



# Mast cells are critical for controlling the bacterial burden and the healing of infected wounds

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**Skin wound infections are a significant health problem, and antibiotic resistance is on the rise. Mast cells (MCs) have been shown to contribute to host–defense responses in certain bacterial infections, but their role in skin wound superinfection is unknown. We subjected 2 MC-deficient mouse strains to *Pseudomonas aeruginosa* skin wound infection and found significantly delayed wound closure in infected skin wounds. This delay was associated with impaired bacterial clearance in the absence of MCs. Engraftment of MCs restored both bacterial clearance and wound closure. Bacterial killing was dependent on IL-6 released from MCs, and engraftment with IL-6–deficient MCs failed to control wound infection. Treatment with recombinant IL-6 enhanced bacterial killing and resulted in the control of wound infection and normal wound healing in vivo. Taken together, our results demonstrate a defense mechanism for boosting host innate immune responses, namely effects of MC-derived IL-6 on antimicrobial functions of keratinocytes.**

mast cells | skin wound infection | bacterial infection | innate immunity | *pseudomonas aeruginosa*

**B**acterial skin wound infection is a serious health risk in which a wound fails to heal, leading to increased patient distress, prolonged hospital stays, and increased mortality and health care costs (1, 2). *Pseudomonas aeruginosa*, a Gram-negative bacterium, is the second most common cause of nosocomial skin wound infections and is associated with high morbidity and mortality (3). *P. aeruginosa* presents a rising therapeutic challenge due to its ability to form biofilms and increasing antibiotic resistance (4). Therefore, effective treatment, combined with rapid wound closure, is essential to restrain such infections.

Mast cells (MCs) are key effector cells of the innate immune system (5, 6). MCs have been shown to be involved in the defense of bacterial infections by releasing soluble factors that recruit or activate immune cells such as neutrophils (7), dendritic cells, and T cells. They also can contribute to antibacterial defense by acting as phagocytes, releasing antimicrobial peptides, and forming extracellular traps (8). By contrast, the role of MCs in skin wound healing is controversial. Whereas some studies indicate that MCs are essential in certain models of wound healing (9–12), others found that skin wound closure may not be affected by the absence of MCs (13–15).

However, there have been no reports investigating the potential roles of MCs during skin wound infection. Therefore, we addressed this clinically relevant topic and identified, and characterized in detail, a mechanism by which MCs can control bacterial skin wound infections and promote wound healing.

## Results

**MC-Deficient Mice but Not WT Mice Exhibit Impaired Skin Wound Healing in Response to Skin Wound Infection with *P. aeruginosa*.** To investigate the role of MCs in a physiological model of skin wound infection, skin wounds of MC-deficient and WT mice were inoculated with *P. aeruginosa* mimicking a naturally occurring infection route. Wound closure was evaluated by lesion size over time. As early

as 1 d after skin wound infection, MC-deficient *Kit<sup>W</sup>/Kit<sup>W-v</sup>* mice exhibited increased wound size compared to infected WT mice, indicative of impaired wound closure (Fig. 1A). This difference in wound size between *Kit<sup>W</sup>/Kit<sup>W-v</sup>* and WT mice was most pronounced between day 2 and day 4 after wound infection with an average of  $111 \pm 9\%$  of the initial wound size in MC-deficient mice as compared with  $61 \pm 5\%$  in WT mice (Fig. 1A). By contrast, wound closure in noninfected mice did not differ in size between the 2 genotypes, indicating that MCs are dispensable for normal closure of noninfected wounds in this model.

To determine whether the impaired wound healing was only associated with the absence of MCs in *Kit<sup>W</sup>/Kit<sup>W-v</sup>* mice, some *Kit<sup>W</sup>/Kit<sup>W-v</sup>* mice were reconstituted with bone marrow-derived cultured MCs (BMCMCs). Local adoptive transfer of BMCMCs to the skin of *Kit<sup>W</sup>/Kit<sup>W-v</sup>* mice 4 wk before wounding restored MC numbers (SI Appendix, Fig. S6A) and normalized wound closure of infected skin wounds (Fig. 1B). The wound closure kinetics of *Kit<sup>W</sup>/Kit<sup>W-v</sup>* + BMCMCs mice closely resembled those of WT mice (Fig. 1B). This demonstrates that MCs are required for normal healing of skin wounds infected with *P. aeruginosa*.

