, San Diego, CA) staining. CD4+ cells were identified by FITC-labeled mAb GK.1.5 (BD Pharmingen) staining. Cells were analyzed by forward scatter, side scatter, and FITC fluorescence using a laser flow cytometer (FACSCan; BD Biosciences, Mountain View, CA). Data are presented as fluorescence histograms of CD4 and CD8 expression on a four-decade log scale.

Histopathology

Age-matched wild-type, E−/−P−/− with and without ulcerative dermatitis, and E−/−P−/−Rag−1−/− mice were euthanized by cervical dislocation under anesthesia for histopathological analysis. After gross examination for any discernable pathology, lung, intestine, kidney, cervical lymph nodes, liver, spleen, and skin samples were fixed in 10% buffered formalin (Sigma-Aldrich), embedded in paraffin, sectioned, and examined after H&E staining.

Microbiological analyses

Tissue samples for microbiological analysis were obtained under sterile conditions from the lung, liver, and spleen from wild-type Rag−1−/−, E−/−P−/− with and without ulcerative dermatitis, and E−/−P−/−Rag−1−/− mice. Tissues were homogenized with a sterile grinder in 0.5 ml PBS. A 100-µl sample from each tissue was plated onto trypticase soy agar with 5% sheep RBCs (Difco, Detroit, MI). Plates were observed at 24 and 48 h for growth.

Statistics

Total leukocyte counts, neutrophil counts, mononuclear cell counts, and serum G-CSF levels between groups were compared using Kruskal-Wallis One-Way ANOVA on ranks and pairwise multiple comparison by Dunn’s method. Spleen weights and lymph node weights between groups were compared using Kruskal-Wallis one-way ANOVA and pairwise multiple comparison by the Tukey test. Statistical significance was set at p < 0.05.

Results

Generation of E−/−P−/−Rag−1−/− mice

Mice deficient in E- and P-selectin and Rag-1 were generated by cross-breeding E−/−P−/− C57BL/6 N6 (1) and Rag−1−/− C57BL/6 N10 (The Jackson Laboratory) mutants. This resulted in all necessary control groups, including E−/−P−/−Rag−1−/− (wild-type), E+/−P−/−Rag−1−/− (Rag−1−/−), E−/−P−/−Rag−1−/− (E−/−P−/−), and E−/−P−/−Rag−1−/− (Fig. IA; Table I). Mice expressing E- and P-selectin (E+/−P+/+), heterozygous for E- and P-selectin (E+/−P+/−), and homozygous for a null mutation in E- and P-selectin (E−/−P−/−) were determined by PCR. Consistent with previous results (1, 4) and the close distance of the E- and P-selectin genes, we found that E- and P-selectin cosegregated in all mice tested.

Mice expressing E- and P-selectin (E+/−P+/+) show a band from the wild-type P-selectin reaction at 260 bp and at 1000 bp from the E-selectin wild-type reaction (Fig. IA, lane 1). No band is detectable either of the knockout reactions in E+/−P+/+ mice (Fig. IA, lane 2). A heterozygous mutation for E- and P-selectin (E+/−P+/+) results in both wild-type and knockout bands from the P-selectin and the E-selectin reactions (Fig. IA, lanes 3 and 4). Mice that have a null mutation in E- and P-selectin (E−/−P−/−) show no band from either wild-type reaction (Fig. IA, lanes 5 and 7) and a single band from the respective knockout reactions (Fig. IA, lanes 6 and 8). Verification for a null mutation for Rag-1 was determined by flow cytometry. Mice deficient in Rag-1 have no CD4+ or CD8+ lymphocytes in blood (Fig. IB, bottom panels).

