

Filamentous Bacteriophage Delay Healing of Pseudomonas-Infected Wounds

Michelle S. Bach^{1,2#}, Christiaan R. de Vries^{1#}, Johanna M. Sweere^{1,2}, Medeea Popescu^{1,2}, Jonas D. Van Belleghem¹, Gernot Kaber¹, Elizabeth B. Burgener³, Dan Liu¹, Quynh-Lam Tran¹, Tejas Dharmaraj¹, Maria Birukova^{1,2}, Vivekananda Sunkari^{1,2}, Swathi Balaji⁴, Sundeep Keswani⁴, Niaz Banaei⁵, Dolly K. Khona⁶, Laurence Nedelec¹, Chandan K. Sen⁶, Venita Chandra⁷, Patrick R. Secor⁸, Gina A. Suh,^{9†} and Paul L. Bollyky^{1†*}

Affiliations:

¹ Division of Infectious Diseases and Geographic Medicine, Department of Medicine, Stanford University, Stanford, CA

² Stanford Immunology, Stanford University, Stanford, CA.

³ Center for Excellence in Pulmonary Biology, Department of Pediatrics, Stanford University, Stanford, CA

⁴ Division of Pediatric Surgery, Department of Surgery, Baylor College of Medicine, Houston, TX

⁵ Division of Pathology, Department of Medicine, Stanford University, Stanford, CA

⁶ Department of Surgery, Indiana University, Indianapolis, IN

⁷ Department of Vascular Surgery, Stanford University, Stanford, CA

⁸ University of Montana, Missoula, MT

⁹ Division of Infectious Diseases, Mayo Clinic, Rochester, MN

Contributed equally to the work

† These authors share senior authorship.

* To whom correspondence should be addressed: pbollyky@stanford.edu. Twitter: @bollyky

Abstract

We have identified a novel role for filamentous bacteriophage in the delayed healing associated with chronic *Pseudomonas aeruginosa* (*Pa*) wound infections. In a mouse model of chronic *Pa*- infected wounds, Pf, a filamentous phage produced by *Pa*, impaired keratinocyte migration, prevented wound re-epithelialization, and delayed healing in both the absence and presence of live bacteria. Mechanistically, the immune response to Pf phage produces soluble factors that impair keratinocyte migration and delay wound re-epithelialization. In a prospective cohort study of 113 human patients, *Pa* was detected in 36 patients and 25 of these (69%) were positive for Pf phage. Pf(+) wounds were significantly older and more likely to increase in size over time than Pf(-) wounds. Together, these data implicate Pf in the delayed wound healing associated with *Pa* infection. We propose that Pf phage may have potential as a biomarker and therapeutic target for delayed wound healing.

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Introduction

Chronic wounds are associated with extensive human suffering and massive economic costs (Järbrink et al., 2016; Kirsner, 2016). Over 6.5 million Americans have chronic wounds, and their care is estimated to cost \$25 billion annually (Kirsner, 2016). Bacterial infections frequently complicate chronic wounds, leading to delayed wound healing (Guo and Dipietro, 2010), increased rates of amputation (Gardner and Frantz, 2008), and extensive morbidity and mortality (Bowler, 2002; Clinton and Carter, 2015; Scali and Kunimoto, 2013).

Wounds infected with the Gram-negative pathogen *Pseudomonas aeruginosa* (*Pa*) are characterized by poor outcomes (Dowd et al., 2008; James et al., 2008; Kirketerp-Møller et al., 2008; Malic et al., 2009). *Pa* is prevalent in infections of burns (Tredget et al., 2004), diabetic ulcers, and post-surgical sites (Percival et al., 2015). The presence of *Pa* is associated with delayed wound closure in both humans as well as animal models (Bryers, 2008; Guo and Dipietro, 2010; Watters et al., 2013; Zhao et al., 2012), and wounds infected with *Pa* tend to be larger than those in which *Pa* is not detected (Gjødsbøl et al., 2006; Jesaitis et al., 2003). This wound chronicity is associated with ineffective bacterial clearance (Sen et al., 2009) and delayed wound healing (Bjarnsholt et al., 2008). In addition to wound infections, *Pa* is a major human pathogen in other contexts (Bryers, 2008; Høiby et al., 2011; Sen et al., 2009) due to increased antibiotic resistance incidence. In recognition of the magnitude of this problem, *Pa* was recently listed as a critical priority pathogen by the World Health Organization (WHO) and the Centers for Disease Control (CDC) (Centers for Disease Control and, 2013; Tacconelli et al., 2018). Therefore, there is great interest in identifying novel biomarkers, virulence factors, and therapeutic targets associated with *Pa* infections.

One such potential target is Pf bacteriophage (Pf), a filamentous virus produced by *Pa* (Dowd et al., 2008; Malic et al., 2009). Unlike lytic bacteriophages used in phage therapy (Gordillo Altamirano and Barr, 2019; Górski et al., 2019), Pf typically does not lyse its bacterial hosts. Instead, Pf phages integrate into the bacterial chromosome as prophage and can be produced without destroying their bacterial hosts (Rakonjac et al., 2011; Roux et al., 2019). Indeed, the production of phage virions is partially under bacterial control (Castang and Dove, 2012). We and others have reported that Pf phage contribute to *Pa* fitness by

serving as structural elements in *Pa* biofilms (Secor et al., 2015b) and contributing to bacterial phenotypes associated with chronic *Pa* infection, including aggregation (Secor et al., 2018) and reduced motility (Secor et al., 2017).

Pf phage also impact host immunity in ways that promote chronic infection. We recently reported that Pf phage directly alter cytokine production and decrease phagocytosis by macrophages (Secor et al., 2017; Sweere et al., 2019b). These effects were associated with endocytosis of Pf by mammalian cells and triggering of maladaptive anti-viral immune responses that antagonize anti-bacterial immunity. Consistent with these effects, *Pa* strains that produced Pf phage (Pf(+) strains) were more likely than strains that do not produce Pf (Pf(-) strains) to establish wound infections in mice (Sweere et al., 2019b). These and other studies implicate Pf phage as a virulence factor in *Pa* infections (Burgener et al., 2019; Rice et al., 2009; Secor et al., 2015b).

We hypothesized that Pf phage might contribute to the delayed wound healing often associated with *Pa* wound infections. To test this, we first examined the effects of Pf phage in a well-established *in vitro* wound healing model (a scratch assay). We then examined the effects of Pf phage on wound healing in an *in vivo* mouse model in the absence or presence of live *Pa* (Sweere et al., 2019a; Sweere et al., 2019b). Finally, we examined the association between Pf phage and delayed wound healing in a cohort of 113 patients seen at the Stanford Advanced Wound Care Center.

Results

Infection with a Pf(+) strain of *Pa* causes worse morbidity in a murine chronic wound infection model.

We first examined the impact of a Pf phage-positive (Pf+) strain of *Pa* (PAO1) versus a Pf phage-negative (Pf-) isogenic strain (PAO1ΔPf4) on morbidity and mortality in a delayed inoculation murine wound infection model in which a provisional wound matrix is allowed to form for 24 hours after wounding before introduction of bacteria. We previously pioneered this model to study the initial establishment of *Pa* wound infections in otherwise healthy C57BL/6 mice in the absence of foreign material, diabetes, or other forms of immune suppression (de Vries et al.; Sweere et al., 2019a; Sweere et al., 2019b). Here, we have adapted this model to study wound healing in the setting of infection over a two-week time period. In brief, we generated bilateral, full-thickness, excisional wounds on the dorsum of mice. These wounds were then inoculated 24 hours later with 1×10^7 /mL PAO1, PAO1ΔPf4, or PBS as a control. We then assessed wound size daily through day 13 post-*Pa* inoculation (day 14 post-wounding). A schematic of this protocol is shown in **Figure 1A**.

All mice survived the 14-day experiment (**Figure S1**) and all wounds were obviously infected on Day 3 by gross appearance (**Figure 1B**), regardless of whether they received PAO1 or PAO1ΔPf4. However, wounds in the PAO1 group were characterized by diffuse inflammation and disintegration of the epithelium of the wounded area. In contrast, PAO1ΔPf4 wounds were characterized by less extensive inflammation and redness limited to the area around the wounded area (**Figure 1B**). Moreover, mice inoculated with PAO1 lost significant weight compared to mice that received PAO1ΔPf4 or PBS (**Figure 1C**). These data implicate Pf phage in worse morbidity in this model.

Pf phage delay wound healing.

