A critical role for neutrophil elastase in experimental bullous pemphigoid

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Bullous pemphigoid (BP) is an autoimmune skin disease characterized by subepidermal blisters and autoantibodies against 2 hemidesmosome-associated proteins, BP180 and BP230. The immunopathologic features of BP can be reproduced in mice by passive transfer of anti-BP180 antibodies. Lesion formation in this animal model depends upon complement activation and neutrophil recruitment. In the present study, we investigated the role of neutrophil elastase (NE) in antibody-induced blister formation in experimental BP. Abnormally high levels of caseinolytic activity, consistent with NE, were detected in extracts of lesional skin and blister fluid of mice injected with anti-BP180 IgG. The pathogenic anti-BP180 IgG failed to induce subepidermal blistering in NE-null (NE−/−) mutant mice. NE−/− mice reconstituted with neutrophils from wild-type mice became susceptible to experimental BP. Wild-type mice given NE inhibitors (α1-proteinase inhibitor and Me-O-Suc-Ala-Ala-Pro-Val-CH2Cl), but not mice given cathepsin G/chymase inhibitors (α1-antichymotrypsin or Z-Gly-Leu-Phe-CH2Cl), were resistant to the pathogenic activity of anti-BP180 antibodies. Incubation of murine skin with NE induced BP-like epidermal-dermal detachment. Finally, NE cleaved BP180 in vitro and in vivo. These results implicate NE directly in the dermal-epidermal cleavage induced by anti-BP180 antibodies in the experimental BP model.


Introduction

Bullous pemphigoid (BP) is an autoimmune skin disease characterized by subepidermal blisters, a dermal inflammatory infiltrate, and in vivo deposition of autoantibodies and complement components along the dermal-epidermal junction (DEJ) (1). Ultrastructural studies have shown that the DEJ separation in BP lesions occurs through the lamina lucida, the electron-lucent region that separates the basal cell plasma membrane from the underlying basal lamina (2, 3). This split is accompanied by an extensive inflammatory infiltrate and destruction of hemidesmosomal and extracellular matrix components (2–4).

One of the main antigenic targets of BP autoantibodies is a 180-kDa transmembrane hemidesmosome-associated glycoprotein designated BP180 (also known as BPAG2 or type XVII collagen; ref. 5–13). The extracellular domain of this protein contains a series of collagen-like triple-helical domains. Structural studies showed that the BP180 ectodomain exists in a multi-meric rod-like conformation (14, 15). BP autoantibodies react with at least 4 distinct antigenic sites on the BP180 ectodomain, all of which are clustered within a 45-amino acid noncollagenous stretch adjacent to the membrane-spanning domain (12, 16).

We have described a mouse model of BP that involves the passive transfer of antibodies directed against mouse BP180 (17). Neonatal BALB/c mice injected with these antibodies develop a blistering skin disease that exhibits all of the key immunopathologic features of BP. Using this animal model, we have shown that the antibody-induced lesion formation is dependent on complement activation (18) and neutrophil infiltration of the upper dermis (19). Blockage of neutrophil recruitment into skin sites resulted in the neutralization of the pathogenic activity of anti-mBP180 (anti-mBP180) antibodies in mice. Proteinases and reactive free radicals from infiltrating inflammatory cells, acting either alone or synergistically, have been implicated as effector molecules contributing to tissue damage in BP lesions (20, 21). Neutrophil granules contain a variety of proteolytic enzymes, including elastase, cathepsin G (CG), collagenase, and gelatinase B (GB), which are known to degrade specific elements of the extracellular matrix (22–24). Upon cell activation, these enzymes are secreted into the pericellular space (22). These and other proteinases, e.g., plasmin and plasminogen activators, have been detected in BP blister fluid and within lesion-
the role of neutrophil elastase in BP blister formation

Methods

Reagents. Human NE, CG, α1-proteinase inhibitor (α1-PI), α1-antichymotrypsin (α1-ACT), and myeloperoxidase (MPO) were from Athens Research and Technology, Inc. (Athens, Georgia, USA). Mouse GB was from Triple Point Biologics (Forest Grove, Oregon, USA). PMSF, 1,10-phenanthroline, chymostatin, DMSO, casein, gelatin, and PMA were obtained from Sigma Chemical Co. (St. Louis, Missouri, USA). Methoxysuccinyl-Ala-Ala-Pro-Val-p-nitroanilide (Met-O-Suc-Ala-Ala-Pro-Val-pNA), methoxysuccinyl-Ala-Ala-Pro-Val-chloromethylketone (MeoSuc-AAPV-CK), succinyl-Ala-Ala-Pro-Phe-p-nitroanilide (Suc-Ala-Ala-Pro-Phe-pNA), and carboxbenzox-Gly-Leu-Phe-chloromethylketone (Z-GLF-CK) were from Enzyme Systems Products Inc. (Dublin, California, USA).

Laboratory animals. Breeding pairs of BALB/c mice were purchased from The Jackson Laboratory (Bar Harbor, Maine, USA) and maintained at the Medical College of Wisconsin Animal Resource Center. NE-null mutants (NE−/−) and the matched control littermates (NE+/+) were generated as described previously by homologous recombination in 129Sv/J embryonic stem (ES) cells and transmitted to germline in 129Sv/J × C57BL/6J background (33). Neonatal mice (24–36 hours old with body weights between 1.4 and 1.6 g) were used for passive transfer experiments.