We further confirmed our finding that the MC deficiency, but not MC-independent defects of the *Kit<sup>W</sup>/Kit<sup>W-v</sup>* mouse model, was responsible for the delayed wound closure by testing a non-c-Kit–dependent model of MC deficiency, the *Cpa3-Cre; Mcl-1<sup>fl/fl</sup>* mouse (Fig. 1C and SI Appendix, Fig. S6B). Upon skin wound infection with *P. aeruginosa*, MC-deficient *Cpa3-Cre; Mcl-1<sup>fl/fl</sup>* mice exhibited larger wounds compared with WT mice, peaking at day 3 with a difference of  $40 \pm 13\%$  between the genotypes. By contrast, no significant differences in wound closure kinetics was noted in the sham-infected mice.

**MC-Deficient Mice Show Higher Bacterial Counts Compared to WT Mice in Response to Topical Wound Infection with *P. aeruginosa*.** To identify the mechanisms of impaired healing of *P. aeruginosa*-infected skin wounds, we analyzed the skin bacterial load during the course of

## Significance

**Mast cells are key effector cells of the innate immune system and have been shown to orchestrate host–defense responses against a variety of pathogens. In this report, we demonstrate the importance of mast cells for the control of bacterial wound infections, by which mast cell-derived IL-6 turns keratinocytes into better bacterial killers.**

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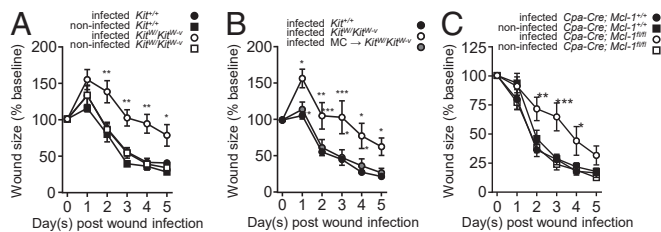
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**Fig. 1.** MCs are required for normal healing of *P. aeruginosa*-infected wounds. MC-deficient mice show delayed wound closure after local *P. aeruginosa* skin wound infection. (A–C) Wound closure kinetics of infected or noninfected PBS-treated skin wounds of *Kit*<sup>+/+</sup> (*n* = 9 PBS, *n* = 7 infected) mice and *Kit*<sup>W/Kit</sup><sup>W-v</sup> mice (*n* = 5 PBS, *n* = 5 infected) (A); *Kit*<sup>+/+</sup> mice (*n* = 9), *Kit*<sup>W/Kit</sup><sup>W-v</sup> mice (*n* = 7), and *Kit*<sup>W/Kit</sup><sup>W-v</sup> mice engrafted intradermally with bone marrow-derived cultured MCs (MCs → *Kit*<sup>W/Kit</sup><sup>W-v</sup>) (*n* = 8) (B), and MC-deficient *Cpa3-Cre; Mcl-1*<sup>fl/fl</sup> mice (*n* = 6 PBS, *n* = 9 infected) compared to *Cpa3-Cre; Mcl-1*<sup>fl/fl</sup> mice (*n* = 7 PBS, *n* = 8 infected) (C). Data are expressed as mean ± SEM and pooled from 3 independent experiments each, analyzed using 2-way repeated measure ANOVA followed by Bonferroni post hoc test. \**P* < 0.05; \*\**P* < 0.01; \*\*\**P* < 0.001; n.s., not significant.