We next examined the effect of Pf phage on wound healing in our *in vivo* model. Healing was defined as the remaining open wound area as a fraction of the original wound area. Inoculation with PAO1ΔPf4 led to significantly improved wound healing over time compared to inoculation with PAO1. Indeed, wound healing in many of the mice inoculated with PAO1 appeared “stuck” and mice failed to resolve their wounds

(**Figure 1D**). By Day 13, most of the mice from the PAO1 Δ Pf4 group resolved their infections and healed their wounds completely, whereas most wounds infected with PAO1 persisted (**Figure 1E**). Similar results were observed when area under curve (AUC), a measure of cumulative change in wound area, was considered (**Figure 1F**).

We next examined wound histology from tissue sections collected on Day 13 post-wounding that were collected from these mice. We observed that wounds infected with PAO1 Δ Pf4 exhibited healthy granulation tissue (**Figure 1G**) whereas wounds infected with PAO1 had substantial necrotic debris, minimal granulation tissue, and obvious infection (**Figure 1H**). When the epithelial gap was measured, wounds infected with PAO1 were found to have significantly less re-epithelialization by day 13 compared to PAO1 Δ Pf4 (**Figure 1I**).

Taken together, these data indicate that the production of Pf phage by *Pa* is associated with increased morbidity, ongoing infection, impaired wound healing, and delayed re-epithelialization in this model.

Pf phage delay re-epithelialization in the absence of bacterial infection.

Because the aforementioned data were collected in the setting of infection, it was difficult to know what to attribute to ongoing infection versus direct effect of Pf on wound healing. To assess this directly, we treated mice with purified Pf phage in the absence of bacterial infection. In particular, we administered heat killed PAO1 (HK-*Pa*) or PBS to C57BL6 mice one day after wounding in the absence or presence of supplemental Pf4 phage. Day 7 rather than Day 13 was used in the studies in Figure 2 because wounds healed more rapidly in the absence of live bacterial infections. Moreover, because in the absence of infection mouse wounds heal with contracture we used a standard splinting procedure that incorporates silicone rings to prevent this (Wang et al., 2013). A schematic detailing this protocol is shown in **Figure 2A**. Both the wound area on Day 1 (**Figure 2B**) and the epithelial gap on Day 7 post-wounding (**Figure 2C**) were assessed morphometrically. Representative area demarcations are shown inset into these images.

We observed that supplementation with Pf4 phage was associated with a larger epithelial gap on Day 7 when normalized to Day 1 wound area (**Figure 2D**). However, overall inflammatory cell counts were not altered by Pf phage supplementation (**Figure S2**). We conclude that supplementation with Pf phage reduced re-epithelialization of murine wounds in the absence of infection.

Pf phage enhance *in vitro* *Pa* biofilm formation.

We next investigated the effects of Pf phage on biofilm structure, given the importance of this in chronic wounds and delayed healing (Zhao et al., 2013). We seeded PAO1 and PAO1ΔPf4 on polycarbonate membrane discs and stained them with Sypro Ruby (biofilm matrix stain) and Syto9 (cell stain). Confocal laser scanning microscopy showed significantly increased biofilm thickness of PAO1 compared to PAO1ΔPf4 (**Figure S3A-B**). Supporting this result, microtiter plate crystal violet assay for biofilm attachment showed a significantly increased absorbance at 24 hours in PAO1 compared to PAO1ΔPf4 (**Figure S3C**). Finally, scanning electron microscopy (SEM) of *Pa* cultures on membrane discs showed enhanced biofilm formation in PAO1 compared to PAO1ΔPf4 (**Figure S3D**). Together, these data indicate that Pf phage promote formation of more robust biofilms.

Supernatant from macrophages exposed to Pf phage impair *in vitro* keratinocyte migration

We next asked whether Pf phage inhibit cell migration in a scratch assay—an *in vitro* model commonly used to study mechanisms related to wound healing (Liang et al., 2007). A schematic of this protocol is shown with representative examples of keratinocyte migration over 24 hours (**Figure 3A**).

Migration of HaCaT keratinocytes was not affected by purified Pf phage (**Figure 3B**). Although a few (~1%) HaCaT cells take up AlexaFluor 488-labeled Pf phage (**Figure 3C**) this is a negligible amount compared to the relatively large numbers of monocytes and macrophages that we previously reported take up Pf both *in vitro* and within the wound bed (Sweere et al., 2019b).

Given our previous reports that internalization of Pf phage suppresses production of TNF α made in response to LPS (Sweere et al., 2019b) as well as a number of other chemokines and cytokines (Secor et

al., 2017), we hypothesized that changes in the wound inflammatory milieu might indirectly impact keratinocyte migration and thus wound healing. In support of this model, we found that supernatant from U937 macrophages stimulated with Pf and LPS slowed migration of HaCaT keratinocytes compared to supernatant from keratinocytes cultured in supernatant from mock controls (**Figure 3D-E**). U937 macrophages stimulated with Pf with or without LPS showed increased uptake of Pf compared to controls (**Figure 3F**). These data point to an indirect role for Pf phage in inhibition of keratinocyte migration. Consistent with this, we previously reported that macrophages as well as a number of other cell types take up Pf phage within the wound bed (Sweere et al., 2019b).

Pf phage alter macrophage cytokine production profiles.

To better understand how macrophage internalization of Pf phage impacts cytokine and chemokine production we performed a Luminex assay on primary human monocytes stimulated with Pf phage and LPS. We find that murine bone marrow-derived macrophages (BMDMs) take up Pf phage (**Figure 4A**) and that this leads to altered production of multiple cytokines and chemokines, including TNF (**Figure 4B**) as well as MIG (CXCL9), SDF1a (CXCL12), NGF (nerve growth factor), and IL-10 (**Figure 4C**).

We then attempted to overcome the inhibitory effects of Pf phage on keratinocyte migration and wound healing in mice via supplementation with several factors shown here or in our previous studies to be suppressed by Pf phage, including TNF, CXCL1, or IL-10 (Secor et al., 2017; Sweere et al., 2019b). However, these agents did not overcome the effect of Pf or restore normal wound healing in this model (data not shown).

Together these data suggest that Pf phage alter the inflammatory milieu in ways that impact keratinocyte migration and re-epithelialization. Further, these effects are either multi-factorial or involve cytokines and chemokines other than those directly tested here.

Pf phage are abundant in human wounds infected with *P. aeruginosa*

In light of the decreased wound healing observed in the presence of Pf4 phage in our *in vivo* model, we investigated the clinical impact of Pf phage on human chronic *Pa* wound infections. A total of 113 consecutive patients referred to the Infectious Disease service at the Stanford Advanced Wound Care Center (AWCC) were enrolled from June 2016 to June 2018. Our protocol for screening and classifying these patients is described in **Figure S4**. Patient characteristics are described in **Table 1**. Three *Pa*-positive patients (one who was Pf(+), and two who were Pf(-)) were excluded from the final analysis due to amputation, exposed bone/tendon, or loss to follow up (**Figure S4, S5**). Microbiological characteristics of the *Pa*-positive wound isolates are described in **Table 2**. Representative images from *Pa*-infected Pf(+) and Pf(-) wounds are shown in **Figure 5A,B**. Additional representative images are shown in **Figure S6**.

We found that samples from 39 unique patients out of 113 were *Pa*-positive (35%) (**Figure 5C**). Of the *Pa*-positive patients included in our analysis, Pf prophage (integrated into the bacterial chromosome) was detected in 69% (25/36) of the wounds. Pf phage levels ranged from 3.55×10^3 - 2.69×10^8 copies/swab, with a mean of 2.16×10^7 copies/swab in Pf(+) samples. Subjects in the Pf(-) subset were older compared to the Pf(+) subset (61.6 +/- 14.7 versus 76.6 +/- 14.5; p=0.008; **Table 1**). Higher numbers of Pf(+) patients were on antimicrobial treatments compared to Pf(-) patients, though this was not statistically significant (56% vs 18%; p=0.067; **Table 2**). There were no significant correlations between Pf phage status and gender, BMI, recurrence of infection, race/ethnicity, co-morbidities, antimicrobial resistance, or co-infection with other bacterial or fungal species (**Table 2**). This excludes co-infection as a confounding variable and highlights the correlation between Pf phage status and clinical outcomes.