Preparation of pathogenic anti-BP180 IgG. The preparation of recombinant murine BP180 and the immunization of rabbits were performed as described previously (17). Briefly, a segment of the ectodomain of the murine BP180 antigen (34) was expressed as a glutathione S-transferase (GST) fusion protein using the pGEX prokaryotic expression system (Pharmacia LKB Biotech Inc., Piscataway, New Jersey, USA). The murine BP180 fusion protein, designated GST-mBP180ABC, was purified to homogeneity by affinity chromatography (35). New Zealand white rabbits were immunized with the purified mBP180 fusion protein, and the IgG fraction from the serum (designated R621) was purified as described previously (17). The IgG fractions were concentrated, sterilized by ultrafiltration, and the protein

Table 1

Summary: the role of neutrophil elastase in BP blister formation

<table>
<thead>
<tr>
<th>Host mice</th>
<th>IgG injected</th>
<th>Treatment</th>
<th>No. of mice</th>
<th>Disease activity</th>
</tr>
</thead>
<tbody>
<tr>
<td>BALB/c</td>
<td>R621</td>
<td>α1-PI</td>
<td>24</td>
<td>3+</td>
</tr>
<tr>
<td>NE−/−</td>
<td>R621</td>
<td>α1-ACT</td>
<td>10</td>
<td>–</td>
</tr>
<tr>
<td>NE−/−</td>
<td>R621</td>
<td>MeoSuc-AAPV-CK</td>
<td>10</td>
<td>–</td>
</tr>
<tr>
<td>NE−/−</td>
<td>R621</td>
<td>Z-GLF-CK</td>
<td>10</td>
<td>3+</td>
</tr>
<tr>
<td>NE+/+</td>
<td>R621</td>
<td>–</td>
<td>8</td>
<td>3+</td>
</tr>
<tr>
<td>NE+/+</td>
<td>R621</td>
<td>–</td>
<td>8</td>
<td>–</td>
</tr>
<tr>
<td>NE−/−</td>
<td>R621</td>
<td>5 × 10⁶ NE−/− PMN</td>
<td>4</td>
<td>3+</td>
</tr>
<tr>
<td>NE−/−</td>
<td>R621</td>
<td>5 × 10⁶ NE−/− PMN</td>
<td>4</td>
<td>–</td>
</tr>
<tr>
<td>NE−/−</td>
<td>R621</td>
<td>2.5 × 10⁶ NE−/− PMN</td>
<td>4</td>
<td>–</td>
</tr>
</tbody>
</table>

aNeonatal BALB/c, NE−/−, and the matched control (NE+/+) mice were injected intradermally with either nonpathogenic anti-mBP180 IgG (R50) or pathogenic antibody (R621). A NE inhibitor (α1-PI or MeO Suc-AAPV-CK) or CG/chymase inhibitor (α1-ACT or Z-GLF-CK) was given intradermally 90 minutes after IgG injection or coinjected with IgG. For neutrophil reconstitution, NE−/− mice were given neutrophils intradermally from NE+/+ or NE−/− mice. α1-PI, α1-proteinase inhibitor; α1-ACT, α1-antichymotrypsin; MeoSuc-AAPV-CK, Met-O-Suc-Ala-Ala-Pro-Val-CH₂Cl; Z-GLF-CK, Z-Gly-Leu-Phe-CH₂Cl. bInjected animals were evaluated 12 hours after IgG injection. Disease activity is scored on a scale of “–” (no detectable skin lesions) to “3+” (intense erythema with frank “epidermal detachment” sign). See Methods for details.
concentrations were determined by OD$_{280nm}$ (E [1%, 1 cm] = 13.6). The titers of anti-mBP180 antibodies in both the unfraccionated rabbit serum and in the purified IgG fraction were assayed by indirect immunofluorescence (IF) using mouse skin cryosections as substrate. The antibody preparations were also tested by immunoblotting against the GST-mBP180ABC fusion protein. The IF and immunoblotting techniques have been reported elsewhere (17). The pathogenicity of these IgG preparations was tested by passive transfer experiments as described below. A nonpathogenic anti-mBP180 IgG preparation (designated R50) was used as a control (19).

\textit{Induction of experimental BP and clinical evaluation of animals:} Neonates were given 1 intradermal injection of a sterile solution of either control IgG or anti-BP180 IgG in PBS (50 µL/vol; 2.5 mg IgG/g body weight), as described elsewhere (17). The skin of neonatal mice from the test and control groups were examined 12 or 24 hours after the injection of the IgG fractions. The extent of cutaneous disease was scored as follows: (−), no detectable skin disease; +, mild erythematic reaction with no evidence of the “epidermal detachment sign” (this sign was elicited by gentle friction of the mouse skin, which, when positive, produced fine, persistent wrinkling of the epidermis); 2+, intense erythema and epidermal detachment sign involving 10–50% of the epidermis in localized areas; and 3+, intense erythema with frank epidermal detachment sign involving more than 50% of the epidermis in the injection site.

