

Quorum sensing and motility mediate interactions between *Pseudomonas aeruginosa* and *Agrobacterium tumefaciens* in biofilm cocultures

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Communicated by E. Peter Greenberg, University of Washington School of Medicine, Seattle, WA, January 3, 2006 (received for review July 6, 2005)

In the environment, multiple microbial taxa typically coexist as communities, competing for resources and, often, physically associated within biofilms. A dual-species cocultivation model has been developed by using two ubiquitous and well studied microbes *Pseudomonas aeruginosa* (*P.a.*) and *Agrobacterium tumefaciens* (*A.t.*) as a tractable system to identify molecular mechanisms that underlie multispecies microbial associations. Several factors were found to influence coculture interactions. *P.a.* had a distinct growth-rate advantage in cocultures, increasing its relative abundance during planktonic and biofilm growth. *P.a.* also demonstrated a slight quorum-sensing-dependent increase in growth yield in liquid cocultures. *P.a.* dominated coculture biofilms, “blanketing” or burying immature *A.t.* microcolonies. *P.a.* flagellar and type IV pili mutant strains exhibited deficient blanketing and impaired competition in coculture biofilms, whereas, in planktonic coculture, these mutations had no effect on competition. In contrast, *A.t.* used motility to emigrate from coculture biofilms. In both planktonic and biofilm cocultures, *A.t.* remained viable for extended periods of time, coexisting with its more numerous competitor. These findings reveal that quorum-sensing-regulated functions and surface motility are important microbial competition factors for *P.a.* and that the outcome of competition and the relative contribution of different factors to competition are strongly influenced by the environment in which they occur.

flagella | interspecies interactions

In theoretical and empirical ecology, competition between species plays a central role in defining community structure and activity. Stable coexistence of diverse organisms in communities is thought to be fostered by individual tradeoffs and optimization of competitive strategies along resource gradients (1). Outside of the laboratory, microorganisms usually coexist in multicellular communities, governed by competition for common nutritional resources with other community members (2). Competitive fitness can be realized simply by occupying a suitable or specialized nutritional niche. Motility provides a mechanism by which microbes continually reposition themselves, adapting to changing nutritional and physical conditions. Another effective competitive strategy is to secrete antimicrobial compounds that kill or impair other species that occupy the same niche. Competitive interactions are most likely to occur when local microbial population densities are high, such as in biofilm communities. At high population densities, the process of quorum sensing is an important mechanism that coordinates and reinforces community behaviors in many bacterial species. Biofilm formation and quorum sensing are two bacterial community behaviors that clearly have significant potential to influence multispecies interactions (3–5). We hypothesize that motility, quorum sensing, and biofilm formation are among the mechanisms by which bacteria compete and persist within microbial communities.

To examine this hypothesis, we have developed a dual-species model system, composed of *Pseudomonas aeruginosa* (*P.a.*) and

Agrobacterium tumefaciens (*A.t.*). *P.a.* is a Gram-negative γ -proteobacterium ubiquitous in soil and aquatic environments. It is also an opportunistic pathogen that causes many nosocomial infections and is frequently associated with the chronic lung infections that plague people suffering from cystic fibrosis. *P.a.* is a paradigm for the study of acyl-HSL-based quorum sensing and the formation of surface-associated communities called biofilms (6, 7). *A.t.* is a Gram-negative α -proteobacterium that causes crown gall disease in plants. This microbe has served as a model for horizontal gene transfer, host–microbe interactions, pathogenesis, and acyl-HSL-based signaling for many years. *P.a.* and *A.t.* have been isolated from the same environment, where they coexist as common residents of freshwater, bulk soil, and the rhizosphere (8–10).

We report here, that *P.a.* manifests a significant competitive advantage over *A.t.*, simply through its rapid growth rate in laboratory culture. We show that other functions can influence competition independently of growth rate. Our studies reveal important roles for quorum sensing and surface motility in the competitive interaction of *P.a.* with *A.t.* in cocultures. Examination of an *A.t.* flagellar-motility mutant suggests a very different role for *A.t.* swimming in biofilm coculture interactions.

