

Pseudomonas aeruginosa Gene Products PilT and PilU Are Required for Cytotoxicity In Vitro and Virulence in a Mouse Model of Acute Pneumonia

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Type IV pili of the opportunistic pathogen *Pseudomonas aeruginosa* mediate twitching motility and act as receptors for bacteriophage infection. They are also important bacterial adhesins, and nonpiliated mutants of *P. aeruginosa* have been shown to cause less epithelial cell damage in vitro and have decreased virulence in animal models. This finding raises the question as to whether the reduction in cytotoxicity and virulence of nonpiliated *P. aeruginosa* mutants are primarily due to defects in cell adhesion or loss of twitching motility, or both. This work describes the role of PilT and PilU, putative nucleotide-binding proteins involved in pili function, in mediating epithelial cell injury in vitro and virulence in vivo. Mutants of *pilT* and *pilU* retain surface pili but have lost twitching motility. In three different epithelial cell lines, *pilT* or *pilU* mutants of the strain PAK caused less cytotoxicity than the wild-type strain but more than isogenic, nonpiliated *pilA* or *rpoN* mutants. The *pilT* and *pilU* mutants also showed reduced association with these same epithelial cell lines compared both to the wild type, and surprisingly, to a *pilA* mutant. In a mouse model of acute pneumonia, the *pilT* and *pilU* mutants showed decreased colonization of the liver but not of the lung relative to the parental strain, though they exhibited no change in the ability to cause mortality. These results demonstrate that pilus function mediated by PilT and PilU is required for in vitro adherence and cytotoxicity toward epithelial cells and is important in virulence in vivo.

Type IV pili of *Pseudomonas aeruginosa* are significant adhesins for interaction with mammalian cells (reviewed in references 15 and 25), contributing to this organism's ability to cause opportunistic infections in humans. They also are required for motility of the bacteria across a solid surface (twitching motility) (8) and for the binding and entry of bacteriophages (6, 7). *P. aeruginosa* pili are polymers of a single gene product, called PilA or pilin (32), but their assembly and function requires the products of at least 30 additional genes (reviewed in reference 1). Three of these genes, *pilB*, *pilT*, and *pilU*, encode proteins that contain motifs common among nucleotide-binding proteins (Walker box A domains [37]) and thus are postulated to contribute energy to pili formation or function (1, 18, 40). Inactivation of *pilB* results in a nonpiliated phenotype (24), while strains with defects in *pilT* or *pilU* overexpress surface pili but are no longer motile on a solid surface (10, 40). These observations, in addition to electron microscopy studies comparing phage binding of *pilT* mutant and wild-type strains (9), suggested that PilB enables pili extension whereas PilT and PilU catalyze pili retraction (15, 23). However, the precise functions of PilT and PilU in this process

remain obscure, and *pilU* mutants, unlike *pilT* mutants, remain sensitive to bacteriophage infection (40). The proposed dynamic nature of pili could promote the entry of bacteriophage into the bacterium and provide the impetus for twitching motility, but its importance in *P. aeruginosa* adherence to mammalian cells and virulence has not been examined. Importantly, since *pilT* and *pilU* mutants retain (hyper)expression of surface pili, such mutants provide an opportunity to dissect the role of pili as adhesins and as the mediators of twitching motility in the process of *P. aeruginosa* infection. This may be also relevant to infection caused by a wide range of other type IV piliated bacteria, including *Neisseria* spp., *Moraxella* spp., *Aeromonas* spp., and *Legionella* spp. (4, 22, 30, 35, 41).

The *pilT* and *pilU* genes are contiguous on the *P. aeruginosa* chromosome but reside in a locus separated from other genes involved in pilus biogenesis and function. Northern blot and complementation analyses suggest that the two genes are transcribed independently and thus are not part of the same operon (40). The encoded proteins are closely related to each other (39% amino acid identity and 61% similarity) but are more distantly related to PilB and to other putative bacterial nucleotide-binding proteins involved in fimbrial biogenesis, protein secretion, or DNA transfer (39, 40). Direct homologs of PilT that are involved in type IV pili function have been identified in enteropathogenic *Escherichia coli* (EPEC) (31), *Myxococcus xanthus* (43), and *Neisseria gonorrhoeae* (11). Examination of the unfinished genome sequences of *N. gonorrhoeae* and *N. meningitidis* indicates that these species at least also contain homologs of PilU (40a). An EPEC mutant defective in the *pilT*-like *bfpF* gene, despite having morphologically wild-type bundle-forming pili, displayed increased adherence to epithelial cells in culture, formed irregular bacterial aggregates,