Pf phage are associated with chronic *Pa* wound infections

We assessed whether the absence or presence of Pf phage was associated with chronic or non-healing human wound infections, as characterized by wound age. Chronic wounds are defined as lasting longer than 3 months in duration without complete healing (32). We therefore asked whether chronic wounds were more likely to be infected with *Pa* strains producing Pf phage. At the end of the study period, all wounds were classified into groups based on the age of the wound, with cutoffs set at ≤ 3 months, ≤ 6

months, ≤ 9 months, ≤ 12 months, or > 12 months. They were then further classified as Pf(+) vs Pf(-). We found that 0% (0/3) of wounds ≤ 3 months were Pf(+); 25% (2/8) wounds ≤ 6 months of age were Pf(+); 41.7% (5/12) of wounds ≤ 9 months of age were Pf(+); 50% (7/14) of wounds ≤ 12 months of age were Pf(+); and 81.8% (18/22) of wounds > 12 months of age were Pf(+) ($n = 36$, $p=0.007$ by Chi-square test) (**Figure 5D**). Wounds infected with Pf(+) strains of *Pa* were significantly older than Pf(-) strains (mean age 3.8 years vs. 2.0 years, respectively; median age 2.1 years vs 0.5 years, respectively; $p=0.046$ by Mann Whitney test) (**Figure 5E**). Moreover, a survival analysis using the endpoint of wound closure showed a significant difference between time to wound closure of the Pf(+) wounds vs Pf(-) wounds ($p = 0.013$ by Gehan-Breslow-Wilcoxon test) (**Figure 5F**). Together, these data demonstrate that Pf phage production is associated with clinically-defined chronic wounds.

Pf phage are associated with wound progression

Given the association between the presence of Pf phage and chronic / non-healing wounds, we next asked whether patients infected with Pf(+) *Pa* strains indeed have impaired wound healing compared to patients infected with Pf(-) *Pa* strains. To this end, we measured changes in wound dimensions over time in the 36 *Pa*-positive patients, starting with measurements taken at the patient's initial clinic visit. We found that Pf(+) patients were more likely to experience an increase in wound size over time than patients who were Pf(-) (8/25 versus 0/11, respectively; $p = 0.033$ by two-sided Chi square test) (**Figure 5G-H**). These data indicate that Pf(+) strains of *Pa* are associated with delayed wound healing.

Discussion

We report that Pf phage promote chronic wounds and impaired healing in mice. Pf phage delayed healing in these animals in both the absence or presence of live *Pa*, indicative of effects on wound healing independent of effects on chronic infection. Further, in a prospective cohort study in humans, Pf phage were associated with more chronic wound infections and wound size progression. Together, these data strongly implicate Pf phage in chronic wounds and delayed wound healing.

These effects of Pf phage on delayed healing were associated with impaired re-epithelialization. Re-epithelialization is critical for wound closure and its failure leads wounds open to reinjury, re-infection, and perpetuation of a chronic wound (Barrientos et al., 2008; Pastar et al., 2014; Sivamani et al., 2007). Our data are consistent with reports that *Pa* infections can cause impaired epithelialization (Roy et al., 2014) although this was not previously attributed to Pf phage.

Our data implicate crosstalk between immune cells and keratinocytes in these effects. Conditioned media from Pf-treated monocytes prevented keratinocyte migration *in vitro*. However, direct treatment of keratinocytes with Pf did not, indicating that Pf effects on re-epithelialization are indirect. We previously reported that Pf phage impact production of multiple cytokines and chemokines (Secor et al., 2017; Sweere et al., 2019b; Van Belleghem et al., 2017; Van Belleghem et al., 2018). We have identified multiple factors in this study as well, including several known to have effects on keratinocyte migration and function (Balaji et al., 2015; Banno et al., 2004). In addition to chemokines, other soluble factors present in the wound environment also impact wound healing and might also contribute to the effects reported here (Tu et al., 2019; Williams et al., 2018). Together, these data point to Pf internalization by immune cells present in the wound bed, alternations in the inflammatory milieu, and indirect effects on keratinocyte migration leading to delayed re-epithelialization. However, we cannot exclude potential roles for other cell types within the wound bed.

These data are consistent with other reports implicating Pf phage in the pathogenesis of *Pa* infections. Pf contributes to *Pa* virulence in lung infections in mice (Rice et al., 2009; Secor et al., 2015b) and in humans (Burgener et al., 2019). We ourselves previously reported that Pf phage contribute to biofilm

formation (Secor et al., 2015a; Secor et al., 2015b) and promote bacterial phenotypes associated with chronic infections (Secor et al., 2017). This work also builds on our recent report that Pf phage promote the establishment of wound infections (Sweere et al., 2019b). However, while that study focused on bacterial clearance and initial wound infections, the present work focuses on re-epithelialization and wound resolution.

It is notable that the incidence of *Pa* in our patient sample (35%) is higher than what is typically reported in other cohorts (Panagea and Galloway, 1998). However, because our patients represent a random cross section of individuals referred to the Infectious Disease service, they are likely to include sicker, more complicated patients than might be seen in a general medicine service. It will be important to assess the prevalence of Pf phage in *Pa* infections in other contexts and at other sites.

A restriction of our study is its limited prospective cohort design, which does not permit us to follow all patients, especially those with more chronic wounds, for longer than two years. Larger longitudinal studies will allow us to confirm our findings. Since we did not find differences in antibiotic resistance or co-infection profiles between Pf(+) and Pf(-) *Pa*-infected wounds, we suspect that these factors do not serve as confounding variables. Lastly, chronic wound infection is a complex disease characterized by multiple factors that contribute to disease severity and progression, which can only be controlled for through longitudinal studies with larger samples. Our ongoing work aims to further investigate Pf phage pathogenicity through a larger prospective longitudinal study.

We conclude that Pf phage may provide a novel biomarker and therapeutic target for the delayed wound healing that occurs in the context of *Pa* wound infections. This finding may have particular relevance given our recent report that a vaccine targeting Pf could prevent initial *Pa* wound infection (Sweere et al., 2019b). Future longitudinal studies in independent cohorts are needed to validate these studies.

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Author Contributions

M.S.B. contributed to the study design, conducted experiments, acquired and analyzed data, and prepared the manuscript. C.D.V. provided methods, conducted experiments, analyzed data, and revised the manuscript. J.M.S. provided methods and assisted in conducting experiments, data analysis, and manuscript revisions. E.B.B. provided methods and assisted in data analysis. S.B. and S.K. provided methods and assisted in conducting experiments and data analysis. M.B. and V.S. assisted in conducting experiments and data analysis. P.L.B. and G.A.S. provided oversight in study design, conducting experiments, data analysis, and manuscript revision.

Declaration of Interests

The authors declare no competing interests.

Figures and Tables

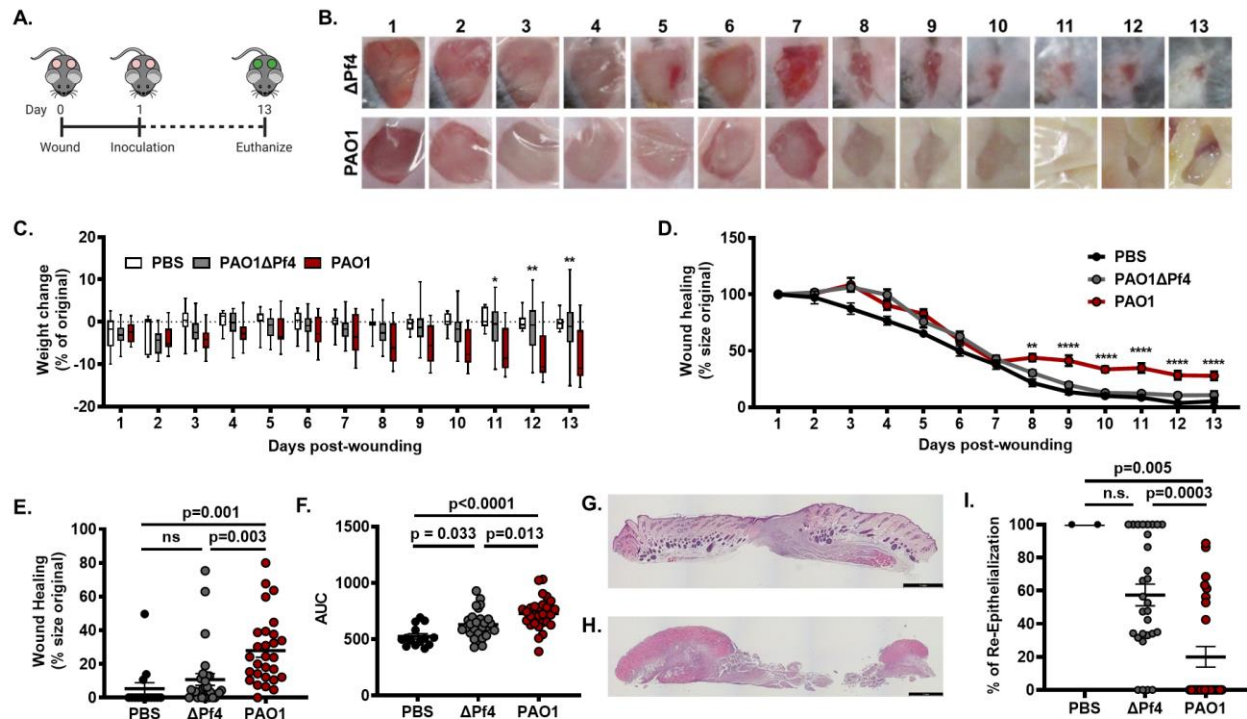


Figure 1. Pf phage lead to decreased wound healing in murine models.