After clinical examination, the animals were sacrificed, and skin and serum specimens were obtained. The skin samples were used for routine histological examination using light microscopy (hematoxylin and eosin [H&E] staining) and direct IF assays to detect rabbit IgG and mouse C3 deposition at the basement membrane zone (BMZ). Other skin samples were used for enzymatic assays described below. The sera of injected animals were tested by indirect IF techniques to determine the titers of rabbit anti-mBP180 antibodies. Direct and indirect IF studies were performed as described previously (17) using commercially available FITC-conjugated goat anti-rabbit IgG (Kirkegaard & Perry Laboratories Inc., Gaithersburg, Maryland, USA). Monospecific goat anti-mouse C3 serum was purchased from Cappel Research Products (Durham, North Carolina, USA).

\textit{Quantification of PMN accumulation at antibody injection sites.} Tissue MPO activity was used as an indicator of polymorphonuclear neutrophils (PMNs) within skin samples of experimental animals, as described elsewhere (36). A standard reference curve was first established by obtaining activity levels on aliquots of known amounts of purified MPO. The mouse skin samples were extracted by homogenization in a buffer containing 0.1 M TrisCl, pH 7.6, 0.15 M NaCl, 0.5% hexadecyltrimethylammonium bromide. MPO activity levels in supernatant fractions were determined by the change in optical density at 460 nm resulting from decomposition of H$_2$O$_2$ in the presence of o-dianisidine. MPO content was expressed as relative MPO activity (Δ OD$_{460nm}$/mg protein). Protein concentrations were determined by the Bio-Rad (Hercules, California, USA) dye binding assay using BSA as a standard.

\textit{Enzyme assays for NE and CG/chymase.} NE activity in skin and neutrophil extracts of mice was measured using the NE-specific substrate Met-O-Suc-Ala-Ala-Pro-Val-pNA according to Nakajima et al. (37). CG/chymase activity in skin and neutrophil extracts of mice were assayed using the CG/chymase substrate Suc-Ala-Ala-Pro-Phe-pNA, as described by Barrett (38).

\textit{Zymographic analysis.} Proteinase profiles were determined by zymography as described previously (32, 39). Briefly, protein extracts of neutrophils and skin samples from IgG-injected animals were subjected to SDS-PAGE on casein-containing acrylamide gels (12% acrylamide and 1% casein) or gelatin-containing acrylamide gels (8% acrylamide and 1% gelatin) under nonreducing conditions. After electrophoresis, gels were washed twice with 2.5% Triton X-100 for 30 minutes to remove SDS. They were then rinsed briefly with H$_2$O and then incubated overnight at 37°C in reaction buffer (50 mM Tris, pH 7.4, 150 mM NaCl, and 5 mM CaCl$_2$). The gels were stained with 0.125% Coomassie brilliant blue. Caseinolytic and gelatinolytic activity appeared as colorless zones against a blue background.

To characterize the proteinases seen on zymograms, test samples were mixed with a panel of inhibitors or negative control solvents and incubated for 15 minutes in room temperature before electrophoresis. The final concentrations of inhibitors were 3 µg/mL α1-PI (H$_2$O as the solvent), 1 µg/mL MeOSuc-AAPV-CK (DMSO as the solvent), 1 µg/mL Z-GLF-CK (DMSO as the solvent), 1 mM PMSF (isopropanol as the solvent), 5 mM EDTA (H$_2$O as the solvent), 10 mM 1,10-phenanthroline (methanol as solvent), and 0.1 mM chymostatin (DMSO as the solvent). After elec-

![Zymogram Image]

\textbf{Figure 2}

Identification of NE, GB, and CG in the blister fluid of experimental BP. Blister fluids (20 µL/lane) from BALB/c mice injected with pathogenic IgG (lanes 2 and 4) or PBS “washout” (20 µL/lane) from BALB/c mice injected with control IgG (lanes 1 and 3) were analyzed by casein (a) and gelatin (b) gel zymography. NE, (lane 2), CG (lane 2), and GB (lane 4) were present in the blister fluid, but not control samples (lanes 1 and 3).
at the lesional site of the dermis. In contrast, neonatal mice showed a subepidermal vesicle with neutrophilic infiltrate.

Figure 3: Clinical, IF, and histological examination of neonatal NE–/– mice injected with rabbit anti-mBP180 IgG. Pathogenic rabbit anti-mBP180 IgG (intradermal injection, 2.5 mg/g body weight) produced extensive epidermal disease in neonatal NE–/– mice. (a) The skin of these animals showed linear deposition of rabbit IgG (b) and mouse C3 (c) at the BMZ by direct IF. H&E-stained sections from these mice showed a subepidermal vesicle with neutrophilic infiltrate (d). The inset, a higher magnification of d, demonstrates neutrophils at the lesional site of the dermis. In contrast, neonatal NE–/– mice injected intradermally with rabbit anti-mBP180 IgG showed no clinical evidence of skin disease (e). Direct IF studies showed rabbit IgG (f) and mouse C3 (g) deposition at the BMZ, but these animals showed no evidence of subepidermal vesiculation at the light microscopic level (h). The inset, a higher magnification of h, exhibits neutrophils in the dermis. Site of blisters (black arrows in a and e), site of basal keratinocytes (white arrowheads in b, c, f, and g; black arrowheads in d and h), dermis (d), epidermis (e), vesicle (v). Original magnification b–d and f–h: ×100. Inset (original magnification ×400): neutrophils (black arrowheads).