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TABLE 1. Strains used

Strain	Relevant characteristics	Source or reference
PAK	Wild type; pili ⁺ and flagella ⁺	J. Mattick
PAK-N1	Tn5G insertion in <i>rpoN</i> ; pili ⁻ and flagella ⁻	19
PAK <i>pilT</i> ::Tn5	Tn5B21 insertion in <i>pilT</i> (mutant R364); hyperfimbriated, phage resistant	40
PAK <i>pilU</i> ::Tn5	Tn5B21 insertion in <i>pilU</i> (mutant S34); hyperfimbriated, phage sensitive	40
PAK <i>pilA</i> ::TcR	Tetracycline resistance cassette inserted into <i>pilA</i> ; nonpilated	38

and had decreased virulence in humans (2, 5). An in-frame deletion of the *M. xanthus pilT* gene did not affect the expression of pili or cell to cell agglutination but did cause a loss of pilus-dependent social gliding motility (43). Furthermore, the natural competence and twitching motility of a *N. gonorrhoea pilT* mutant were abolished, but this strain retained wild-type pilus expression and adherence to epithelial cells (41–43). Inactivation of *pilT* was also shown to rescue the pilus production defect of *pilC* mutants, further demonstrating a role for *N. gonorrhoea* PilT in pilus function (42).

A number of studies have implicated *P. aeruginosa* pili as factors important for adherence to epithelial cells in vitro (reviewed in references 15 and 25) and for virulence at several sites of infection (13, 14, 27). These studies have all used nonpilated mutants of *P. aeruginosa* except for one investigation which demonstrated that a *pilT* mutant of *P. aeruginosa* PAO (DB2) and two other hyperpilated isolates were noninfectious in a mouse corneal infection model (17). This study suggested that the presence of pili (as adhesins) per se is insufficient to support virulence in vivo. Here we sought to examine further the importance of PilT and PilU in causing epithelial cell injury in vitro and in vivo by comparing the adherence, cytotoxicity, and virulence of the parental strain PAK to those of isogenic *pilT* and *pilU* mutants and of isogenic nonpilated strains.

PilT and PilU contribute to PAK-mediated cytotoxicity toward multiple epithelial cell lines. Previous studies have demonstrated that the addition of *P. aeruginosa* PAK to the apical surface of a highly polarized epithelial cell monolayer consisting of Madin-Darby canine kidney (MDCK) cells resulted in significant cytotoxicity in vitro (3). Comparison of isogenic mutants and different strains demonstrated that the amount of cell damage observed in vitro correlated well to relative virulence in an animal model of acute pneumonia (16, 28). We used this MDCK system to determine if PilT or PilU were required for cytotoxicity in vitro. MDCK type II cells (5×10^6) were grown as a confluent monolayer on Transwell filters (Corning) for 3 days as previously described (20), washed, and placed in minimal essential medium Eagle (MEM) supplemented with 20 mM HEPES buffer pH 7.4 (MEM-lite). Approximately 10^7 CFU (as determined by dilution plating) of a stationary-phase culture of PAK wild-type or isogenic mutant strain (Table 1) grown for 16 h in Luria broth without shaking at 37°C was added to the apical surface of the MDCK cell monolayer (multiplicity of infection of 2). After incubation for 9 h at 37°C in room air, aliquots of the apical and basal medium were removed and assayed for lactate dehydrogenase (LDH) activity as instructed by the manufacturer (Sigma Chemical Co.). The percentage of cell death was calculated by comparison of the total LDH released to that released from