Abbreviations used in this paper: Rag, recombination activation gene.
fltin genotype was identified by a 650-kb mutant fragment and a 1000-kb
epithelium, and a dermal in
mononuclear cells in the peripheral circulation compared with
wild-type mice (Table II). Rag1-deficient mice had neutrophil
counts similar to wild-type mice (Table II). E⁻/⁻P⁻/⁻ mice dis-
played significantly elevated leukocyte counts compared with
wild-type and Rag-1⁻/⁻ mice. E⁻/⁻P⁻/⁻ mice were severely neu-
trophilic (Table II). Neutrophil counts in E⁻/⁻P⁻/⁻Rag-1⁻/⁻ mice were almost as high as in E⁻/⁻P⁻/⁻ mice. However, E⁻/⁻
P⁻/⁻Rag-1⁻/⁻ mice had significantly fewer circulating mononu-
clear cells than E⁻/⁻P⁻/⁻ mice (Table II).

Consistent with elevated circulating neutrophil levels, E⁻/⁻P⁻/⁻ mice showed increased serum G-CSF levels compared with
wild-type and Rag-1⁻/⁻ mice (Table II). E⁻/⁻P⁻/⁻Rag-1⁻/⁻ mice showed G-CSF levels similar to E⁻/⁻P⁻/⁻ mice (Table II).
Similar increases in neutrophil counts and G-CSF levels in E⁻/⁻P⁻/⁻Rag-1⁻/⁻ mice compared with E⁻/⁻P⁻/⁻ mice show that the level of neutrophilia in these mice is not dependent on
lymphocyte function (Table II).

Clinical and pathological findings
E⁻/⁻P⁻/⁻Rag-1⁻/⁻ mice were generally healthier than E⁻/⁻P⁻/⁻ mice. E⁻/⁻P⁻/⁻Rag-1⁻/⁻ breeders consistently produced litters for ~1 year. E⁻/⁻P⁻/⁻ breeders only occasionally generated litters until 5–6 mo of age in the same specific pathogen-free barrier facility. Therefore, the E⁻/⁻P⁻/⁻ colony was maintained with heterozygous breeders.

Approximately 50% (18 of 35) of the E⁻/⁻P⁻/⁻ mice 6–12 mo old developed conjunctivitis and/or facial and submandibular ulcerative dermatitis (Fig. 2C). Initially, these mice showed hair loss around the eyes and the development of conjunctivitis. This was followed by hair loss on the nose and neck and subsequent skin lesion development (Fig. 2A). These lesions have primarily been shown to be colonized by commensal bacteria, but not specific pathogens (1, 4). E⁻/⁻P⁻/⁻Rag-1⁻/⁻ mice were completely protected from the development of ulcerative dermatitis (Fig. 2B).

None of the 62 E⁻/⁻P⁻/⁻Rag-1⁻/⁻ mice developed facial or sub-
mandibular ulcerative dermatitis, including 25 mice older than 12
mo (Fig. 2C).

Microbiological cultures
To investigate possible bacteremia and colonization of internal or-
gans by commensal bacteria in E⁻/⁻P⁻/⁻ mice, liver, lung, and
spleen samples were analyzed for bacterial growth. No positive
cultures were obtained in wild-type (n = 4), Rag-1⁻/⁻ (n = 5),
E⁻/⁻P⁻/⁻ showing ulcerative dermatitis (n = 5), E⁻/⁻P⁻/⁻ with
no signs of ulcerative dermatitis (n = 5), or E⁻/⁻P⁻/⁻Rag-1⁻/⁻ (n = 8) mice.

Histopathology
Cervical skin from E⁻/⁻P⁻/⁻Rag-1⁻/⁻ mice showed an intact epi-
dermis, no dermal infiltrate, and no bacterial colonization (Fig.
3C). Tissue samples from E⁻/⁻P⁻/⁻ mice with ulcerative skin
lesions showed areas of epidermal loss as well as hyperplastic
epithelium, and a dermal infiltrate consisting of mixed inflamma-

White blood cell counts and serum G-CSF levels
Because mice deficient in Rag-1 do not develop functional T or B
lymphocytes, Rag-1⁻/⁻ mice show a significant reduction of
mononuclear cells in the peripheral circulation compared with
wild-type mice (Table II). Rag1-deficient mice had neutrophil