A) Schematic of the full thickness delayed-inoculation chronic *Pa* wound infection murine model. B) Images of representative mice from PAO Δ Pf4 and PAO1 groups from Day 1-13 post-wounding. C) Weight loss of mice across all days post-wounding (For PAO1 vs PAO1 Δ Pf4, Day 11: $p < 0.05$; Day 12: $p < 0.01$; Day 13: $p < 0.01$; two-way ANOVA). D) Wound healing rates for all wounds across all days from Day 1-13 post-wounding. $n=28$ wounds in PAO Δ Pf4 and PAO1 groups; $n=14$ wounds in PBS group. Results are mean % size of original wound \pm SE. $**p < 0.005$ $****p < 0.0001$ by one-way ANOVA E) Wound healing at Day 13 post-wounding measured as percentage of original wound area. F) Area-under-curve (AUC) analysis for wound healing rates compiled for all wounds for all Days 1-13 post-wounding ($n=28$ wounds in PAO Δ Pf4 and PAO1 groups; $n=14$ wounds in PBS group; comparison by one-way ANOVA). G) Representative H&E stain of Day 13 wound infected with PAO1 Δ Pf4. H) Representative H&E stain of Day 13 wound infected with PAO1. I) Percent re-epithelialization calculated as ((Day 1 epithelial gap – Day 13 epithelial gap)/Day 1 epithelial gap). ($p=0.005$; $p=0.0003$ by one-way ANOVA).

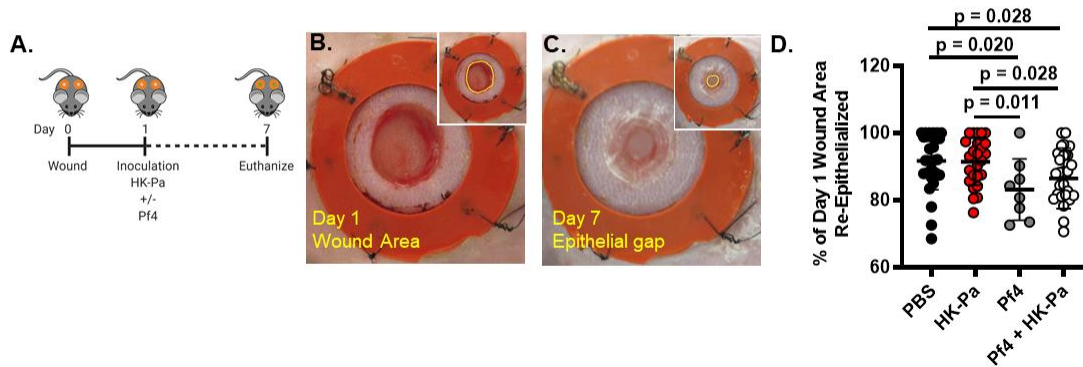


Figure 2. Pf phage delay re-epithelialization in the absence of bacterial infection in murine models.

A) Schematic of full-thickness chronic *Pa* wound infection murine model using ring system. B) Image of a representative wound area inoculated with heat-killed *Pa* on Day 1. C) Image of the same wound area inoculated with heat-killed *Pa* on Day 7. D) Epithelial gap on Day 7 normalized to Day 1 wound area. Combined results from multiple experiments with 8-30 wounds in total per group. PBS vs Pf4 $p=0.028$, PBS vs Pf4 + HK-Pa $p=0.020$, HK-Pa vs Pf4 $p=0.011$, HK-Pa vs Pf4 + HK-Pa $p=0.028$ by unpaired Student's T-test.

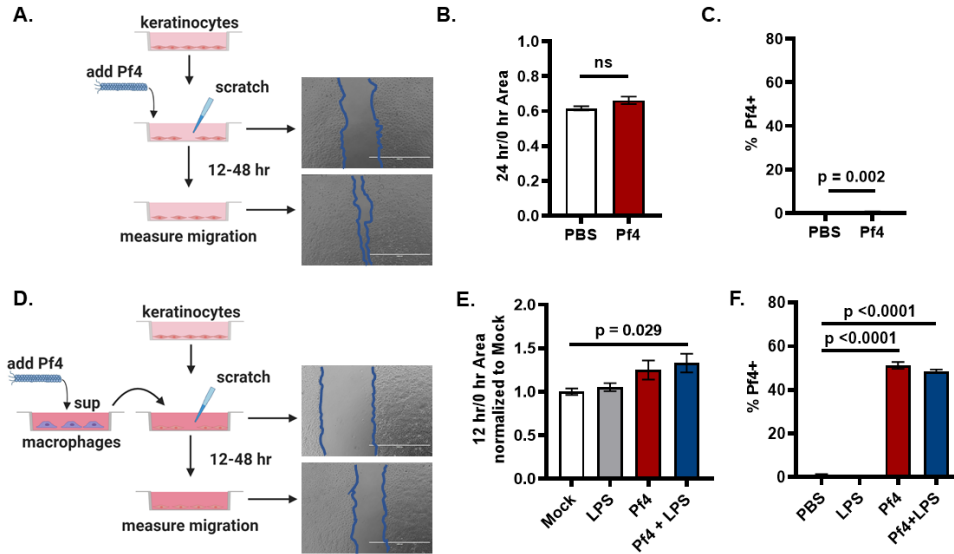


Figure 3. Supernatant from macrophages stimulated with Pf phage slows keratinocyte migration.

A) Schematic of fibroblast and keratinocyte migration assays with representative images of migration assay of HaCaT keratinocytes at 0 hour and 24 hours. B) Effect of Pf4 (1×10^{10} copy #/ml) on HaCaT keratinocyte migration, presented as area at 24 hours/area at 0 hours. Results representative of 3 independent experiments. C) Percentage of Pf4 phage uptake of HaCaT keratinocytes after culture for 3 hours with 1×10^{11} copy #/ml AlexaFluor 488 labeled Pf4 phage. Combined data from 2 independent experiments. D) Schematic of keratinocyte migration assays in which HaCaT keratinocytes were incubated with supernatant from U937-derived macrophages primed with Pf4 and stimulated with LPS. E) Effect on keratinocyte migration of supernatant from U937-derived macrophages that had been primed with Pf4 (1×10^{10} copy #/ml) overnight and stimulated LPS (1 ug/ml) for 4 hours. Results are combined from 6 independent experiments presented as area at 12 hours/area at 0 hours normalized to the mock phage prep condition. F) Percentage of Pf4 phage uptake by U937-derived macrophages primed with Pf4 (1×10^{10} copy #/ml) overnight and stimulated LPS (1 ug/ml) for 4 hours. $P < 0.0001$ by one-way ANOVA. Representative of 2 independent experiments.

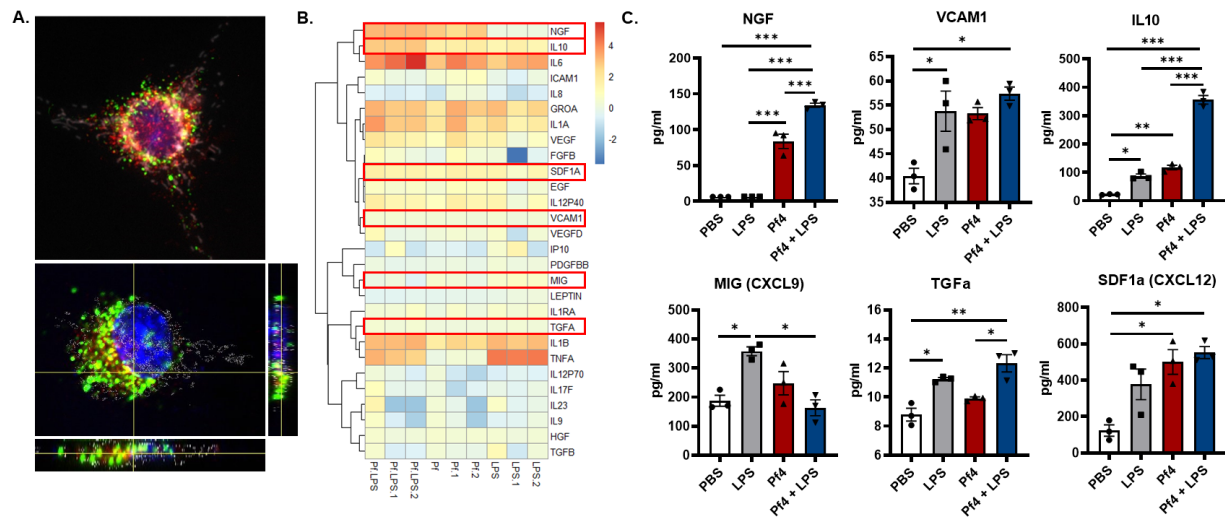


Figure 4. Pf phage uptake by macrophages alters cytokines that affect keratinocyte migration