infection. Using *in vivo* live imaging of fluorescently labeled bacteria, the bacterial load within skin wounds was monitored over 5 d following wound infection. *Kit*<sup>W/Kit</sup><sup>W-v</sup> mice exhibited a 14-fold higher bacterial load in skin wounds than the corresponding WT mice on day 1 after skin wound infection (Fig. 2*A* and *B*). Although bacterial numbers decreased in both genotypes over time, *Kit*<sup>W/Kit</sup><sup>W-v</sup> mice showed a 20-fold higher bacterial load at day 5. To ascertain whether the increased bacterial load consisted of viable bacteria, we conducted plate counts at day 1 after skin wound infection. Lysates of infected skin wounds of *Kit*<sup>W/Kit</sup><sup>W-v</sup> mice contained 7 times more *P. aeruginosa* compared with the WT mice. To assess whether this antibacterial effect required MCs, intradermal reconstitution of *Kit*<sup>W/Kit</sup><sup>W-v</sup> with MCs was carried out, which resulted in a reduced bacterial load to approximately 2-fold WT levels (Fig. 2*C*). In accord with these findings, MC-deficient *Cpa3-Cre; Mcl-1*<sup>fl/fl</sup> mice exhibited a 5-fold increase in bacterial load compared with WT mice (Fig. 2*D*).

Next, we analyzed whether this antibacterial effect of MCs is dependent on immune cell recruitment. For this purpose, 4-mm skin biopsies derived from *Kit*<sup>W/Kit</sup><sup>W-v</sup> vs. WT mice were infected *ex vivo* with *P. aeruginosa*. Infection of skin explants derived from *Kit*<sup>W/Kit</sup><sup>W-v</sup> mice showed 12 times reduced bacterial clearance compared to WT skin explants (Fig. 2*E*), suggesting that MC-mediated bacterial clearance in this setting is not dependent on immune cell recruitment. Impaired bacterial elimination in the absence of MCs was also detected for Gram-positive *Staphylococcus aureus* (Fig. 2*F*). Supporting the evidence that reduced neutrophil recruitment in *Kit*<sup>W/Kit</sup><sup>W-v</sup> mice was not the underlying cause of the impaired bacterial clearance, we analyzed neutrophil numbers and myeloperoxidase (MPO) activity within skin wounds upon infection. In both MC-deficient and WT mice, neutrophil numbers as well as MPO activity (a marker of neutrophil activity) were increased but not different between the genotypes (Fig. 2*G* and *H*). These findings point toward an important role of MCs in bacterial clearance that appears to be independent of immune cell recruitment and the type of bacteria.

**Antimicrobial Response to *P. aeruginosa* Infection Depends on MC–Keratinocyte Interaction.** As our results pointed to a MC-mediated antibacterial effect, we investigated the direct antibacterial effects of MCs *in vitro*. However, MCs infected with *P. aeruginosa* were not able to reduce numbers of bacteria (Fig. 3*A*). Therefore, we hypothesized that MCs might stimulate epidermal cells to elicit the antimicrobial activity. We cocultured mouse MCs together with mouse keratinocytes (KCs) and analyzed the effects on *P. aeruginosa*. Upon their infection with *P. aeruginosa*, an increase of 56 ± 10% in the antibacterial effect within cocultures of MCs and KCs was

observed compared with single cultures of MCs or KCs (Fig. 3*A*). This antibacterial effect was amplified by increasing the KC/MC ratio of the coculture (Fig. 3*B*), supporting the conclusion that the antimicrobial effects were KC-derived.

Since KCs are a major source of antimicrobial peptides (AMPs), we analyzed the production and release of AMPs by KCs. Upon infection of the coculture, KCs increased expression of the mBD-14 gene (*Defb14*) 4-fold as well as CRAMP (*Camp*) 2-fold. The expression level of Psoriasin (*S100A7*), however, remained unchanged (Fig. 3*C*). To confirm that these AMPs are in fact secreted, we measured protein levels of mBD-14 and CRAMP. Analysis of the supernatants of previously infected MC–KCs cocultures by enzyme-linked immunosorbent assay (ELISA) allowed for the detection of mBD-14 and CRAMP protein. While mBD-14 showed a moderate increase in protein levels (of 385 ± 118 pg/mL) a significant release of CRAMP (2,140 ± 348 pg/mL) was observed (Fig. 3*D*). Moreover, we analyzed AMP expression after skin wound infection in the MC-deficient mouse model. Upon *P. aeruginosa* skin wound infection, MC-deficient *Kit*<sup>W/Kit</sup><sup>W-v</sup> mice showed consistently lower levels of mBD-14, CRAMP, and Psoriasin compared to WT or MC reconstituted mice at day 1 after skin wound infection (Fig. 3*E*).