![FIGURE 1. E⁻/⁻P⁻/⁻Rag-1⁻/⁻ mice were identified by PCR and flow cytometry. A, PCR was used to determine mice expressing E- and P-selectin (E⁻/⁻P⁻/⁻), heterozygous for E- and P-selectin (E⁻/⁻P⁺/⁻), and homozygous null mutations in E- and P-selectin (E⁻/⁻P⁻/⁻). The E-selectin genotype was identified by a 650-kb mutant fragment and a 1000-kb wild-type fragment. The P-selectin genotype was identified by a 1300-kb mutant fragment and a 260-kb wild-type (WT) fragment. B, Rag-1-deficient mice were identified by absence of CD4⁺ and CD8⁺ expressing cells in whole blood using flow cytometry. KO, Knockout.](image-url)

<table>
<thead>
<tr>
<th>Table I. No. of mice in study</th>
<th>No. of Mice</th>
</tr>
</thead>
<tbody>
<tr>
<td>Wild type</td>
<td>67</td>
</tr>
<tr>
<td>Rag-1⁻/⁻</td>
<td>33</td>
</tr>
<tr>
<td>E⁻/⁻P⁻/⁻</td>
<td>35</td>
</tr>
<tr>
<td>E⁻/⁻P⁻/⁻Rag-1⁻/⁻</td>
<td>62</td>
</tr>
</tbody>
</table>
body weights in E/P \(^{-/-}\) mice, the spleen weights normalized for these mice showed an absence of white pulp (Fig. 4). As a result, mice exhibited a similar pattern of extramedullary hemopoiesis; however, 50% of E\(^{-/-}\)/P\(^{-/-}\) mice generally had less inflammatory cell infiltrate in the lungs compared with E\(^{-/-}\)/P\(^{-/-}\) mice (Fig. 5). Where seen, infiltrates appeared to be more focal than diffusely distributed.

The splenic architecture of wild-type mice showed normal, well-defined regions of red and white pulp (Fig. 4). As expected, the lymphoid follicles (white pulp) were absent in Rag-1\(^{-/-}\) mice (Fig. 4), but some myeloid and erythroid precursor cells were present. E\(^{-/-}\)/P\(^{-/-}\) mice showed a profound expansion of red pulp due to extramedullary hemopoiesis, disrupting the normal splenic architecture (Fig. 4). Large numbers of megakaryocytes, erythroid, myeloid precursors, neutrophils, and plasma cells were present throughout the spleen. The white pulp in these mice was smaller, distorted, and less distinct (Fig. 4). E\(^{-/-}\)/P\(^{-/-}\)/Rag-1\(^{-/-}\) mice exhibited a similar pattern of extramedullary hemopoiesis; however, these mice showed an absence of white pulp (Fig. 4). As a result of extramedullary hemopoiesis, the spleen weights normalized for body weights in E\(^{-/-}\)/P\(^{-/-}\) and E\(^{-/-}\)/P\(^{-/-}\)/Rag-1\(^{-/-}\) mice were increased ~3- and 2-fold, respectively, compared with wild-type and Rag-1\(^{-/-}\) mice (Table III).

Lung sections from Rag-1\(^{-/-}\) mice (Fig. 5) were indistinguishable from wild-type mice (data not shown). Mice lacking E- and P-selectin showed degrees of lung pathology varying from mild to severe. Interstitial cellularity was increased secondary to leukocyte infiltration (Fig. 5), and in more severe cases evidence of interstitial edema and hemorrhage were present. E\(^{-/-}\)/P\(^{-/-}\)/Rag-1\(^{-/-}\)

### Table II. Leukocyte counts and serum G-CSF levels

<table>
<thead>
<tr>
<th></th>
<th>Total Leukocyte Counts (cells/µl)</th>
<th>Neutrophils/µl</th>
<th>Mononuclear Cells/µl</th>
<th>Serum G-CSF (pg/ml)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Wild type</td>
<td>15,000 ± 1,000</td>
<td>2,000 ± 200</td>
<td>12,000 ± 1,000</td>
<td>24 ± 4</td>
</tr>
<tr>
<td>Rag-1(^{-/-})</td>
<td>3,500 ± 300(^{a})</td>
<td>1,700 ± 200</td>
<td>3,000 ± 600(^{a})</td>
<td>43 ± 6</td>
</tr>
<tr>
<td>E(^{-/-})/P(^{-/-})</td>
<td>41,000 ± 2,000(^{e})</td>
<td>23,000 ± 1,000</td>
<td>18,000 ± 1,000(^{e})</td>
<td>72 ± 11(^{e})</td>
</tr>
<tr>
<td>E(^{-/-})/P(^{-/-})/Rag-1(^{-/-})</td>
<td>22,000 ± 2,000(^{d})</td>
<td>18,000 ± 1,000</td>
<td>4,100 ± 300(^{d})</td>
<td>91 ± 16(^{d})</td>
</tr>
</tbody>
</table>