A) Confocal imaging of murine bone marrow derived macrophage (BMDMs) showing internalization of AlexaFluor 488 labeled Pf4 phage. B) Heat map from Luminex Immunoassay performed on supernatant from human primary monocytes stimulated with Pf4 and LPS. Results are presented as log₂ fold change of concentration compared to PBS control. C) Interpolated values (pg/ml) of select cytokines and growth factors from Luminex assay of human primary monocytes stimulated with Pf4 and LPS. *p<0.05, **p<0.01, ***p<0.001 by one-way ANOVA.

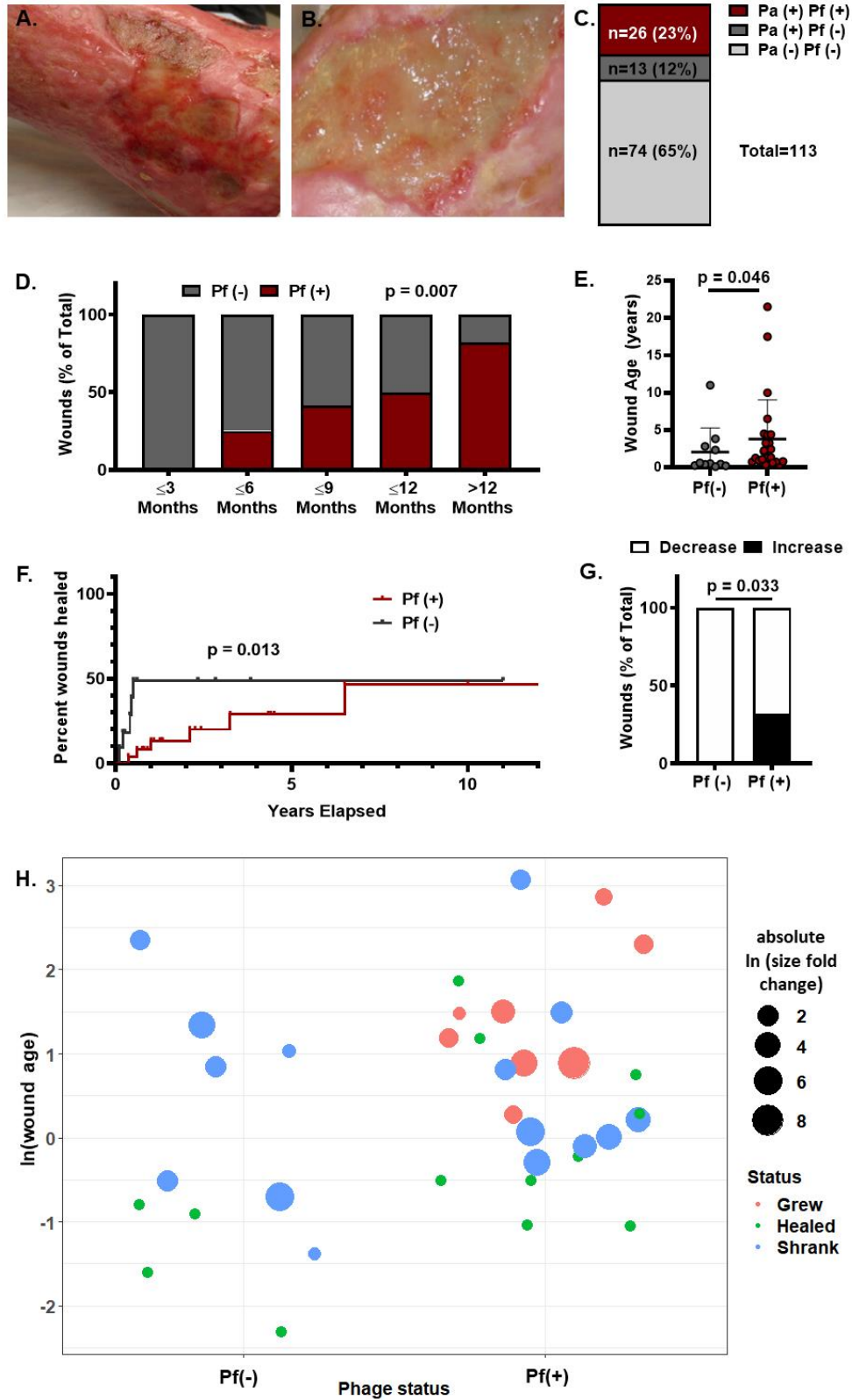


Figure 5. Pf phage are associated with more chronic wound infections and increased wound size in humans.

A) Representative image of Pf(+) wound. B) Representative image of Pf(-)wound. C) *Pa* and Pf phage prevalence in a cohort of 113 patients seen at the Stanford Advanced Wound Care Center. D) Prevalence of Pf(+) wounds when grouped by age of wound into ≤ 3 months, ≤ 6 months, ≤ 9 months, ≤ 12 months, or > 12 months. Results presented as % of total wounds in each age group. $n=36$; $p=0.007$ by Chi-square test. E) Pf(-) ($n=11$) and Pf(+) ($n=25$) wound age presented in years. Mean wound age Pf(-) 2.0 years vs Pf(+) 3.8 years. Median wound age Pf(-) 0.5 years vs Pf(+) 2.1 years, $p=0.046$ by Mann-Whitney test. F) Survival analysis of Pf(+) and Pf(-) *Pa* wound infections, with wound healing as end point. Wounds that were not healed by the end of the study were censored. $p=0.013$ by Gehan-Breslow-Wilcoxon test. G) Pf(+) status is associated with increases in wound size. Pf(-): $n=11$; Pf(+): $n=25$; $p=0.033$ by Chi square test. I) Plot showing wound age and change in size. Wounds were plotted based on Pf(-) or Pf(+) phage status and wound age (calculated as $\ln(\text{wound age in years})$). Wound size change is represented as absolute \ln fold change, with color indicating whether wound grew (red), healed (green), or shrank (blue).

Table 1. Patient Demographic Data.

	Pa(+)/Pf(+)	Pa(+)/Pf(-)	p-value
Samples (n, %)	25, 69%	11, 31%	-
Age of Patients (years)			
Mean (SD)	61.6 (14.7)	76.6 (14.5)	<u>0.008</u>
Gender (n)			
Male	9 (36%)	6 (55%)	0.465
Female	16 (64%)	6 (45%)	
Race/Ethnicity			
Caucasian	17 (68%)	5 (45%)	0.144
Asian	3 (12%)	1 (9%)	
Hispanic	3 (12%)	5 (45%)	
African-American	2 (8%)	0 (0%)	
Body Mass Index (kg/m ²)			
Mean (SD)	27.95 (11.9)	30.28 (6.9)	0.551
Age of Wound (years)			
Median	2.1	0.5	<u>0.046</u>
Recurrence of Infection	9 (43%)	3 (43%)	0.756
Co-Morbidities			
Diabetes Mellitus	12 (48%)	3 (27%)	0.295
Renal Disease	8 (32%)	1 (9%)	0.223

Clinical information on patients from the AWCC with culture-positive and qPCR-positive *Pa*-infected non-healing wounds who participated in the wound swab study. Renal Disease was defined as patients with Chronic Kidney Disease (CKD) or End Stage Renal Disease (ESRD). Statistical significance was measured using Fisher's Exact Test for the following parameters: Gender, Infection Recurrence, and Co-Morbidities. Statistical significance was measured using an Unpaired T-Test for Age and BMI, Chi-Square Test for Race/Ethnicity, and Unpaired Two-Tailed Mann-Whitney Test for Age of Wound because the data was nonparametric. 9 patients were excluded from the Recurrence of Infection analysis because they did not provide an answer on the patient intake form provided by the AWCC.

Table 2. Human Wound Microbiology Data.