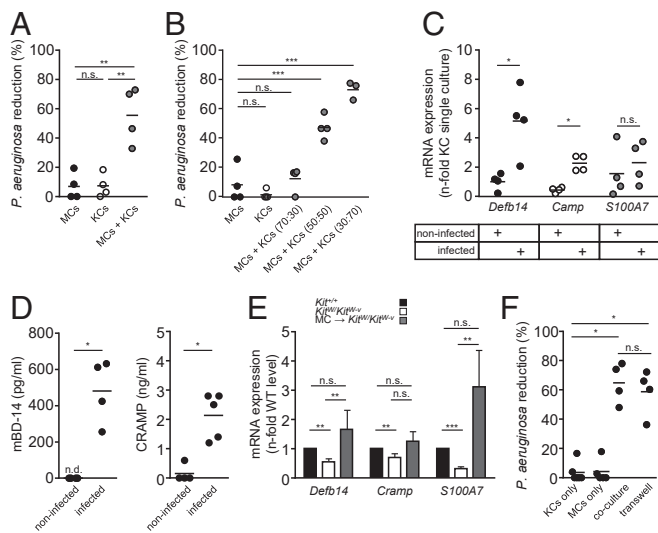
We next determined whether the collaborative antibacterial activity of MCs and KCs required direct cell contact or was mediated by a soluble factor. We used transwell inserts to separate MCs and KCs and found that the transwell inserts did not hinder the bacterial reduction ability of the coculture. We concluded that direct cell-cell contact between the 2 cell types was not essential for the observed antibacterial effect (Fig. 3*F*). These findings suggest that MCs protect mice from *P. aeruginosa* skin wound infection by secreting mediators that can induce antibacterial effects of KCs.

**MC-Derived IL-6 Is Required for a KC Antimicrobial Response.** The antibacterial effect of KCs upon infection with *P. aeruginosa* only became apparent when they were cocultured with MCs. To identify the MC mediators involved in the induction of antibacterial KC effects, we analyzed the supernatants of infected MC–KC cocultures and found high levels of interleukin (IL)-6 (*SI Appendix, Fig. S1A*). To identify the source of IL-6 secretion, we performed expression analysis of IL-6 mRNA transcripts from MCs and KCs derived from infected cocultures. IL-6 mRNA expression levels were 36-fold higher in MCs as compared with KCs, suggesting that MCs, but not KCs, predominantly produce and release IL-6 when cocultures are infected with *P. aeruginosa* (*SI Appendix, Fig. S1B*). Notably, IL-6 production by MCs was dependent on coculture with KCs, indicating that KC-derived signals are required to induce IL-6 production by MCs (*SI Appendix, Fig. S1B*). Further characterization of such KC-derived signals suggests that the release of IL-1 family members by KCs infected with PA is able to stimulate IL-6 production in MCs by a MyD88-dependent pathway (*SI Appendix, Fig. S2A–F*), thereby augmenting bacterial clearance (*SI Appendix, Fig. S2G*). Although IL-33 stimulation of MCs results in high levels of IL-6 (*SI Appendix, Fig. S2E*), bacterial clearance and the release of IL-6 by MCs seems to be independent from their expression of the IL-33/ST2 receptor (*SI Appendix, Fig. S2H and J*).

Next, we investigated the importance of MC-derived IL-6 for the collaborative antimicrobial effect of MC–KC cocultures using MCs derived from bone marrow of IL-6-deficient mice. Cocultures of *Il6*<sup>-/-</sup> MCs and KCs, in contrast to cocultures with WT MCs, were unable to reduce numbers of *P. aeruginosa* in MC–KC cocultures (*SI Appendix, Fig. S1C*). Moreover, we analyzed the AMP mRNA expression levels of KCs in MC–KC cocultures containing *Il6*<sup>-/-</sup> MCs and found no up-regulation of AMP expression in KCs upon infection with *P. aeruginosa* compared with cocultures containing WT MCs (*SI Appendix, Fig. S1D*).