*All values expressed as mean ± SEM.

Different from wild-type and Rag-1\(^{-/-}\) groups.

Different from wild-type and E\(^{-/-}\)/P\(^{-/-}\) groups.

Different from Rag-1\(^{-/-}\) and E\(^{-/-}\)/P\(^{-/-}\)/Rag-1\(^{-/-}\) groups.

Different from wild-type group (p < 0.05).

#### FIGURE 3. Histopathology of ventral cervical skin sections. A and B. Cervical skin from E\(^{-/-}\)/P\(^{-/-}\) mice was ulcerated (arrowhead) and showed a chronic hyperplastic epithelial response in areas bordering epithelial denudation (arrows). A and B. The dermis is characterized by a mixed inflammatory infiltrate, noticeably lacking in neutrophils but containing plasma cells, macrophages, and other mononuclear cells. C. Skin from E\(^{-/-}\)/P\(^{-/-}\)/Rag-1\(^{-/-}\) mice showed none of the changes described in the E\(^{-/-}\)/P\(^{-/-}\) mice. The epithelium has a normal uniform thickness, and the dermis contains normal connective and muscular tissue with very few inflammatory cells present. Sebaceous glands and hair follicles also demonstrate normal architecture. Original magnification: A and C, ×10; B, ×40.

#### FIGURE 2. E\(^{-/-}\)/P\(^{-/-}\)/Rag-1\(^{-/-}\) mice do not develop ulcerative dermatitis. A. Mice lacking E- and P-selectin often develop conjunctivitis that progresses to ulcerative dermatitis on the nose and neck. C. Approximately 50% of E\(^{-/-}\)/P\(^{-/-}\) mice displayed ulcerative dermatitis or conjunctivitis by 10 mo of age. B. E\(^{-/-}\)/P\(^{-/-}\)/Rag-1\(^{-/-}\) mice are protected from the development of ulcerative dermatitis. None of the 62 E\(^{-/-}\)/P\(^{-/-}\)/Rag-1\(^{-/-}\) mice developed lesions, including 25 mice >12 mo old (C).
Rag-1-deficient mice had small cervical lymph nodes (5.0 ± 0.4 mg) with few mature lymphoid cells (Fig. 6) compared with wild-type mice (11.0 ± 0.1 mg; Fig. 6; Table III). E−/−P−/− mice displayed cervical lymphadenopathy (19.6 ± 2.4 mg), the normal corticomedullary architecture was disrupted, and lymph nodes were infiltrated by inflammatory mononuclear and plasma cells (Fig. 6). In some cases, particularly in mice with severe skin lesions overlying the submandibular lymph nodes, lymphoid follicles were completely effaced by plasma cells. E−/−P−/−Rag-1−/− mice had very small lymph nodes that were not different from Rag-1-deficient mice (3.7 ± 0.7 mg; Table III). These lymph nodes contained few lymphocytes, no plasma cell infiltrate, and more connective tissue and fibroblasts (Fig. 6). Colon, small intestine, liver, and kidney from all groups showed no significant pathology.

Discussion

The development of conjunctivitis and ulcerative dermatitis in E−/−P−/− mice is associated with an active immune response and plasma cell infiltration of cervical lymph nodes. In this study, we show that E−/−P−/−Rag-1−/− mice lacking this immune response did not develop ulcerative dermatitis, showed small cervical lymph nodes, and decreased lung pathology compared with E−/−P−/− mice. This indicates that an altered or enhanced immune response in E−/−P−/− mice is, at least in part, causative of the phenotype.