	Pa(+) Pf(+)	Pa(+) Pf(-)	p-value
Pf Phage, copies/swab, mean (range)	2.16x10 ⁷ (3.55x10 ³ - 2.69x10 ⁸)	0	-
<i>P. aeruginosa</i> , copies/swab, mean (range)	7.55x10 ⁷ (6.82x10 ² – 9.2x10 ⁸)	2.74x10 ⁷ (5.56x10 ³ – 1.79x10 ⁸)	0.395
Antimicrobials at Time of Swab			
Antimicrobials	14 (56%)	2 (18%)	0.067
Anti-Pseudomonal	6 (24%)	2 (18%)	>0.999
Antibiotic Resistance			
Levofloxacin	8 (32%)	1 (9%)	0.223
Ciprofloxacin	5 (20%)	1 (9%)	0.643
Imipenem	4 (16%)	4 (36%)	0.214
Meropenem	4 (16%)	3 (27%)	0.650
Gentamicin	2 (8%)	1 (9%)	>0.999
Piperacillin	1 (4%)	1 (9%)	0.524
Co-Infection			
Presence of Co-Infection	19 (76%)	7 (64%)	0.454
<i>Staphylococcus aureus</i>	10 (40%)	4 (36%)	>0.999
Gram-Positive	8 (32%)	2 (18%)	0.690
Other Gram-Negative	11 (44%)	5 (45%)	>0.999
<i>Candida</i> Species	0 (0%)	1 (9%)	0.306

Statistical significance for *Pa* copies/swab was analyzed using an Unpaired T-Test. Statistical significance for Antimicrobials at Time of Swab, Antibiotic Resistance, and Co-Infection was analyzed

using Fisher's Exact Test. Anti-Pseudomonal antibiotics included in the analysis were ciprofloxacin and levofloxacin. 0 patients were resistant to Tobramycin. None of the Co-Infection parameters were significant, which strengthens the correlation between Pf phage and clinical outcomes.

Methods

LEAD CONTACT AND MATERIALS AVAILABILITY

- Further information and requests for resources and reagents should be directed to and will be fulfilled by the Lead Contact, Paul L. Bollyky (pbollyky@stanford.edu).
- This study did not generate new unique agents.

EXPERIMENTAL MODEL AND SUBJECT DETAILS

Bacterial strains and culture conditions

P. aeruginosa strain PAO1 was used for all experiments. Isogenic phage-free strain PAO1 Δ Pf4 is derived from strain PAO1, but PAO1 Δ Pf4 lacks the genomic copy of Pf4 phage entirely (Rice et al., 2009). This strain can still be re-infected by Pf4 phage. In general, bacteria were prepared as follows. Frozen glycerol stocks were streaked on Luria-Bertani (LB) agar containing selective antibiotics (PAO1 and clinical strains: none; PAO1 Δ Pf4: 10 μ g/mL gentamicin) and grown overnight at 37°C. An isolated colony was picked and grown overnight at 37°C in LB medium, pH 7.4 under shaking, aerobic conditions. The next day, cultures were diluted to OD₆₀₀ = 0.05 in 75 mL LB media and cultures were grown until early exponential phase (OD₆₀₀ \pm 0.3). OD₆₀₀ was measured and the required number of bacteria was calculated, washed, and prepared as described in the experiments.

Cell Culture Conditions

Human primary monocytes, U937 cell line (ATCC CRL-1593.2), and other immune cells were cultured in RPMI with 10% FBS, 100 IU penicillin, 100 μ g/mL streptomycin and 1 mM sodium pyruvate. HaCaT human keratinocyte cell line (AddexBio, Cat. No. T0020001) was cultured in DMEM with 10% FBS, 100 IU penicillin, and 100 μ g/mL streptomycin in T-75 flasks (Corning, Cat. No. 430641). All cells were cultured on sterile tissue culture- treated plates (Falcon). Cells were cultured at 37°C, 5% CO₂ at a 90% humidified atmosphere.

Mice

Mice were bred and maintained under specific pathogen-free conditions, with free access to food and water, in the vivarium at Stanford University. Mice that underwent surgery received additional Supplical Pet Gel (Henry Schein Animal Health, Cat. No. 029908) and intraperitoneal injections of sterile saline (Hospira, Cat. No. 0409-4888-10). All mice used for *in vivo* infection experiments were littermates. Conventional C57BL/6J mice were purchased from The Jackson Laboratory (Bar Harbor, ME). All experiments and animal use procedures were approved by the Institutional Animal Care & Use Committee at the School of Medicine at Stanford University.

Collection of wound swabs from human patients

From 06/2016 to 06/2018, patients visiting the Stanford Advanced Wound Care Center (AWCC) in Redwood City, California with open wounds were swabbed in duplicate over a one square inch area using Levine's technique, using nylon-flocked wet swabs (Copan Diagnostics, Cat. No. 23-600-963). Swabs were collected in PBS and stored in -80°C before transport on dry ice. In the laboratory, the swabs in PBS were thawed, vortexed vigorously for 15 seconds, and the contents were aliquoted for quantitation of *Pa* rpIU gene and Pf prophage gene PAO717, as detailed below. Patients at the Wound Care Center were also swabbed for confirmation by diagnostic laboratory culture for the presence of *Pa*. Patients were subsequently followed until wounds completely healed or until August 2018. Patients were considered *Pa*-positive if their swabs had detectable *Pa* rpIU and their diagnostic cultures were positive. Patients were considered Pf phage-positive if both duplicate wound swabs had detectable levels of Pf phage genes. None of the *Pa*-negative patients had detectable Pf phage. For patients with multiple wounds, the dominant wound was selected for analysis. Patient enrollment and swab collection were done in compliance with the Stanford University Institutional Review Board for Human Research. Written informed consent was obtained from each patient before swab collection.

Human wound closure analysis

Length, width, and depth measurements of the wounds were taken and recorded into the patient's flowsheet at the start of each visit by the intake nurse. Additional measurements of undermining and tunneling were also taken when applicable. Wound measurements were taken for all patients in the study each time they visited the Stanford Advanced Wound Care Center (AWCC). The AWCC nursing staff is trained in collecting and recording the longest length, width, and depth for each wound. Total volume was calculated for each wound for analysis. Wound healing was defined as a reduction in size compared to the first recorded measurement, whereas wound size increase was defined as an increase in wound volume compared to the first recorded measurement. Wound images were obtained directly from the EMR and uploaded at each clinic visit by the AWCC staff from the designated clinic camera used for this purpose. Images were cropped and resized to protect patient identity. Informed consent was obtained from patients before images were taken. Patient data was collected from EMR chart records, including patient age, gender, co-morbidities, wound age, and other variables. This included history and physicals, progress notes, and documents uploaded into the EMR, such as the AWCC patient intake questionnaire. Patient flowsheet review was accessed for precise wound measurements and laboratory results were accessed to assess renal function and glycemic control. Microbiologic data were reviewed for antibiotic resistance profiles.

METHOD DETAILS

Materials and Reagents

The following chemicals, antibiotics, and reagents were used: RPMI (HyClone, Cat. No. SH30027.01); DMEM (Hyclone, Cat. No. SH30243.01); PBS (Corning Cellgro, Cat. No. 21-040-CV); tryptone (Fluka Analytical, Cat. No. T7293); sodium chloride (Acros Organics, Cat. No. 7647-14-5); yeast extract (Boston BioProducts, Cat. No. P-950); agar (Fisher BioReagents, Cat. No. BP9744); gentamicin (Amresco, Cat. No. E737).

Preparation of heat-killed bacteria

Frozen glycerol stocks were streaked on LB agar as described above. Individual colonies were grown in 5 mL of LB broth the next day for 2 hours to approximately 2×10^8 CFU/mL. The bacterial cultures were centrifuged at $6,000 \times g$ for 5 minutes, and the pellet was washed in 1 mL of PBS three times. Finally, the pellet was resuspended in 1 mL of PBS and heated for 30 minutes at 90°C under shaking conditions. The preparation was checked for sterility by plating.

Phage Purification

This was performed as previously reported. In brief, phage were harvested following treatment with 1 $\mu\text{g}/\text{mL}$ DNase I (Roche, Cat. No. 4716728001) and PEG8000-precipitation as described previously (Sweere et al., 2019b). In some experiments, supernatant was treated with 250 $\mu\text{g}/\text{mL}$ of RNase A (Thermo Fisher Scientific, Cat. No. EN0531) or 85 U/mL of benzonase (Novagen, Cat No. 70746) for 4 hours at 37°C before sterilization. The purified phage pellets were suspended in sterile PBS and dialyzed in 10-kDa molecular weight cut-off tubing (FisherScientific, Cat. No. 88243) against PBS, quantified by qPCR, diluted at least $10,000\times$ to appropriate concentrations in sterile PBS and filter-sterilized. We sent three different Pf4 preparations diluted in PBS to working concentrations (1×10^8 Pf4/mL) to Nelson Labs (Salt Lake City, UT), where they were tested for endotoxin by *Limulus* amoebocyte lysate testing. All three preps had endotoxin levels under the test sensitivity level of 0.05 EU/mL.