In vitro degradation of recombinant murine BP180 by NE. Protein extracts (55 μg/lane) of lesional skin and nonlesional skin were resolved by 7% SDS-PAGE gel under denaturing conditions and then immunoblotted using rabbit anti-mBP180 IgG.

In vitro treatment of mouse skin in organ culture. Mouse skin sections were obtained from neonatal BALB/c mice (36–48 hours old) and cut into 2 × 2-mm strips with a razor blade. The skin strips were then incubated in MEM either with or without 100 μg/mL human NE at 37°C for various periods of time. At the end of the incubation, the skin strips were rinsed in fresh MEM, fixed in 10% formalin, and embedded in paraffin, after which sections were cut and stained with H&E.

Neutrophil isolation and extraction. Mouse neutrophils were isolated from heparinized blood by dextran sedimentation followed by separation on a density gradient as described (40). The purified neutrophils were homogenized and extracted in the same manner as described for skin samples.

In vitro neutrophil degranulation. In vitro neutrophil degranulation assays were performed as described (41). Purified neutrophils from NE–/– and NE+/+ mice were suspended in HBBS (GIBCO BRL, Grand Island, New York, USA) at a final concentration of 107/mL and triggered with 50 ng/mL PMA in the absence or presence of 5 μg/mL MeOSuc-AAPV-CK or Z-GLF-CK for 15 minutes at 37°C. The cells were then pelleted by centrifugation (1,000 g, 5 minutes), and the supernatant was analyzed by gelatin gel zymography for GB activity and enzyme assays for NE and CG as described above.

Identification of NE, GB, and CG in blister fluids. One hundred microliters of PBS was injected into the skin blisters (formed 12 hours after pathogenic IgG injection) and nonlesional sites, and withdrawn 1 minute later. The “washout” PBS was centrifuged at low speed (1,000 g) for 5 minutes to remove cells and then high speed (12,000 g) for 5 minutes to remove cell debris. The supernatant was analyzed by zymography and enzyme assays as given above.

Identification of BP180 degradation products in blisters. Purified GST-mBP180ABC fusion protein (2 μg, as described above) was incubated with highly purified human NE (0.125 μg) in the in vitro degradation buffer containing 50 mM Tris-Cl, pH 7.5, and 1 mM DTT. Reactions were carried out at 37°C for 0, 0.25, 0.5, 1, or 2 hours and terminated by adding an equal volume of SDS-PAGE sample buffer and heating at 100°C for 5 minutes. Reaction mixtures were then resolved by electrophoresis through 15% SDS-PAGE gels under denaturing conditions. The GST-
mBP180ABC and its degraded fragments were detected using Coomassie blue staining and immunoblotting using rabbit anti-mBP180 IgG.

Statistical analysis. The data were expressed as mean ± SEM and were analyzed using Student’s t test. A P value less than 0.05 was considered significant.

Results
Significantly elevated levels of NE were present in experimental BP lesions and blister fluids. To identify NE, skin samples were analyzed by casein gel zymography. As shown in Figure 1, caseinolytic bands ranging from 24 to 72 kDa were seen in the extracts of lesional/perilesional skin of anti-BP180 IgG-injected mice (Figure 1, lane 3) in addition to the 24-kDa band observed in extracts of control skin samples obtained from mice injected with either PBS (Figure 1, lane 1) or normal rabbit IgG (Figure 1, lane 2). The caseinolytic activity associated with these bands was inhibited by PMSF (Figure 1, lane 8), a general serine proteinase inhibitor, but not by the metalloproteinase inhibitor, EDTA (Figure 1, lane 6). The 29-kDa band comigrated with murine NE (Figure 1, lanes 4 and 7). Inhibition experiments confirmed the activity as NE, because its activity could be completely blocked by the natural NE inhibitor, α1-PI (Figure 1, lane 11) and by the synthetic peptide NE inhibitor, MeOSuc-AAPV-CK (Figure 1, lane 12), but not by the synthetic peptide CG/chymase inhibitor, Z-GLF-CK (Figure 1, lane 13). The 24-kDa band, which was found in the skin of both anti-mBP180 IgG-injected and control mice, was identified as a CG/chymase-like proteinase, because its activity could be blocked by Z-GLF-CK (Figure 1, lane 13), but not by MeOSuc-AAPV-CK (Figure 1, lane 11). These results demonstrate that elevated levels of NE are associated with lesional/perilesional tissue in experimental BP.

For NE to be relevant pathologically, NE would need to be released from neutrophils in the skin inflammatory site. To evaluate this, we determined the presence of NE in the skin blister fluid. Significantly high levels of NE activity were detected in the blister fluid of IgG-injected NE+/+ mice (Figure 2a). Similarly, CG (Figure 2a) and GB (Figure 2b) were also present in the blister fluid. These results demonstrate that NE is released from infiltrating neutrophils at lesional sites.