Table III. Spleen and lymph node weights

<table>
<thead>
<tr>
<th></th>
<th>Spleen wt (mg)</th>
<th>Spleen wt/Body wt (mg/g)</th>
<th>One Cervical Lymph Node wt (mg)</th>
<th>Cervical Lymph Node wt/Body wt (mg/g)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Wild type</td>
<td>90 ± 6</td>
<td>3.0 ± 0.2*</td>
<td>11.0 ± 0.1*</td>
<td>0.36 ± 0.02*</td>
</tr>
<tr>
<td>Rag-1−/−</td>
<td>62 ± 19*</td>
<td>2.3 ± 1.0*</td>
<td>5.0 ± 0.4*</td>
<td>0.18 ± 0.04*</td>
</tr>
<tr>
<td>E−/−P−/−</td>
<td>175 ± 21*</td>
<td>8.3 ± 1.2*</td>
<td>19.6 ± 2.4*</td>
<td>0.93 ± 0.09*</td>
</tr>
<tr>
<td>E−/−P−/−Rag-1−/−</td>
<td>130 ± 20</td>
<td>5.0 ± 0.9</td>
<td>3.7 ± 0.7*</td>
<td>0.14 ± 0.03*</td>
</tr>
</tbody>
</table>

* All values expressed as mean ± SEM.
* wt, weight.
* Different from E−/−P−/− group.
* Different from Rag-1−/− group.
* Different from wild-type group.
* Different from E−/−P−/−Rag-1−/− group (p < 0.05).
E−/−P−/− mice develop ulcerative dermatitis of the nose and cervical regions at all institutions where they are bred. Mice lacking all three selectins (E−/−P−/−L−/−) develop ulcerative dermatitis in some facilities (University of Washington, Seattle, WA; and Harvard Medical School, Boston, MA), while they are largely protected from disease development in other facilities (Baylor College of Medicine, Houston, TX; University of Virginia, Charlottesville, VA). The causative trigger of ulcerative dermatitis is not known but appears to be related to environmental and/or housing conditions. We hypothesized that the skin pathology results from a compensatory immune response to commensal bacteria, including plasma cell hyperproliferation, and that a diminished immune response in the absence of ICAM-1 (E−/−P−/−I−/−) and L-selectin (E−/−P−/−L−/−) protects these mice from disease development. In this study, we show that an absence of functional T and B cells (E−/−P−/−Rag-1−/−) completely protects these mice from the development of ulcerative dermatitis, although housed in the same environment.

Histological analysis of skin sections of E−/−P−/− mice reveals a complete absence of neutrophils in skin lesions, showing that neutrophils are not recruited to these areas of infection in the absence of E- and P-selectin. Interestingly, significant levels of lymphocyte recruitment occur despite the lack of E- and P-selectin. Lymphocyte recruitment to skin lesions is likely due to the critical involvement of α4 integrins in the recruitment of lymphocytes to sites of chronic inflammation. We have shown that lymphocyte recruitment and adhesion were not affected in chronic models of inflammation (thioglycolate-induced peritonitis and 6-h TNF-α-induced inflammation of the mouse cremaster muscle, respectively) in mice lacking E-, P-, and L-selectin and ICAM-1 (E−/−P−/−I−/−−/−; Ref. 15). Leukocyte adhesion was almost completely inhibited only with anti-α4 integrin mAb pretreatment.

T lymphocyte migration to dermal inflammation induced by IFN-γ, TNF-α, LPS, or polyinosine-cytosine was markedly reduced by anti-α4 integrin treatment (50–60%). T lymphocyte migration was inhibited by only 30–60% with anti-E- and P-selectin treatment compared with 10–40% with either anti-E- or P-selectin treatment. Blockade of both E- and P-selectin and α4 integrin function severely impaired lymphocyte recruitment in these models of dermal inflammation (>90%; Ref. 16). Although E- and P-selectin have a role in lymphocyte recruitment, they are not strictly required in some models of inflammation.