Phages were purified by PEG precipitation only unless noted otherwise. Bacteria were infected with stocks of Pf4 phage at mid-log phase and cultured in 75 mL of LB broth for 48 hours at 37°C under

shaking conditions. Bacteria were removed by centrifugation at $6,000 \times g$ for 5 minutes, and supernatant was treated with $1 \mu\text{g/mL}$ DNase I (Roche, Cat. No. 4716728001) for 2 hours at 37°C before sterilization by vacuum filtration through a $0.22 \mu\text{m}$ filter. Pf phage were precipitated from the supernatant by adding 0.5 M NaCl and 4% polyethylene glycol (PEG) 8000 (Milipore Sigma, Cat. No. P2139). Phage solutions were incubated overnight at 4°C . Phage were pelleted by centrifugation at $13,000 \times g$ for 20 minutes, and the pellet suspended in sterile TE buffer (pH 8.0). The suspension was centrifuged for $15,000 \times g$ for 20 minutes, and the supernatant was subjected to another round of PEG precipitation. The purified phage pellets were suspended in sterile PBS and dialyzed in 10- kDa molecular weight cut-off tubing (FisherScientific, Cat. No. 88243) against PBS, quantified by qPCR, diluted at least $10,000\times$ to appropriate concentrations in sterile PBS and filter-sterilized.

For migration experiments with U937 supernatants, phage purification was performed by anion exchange. The phage filtrate was subsequently concentrated to 19 ml using Amicon Ultra 15 ml 50kDa Ultracel spin filters (Millipore). The pH of the phage concentrate was adjusted to pH 5.5 using a 250 mM MES buffer (pH 5.5) to a final concentration of 25 mM MES. The phage concentrate was passed through a Pierce Strong Cation Exchange maxi prep columns (Thermo Fisher) as instructed by the manufacturer. The flow through containing the phage was dialyzed overnight against 4 l 25 mM Tris-HCl (pH 8). The following day the dialyzed phage preparation was further purified through a Pierce Strong Anion Exchange (SAX) maxi prep columns (Thermo Fisher) as instructed by the manufacturer. Briefly, the phage preparation was transferred to the SAX column and centrifuged at $500 g$ for 5 min. the flow through was discarded and the column was washed once with 25 mM Tris-HCl (pH 8). Following the first wash, the column was washed three additional times using 25 mM Tris-HCl (pH 8) supplemented with 0.3 M NaCl. In a final step the phage was eluted from the column in three steps using 10 ml of 25 mM Tris-HCl (pH 8) supplemented with 1 M NaCl. The purified phage Pf4 was further concentrated using Amicon Ultra 15 ml 50kDa Ultracel spin filters (Millipore) to a final volume of 2.5 ml. The purified phage Pf4 was desalted in a final step using a PD-10 Desalting Columns with Sephadex G-25 Medium (GE Healthcare) and eluted with PBS. The phage concentration was determined through qPCR and Tris-Tricine protein gel.

Quantification of Pf phage

As several factors can produce plaques on bacterial lawns (other species of phage, pyocins, host defensins, etc.), we quantitated Pf phage using a qPCR assay as previously described (Secor et al., 2015b). In brief, to quantitate Pf prophage in human or mouse wound homogenates and purified Pf phage preparations, bacterial cells and debris were removed by centrifugation at 8,000× g for 10 min. Supernatants were boiled at 100°C for 20 min to denature any phage particles, releasing intact Pf phage DNA. 2 μL was used as a template in 20 μL qPCR reactions containing 10 μL SensiFAST™ Probe Hi-ROX (Bioline, Cat. No. BIO-82020), 200 nM probe, and forward and reverse primers with concentrations depending on the target. Cycling conditions were as follows: 95°C 2 min, (95°C 15 sec., 60°C 20 sec.) x 40 cycles on a StepOnePlus Real-Time PCR system (Applied Biosystems). For a standard curve, the sequence targeted by the primers and probe were inserted into a pUC57 plasmid (Genewiz) and tenfold serial dilutions of the plasmid were used in the qPCR reactions. For the human wound swabs, the primers and probe were designed to recognize PAO717, a gene conserved across the Pf phage family (F at 600 nM: TTCCCGCGTGGAATGC; R at 400 nM: CGGGAAGACAGCCACCAA; probe: AACGCTGGGTCGAAG); the *Pa* 50S ribosomal gene *rpIU* was used to confirm *Pa* infection (F at 200 nM: CAAGGTCCGCATCATCAAGTT; R at 200 nM: GGCCCTGACGCTTCATGT; probe: CGCCGTCGTAAGC). For the Pf4 phage purifications, the primers and probe were designed to recognize a Pf4 phage-specific intergenic region between PAO728 and PAO729 (F at 500 nM: GGAAGCAGCGCGATGAA; R at 500 nM: GGAGCCAATCGCAAGCAA; probe: CAATTGCGCTGGTGAA). For the Pf phage purification preparations, levels of *Pa* 50S ribosomal gene *rpIU* were measured to correct for contaminating genomic *Pa* DNA, but those levels were usually negligible.

Fluorescent labeling of Pf phage

Purified Pf4 preparations were labeled with AlexaFluor 488-labeled TFP ester (Molecular Probes, Cat. No. A37570) following the manufacturer's protocol. Following labeling, labeled Pf4 viral particles were

separated from unincorporated dye using PD10 Sephadex G-25 desalting columns (GE Healthcare, Cat. No. 17085101) according to manufacturer's instructions. The labeled viral particles were then quantified using qPCR.

Uptake assays of labeled Pf phage by HaCaT keratinocytes

HaCaT cells were seeded at 1.2×10^5 cells per well in 24 well plates and incubated overnight at 37°C. AlexaFluor 488-labeled Pf4 phage was added to wells for a final concentration of 1×10^{11} labeled Pf4/ml. Cells were incubated for 3 hours at 37°C for uptake analysis or 4°C for adsorption analysis. Cells were removed from culture plate with 0.25% trypsin-EDTA solution, washed in cold flow cytometry buffer twice, stained for live-dead discrimination, and fixed according to our flow cytometric protocol. Acquisition was performed on a LSR II flow cytometer (BD Biosciences) or Aurora spectral flow cytometer (Cytex). Analysis was performed using FlowJo (BD Life Sciences).

***In vivo* murine full-thickness wound infection model**

This was done as recently described (Sweere et al., 2019a; Sweere et al., 2019b). In brief, ten-to-twelve-week old male mice were anesthetized using 3% isoflurane, and their backs were shaved using a hair clipper and further depilated using hair removal cream (Nair). The shaved area was cleaned with sterile water and disinfected twice with Betadine (Purdue Fredrick Company, Cat. No. 19-065534) and 70% ethanol. Mice received 0.1-0.5 mg/kg slow-release buprenorphine (Zoopharm Pharmacy) as an analgesic. Mice received two dorsal wounds by using 6-mm biopsy punches to outline the wound area, and the epidermal and dermal layer were excised using scissors. For certain experiments as indicated in the text, a silicone ring was sutured in place around the wound. The wound area was washed with saline and covered with Tegaderm (3M, Cat. No. 1642W). Bacteria were grown as described above and diluted to 1×10^7 CFU/mL in PBS. Mice were inoculated with 40 μ L per wound 24 hours post-wounding, and control mice were inoculated

with sterile PBS. Mice were weighed daily and given Supplical Pet Gel and intraperitoneal injections of sterile saline. Upon takedown, wound beds were excised and processed for histological analysis.