Elastase-deficient mice are resistant to the pathogenic effects of anti-BP180 antibodies. To determine directly whether NE is involved in subepidermal blister formation in experimental BP, passive transfer experiments were performed on the following 3 groups of mice: NE+/+ mice, matched control littermates (NE−/−), and positive control BALB/c mice. Mice from these 3 groups were injected intradermally with identical doses of pathogenic anti-BP180 IgG (2.5 mg/g body weight). As expected, the NE+/+ (n = 8) (Figure 3a; Table 1) and BALB/c (n = 9) mice developed extensive blisters within 12 hours after injection with anti-BP180 IgG (Table 1). The skin of these animals was markedly erythematous and developed persistent epidermal bullous lesions. Direct IF analysis of lesional/perilesional skin of the mice showed in vivo deposition of rabbit IgG (Figure 3b) and mouse C3 (Figure 3c) at the BMZ. H&E-stained skin sections from these mice showed DEJ separation with a neutrophil infiltration (Figure 3d). In contrast, the NE−/− (n = 8) mice (Figure 3e) exhibited no blisters 12 hours after injection with anti-BP180 IgG. Indirect IF demonstrated that the titers of circulating rabbit anti-BP180 IgG in NE−/− mice were comparable to those of the control mice (average titers were greater than 1:5260 for all injected mice). Direct IF of the skin of NE−/− mice also showed deposition of rabbit IgG (Figure 3f) and mouse C3 (Figure 3g) at the BMZ. Histologic examination of the mice skins showed no signs of dermal-epidermal detachment, but did reveal a neutrophil infiltrate in the upper dermis (Figure 3h). This inflammatory infiltrate was less prominent than the infiltrates in the skin of NE+/+ mice.

Quantification of neutrophil infiltration by the MPO assay showed a significant difference between NE−/− mice (0.41 ± 0.06 OD460nm/mg protein) and the control groups (1.27 ± 0.21 OD460nm/mg protein for NE+/+ mice; 1.06 ± 0.14 OD460nm/mg protein for NE+/+ mice). The MPO and NE activities (mean ± SEM) of skin extracts from NE−/− mice injected intradermally with pathogenic rabbit anti-mBP180 IgG. Neonatal NE−/− (bars 1–3) and NE+/+ (bars 4–6) mice received 2.5 mg/g body weight anti-mBP180 IgG. MPO (a) and NE (b) activities in skin in the injection sites were assayed 4 hours (bars 2 and 5 in a; bars 1 and 3 in b) and 12 hours (bars 1, 3, 4, and 6 in a; bars 2 and 4 in b) after IgG administration. n = 8 for NE−/− and NE+/+ groups. *P < 0.01; **P < 0.001. Student t test for paired samples (bar 2 vs. 5, bar 3 vs. 6 in a; bar 1 vs. 3, bar 2 vs. 4 in b).
In vitro neutrophil degranulation. Neutrophils were purified from NE+/+ (lane 1) and NE−/− (lanes 2 and 3) mice and triggered at a final concentration of 10⁷ neutrophils/mL with 50 ng/mL PMA in the absence (lanes 1 and 2) or presence (lane 3) of 5 μg/mL MeOSuc-AAPV-CK (a synthetic peptide NE inhibitor) for 15 minutes at 37°C. Cell-free releases (equivalent to 10⁷ neutrophils) from triggered neutrophils were analyzed by casein (a) and gelatin (b) zymography. Upon stimulation with PMA, NE−/− (lane 1), NE+/+ (lane 2), and inhibitor-treated NE−/− (lane 3) neutrophils released almost the same levels of GB and CG. NE−/− neutrophils also secreted NE, whereas NE+/+ neutrophils did not. These results show that NE−/− and inhibitor-treated NE−/− neutrophils respond normally to PMA stimulation as determined by releasing GB and CG.

BALB/c mice; P < 0.01, 12 hours after injection (Figure 4a). In contrast, at the 4-hour time point, there were no significant differences in tissue MPO activity (Figure 4a) and CG/chymase activity (data not shown) between the NE+/+ and control mice, indicating that the absence of NE activity did not interfere with the early stages of neutrophil migration from the circulation into the tissue inflammatory site.

As expected, NE activity levels (expressed as relative OD₄⁶₀nm reading/min per milligram of protein) were not above background in the antibody-treated NE−/− mice, but were significantly elevated in normal control mice both at 4 hours and 12 hours after injection of anti-BP180 IgG (Figure 4b). At 4 hours after IgG injection, the relative mean NE activity levels were: 0.35 ± 0.03 (BALB/c), 0.31 ± 0.04 (NE+/+), and 0.11 ± 0.01 (NE−/−; P < 0.01) OD₄⁶₀nm/min per milligram of protein. The relative mean NE levels at the 12-hour time point were: 1.47 ± 0.15 (BALB/c), 1.26 ± 0.18 (NE+/+), and 0.14 ± 0.02 (NE−/−; P < 0.001) OD₄⁶₀nm/min per milligram of protein. These results suggest that NE plays a direct role in subepidermal blistering in experimental BP. The results of in vitro neutrophil degranulation experiments (see below) lend further support to this conclusion.