Results

***P.a.* Dominates Planktonic Coculture Interactions.** Because growth rate is a key variable for interpreting subsequent experiments, the doubling times of *P.a.* and *A.t.* were determined in a defined growth medium supplemented with three distinct carbon sources: glucose, succinate, and glutamate. For all three carbon sources tested, *P.a.* grew at least twice as quickly as *A.t.* (see Table 2, which is published as supporting information on the PNAS web site). The carbon source succinate was used in subsequent experiments, where the doubling times of *P.a.* and *A.t.* were estimated to be 29 and 59 min, respectively.

We initially examined the population dynamics of wild-type *P.a.* and *A.t.* planktonic cocultures. Cocultures were inoculated with different ratios (ranging from 10:1 to 1:10) of wild-type *P.a.* and *A.t.* The relative percentage of *A.t.* was determined in the coculture at three points during the growth curve. In all tested cocultures, the percentage of *A.t.* decreased over time (Fig. 1 and data not shown). In a coculture inoculated with a 1:1 ratio, *A.t.* dropped from 50% of total cells to $\approx 16\%$ in late stationary phase. To determine whether our laboratory strain of *P.a.* (originally a clinical isolate) exhibited coculture trends generally representative of other *P.a.* strains, four environmental isolates (see Table 2) were also examined in coculture with *A.t.* Both pure

Conflict of interest statement: No conflicts declared.

Abbreviations: *A.t.*, *Agrobacterium tumefaciens*; CFU, colony-forming unit; *P.a.*, *Pseudomonas aeruginosa*.

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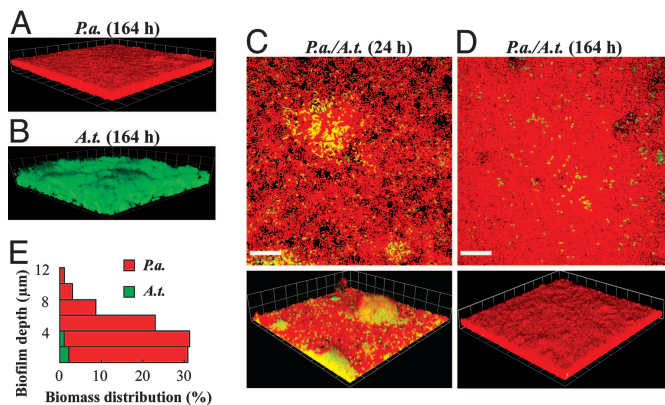


Fig. 2. *P.a.* blankets *A.t.* in coculture biofilms. (A and B) Three-dimensional views of *P.a.* and *A.t.* wild-type pure-culture flow-cell biofilms. (C and D) Coculture biofilms at 24 and 164 h after inoculation. (Top) An *x-y* slice close to the attachment surface. (Bottom) Three-dimensional views. Red cells, *P.a.*, green cells, *A.t.* (Scale bars, 20 μm ; 1 unit in 3D view = 13.4 μm .) (E) A quantitative determination of biomass distributions along biofilm depth in a coculture biofilm at 96 h.

A.t. biomass proceeded to decline to $\approx 1\%$ of the total coculture biomass, remaining at that level for extended periods (≈ 10 d) (Fig. 3D; and see Table 5, which is published as supporting information on the PNAS web site; and data not shown). Viability staining of the 164-h coculture biofilm showed that the majority of cells in the biofilm were alive, indicating that the blanketed *A.t.*, although spatially confined, remain viable (data not shown). Complete blanketing was not observed when *P.a.* was added after *A.t.* had established a biofilm. The addition of planktonic *P.a.* to preestablished pure-culture *A. t.* biofilms (164