Wound Healing Analysis in Murine Models

This was done as previously described (Balaji et al., 2014). In brief, using a dot ruler for standardization, images of both wounds for all mice were taken on each day from Day 1 to Day 13 post-wounding. Images were taken using Canon PowerShot Camera (Item Model No. 1096C001) mounted on a fixed tripod for consistency in the height and angle at which images were taken. Using the images, wound area was measured using Fiji (Image J, National Institutes of Health) (Schindelin et al., 2012) by tracing the border of the wounded area, as well as the border of a standardized dot measure. For histologic analyses, wounds were harvested, bifurcated, fixed in 10% neutral buffered formalin, and embedded in paraffin. Wound sections of 5 μm were cut from paraffin-embedded blocks. Immune cell influx, general tissue morphology, epithelial gap and granulation tissue area were measured from hematoxylin and eosin (H&E) stained sections using morphometric image analysis (Image-Pro, Media Cybernetics, Silver Spring, MD, and Metamorph, Molecular Devices, Downingtown, PA). Images were obtained using a Nikon Eclipse microscope, and image analysis was performed using the Nikon Elements software (Nikon Instruments, Melville, NY, USA). The percentage of CD45+ cells to the total cell infiltrate was calculated in 6 high-powered fields (HPFs, 64X) for each wound section. The HPFs were chosen just above the panniculus carnosus and were evenly distributed between the two wound epithelial margins. On a 4 \times edge-to-edge wound section image, epithelial gap was measured as the distance (in millimeters) between encroaching epithelial margins. The granulation tissue area was measured (in mm^2) as the entire cellular region within the epithelial margin. Any sample that did not yield reliable counting due to sample quality was excluded from the analysis for that particular variable. All scoring was done by an independent pathologist.

Microtiter plate crystal violet assay

Bacterial cells were inoculated in a 96-well polystyrene plate. Plate was incubated at 37C, static condition for 24 and 48 hours. Post incubation, media was discarded and the plate was washed to remove unattached or planktonic bacteria. Wells were stained in crystal violet. Post staining, the plate was washed to remove excessive dye with ethanol and absorbance was measured at 590 nm.

Confocal laser scanning microscopy and Scanning Electron Microscopy of *Pa* biofilms

Bacterial cultures were seeded at 37C in sterile tryptic soy broth. Polycarbonate membrane (PCM) discs were cut at 1 cm diameter and placed on tryptic soy agar plates. Overnight broth cultures were spotted at 10 μ l per disc and culture spots were allowed to dry. Plates were incubated at 37C for 24, 48, and 72 hours to allow biofilm formation. For the 48 and 72 hour biofilms, the discs were transferred to new tryptic soy discs every 24 hours to replenish nutrients. After respective time intervals, PCM discs were transferred to glass slides. Biofilms were stained with Syto9 (cell stain) and Sypro Ruby (biofilm matrix stain). Post staining, discs were mounted with a coverslip and observed on a confocal laser scanning microscopy at 63 \times . Thickness measurements were derived from 'z' values of images. For SEM, biofilms were sputter coated with gold.

Luminex immunoassay

Human peripheral blood mononuclear cells (PBMCs) were isolated from whole blood using lymphoprep (Stem Cell technologies, Cat no. 07851). Monocytes were subsequently isolated from the PBMC mixture using a human monocyte isolation kit without CD16 depletion (Stem Cell Technologies, Cat no. 19058), as described by the manufacturer. The isolated monocytes were resuspended at 5×10^7 cells/ml in RPMI supplemented with MEM (non-essential amino acids, Sodium pyruvate, Penicilline/Streptomycine, L-glutamine (200 mM), 2-mercaptoethanol (50 mM) and 10 % heat inactivated fetal bovine serum. A 100 μ l of the monocyte suspension was transferred to a 96 well plate and stimulated for 15 h with 10^9 phage particles. Subsequently LPS (1 μ g/ml) was added for 4 h. Supernatant was collected and used for cytokine

profiling through Luminex (The Human Immune Monitoring Center, Stanford). Human 63-plex kits were purchased from eBiosciences/Affymetrix and used according to the manufacturer's recommendations with modifications as described below. Briefly: Beads were added to a 96 well plate and washed in a Biotek ELx405 washer. Samples were added to the plate containing the mixed antibody-linked beads and incubated at room temperature for 1 hour followed by overnight incubation at 4°C with shaking. Cold and Room temperature incubation steps were performed on an orbital shaker at 500-600 rpm. Following the overnight incubation plates were washed in a Biotek ELx405 washer and then biotinylated detection antibody added for 75 minutes at room temperature with shaking. Plate was washed as above and streptavidin-PE was added. After incubation for 30 minutes at room temperature wash was performed as above and reading buffer was added to the wells. Each sample was measured in duplicate. Plates were read using a Luminex 200 instrument with a lower bound of 50 beads per sample per cytokine. Custom assay Control beads by Radix Biosolutions are added to all wells.

Migration Assay

For HaCaT cell migration assays a standard scratch assay was used (Cory, 2011). In brief, cells were seeded at a density of 1.5×10^5 cells per well in DMEM 10% FBS in collagen-coated 24-well tissue culture plates. Media was removed and HaCaT cells were then serum starved in DMEM containing only 2% FBS for 24 hours prior to the treatment. A scratch defect was created in the cell monolayer along the diameter in each group using a sterile pipette tip (200 μ l tip). Treatment groups had Pf4 bacteriophage added to media with final concentration of 1×10^{10} copy #/ml. Photographic images were obtained at 0 and 24 hours incubation using an Invitrogen EVOS FL Imaging System. The unfilled scratch defect area was measured at each reference point per well. Data was presented as extent of wound closure, that is, the percentage by which the scratch area has decreased at a given time point for each treatment as compared to the original defect (at 0 hours). All experiments were carried out at minimum in triplicate and the passage number was similar amongst the different groups. For experiments utilizing supernatant from U937 macrophages, media was replaced after serum starvation with supernatant prepared as follows. U937 cells were differentiated into

macrophages by stimulation for 3 days with 100 n g/ml Phorbol 12-myristate 13-acetate (Sigma, Cat. No. P1585). Resulting macrophages were seeded in 6-well plates at 2×10^6 cells/well and stimulated with 1×10^{10} #/ml Pf4 overnight followed by LPS 1 ug/ml for 4 hours at 37°C. Supernatant was then centrifuged to remove debris and cells and frozen at -80°C.

Generation of murine bone marrow-derived macrophages (BMDMs)

Femurs were isolated from male or female mice of >10 weeks of age and bone marrow was flushed out using a 27 G PrecisionGlide needle (BD Biosciences, Cat. No. 305109). Cells were plated at 1×10^6 cells in 10 ml of media per Petri Dish (Fisherbrand, Cat. No. FB0875711). Media was supplemented with 20 ng/ml of M-CSF (Invitrogen, Cat. No. 14-8983-62). An equivalent amount of fresh media containing cytokines was added three days after plating, and 50% of media was changed 5 and 7 days after plating. BMDMs were harvested on day 6 after plating. Supernatant containing non-adherent cells was collected, and adherent cells were lifted by the addition of 10% Accumax in Accutase for 5 min. Both cell populations were pooled to obtain a mixture of mature and immature cells, washed in fresh media at 300g for 5 min, and prepared for experiments.

Confocal microscopy

BMDMs were seeded on glass 12-mm #1.5H precision coverslips (Thorlabs, Cat. No. CG15CH) coated with poly-L-lysine (Sigma, Cat. No. P8920) or 0.2% poly-L-ornithine (Sigma, Cat. No. P3655) according to the manufacturer's instructions at a density of 1×10^6 cells/well in a 6-well plate. Cells were either stimulated with coated 1×10^9 Pf4/ml labeled Pf4 or soluble 1×10^{10} Pf4/ml labeled Pf4 for 3 hours. Cells were washed three times with PBS and subsequently fixed with 10% neutral buffered formalin for 5 minutes at room temperature. After permeabilization with 0.5% Triton X-100, coverslips were washed twice with PBS and excess formalin was quenched by incubation with 150 mM glycine, pH 8.0 (Fisher BioReagents, Cat. No. BP381) for 30 minutes. Coverslips were blocked with Sea Block blocking buffer (Thermo Scientific, Cat. No. 37527) followed by incubation with primary antibodies in Sea Block overnight at 4°C.

Coverslips were washed three times with PBS and incubated with the appropriate fluorescently labeled secondary antibodies. Coverslips were mounted in ProLong Diamond Antifade Mountant (Invitrogen, Cat. No. P36970) with DAPI to label nuclei. Fluorescent specimens were examined using a confocal system (DMX BLAZE 3D SIM, GE Healthcare). Images were acquired using Leica DFC365 FX camera and the Leica Application Suite X (Leica). Image analysis was performed using FIJI (NIH).

QUANTIFICATION AND STATISTICAL ANALYSIS

All statistical analyses were done using GraphPad Prism (GraphPad Software, Inc. La Jolla, CA). All Unpaired T-Tests, Mann-Whitney Test, Fisher's Exact Tests, and Chi-Square Test were two-tailed. Depicted are Means with Standard Error or Standard Deviation of the population unless otherwise stated. Statistical significance was considered $p < 0.05$.

DATA AND CODE AVAILABILITY

Original/source data for figures and tables in this paper is available by request

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