NE inhibitors block blister formation in the BP mouse model. If NE contributes directly to DEJ separation in experimental BP, then blocking NE activity should inhibit the disease phenotype. Thus, neonatal BALB/c mice were given injections of pathogenic anti-mBP180 IgG intradermally and 90 minutes later were injected at the same site with NE inhibitors (α1-PI or MeOSUC-AAPV-CK) or CG/chymase inhibitors (α1-ACT or Z-GLF-CK). Control mice (receiving either PBS or DMSO; n = 5) developed intense blisters 12 hours after antibody administration (Table 1). Mice coinfected with anti-mBP180 IgG and CG/chymase inhibitors (n = 5) also developed subepidermal blisters (Table 1). In contrast, neonatal mice that were treated with R621 IgG followed by either α1-PI or MeOSuc-AAPV-CK (n = 5 for each group) did not develop skin lesions (Table 1). The skin of these animals showed no evidence of subepidermal vesiculation, despite the presence of high levels of circulating rabbit anti-BP180 antibodies, in situ deposition of rabbit IgG and murine C3 at the BMZ, and a neutrophil infiltrate in the upper dermis (data not shown). Coinjecting the NE inhibitor (α1-PI or MeOSuc-AAPV-CK) with the pathogenic anti-BP180 IgG also yielded inhibition of subepidermal blistering (data not shown).

Neutrophil infiltration into the antibody injection sites of the animals listed above was quantified using the MPO assay. A significant difference in extractable MPO activity levels between these groups became apparent 12 hours after injection. The relative mean MPO level for mice treated with both anti-BP180 IgG and an elastase inhibitor, α1-PI, was 0.48 ± 0.04 OD₄⁶₀nm/mg protein, whereas the MPO level for the control groups was 0.90 ± 0.10 OD₄⁶₀nm/mg protein (P < 0.01). To rule out the possibility that the reduction in MPO activity in NE inhibitor-treated mice is due to the direct inhibition of neutrophil recruitment by the inhibitors, MPO activities of skin extracts in the injection sites were analyzed 4 hours after IgG injection. No change in tissue MPO activity was observed in either NE inhibitor–treated or control mice 4 hours after injection of anti-BP180 IgG (0.25 ± 0.03 for NE inhibitor–treated mice versus 0.29 ± 0.03 for control). The higher MPO values at 12 hours after mice were injected with pathogenic IgG alone are due to the development of tissue injury in the DEJ causing additional neutrophil recruitment between 4 and 12 hours after injection.

The experiments discussed above demonstrate that anti-BP180 antibodies are not capable of inducing a BP-like skin disease in NE−/− mice and in BALB/c mice treated with NE inhibitors, suggesting that NE plays a key role in the pathogenic mechanism of experimental BP. The possibility remained, however, that the release and/or activity of other neutrophil products, e.g., GB and CG, are adversely affected in the NE−/− and inhibitor-treated animals. To address this important issue, we performed in vitro neutrophil degranulation experiments following the protocol by...
Desrochers et al. (41). Upon stimulation with PMA, NE–/– (Figure 5, lane 1), NE+/+ (Figure 5, lane 2), and inhibitor-treated NE+/+ (Figure 5, lane 3) neutrophils released similar levels of CG and GB, as determined by zymography. NE was found to be released by NE+/+ (Figure 5a, lane 2), but not NE–/– (Figure 5a, lane 1) and inhibitor-treated NE+/+ (Figure 5a, lane 3) neutrophils. These results demonstrate that NE–/– and NE inhibitor–treated NE+/+ neutrophils show the normal stimulation response.

NE is required for BP180 degradation in the lesional skin of experimental BP and produces dermal-epidermal separation in a mouse skin organ culture system. The appearance of NE in the skin correlated with subepidermal blistering in response to pathogenic IgG, thus suggesting that NE may cleave BP180 and other extracellular matrix proteinase directly. To test this hypothesis, protein extracts from lesional and nonlesional skin of IgG-injected NE+/+ and NE–/– mice were analyzed by immunoblotting using rabbit anti-mBP180 IgG. As shown in Figure 6, both full-length (180 kDa) and degraded (100 kDa) mBP180 were identified in lesional skin of IgG-injected NE+/+ mice (Figure 6, lane 2), whereas the nonlesional skin extracts of NE–/– mice injected with pathogenic IgG (Figure 6, lane 3) and NE+/+ mice injected with control IgG R50 (Figure 6, lane 1) showed only intact mBP180.

Tissue damage caused by exogenous NE was also assessed using an in vitro system. Neonatal mouse skin sections (2 × 2 mm) were incubated with purified human NE (100 μg/mL in MEM) and examined histologically at various times over a 24-hour period (Figure 7). The NE-treated skin sections showed signs of early DEJ separation at 6 hours (Figure 7b) and broad subepidermal vesiculation at 12 hours (Figure 7c). Complete detachment of the epidermis from the dermis was observed at 24 hours (Figure 7d). The control mouse skin incubated with MEM alone did not show any signs of DEJ separation at any of the time points tested (Figure 7e). The NE-induced dermal-epidermal detachment was completely blocked by the NE inhibitors MeOSuc-AAPV-CK (Figure 7f) or α1-PI (not shown) and by coinubcation with a 5-fold molar excess of purified GST-mBP180ABC fusion protein (Figure 7g). CG/chymase inhibitors (Z-GLF-CK or α1-ACT) had no effect on the NE-induced tissue changes (Figure 7h). These results demonstrate that NE can produce DEJ separation of mouse skin.