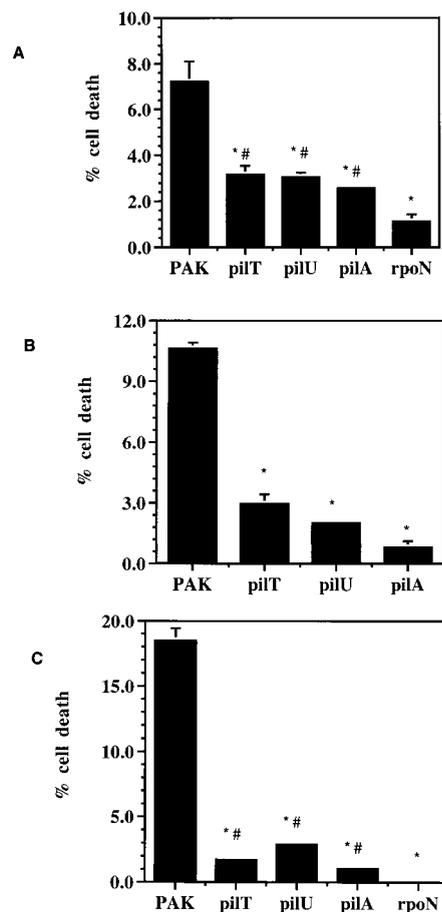


FIG. 1. *P. aeruginosa pilT* and *pilU* mutants have decreased ability to damage epithelial cells. Cytotoxicity of wild-type PAK, hyperpilated *pilT* or *pilU* mutants, or nonpilated *pilA* or *rpoN* mutants was assayed by incubation of the bacteria with the cell type indicated. Cell death was quantitated by LDH release and is expressed as a percentage of the total LDH released by lysis of cells not exposed to bacteria. Each assay was performed in triplicate, and error bars represent SEM. (A) Cytotoxicity of MDCK cells after incubation with bacteria for 9 h. *, $P < 0.003$ compared to PAK; #, $P < 0.003$ compared to *rpoN*. (B) Cytotoxicity observed on A549 cells after incubation with bacteria for 8 h. *, $P < 0.001$ compared to PAK. (C) Cytotoxicity observed on HeLa cells after incubation with bacteria for 5 h. *, $P < 0.003$ compared to PAK; #, $P < 0.05$ compared to *rpoN*. Statistical analysis was performed by using Student's two-tailed *t* test with unequal variance.

uninfected cells lysed with 0.25% Triton X-100 in MEM-lite. The strains lacking PilT or PilU showed reduced cytotoxicity relative to PAK, causing 44 and 42%, respectively, as much cytotoxicity as the wild type, while a *pilA* mutant damaged 36% as many cells as the wild type did (Fig. 1A). A strain with a defect in *rpoN* had even less cytotoxic capability, 16% of the wild-type level, presumably because of pleiotropic effects on a range of genes (36). These findings are not the result of differences in bacterial growth rates since the mutant and wild-type strains had similar rates of growth in minimal or complex medium (data not shown). Furthermore, the mutant strains did not achieve wild-type cytotoxicity even with longer incubation times (up to 11 h; data not shown).

These strains were tested for cytotoxicity toward two additional cell lines, A549 lung pneumocytes and HeLa cells, with similar results (Fig. 1B and C). In preparation for this assay, 2×10^6 A549 cells were grown in Waymouth's 752/1 medium without insulin (Gibco BRL) containing 10% fetal calf serum

for 3 days on Transwell filters at 37°C with 5% CO₂. HeLa cells (2 × 10⁶) were maintained in Dulbecco's modified essential medium containing 5% glucose (UCSF cell culture facility) and 5% fetal calf serum for 2 days in a tissue culture-treated 24-well plate at 37°C with 5% CO₂. As with MDCK cells, cytotoxicity assays were performed by the addition of approximately 10⁷ CFU of the appropriate strain (prepared as described above), but incubation times with these cell types were shortened to 8 h for A549 cells and to 5 h for HeLa cells due to the greater susceptibility of these cells to *P. aeruginosa*-mediated cytotoxicity. Cell death was calculated as described above. When incubated with A549 or HeLa cells, the PAK strains with either the *pilT* or *pilU* gene inactivated were less cytotoxic than the wild type (Fig. 1B and C). Cell death caused by the *pilT* and *pilU* mutants was reduced to 28 and 19%, respectively, of the level for the parental strain with respect to A549 cells and to 9 and 16% with respect to HeLa cells. The *pilA* mutant exhibited 8 and 6% of the wild-type cytotoxicity toward A549 and HeLa cells, respectively. As with MDCK cells, the *rpoN* mutant had the most severe phenotype, causing only 0.2% of the wild-type cytotoxicity toward HeLa cells. Even with extended incubation (up to 11 h [data not shown]), the *pilT* and *pilU* mutants did not achieve wild-type cytotoxicity levels.