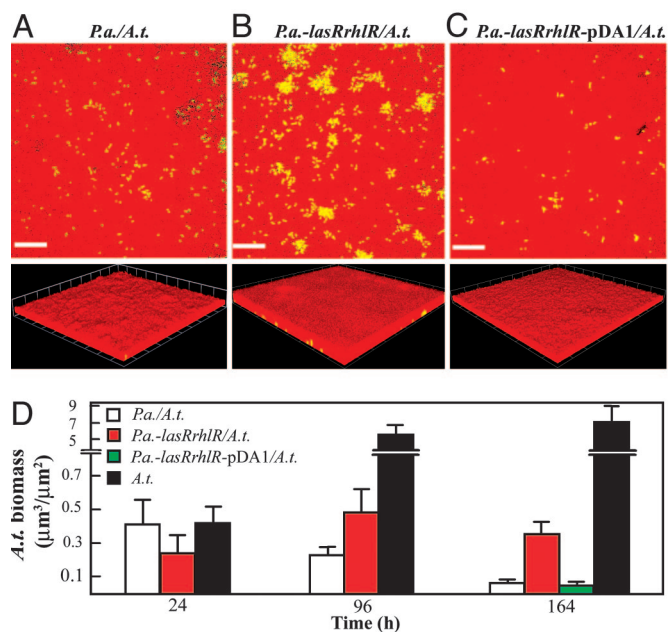


Fig. 3. Quorum sensing provides an advantage to *P.a.* in coculture biofilms. (A–C Top) An *x-y* slice close to the attachment surface. (A–C Bottom) Three-dimensional views. (A) A coculture biofilm of wild-type *P.a.* and *A.t.* (B) A coculture biofilm of *P.a.-lasRrhIR* and wild-type *A.t.* (C) A coculture biofilm of *P.a.-lasRrhIR* complemented with pDA1 and wild-type *A.t.* Red cells, *P.a.*, green cells, *A.t.* (Scale bars, 20 μm ; 1 unit in 3D view = 13.4 μm .) All micrographs were taken at 164 h. (D) COMSTAT determination of the relative amount of *A.t.* biofilm biomass present in pure and coculture biofilms.

h) resulted in significant *P.a.* colonization, although complete blanketing was never achieved (data not shown).

***P.a.* Quorum Sensing Plays a Role in Older Coculture Biofilms.** We next examined whether *P.a.* quorum sensing played a role in biofilm cocultures. The *P.a.-lasRrhIR* mutant strain was used for these studies, because the *P.a.-lasIrhlI* strain has a known defect in twitching motility unrelated to quorum sensing (19). In pure culture, *P.a.-lasRrhIR* had a growth rate similar to wild-type and formed biofilms with similar structure (Table 4). In 24-h biofilms, the amount of *A.t.* biomass present in the coculture with *P.a.-lasRrhIR* was comparable with the wild-type *P.a./A.t.* coculture (Fig. 3D). However, at later time points, the amount of *A.t.* biomass in the *P.a.-lasRrhIR* coculture biofilm remained constant, in contrast to the wild-type *P.a.* coculture, where *A.t.* biomass continued to decrease (compare Fig. 3A and B; and see Fig. 3D and Table 5). Complementation of the *P.a.-lasRrhIR* double mutant with pDA1 restored wild-type biofilm coculture phenotypes (Fig. 3C).

Motility Confers a Competitive Advantage to *P.a.* in Coculture Biofilms. We hypothesized that blanketing by *P.a.* was mediated by surface motility. *P.a.* is capable of moving on a surface using two separate types of motility. Twitching motility is mediated by type-IV pili, whereas swarming motility involves type IV pili, flagella, and secreted surfactants called rhamnolipids (20, 21). Both swarming and twitching motilities have been implicated in *P.a.* biofilm development (17, 22, 23). *A.t.* can swim via flagella, but has not been reported to exhibit surface motility. Two *P.a.* motility mutants were studied in both pure and coculture biofilms. A mutant with a nonfunctional *pilA* gene, encoding the pilin subunit, is unable to produce a type IV pilus and shows no twitching and reduced swarming motility. The *P.a. flgK* gene encodes a flagellar hook protein. A *flgK* mutant is unable to produce flagella and is defective in both swarming and swimming motilities.