To provide proof of concept that IL-6 is required for an optimal defense response from KCs upon infection, we stimulated





**Fig. 3.** *P. aeruginosa* reduction involves MC-KC interaction. Reduction of *P. aeruginosa* numbers in vitro requires MC-KC interaction. (A and B) *P. aeruginosa* CFU from cell culture supernatant after 3 h in vitro infection of either KCs or MCs alone compared with MC-KC cocultures (A) or MC-KC cocultures with different cell ratios (B). Each data point represents an individual experiment, data are expressed as means, pooled from 4 independent experiments and analyzed using one-way ANOVA followed by Tukey's post hoc test. (C) mRNA of AMPs expressed in KCs during coculture with MCs with or without infection with *P. aeruginosa*. Values are normalized to those of the internal control gene beta-actin and represented as fold change relative to that of noninfected KC controls, which was converted to 1. Each data point represents an individual experiment, data are expressed as means, pooled from 4 independent experiments and analyzed by Student's *t* test. (D) Protein levels of secreted AMPs measured in supernatants of infected or noninfected MC-KC cocultures. Each data point represents an individual experiment, data are expressed as means, pooled from 4 independent experiments and analyzed by Student's *t* test (mBD-14) and Mann-Whitney *U* test for CRAMP. (E) AMP expression analysis of skin wound samples 24 h after skin wound infection in *Kit*<sup>+/+</sup> mice, *Kit*<sup>W</sup>/*Kit*<sup>W-v</sup> (*n* = 7) mice, and *Kit*<sup>W</sup>/*Kit*<sup>W-v</sup> mice, which had been intradermally engrafted with MCs (MCs → *Kit*<sup>W</sup>/*Kit*<sup>W-v</sup>) (*n* = 5). Values are normalized to those of the internal control gene beta-actin and represented as fold change relative to *Kit*<sup>+/+</sup> mice, which was converted to 1. (F) *P. aeruginosa* CFU from culture supernatant of MC-KC cocultures and transwell cultures. Each data point represents an individual experiment, data are expressed as means, pooled from 4 independent experiments and analyzed by Mann-Whitney *U* test. Data are shown as means ± SEM and analyzed with Mann-Whitney *U* test. \**P* < 0.05; \*\**P* < 0.01; \*\*\**P* < 0.001; n.s., not significant; n.d., not detectable.

of WT MCs restored IL-6 levels in *P. aeruginosa*-infected skin wounds of *Kit*<sup>W</sup>/*Kit*<sup>W-v</sup> mice to WT levels. In contrast, reconstitution with *Il6*<sup>-/-</sup> MCs resulted in 4-fold lower levels of IL-6 within the *P. aeruginosa*-infected skin wounds, comparable to the IL-6 levels detected in MC-deficient *Kit*<sup>W</sup>/*Kit*<sup>W-v</sup> mice (SI Appendix, Fig. S3A). This indicated that the increase of IL-6 levels in *P. aeruginosa*-infected skin wounds depends on IL-6 released by MCs.

We next determined the bacterial load in *P. aeruginosa*-infected wounds of MC-deficient mice and MC-deficient mice that had received IL-6-deficient or WT MCs. Compared to MC-deficient *Kit*<sup>W</sup>/*Kit*<sup>W-v</sup> mice, WT mice and MC-deficient *Kit*<sup>W</sup>/*Kit*<sup>W-v</sup> mice reconstituted with WT MCs exhibited a similar reduction in bacterial numbers (SI Appendix, Fig. S3B). In contrast, adoptive transfer of MCs lacking IL-6 resulted in a 14-fold higher bacterial burden within the skin wounds compared with WT mice (SI Appendix, Fig. S3B). The bacterial load of MC-deficient *Kit*<sup>W</sup>/*Kit*<sup>W-v</sup> reconstituted with *Il6*<sup>-/-</sup> MCs was comparable to that in MC-deficient *Kit*<sup>W</sup>/*Kit*<sup>W-v</sup> mice (SI Appendix, Fig. S3B).