These results indicate that the putative nucleotide-binding proteins PilT and PilU are required for full *P. aeruginosa*-mediated epithelial cell damage in vitro. In this assay, the result of inactivation of *pilT* or *pilU* on bacterially mediated cytotoxicity is not reproducibly distinguishable. It should be noted that the transposon insertion into *pilT* is not polar to *pilU* (40), and pilus-related defects in each mutant could be rescued by complementation with the appropriate gene (40). Importantly, these results show that the presence of pili on the bacterial surface of the hyperpilated *pilT* and *pilU* mutants is not sufficient to enable normal *Pseudomonas*-epithelial cell interaction and subsequent wild-type levels of cytotoxicity. As more cytotoxicity is caused by either the *pilT* and *pilU* mutant than by the nonpilated *pilA* mutant, the overexpression or presence of abnormally functioning pili on the surface of these strains appears to make a minor contribution to epithelial cell injury. RpoN-dependent gene products, such as the previously identified nonpilus adhesins (26), also appear to play a role in cytotoxicity since the *rpoN* mutant caused significantly less cell damage than the *pilA* mutant in MDCK cells and HeLa cells (Fig. 1A and C).

PilT and PilU contribute to the association of PAK with epithelial cells. To determine if the loss of the cytotoxic capability of the *pilT* or *pilU* mutant corresponded to a defect in adherence to epithelial cells, the association of the mutant strains to the three cell types used was measured. Approximately 10⁷ CFU of each of the various strains (grown and prepared as for the cytotoxicity assays) was incubated with MDCK, A549, or HeLa cells cultured and plated on Transwell filters exactly as described above. Association was assayed by excising the filters (to eliminate bacteria that adhered to plastic) followed by washing the filter-bound cells three times in MEM-lite. The cells were then lysed by treatment with 0.25% Triton X-100 in MEM-lite for 30 min followed by vortexing with glass beads. The bacteria contained within the lysate were quantified by dilution plating on LB agar. Association assays were performed at time points prior to detectable cytotoxicity (3 h for MDCK and A549 cells and 1 h for HeLa cells) to avoid potential bacterial adherence to cell debris. Adherent bacteria were not distinguished from those internalized, but the fraction of internalized bacteria was less than 1% of the fraction of adherent bacteria with all cell types (data not shown). Using