The molecular mechanisms linking a dysregulated immune response to conjunctivitis and cervical ulcerative dermatitis remain unknown and were not addressed in this study. Decreased neutrophil trafficking in E−/−P−/− mice may result in expansion...
of commensal bacteria and an increased antigenic load which may lead to the release of proinflammatory cytokines which potentiate host tissue damage and ulceration. Others have shown that T cells are actively involved in inflammatory skin disorders, including allergic contact dermatitis, atopic dermatitis, and psoriasis (17–19). Cytokines released by activated skin infiltrating T lymphocytes, such as TNF-α, IFN-γ, and IL-4, induce the activation of keratinocytes to produce proinflammatory chemokines, including inducible protein-10, IFN-inducible T cell chemoattractant, and monokine induced by IFN-γ which results in the continued recruitment of CXCR3+ T cells (17). Chronic abnormal chemokine and proinflammatory cytokine production from activated infiltrating T cells and keratinocytes may lead to host tissue damage and lesion development. Consistent with this interpretation, removing lymphocyte function in E–/–P–/– Rag-1–/– mice eliminates this overshooting response to normal flora Ags and protects these mice from tissue damage and ulceration despite defective neutrophil trafficking. Remarkably, the same commensal bacteria colonizing dermal lesions in E–/–P–/– mice do not invade E–/–P–/–Rag-1–/– mice. This suggests that the dysregulated immune response in E–/–P–/– mice is pathogenic rather than protective under vari-um conditions.

Secretion of various cytokines from T cells promotes neutrophil recruitment and stimulates granulopoiesis, linking the acquired immune system with the innate system. IL-17 released from activated T cells has been shown to stimulate neutrophil recruitment through the release of keratinocyte-derived chemokine and macrophage-inflammatory protein-2 (20, 21). T cells regulate hemopoiesis through secretion of various cytokines, including IL-2, IL-3, IL-6, IL-8, IL-17, and GM-CSF (22, 23). IL-17 stimulates granulopoiesis in vivo through G-CSF release (22). E–/–P–/–Rag-1–/– mice show elevated serum G-CSF and levels of neutrophilia similar to E–/–P–/– mice, suggesting that lymphocyte-independent mechanisms completely regulate the elevated neutrophil levels seen in these mice (~20,000 polymorphonuclear cells/μl). Surprisingly, E–/–P–/–Rag-1–/– mice do not show neutrophil levels higher than E–/–P–/– mice. Although granulopoiesis can be enhanced or modulated by the acquired immune system (22), lymphocyte-independent granulopoiesis in E–/–P–/–Rag-1–/– mice is sufficient to compensate for the loss of the acquired immune system. This suggests there is an independence of the acquired and innate immune systems in regulating this level of neutrophilia.

In conclusion, we show that E–/–P–/– Rag-1–/– mice which lack functional lymphocytes display a healthier phenotype than E–/–P–/– mice, although circulating neutrophil counts and serum G-CSF levels are similar to E–/–P–/– mice. E–/–P–/–Rag-1–/– mice did not have hypercellular cervical lymph nodes and showed reduced leukocyte infiltrate in the lungs. Most importantly, E–/–P–/–Rag-1–/– mice were completely protected from the ulcerative dermatitis that develops in E–/–P–/– mice. We conclude that a dysfunctional immune response in E–/–P–/–Rag-1–/– mice resulting from impaired leukocyte trafficking leads to ulcerative dermatitis in E–/–P–/– mice.

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References


CORRECTIONS


An error was made in describing the protocol used for experiments shown in Figs. 5, 6, and 7. Text in the Materials and Methods and Results sections, along with figure legends for Figs. 5, 6, and 7 implied that antigenic readout peptide was only added subsequent to the incubation of macrophages in citrate-buffered saline at varying pH, but peptide was actually added at the indicated concentrations both during the incubation in citrate-buffered saline at varying pH and in the subsequent incubation for 2 h in normal medium. This correction does not change the conclusion of the paper, but is important for executing future studies.


In Fig. 1A, the last lane in the lower right panel was misplaced. It should have appeared as the first lane in the left side of the lower panel. The corrected figure is shown below.