Similar to previous reports, the *P.a.-pilA* pure-culture biofilm was clearly heterogeneous and thick and had a high degree of surface roughness compared with the wild-type parent (Fig. 4A and Table 4) (17, 24). In coculture biofilms, *P.a.-pilA* did not exhibit complete blanketing but, instead, colonized discrete patches on top of *A.t.* microcolonies (Fig. 4B). *A.t.* biomass present in the *P.a.-pilA* coculture biofilm was slightly greater than with the wild-type *P.a.* at 96 h; however, at 164 h, levels of *A.t.* biomass were similar in both cocultures (Fig. 4E and Table 5). In pure culture, the *P.a.-flgK* strain formed flat biofilms very similar to wild-type (compare Figs. 2A and 4C; Table 4). In coculture biofilms, *P.a.-flgK* exhibited reduced blanketing, with much of the *A.t.* exposed to the overlying bulk liquid (Fig. 4D). *A.t.* biomass present in the *P.a.-flgK* coculture biofilm was significantly higher than the *P.a.* coculture at 164 h (Fig. 4E and Table 5). Analysis of sagittal profiles of the *P.a.-flgK/A.t.* coculture revealed that *A.t.* represented a significant portion of the biomass (30–50%) throughout the biofilm (Fig. 4F). The *pilA* and *flgK* mutants were successfully complemented *in trans* with pDA2 (*P_{tac}::pilA*) and pDA3 (*P_{tac}::flgK*), respectively (data not shown). Unlike biofilm cocultures, both *flgK* and *pilA* mutant strains were indistinguishable from wild-type *P.a.* in planktonic coculture (data not shown).

Influence of *A.t.* Flagellar Motility on Coculture Biofilms. To compare the role of *A.t.* and *P.a.* motility functions for liquid and biofilm coculture interactions, *A.t.* bearing a mutation in the *fliR* gene (defective in swimming motility) was examined (25). In liquid culture, this strain had an identical growth rate and the same coculture phenotype as the wild-type *A.t.* strain (data not shown). In pure-culture biofilms, this strain exhibited an attachment-deficient phenotype relative to wild-type *A.t.*, although it

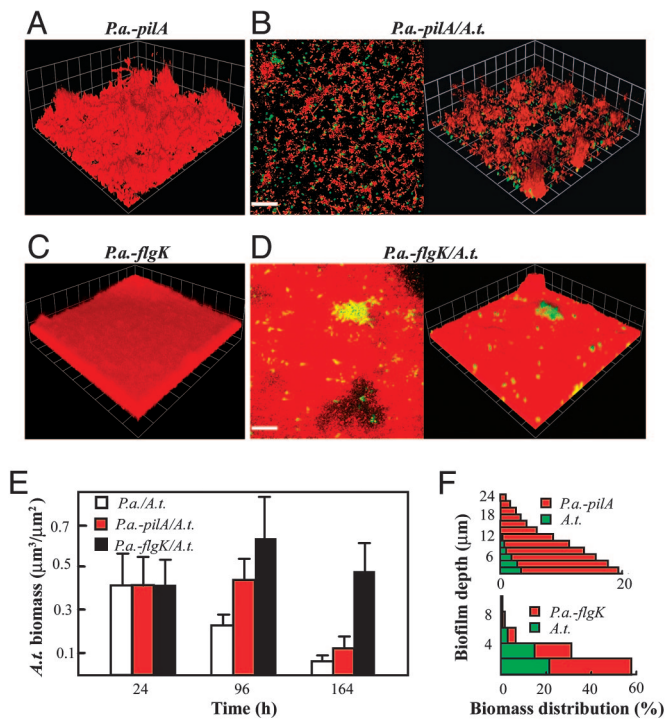


Fig. 4. *P.a.* motility is required for blanketing. (A and C) Three-dimensional views of a *P.a.-pilA* (A) and *P.a.-flgK* (C) pure-culture biofilms. (B and D) *x-y* slices close to the attachment surface and 3D views of a *P.a.-pilA/A.t.* (B) and *P.a.-flgK/A.t.* (D) coculture biofilms. Red cells, *P.a.*; green cells, *A.t.* (Scale bars, 20 μm ; 1 unit in 3D view, 13.4 μm .) All micrographs were taken at 164 h. (E) COMSTAT determination of the relative amount of *A.t.* biofilm biomass present in cocultures. (F) A quantitative determination of biomass distributions along biofilm depth in *P.a.-pilA/A.t.* and *P.a.-flgK/A.t.* coculture biofilms at 96 h.

ultimately formed biofilms with only slightly less biomass than wild-type at 164 h (compare Fig. 2B with Fig. 5A and C and Fig. 8, which is published as supporting information on the PNAS web site). In coculture, *P.a.* blanketed the *A.t.-fliR* mutant and