Furthermore, we investigated the impact of MC-derived IL-6 on skin wound closure upon *P. aeruginosa* infection. In contrast

to the adoptive transfer of WT MCs, MC-deficient *Kit*<sup>W</sup>/*Kit*<sup>W-v</sup> mice reconstituted with *Il6*<sup>-/-</sup> MCs exhibited larger wounds compared to WT mice. Reconstitution with *Il6*<sup>-/-</sup> MCs did not lead to an improvement in wound healing compared with nonreconstituted MC-deficient mice at any time point during wound healing (SI Appendix, Fig. S3C). To further support the importance of MC-derived IL-6 during *P. aeruginosa* skin wound infection, we generated a MC-specific *Il-6* knockout mouse model (*Cpa-Cre; Il-6*<sup>fl/fl</sup>). Subjecting these mice to skin wound infection resulted in significantly impaired wound healing, which confirmed, in mice with normal c-Kit, the relevance of MC-derived IL-6 for normal healing of infected skin wounds (SI Appendix, Fig. S3D).

**Recombinant IL-6 Treatment Stimulates the Antibacterial Response in Mice and Humans.** We next tested whether the topical application of recombinant IL-6 (recIL-6) can compensate for MC deficiency in the promotion of antibacterial responses to wound infection with *P. aeruginosa*. We showed that pretreatment with IL-6 improved wound closure of infected skin wounds in MC-deficient *Kit*<sup>W</sup>/*Kit*<sup>W-v</sup> mice and normalized wound healing to WT levels (SI Appendix, Fig. S4A). By contrast, vehicle pretreatment of MC-deficient mice did not improve wound closure of infected skin wounds. However, IL-6 pretreatment of WT mice did not further improve the closure of infected skin wounds. Moreover, pretreatment of MC-deficient mice with recIL-6 also significantly reduced the bacterial load of skin wounds, as compared with vehicle-treated littermates (SI Appendix, Fig. S4B). RecIL-6 pretreatment normalized the bacterial load of skin wounds in *Kit*<sup>W</sup>/*Kit*<sup>W-v</sup> mice to 1.4-fold that of WT mice. Interestingly, recIL-6 treatment also reduced the bacterial load of skin wounds of MC-competent WT mice 4-fold compared with vehicle-treated WT mice.

Since pretreatment with recIL-6 significantly improved wound healing of MC-deficient mice, and also reduced bacteria within skin wounds of both MC-deficient and WT mice, we aimed to translate these findings into the human system. Therefore, we performed in vitro infection experiments using human KCs. We observed a 13-fold increase in the reduction of *P. aeruginosa* after recIL-6 pretreatment compared to vehicle-treated controls (SI Appendix, Fig. S4C). Furthermore, we could show that IL-6 pretreatment also improves host-defense responses in human skin explants as compared with vehicle-treated controls (SI Appendix, Fig. S4D). Taken together, these data demonstrate that recIL-6 pretreatment can induce antibacterial effects in mice and human KCs, as well as in human skin explants.

## Discussion

In this study, we demonstrate a previously unreported role of MCs in the control of skin wound infection and identify MC-derived IL-6 as a crucial driver of an antibacterial innate immune response in infected skin wounds (SI Appendix, Fig. S5). IL-6 has been previously implicated in wound healing, and increased IL-6 levels have been reported in different models of mouse and human skin injury (16–19). The role of MCs in defense against pathogens also has been well established in several infection models (8). In response to various pathogens, MCs have been shown to exhibit an important function in facilitating the recruitment of other immune cells, i.e., neutrophils, macrophages, and T cells, to the site of infection (20–24). Interestingly, in the model of bacterial skin wound infection reported herein, the control of bacterial burden appeared to be independent from the influx of neutrophils (Fig. 2 G and H) but was mediated by MC-derived IL-6-driven production and release of antimicrobial peptides by KCs (Fig. 3 C–E). In addition, this effect was not limited to a specific pathogen, since both Gram-negative *P. aeruginosa* and Gram-positive *S. aureus* provoked a similar response (Fig. 2 E and F).