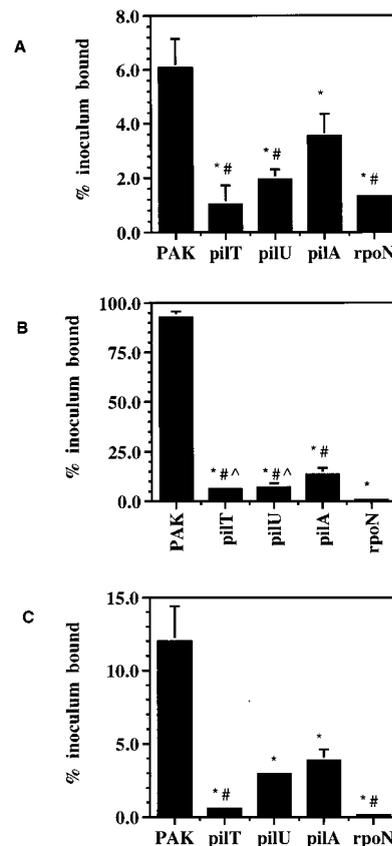


FIG. 2. Association of *P. aeruginosa* with epithelial cells is reduced by inactivation of *pilT* or *pilU*. Wild-type PAK, hyperpilated *pilT* or *pilU* mutants, or nonpilated *pilA* or *rpoN* mutants were incubated with the indicated cell type, and cell-associated bacteria were quantitated after the amount of time indicated. Each assay was performed in triplicate and normalized to the number of CFU initially added to the assay. SEM is represented by the error bars. (A) Fraction of input bacteria associated with MDCK cells after 3 h. *, $P < 0.04$ compared to PAK; #, $P < 0.04$ compared to *pilA*. (B) Fraction of input bacteria associated with A549 cells after 3 h. *, $P < 0.0001$ compared to PAK; #, $P < 0.003$ compared to *rpoN*; ^, $P < 0.05$ compared to *pilA*. (C) Fraction of input bacteria associated with HeLa cells for 1 h. *, $P < 0.006$ compared to PAK; #, $P < 0.003$ compared to *pilA*. Statistical analysis was performed by using the Student's two-tailed *t* test with unequal variance.

this measure, we calculated that 6% of the added wild-type bacteria associated with MDCK monolayers (Fig. 2A), 92% associated with A549 monolayers (Fig. 2B), and 12% associated with HeLa cells (Fig. 2C). The relative binding of PAK to these cell lines is consistent with that previously reported for these cell types (12).

The two nonpilated and two hyperpilated mutants exhibited reduced adherence to each cell type compared to the parental strain (Fig. 2A to C). Of the nonpilated mutants, the *rpoN*-defective strain adhered less than the *pilA* mutant to each cell type, again reflecting a likely role of additional gene products in this interaction. Unexpectedly, the hyperpilated *pilT* or *pilU* mutant also showed reduced binding relative to the *pilA* mutant. The effect was most marked with MDCK monolayers, where the *pilA* mutant bound approximately threefold more effectively than the *pilT* mutant and approximately twofold better than the *pilU* mutant (Fig. 2A). The adherence of each of the *pilT* and *pilU* mutants to A549 cells was roughly twofold less than that of the *pilA* mutant (Fig. 2B), while the *pilT* strain appeared to bind to HeLa cells significantly less than the *pilA*

mutant (Fig. 2C). This trend in the adherence assays is in obvious contrast that observed with in vitro cytotoxicity, where the *pilT* and *pilU* mutant strains displayed a greater ability to kill epithelial cells than the *pilA* mutant. Importantly, the greater cytotoxicity of the hyperpiliated mutants cannot be accounted for simply by increased adherence due to the over-expression of abnormally functioning pili on the surface of these strains. Although the explanation for the greater adherence but reduced cytotoxicity of the *pilA* mutant is unclear, it is possible that the lack of pili on the *pilA* mutant uncovers alternate adhesins. Alternatively, the nonfunctional pili on the surface of the *pilT* or *pilU* strains may act in a dominant negative manner and interfere with these other adhesins, thus inhibiting intimate contact between the bacteria and the host cells.

Since both hyperpiliated and nonpiliated mutants of *P. aeruginosa* show markedly reduced adherence to epithelial cells relative to the wild type (12, 41), the expression of surface pili cannot solely account for the observed cell adherence, suggesting that the extension and retraction of pili or surface translocation by twitching motility may be a relevant factor in adherence to and colonization of host cells. Additional, uncharacterized functions of pili may also contribute to these processes.

In summary, the inactivation of *pilT* or *pilU* decreased both bacterial adherence and cytotoxicity, indicating their importance to these processes. Compared to the *pilA* mutant, however, adhesion of the *pilT* and *pilU* mutant strains was decreased to a greater extent than cytotoxicity. The effect of inactivation of *pilT* in *P. aeruginosa* differs from that in *N. gonorrhoeae* or EPEC with regard to their adherence to epithelial cells. Although not without a phenotype, an *N. gonorrhoeae pilT* mutant bound as well as the wild type to epithelial cells though in a more localized fashion (41), while an EPEC *bfpF* mutant adhered better to mammalian cells than the parental strain (2). This finding suggests that although all of these bacteria possess homologous proteins related to pili function, their absence has slightly different outcomes in each system.

PilT and PilU are required for full virulence in a mouse model of acute pneumonia. To assess the contribution of PilT or PilU to *P. aeruginosa*-induced acute pneumonia, we assayed the virulence of the wild-type and mutant strains in a mouse model of acute pneumonia. Virulence was assessed by two methods, the first using mortality caused by the different isogenic strains as an endpoint and the second measuring the relative ability of the strains to colonize the lung and liver.