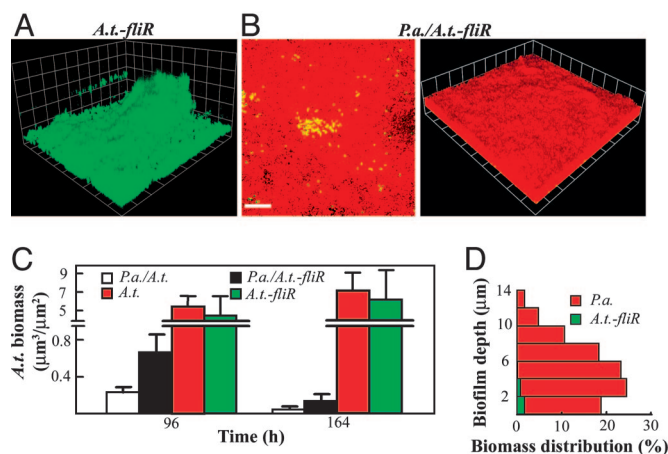


Fig. 5. Biofilm coculture phenotypes of an *A.t.-fliR* mutant strain. (A) A 3D-view of an *A.t.-fliR* pure-culture biofilm; (B) An *x-y* slice close to attachment surface and a 3D-view of a *P.a./A.t.-fliR* coculture biofilm; Red cells, *P.a.*; green cells, *A.t.* (Scale bars, 20 μm ; 1 unit in 3D view, 13.4 μm .) All micrographs were taken at 164 h. (C) COMSTAT determination of the relative amount of *A.t.* biofilm biomass present in pure cultures and cocultures. (D) A quantitative determination of biomass distributions along biofilm depth in a *P.a./A.t.-fliR* coculture biofilm at 96 h.

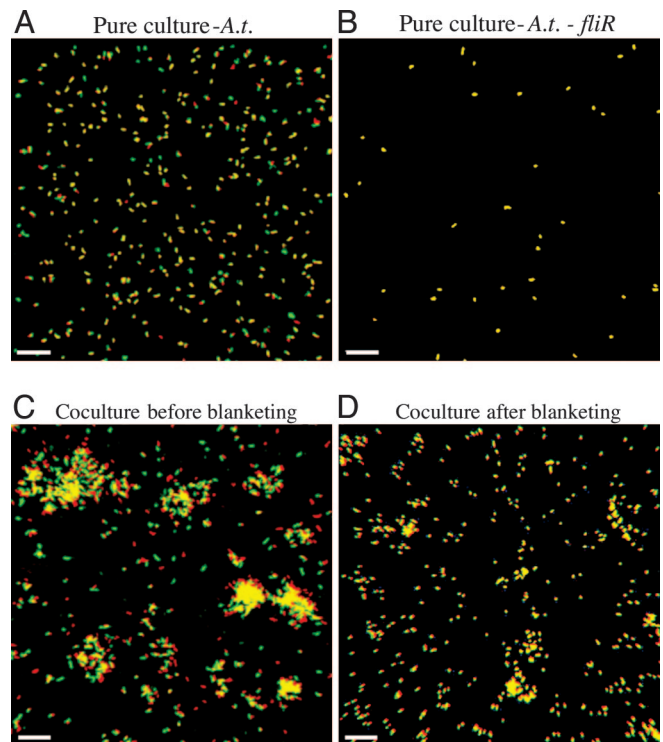


Fig. 6. Different biofilm time-lapse microscopy series. Yellow cells indicate bacteria that have not moved during the course of the time series. Green cells indicate bacteria that were present in the field of view at the beginning of the time course but not at the end. Red cells are bacteria present at the end of the time series that were not present at the start. (A and B) A 1-h time course of a newly inoculated pure-culture biofilm of wild-type *A.t.* (A) or *A.t.-fliR* (B). (Scale bar, 12 μm .) (C and D) A time course of wild-type *A.t.* in coculture with *P.a.* (*P.a.* cells are not visible). (C) A 3-h time course before blanketing has occurred. (D) A 3-h time course after complete blanketing has occurred. (Scale bar, 20.2 μm .)

eventually formed a biofilm with similar structure and biomass distribution as with wild-type *A.t.* (compare Figs. 2D and 5B; Figs. 2E and 5D). Interestingly, at earlier time points, there is more *A.t.-fliR* biomass in the coculture biofilms than observed for *A.t.* (Fig. 5C). We hypothesized that this difference might reflect the repulsion or emigration of wild-type *A.t.* away from surfaces colonized by *P.a.* in earlier coculture biofilms, for which the *A.t.-fliR* strain would be deficient.