NE degrades the ectodomain of recombinant murine BP180 antigen in vitro. To determine whether NE is capable of directly degrading the BP180 antigen, we set up in vitro degradation assays using a 42-kDa recombinant fusion protein, GST-mBP180ABC (Figure 8a) as a substrate for NE. As shown in Figure 8b, when recombinant GST-mBP180ABC was incubated with NE, the 42-kDa fusion protein was degraded into 3 distinct smaller fragments, with molecular weights of 34, 29, and 26 kDa as...
determined by SDS-PAGE (Figure 8b, lanes 2–5). No degradation products were seen in the reaction sample containing no NE (Figure 8b, lane 6) or containing NE at 0 time (Figure 8b, lane 1). Recombinant GST was resistant to NE digestion (Figure 8b, lane 7), indicating that the GST moiety of the GST-mBP180ABC was not a target of NE. To further characterize the NE degradation products, the digestion mixtures described above were subjected to immunoblot analysis using an anti-mBP180 antiserum. The anti-mBP180ABC antibodies labeled the 3 bands that had been detected by Coomassie blue staining and, in addition, labeled 3 additional bands of 18, 15, and 9 kDa (Figure 8b, lanes 9–11). This immunoblot staining pattern was not affected by preadsorption of the anti-BP180 antibody with a 5-fold molar excess of GST-mBP180ABC fusion protein, but not with GST alone (data not shown).

Pathogenic anti-mBP180 IgG induces subepidermal blistering in NE−/− mice reconstituted with neutrophils from NE+/+ but not NE−/− mice. Our data show that NE produces DEJ separation and cleaves BP180 in vitro. However, incubation of human skin sections with human CG also produces DEJ separation (42), and GB also degrades the human recombinant BP180 ectodomain in an in vitro system (31). In addition, although in the early stage (at 4 hours after IgG injection) there is no difference in the number of infiltrating neutrophils between NE+/+ and NE−/− mice, recruitment of neutrophils at the 12-hour time point is reduced in NE−/− mice (Figure 4a). Thus, resistance of NE−/− mice to experimental BP could be due to reduced levels of GB and/or CG. To test this possibility NE−/− mice were injected with pathogenic IgG and reconstituted with neutrophils from either NE+/+ or NE−/− mice. Twelve hours after IgG injection, the NE+/+ mice (n = 4) reconstituted with 5 × 10^5 NE+/+ neutrophils developed subepidermal blisters (Figure 9a, panels A and B) and cleaved BP180 antigen was detected in the lesional skin

Figure 8
In vitro degradation of mBP180ABC by neutrophil elastase. (a) The schematic diagram is a structural representation of mBP180. The vertical black bar designates the transmembrane domain. The COOH-terminal extracellular region is made up of 13 collagen triple-helical (open bar) and 14 noncollagenous (filled bar) domains. The purified recombinant GST-mBP180ABC fusion protein is 42 kDa, containing 26 kDa GST and a 145-amino acid fragment of the mBP180 extracellular domain. The mBP180ABC is made up of an 8-kDa noncollagenous NC14A (filled bar) and an 8-kDa collagenous region (open bar). One pathogenic (oval) and 2 nonpathogenic epitopes (circles) are located within the mBP180ABC (see ref. 51). (b) The GST-mBP180ABC fusion protein (lanes 1–6, 8–11) or GST alone (lane 7) was incubated with (lanes 1–5, 7–11) or without (lane 6) NE at 37°C for 0 minutes (lanes 1 and 8), 0.25 hours (lanes 2 and 9), 0.5 hours (lanes 3 and 10), 1 hour (lanes 4 and 11), or 2 hours (lanes 5–7). The products were then resolved by SDS-PAGE and detected by Coomassie blue staining (lanes 1–7) or immunoblot (lanes 8–11). Fragments of 34 kDa, 29 kDa, and 26 kDa were generated from the cleavage of the fusion protein by elastase (lanes 2–5 and 9–11). No degradation products were seen after incubation without NE (lane 6). GST was not degraded by NE (lane 7). Immunoblot using anti-mBP180ABC IgG identified 3 additional fragments with molecular weights of approximately 18 kDa, 15 kDa, and 9 kDa (lanes 9–11). The immunoreactivity of these bands with anti-mBP180ABC was abolished by preadsorption of the antibody with a 5-fold molar excess of GST-mBP180ABC fusion protein, but not with GST alone (data not shown).
extracts (Figure 9b, lane 2). In contrast, NE+/− mice (n = 4) reconstituted with either 5 × 10⁶ or 2.5 × 10⁶ NE−/− neutrophils showed no sign of blistering (Figure 9a, panels C and D) and only intact BP180 antigen was seen in the skin extracts (Figure 9b, lanes 1 and 3). These results clearly demonstrate that NE plays a major role in subepidermal vesiculation in experimental BP.

**Discussion**

NE has been implicated in numerous inflammatory diseases (22, 43–47). Using NE-null mutant mice, we now provide evidence that NE plays a destructive role in experimental BP, an inflammatory skin blistering disease. Our data show (a) abnormally high levels of NE in the lesional/perilesional skin and blister fluid of experimental BP mice; (b) NE−/− mice are resistant to experimental BP; (c) NE+/− mice reconstituted with NE−/− neutrophils become susceptible to experimental BP; (d) NE inhibitors (1-PI and AAPV) block subepidermal blistering in experimental BP; (e) NE degrades BP180 and extracellular matrix components of the BMZ, causing DEJ separation in the skin culture system; and (f) NE cleaves the recombinant mBP180. These results suggest that NE is directly involved in the disruption of the DEJ leading to subepidermal vesiculation in experimental BP.