For *P. aeruginosa*-induced mouse mortality, approximately 5×10^7 CFU of bacteria (grown for 17 h in MINS medium [16] at 37°C with shaking and then washed and resuspended in 50 ml of phosphate-buffered saline) were instilled into the nares of methoxyfluorane-anesthetized 6- to 12-week-old BALB/c mice (B&K Laboratories), using a pipette tip. Viable counts were determined by dilution plating on LB plates. For 50% lethal dose calculations, five mice each were infected with twofold dilutions of bacteria. Mice were monitored over the subsequent 7 days in compliance with guidelines of the Animal Care Committee of the University of California, San Francisco. Statistical significance was assigned by the chi-square test. As shown in Fig. 3, none of the 15 mice infected with the PAK wild type survived to the 7-day time point. In contrast, the *pilA* mutant strain showed little virulence, and 13 of 15 animals survived for the length of the experiment ($P < 0.0001$). Results with the *rpoN* mutant were similar in that 14 of 15 mice survived for 7 days ($P < 0.0001$). These data are in agreement with previously published results for a neonatal mouse model of

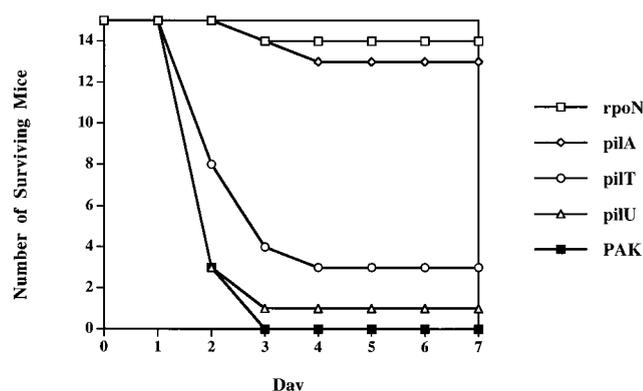


FIG. 3. Virulence of wild-type PAK, hyperpiliated *pilT* or *pilU* mutants, or nonpiliated *pilA* or *rpoN* mutants in a mouse model of acute pneumonia. Approximately 5×10^7 CFU of each strain was used to intranasally infect 15 mice, each of which was monitored for viability over 7 days.

acute pneumonia which demonstrated a loss of virulence associated with a lack of surface pili (33). Surprisingly, in this assay the virulence of the *pilT* and *pilU* mutant strains was statistically indistinguishable from that of the wild type; only 3 of 15 mice receiving *pilT* and 1 of 15 mice inoculated with *pilU* survived for the duration of the experiment. The 50% lethal dose of the *pilU* mutant, the *pilT* mutant, and PAK differed twofold at most (data not shown).

Using colonization as a more sensitive measure of virulence, we performed a competition assay (34) comparing the efficiency of colonization of the lung or liver of the hyperpiliated strains to that of the wild type. Individual strains were grown and prepared as described above for the mortality experiments. Equal numbers (as determined measuring A_{600} and by plating for viable counts) of PAK and *pilU* mutant (either 2.5×10^7 or 5.0×10^7 CFU of each strain), PAK and *pilT* mutant (either 5.0×10^7 or 7.5×10^7 CFU of each strain), or PAK and *pilA* mutant (7.5×10^7 CFU of each strain) were resuspended in a total of 50 μ l of phosphate-buffered saline and instilled into the nares of anesthetized 6-to-12-week old BALB/c mice. Twenty-four hours later, the animals were sacrificed. The right lobes of the lung and the liver were removed under sterile conditions and placed in 1 ml of LB, and serial dilutions were plated onto LB plates containing no antibiotic or 50 μ g of tetracycline per ml (PAK is tetracycline sensitive, whereas each of the mutants carried a tetracycline resistance cassette). A competitive index was calculated by generating the ratio of mutant strain to wild-type bacteria recovered from the lung or liver and comparing it to the same ratio in the infecting inoculum (roughly 1.0 but calculated precisely for each experiment). A competitive index greater than 1.0 indicates that the mutant strain colonized better than wild type, and a competitive index less than 1.0 indicates that the mutant strain was less efficient than the wild type in colonization. The results of two to three experiments using a total of 5 to 10 mice were combined for each sample.