To test this hypothesis, various experimental approaches were tried. Initially, collection of effluent from the flow-cell system failed to show significant emigration of *A.t.* from the coculture biofilm (data not shown). However, time-lapse microscopy revealed that, at early time points in pure and coculture biofilms, *A.t.* has the capacity to detach and leave (Fig. 6A and C; and see Movies 1 and 2, which are published as supporting information on the PNAS web site). Cells from the bulk liquid were also observed to attach to the surface after the time series was initiated. On the other hand, once *A.t.-fliR* attached to the surface, no detachment was observed in pure and cocultures (Fig. 6B; and see Movie 3, which is published as supporting information on the PNAS web site; and data not shown). In coculture with *P.a.*, wild-type *A.t.* was distinctly motile before blanketing and was observed to gradually leave the biofilm. Analysis of time-lapse series revealed an ≈ 10 –15% decrease of *A.t.* biofilm biomass over a 3-h period before complete blanketing (Fig. 6C and data not shown). It is not clear whether *P. a.* induces detachment of *A.t.* in biofilm cocultures. The loss in *A.t.* biofilm biomass in cocultures may simply be due to normal

detachment rates coupled with a decrease or halt in *A.t.* biofilm growth.

In older coculture biofilms, after blanketing had occurred, both *A.t.* and *A.t.-fliR* were immobilized (Fig. 6D; and see Movie 4, which is published as supporting information on the PNAS web site; and data not shown). Ultimately, at later time points, the amount of *A.t.-fliR* present in the coculture biofilm was only slightly greater than the wild-type *A.t.* (Fig. 5C and Table 5).

Discussion

In this study, we have begun to examine the complex interactions between two common environmental microorganisms, *P.a.* and *A.t.*, in planktonic and biofilm growth modes. During exponential growth in dispersed, liquid culture, *P.a.* dominated *A.t.* because of a higher growth rate. Within biofilms grown on glass surfaces in the same defined medium, *P.a.* was also found to numerically dominate the population and to cover adherent *A.t.*, a process requiring motility via flagella and type IV pili. Quorum-sensing mutants displayed an impaired competition phenotype in both liquid and flow-cell biofilm cultures. Motility was found to be important for both species in coculture biofilms. Although *A.t.* was outnumbered after the rapid-growth phase in both growth formats, its population remained viable, leading to a period of coexistence of these two microbes.

Quorum sensing appears to allow *P.a.* to achieve a slightly higher growth yield in liquid cocultures. Several quorum-sensing-regulated secreted functions related to nutrient acquisition might explain this observation. Quorum sensing regulates multiple secreted proteases, which might act to degrade *A.t.* exoproducts, which could then serve as a nutrient source (26). An alternative possibility is that *P.a.* may have a lower K_s value than *A.t.* for key limiting nutrients in coculture. Quorum-sensing-regulated functions may affect the K_s values for these substrates. An example is iron acquisition by the quorum-sensing-controlled siderophore pyoverdine, which is used to secure iron in a form less accessible to other species (although the defined medium used in this study is iron replete).

A *P.a.* quorum-sensing mutant strain was impaired in biofilm cocultures. The quorum-sensing mutant strain coculture biofilm displayed a modest increase in the amount of *A.t.* biomass. One potential explanation is that the dense carpet of *P.a.* cells covering *A.t.* produces a quorum-sensing-regulated toxic compound(s) that kills or inhibits its growth. A similar role for quorum sensing has been shown for *Pseudomonas aureofaciens*, which uses quorum-sensing-regulated phenazine antibiotics to compete with the local flora of the wheat rhizosphere (27). Filtered supernatants of *P.a.* liquid cultures failed to inhibit or kill *A.t.* However, we cannot rule out the possibility that direct cell contact is necessary for killing/inhibition or the high cell densities of *P.a.* present in biofilms produced an elevated, lethal concentration of a secreted antimicrobial(s).