In a recent study, MC-derived IL-6 was also found to augment host defense to herpes simplex virus (HSV) infection. HSV was

not able to directly induce the production of IL-6 by MCs but did so by an IL-33–dependent mechanism, most likely initiated by infected KCs. Blocking the IL-33 receptor T1/ST2 on BMCMCs significantly reduced IL-6 production by BMCMCs (25). Notably, we also observed an early increase of IL-33 in MC-KC cocultures infected with *P. aeruginosa*. However, IL-6 release was not reduced in T1/ST2<sup>-/-</sup> MCs stimulated with the coculture supernatant. This may be explained by the fact that in addition to IL-33, other members of the IL-1 family, i.e., IL-1 $\alpha/\beta$  and IL-36, were simultaneously released, suggesting a biologically redundant response compensating for the lack of IL-33 signaling via the T1/ST2 receptor (*SI Appendix, Fig. S2*).

The relevance of MCs in wound healing is still regarded as controversial. Our group has previously shown that the healing of skin wounds is delayed in the absence of MCs (10). As demonstrated by the results of the current study, this effect was likely due to the control of the bacterial burden on skin wounds, since we found that the healing of noninfected wounds was not delayed in either Kit-dependent or Kit-independent MC-deficient mice. Our finding that MCs are not significantly involved in the closure of noninfected wounds is in accord with several other studies investigating wound healing in Kit-independent MC-deficient mouse models (13–15).

IL-6 previously has been described to augment the production of antimicrobial peptides in a skin graft model (26). We, therefore, hypothesized that MC-derived IL-6 may be crucial for the induction of antimicrobial peptides in KCs. In fact, we were able to show that MC-derived IL-6 induces antimicrobial peptide production and enhances the antibacterial capacity of KCs in vitro and in vivo.

IL-6 has also been reported to be expressed in human wounds (27). Interestingly, patients treated with the humanized neutralizing anti-IL6 receptor antibody Tocilizumab exhibit a noticeable increase in the frequency of skin and s.c. infections (28). In our study, the ability of IL-6 to improve bacterial clearance was also not limited to the mouse system as treatment of human KCs and human skin explants with recombinant human IL-6 significantly

reduced bacterial numbers in the supernatant (*SI Appendix, Fig. S4 C and D*).

In conclusion, our findings show that MC-derived IL-6 is crucial for the control of skin wound infections and for the healing of infected wounds. These results may allow for the development of novel and effective approaches for the prevention and treatment of bacterial skin, and other, infections, which are urgently needed in this time of increasing antibiotic resistance.

## Materials and Methods

**Animals.** Mice were obtained from breeding colonies of the animal facilities of Charité–Universitätsmedizin Berlin and all animal experimentation were approved by the ethics committee of the state of Berlin, Landesamt für Gesundheit und Soziales (LAGeSo). Where indicated, engraftment of back skin was performed by intradermal injection of 10<sup>6</sup> BMCMCs (29, 30). Full details can be found in *SI Appendix, SI Materials and Methods*.

**Model of Skin Wound Infection with *P. aeruginosa*.** Skin wounds were created on the lower back using a 6-mm biopsy punch as described previously (10). Full details can be found in *SI Appendix, SI Materials and Methods*.

**In Vitro Bacterial Infection Assays.** Pam212 mouse KCs, human HaCaT cells, and human skin explants were infected with *P. aeruginosa*. For IL-6 treatment: 50 ng/mL (or as indicated) reIL-6 was added 1 h prior to infection. Full details can be found in *SI Appendix, SI Materials and Methods*.

**Quantitative Real-Time PCR, ELISA, and MPO Assay.** Full details can be found in *SI Appendix, SI Materials and Methods*.

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