The competitive index of the *pilU* mutant in the lung was 2.3 (standard error of the mean [SEM] 0.73, $n = 5$), suggesting that this mutant strain may colonize the lung better than the wild type. In contrast, the liver competitive index was 0.10 (SEM 0.04, $n = 10$), indicating that the *pilU* mutant exhibited a 25-fold decrease in its ability to spread to and colonize the liver 24 h postinoculation. Likewise, the *pilT* mutant was approximately 20-fold less efficient in colonizing the liver compared to the lung; the lung index was 1.08 (SEM 0.79, $n = 6$),

while in the liver it was 0.07 (SEM 0.03, $n = 8$). For either the PAK-*pilU* or PAK-*pilT* mixture, the total CFU recovered from the livers was at least 2 orders of magnitude less than the quantity of bacteria in the lungs (data not shown). The recovered *pilU* and *pilT* strains were indistinguishable from the input bacteria and failed to exhibit twitching motility on Vogel-Bonner medium plates. When mice were infected with a combination of PAK and the *pilA* mutant, no bacteria were recovered from the liver, even when a larger inoculum of 7.5×10^7 was used. This precluded calculating a competitive index for the liver, though in the same experiment, the competitive index in the lungs was 2.14 (SEM 0.70, $n = 5$). Similarly, recovery of bacteria (wild type or mutant) from the lungs and livers of PAK-*pilT*-infected mice required use of a larger inoculum of each strain compared to mice inoculated with PAK-*pilU*. The negative effect of the mutant strains on the wild-type infection implies that the bacteria present in a mixture act synergistically to colonize the lung or the liver, possibly at the stage of initial interaction with the nasal epithelium and upper airways. It may be that the *pilU* mutant strain is more efficient at this process than either the *pilT* or *pilA* mutant. Alternatively, the mutant strains may differ in the ability to provide protection from simultaneous infection with the parental strain.

Thus, in the mouse model of acute pneumonia, the virulence of the *pilT*, *pilU*, and *pilA* mutants as measured by the competition assay mirrored their *in vitro* cytotoxicity. Although the *pilT* and *pilU* strains colonize the lungs as well as or better than the parental strain, the hyperpiliated mutants appear defective in the ability to spread to or colonize the liver. This observation suggests that death caused by the *pilU* or *pilT* strain in this animal model results from localized lung damage, although our data do not rule out bacteremia and colonization of distant organs at later times during infection.

In summary, inactivation of *P. aeruginosa pilT* and *pilU* genes caused a reduction in adherence to and cytotoxicity toward three epithelial cell types and in ability to colonize the liver following intranasal inoculation in an animal model of acute pneumonia. PilT and PilU thus contribute to *P. aeruginosa*-induced cell damage *in vitro* and *in vivo*. As one probable function of PilT and PilU is to catalyze the retraction of pili, the results indicate that this process is likely necessary for full cytotoxic capability. Since pili are adhesins, they may be necessary to cause initial interactions between the bacteria and epithelial cells, and the proposed retraction of pili could then allow a formation of a more intimate association via other nonpilus adhesins. Evidence suggests that EPEC use such a mechanism of retraction of their bundle-forming pili, catalyzed by the PilT homolog BfpF, to interact with host cells. The bundle-forming pili of EPEC allow an initial, weak interaction of the bacteria with epithelial cells (29) that may be necessary for intimate association involving the adhesin intimin and the bacterium-produced receptor Tir (21). In EPEC, and possibly in *P. aeruginosa*, this intimate association may be necessary for subsequent translocation of effectors by type III secretion. It should be noted that the hyperpiliated *P. aeruginosa pilT* and *pilU* mutants still retain more cytotoxicity and virulence than nonpiliated strains. This finding implies that the overexpression of abnormally functioning pili can partially correct for their lack of retraction. However, our results also indicate that this compensation is due to an uncharacterized mechanism other than increased adherence to host cells.

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