In flow-cell biofilms, *P.a.* formed confluent mats on the biofilm surface that eventually submerged *A.t.* microcolonies. The blanketing phenotype was impaired in both *flgK* and *pilA* mutants, suggesting that both of these surface appendages and, by extension, the motility they provide, are required for this phenomenon. However, the severity of the competitive deficiency was different between the two mutants. The amount of *A.t.* biofilm biomass remained relatively constant in coculture with the *flgK* mutant, whereas, in *pilA* cocultures, the amount of *A.t.* biomass decreased over time. In the *flgK* coculture *A.t.* was present at the surface of the biofilm, whereas, in the *pilA* coculture, it localized to the biofilm depths. Therefore, the success of *A.t.* in the *flgK* coculture may result from its exposure to nutrients present in the overlying liquid medium. The pure culture biofilms of wild-type *P.a.* and the *flgK* mutant were structurally indistinguishable, whereas the *flgK* mutant differed substantially from the wild-type in the coculture biofilms, sug-

gesting that some functions have little impact on the development of pure-culture biofilms, but can have a significant impact in the development of biofilms in a mixed-species environment. Another important point is that *flgK* and *pilA* mutants did not exhibit a competitive defect in planktonic coculture. This observation highlights the importance of the microenvironment in dictating key competitive factors.

Surprisingly, disruption of flagellar motility in *A.t.* did not augment the dominance of *P.a.* in flow-cell biofilm coculture. In pure culture, such a mutation has a distinct adherence defect (Fig. 8). Instead, in biofilm cocultures, the *A.t.-fliR* mutant accumulated more initial biomass on the surface (Fig. 5C), although this biomass was eventually subsumed, as with the wild type, by the rapidly moving *P.a.* cells. Our time-lapse microscopy observations of early stages of surface colonization suggest that a significant fraction of *A.t.* cells contacting and transiently adhering to the surface is highly motile, and many *A.t.* cells tend to swim away from the surface. In contrast, once bound to the surface, the *fliR* mutant is much more stable, with few emigrating cells. It is intriguing to speculate that, in coculture biofilms with *P.a.*, *A.t.* flagellar motility may be more important as a mechanism of escape than as a factor that enhances surface colonization. These data suggest that there may be two distinct mechanisms that result in the diminished *A.t.* biomass observed in older coculture biofilms. The first could be the erosion of biomass because of emigration of *A.t.* from the biofilm before blanketing. The second is the loss of *A.t.* biomass after blanketing (not attributable to emigration from the biofilm), perhaps because of a quorum-sensing-regulated function (as seen in Fig. 3).

The dual-species format used in this study is a powerful approach for identifying key attributes that allow specific microbes to effectively compete and coexist in different environments. Indeed, other laboratories are increasingly taking advantage of reconstituting simple, defined multispecies laboratory model systems to gain insight into microbial interactions (28–30). Our findings identified *P.a.* quorum sensing and motility via pili and flagella as functions that contribute to its competitive interactions with *A.t.* in our binary coculture system. We are currently exploring whether our observations in laboratory coculture represent general features of common competitive interactions that occur in the environment. For example, do other species use quorum sensing to regulate their competitive interactions with cohabiting microbes in dense, polymicrobial environments? If so, the degradation or sequestration of acyl-HSLs within these environments, now well documented for several diverse bacteria, could be particularly influential (31). Ultimately, by using observations extrapolated from simple, tractable coculture model systems, we hope to identify important aspects of microbial interactions in complex systems.

Materials and Methods

Bacterial Strains and Media. The *P.a.* and *A.t.* strains used in this study are described in Table 2 (and see *Supporting Text*, which is published as supporting information on the PNAS web site). Bacterial cultures were grown in AT minimal medium (32). The carbon sources were 0.5% glucose, 0.4% succinate, or 0.5% glutamate. Where indicated, before coculture assembly, acyl-HSL was added and dried in the culture flask for a working concentration of 5 μ M for C4-HSL or 2 μ M for 3OC12-HSL. For supernatant killing assays, 5 ml of a 24-h-old stationary-phase culture was passed through 0.22- μ m filters (Millex-GV; Millipore, Bedford, MA), and the filtrate was added to washed stationary phase *A.t.* cells. *A.t.* viability was measured by plate counts.

Planktonic Pure-Culture and Coculture Growth Curves. All growth curves were conducted at 30°C and performed in triplicate. Cocultures were inoculated by growing pure cultures to log phase