We also rule out the possibility that the protective effects observed in NE−/− mice are only transient in nature because the protection continues to be observed up to 48 hours after IgG injection (data not shown).

The BP180 ectodomain exists in an extended trimeric conformation consisting of a series of articulated collagen-like rigid rods (14, 15) spanning the lamina lucida and inserting into the lamina densa of the basement membrane (13, 48). It has been hypothesized that the BP180 ectodomain interacts with one or more components of the extracellular matrix, thereby stabilizing anchorage of the basal keratinocyte. Consistent with this hypothesis are the recent findings that mutations within the BP180 gene lead to the inherited blistering disease, generalized atrophic benign epidermolysis bullosa (49). NE degrades collagens (50–53), fibronectin (54), and proteoglycans (55, 56). The results of our in vitro enzyme digestion experiments demonstrate that NE also cleaves the extracellular domain of the BP180 protein. Thus, it is reasonable to hypothesize that degradation of cell-matrix adhesion molecules, including BP180, by NE and/or other proteolytic enzymes is an important step leading to the detachment of the basal keratinocytes from the basement membrane.

In previous studies we have shown that neutrophil migration to sites of infection was not impaired in NE−/− mice (33). The present results further demonstrate that NE deficiency does not impair transvascular migration of neutrophils into the skin triggered by pathogenic anti-mBP180 antibodies. There was no difference in neutrophil infiltration between NE−/− mice and wild-type control mice 4 hours after the injection of pathogenic IgG, when subepidermal blistering has already begun in control mice. We hypothesize that NE-mediated degradation product(s) are chemotactic for neutrophils, which would account for the higher levels of neutrophil accumulation in the later stages of the disease process in control mice compared with NE-deficient mice. This is similar to findings in macrophage elastase-deficient mice (MME−/−) that fail to accumulate macrophages in response to cigarette smoke (57). In this condition there is evidence that MME cleavage of elastin is responsible for monocyte chemotaxis (S.D. Shapiro, unpublished data).

We recently found that GB-deficient mice are also resistant to the pathogenic activity of anti-mBP180 IgG (32). GB is present in BP lesional skin and blister fluid (31). GB cleaves a number of extracellular matrix proteins (58–62), α1-PI (58), and recombinant BP180 (31). GB could contribute to subepidermal blistering in BP directly by cleaving structural proteins in the
DEJ or indirectly by inactivating α1-PI, the principal NE inhibitor. CG also produces DEJ separation when incubated with skin sections (42). Therefore, the DEJ separation triggered by pathogenic anti-BP180 IgG could be due to the proteolytic activity of NE alone or in concert with GB, CG, and/or other proteolytic enzymes. In fact, we found that neutrophil extracts from NE−/− mice incubated with mouse skin sections produce DEJ separation and cleave the recombinant mouse BP180 ectodomain. These data indicate that other proteinases may participate in these events, at least in vitro (data not shown). The exact relationship between NE, GB, CG, and other proteinases in experimental BP is under investigation.

Despite the striking similarities in the immunopathological features of the experimental model of BP and the human disease, there is one conspicuous difference. The human disease is characterized by an inflammatory infiltrate predominated by eosinophils (4), whereas in the mouse model, the neutrophil is the major inflammatory cell type at the lesion site (17). This discrepancy might signify a key difference in the pathogenic mechanisms operating in humans and mice. Whereas it is clear that the eosinophil plays only a secondary role, if any, in subepidermal blistering in the mouse model, it might be the eosinophil, rather than the neutrophil, that plays an essential role in the initiation and/or progression of human BP. However, a causal link between the presence of eosinophils and subepidermal blistering in humans with BP has not been established. On the contrary, several observations implicate neutrophils in human BP. First, neutrophil-predominant forms of BP have been described in which eosinophils are absent in the inflammatory cell infiltrate, suggesting that eosinophils are not essential for blister formation (63). Second, neutrophils were an essential component in achieving dermal-epidermal detachment in an in vitro model of BP described by Gammon et al. (64). In this model system, after incubation of cultured human skin with a BP serum, complement, and peripheral blood leukocytes, neutrophils lined up along the BMZ and breakdown of dermal-epidermal cohesion occurred. Further studies of inflammatory cells in human BP are necessary to assess the fidelity with which the BP mouse model mimics the human disease.

In summary, NE activity is essential for the generation of subepidermal blisters in experimental BP (see Table 1). It is presumed that this enzyme, acting alone or with other effector molecules, carries out this pathogenic role in BP by degrading components of the BMZ that function in maintaining dermal-epidermal cohesion. Degradation of one such adhesion-related molecule, BP180, appears to be a key in the pathogenic cascade leading to blister formation; i.e., the extracellular domain of this transmembrane protein is the primary antigenic target of pathogenic antibodies and is a substrate for NE. These findings are important in terms of the molecular pathogenesis of BP and point to new directions for the development of therapies for BP.

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