

1 **Virulence factor expression patterns in *Pseudomonas aeruginosa* strains from infants with**  
2 **cystic fibrosis**

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23

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26

1 **ABSTRACT**

2 *Pseudomonas aeruginosa* is the leading cause of morbidity and mortality in cystic fibrosis (CF).  
3 This study examines the role of organism-specific factors in the pathogenesis of very early *P.*  
4 *aeruginosa* infection in the CF airway. One-hundred-and-sixty-eight longitudinally collected *P.*  
5 *aeruginosa* isolates from children diagnosed with CF following newborn screening were genotyped  
6 by pulsed-field gel electrophoresis and phenotyped for 13 virulence factors. Associations between  
7 virulence factors and gender, exacerbation, persistence, timing of infection and infection site were  
8 assessed using multivariate regression analysis. Ninety-two strains were identified. Persistent  
9 strains showed significantly lower levels for pyoverdine, rhamnolipid, haemolysin, total protease,  
10 and swimming and twitching motility, than strains eradicated by aggressive antibiotic treatments.  
11 Initial strains had higher levels of virulence factors, and significantly higher phospholipase C, than  
12 subsequent genotypically-different strains at their initial isolation. Strains from males had  
13 significantly lower pyoverdine and swimming motility than females. Colony size was significantly  
14 smaller in strains isolated during exacerbation than those isolated during non-exacerbation periods.  
15 All virulence factors were higher and swimming motility significantly higher, in strains from  
16 bronchoalveolar-lavage (BAL) and oropharyngeal sites than BAL-alone. Using unadjusted  
17 regression modelling, age at initial infection and age at strain isolation showed U-shaped profiles  
18 for most virulence factors. Among subsequent strains, the longer the time since initial infection, the  
19 lower the level of most virulence factors. This study provides new insight into virulence factors  
20 underpinning impaired airway clearance seen in CF infants despite aggressive antibiotic therapy.  
21 This information will be important in the development of new strategies to reduce the impact of *P.*  
22 *aeruginosa* in CF.

23

## 1 INTRODUCTION

2 *Pseudomonas aeruginosa* airway infection of young children with cystic fibrosis (CF) is typically  
3 intermittent, involving multiple non-mucoid strains that are sensitive to anti-pseudomonal  
4 antibiotics. However, one strain eventually becomes established and by adolescence almost 80% of  
5 CF patients are chronically-infected, resulting in increased morbidity and mortality [1].

6  
7 Many host factors affect the course of infection in the CF lung [2], but there is limited  
8 understanding about organism-specific traits affecting infectivity, severity of infection and  
9 persistence. The environment is the main source for acquiring *P. aeruginosa*, suggesting that  
10 environmental isolates have all the virulence determinants to initiate infection and cause disease [3].  
11 *P. aeruginosa* adapts to long-term survival in the CF lung by overproducing alginate, forming  
12 biofilms and down-regulating virulence factors [4-5]. Tracking of adaptive mutations in isolates  
13 from chronically-infected adults showed defects in swimming and twitching motility, pyocyanin  
14 secretion and biofilm formation [6]. However, *P. aeruginosa* strains infecting infants and young  
15 children have not been systematically examined to establish the characteristics underpinning  
16 pathogenesis of early infection in the CF lung.

17  
18 Here we present the virulence factor phenotype of 168 *P. aeruginosa* isolates collected  
19 longitudinally over the first five-years of life from infants diagnosed with CF after newborn  
20 screening and classified by genotyping as 92 distinct strains. The Type 3 Secretion System profiles  
21 of this cohort were published previously [7]. Relationships were sought between virulence factor  
22 expression and clinical factors.

## 1 MATERIALS AND METHODS

### 2 Clinical isolates

3 Investigations were conducted on 168 *P. aeruginosa* isolates from 58 children aged 3-63 (mean  
4 26.8) months collected within the Australasian Cystic Fibrosis Bronchoalveolar Lavage (ACFBAL)  
5 randomised controlled trial [8]. Isolates came from 150 specimens: 57 bronchoalveolar lavage  
6 (BAL) fluid (35 children); 82 oropharyngeal (OP) swabs (44 children) and 11 sputa (9 children  
7 post-ACFBAL study). On average, 2.5 (range 1-18) isolates were tested per child. Ethics  
8 Committees of all participating hospitals approved the study.

9  
10 The ACFBAL study randomised infants diagnosed with CF after newborn screening to either BAL-  
11 directed therapy or standard care using clinical judgment and OP cultures [8]. BAL-directed infants  
12 underwent BAL at enrolment before six-months of age, and then with exacerbations requiring  
13 hospitalisation, if OP cultures grew *P. aeruginosa*, and after *P. aeruginosa* eradication therapy. All  
14 subjects had OP cultures taken during exacerbations and after eradication therapy, while each  
15 underwent BAL at age five. Children with positive *P. aeruginosa* BAL ( $\geq 10^3$  colony-forming  
16 units/ml) or OP (standard group only) cultures received two-weeks of intravenous tobramycin with  
17 either ticarcillin-clavulanate, or ceftazidime, followed by four-weeks of oral ciprofloxacin and eight  
18 weeks of nebulised tobramycin inhalation solution [8]. After treatment, BAL-directed children had  
19 further BAL and OP cultures, while those receiving standard care had OP cultures alone. If *P.*  
20 *aeruginosa* persisted, the eradication protocol was repeated, but if this also failed the child was  
21 categorised as chronically-infected. Samples were processed by hospital laboratories where *P.*  
22 *aeruginosa* identification was undertaken using standard techniques and interpretative criteria [9].

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## 1 **Genotypic testing**

2 Isolates were genotyped using pulsed-field gel electrophoresis (PFGE) following *SpeI* digestion  
3 [10]. Analyses were performed using cluster analysis software (GelComparII™, Applied Maths,  
4 Belgium) and the criteria of Tenover et al [11].

## 5 **Phenotypic tests**

- 6 i. **Pyocyanin** Overnight *P. aeruginosa* grown in Cation-Adjusted Mueller-Hinton Broth  
7 (CAMHB) (Oxoid Ltd, Australia) were scored (0-4) for intensity of green pigment: 0 = no pigment,  
8 1 = light green, 2 = moderate green, 3 = dark green, 4 = very dark green [12].
- 9 ii. **Pyoverdine** 5µl of overnight CAMHB cultures were spotted onto Kings B agar (Oxoid),  
10 dried and incubated at 37°C. Plates were viewed under a transilluminator after 24h and scored on  
11 the intensity of the fluorescent yellow pigment produced by pyoverdine using a graded system: 0=  
12 no fluorescence, 1 = lightly fluorescent yellow, 2 = moderately fluorescent yellow, 3 = strongly  
13 fluorescent yellow, 4 = very strongly fluorescent yellow [13].
- 14 iii. **Haemolysin** 5µl of overnight CAMHB culture was spotted onto Columbia Horse Blood  
15 Agar (Oxoid), dried and incubated for 20h at 37°C. Two perpendicular diameters of clearance zones  
16 around colonies were measured and the area ( $\pi r^2$ ) calculated (mm<sup>2</sup>). This method was utilised in iv  
17 to viii below.
- 18 iv. **Total Protease** 5µl of overnight CAMHB culture were spotted onto a 15%v/v skim milk  
19 agar plate dried and incubated for 24h at 37°C.
- 20 v. **Elastase** 5µl of overnight CAMHB culture were spotted onto 10ml elastin agar plates, dried  
21 and incubated for 48h at 37°C.
- 22 vi. **Phospholipase C** 5µl of overnight CAMHB culture were spotted onto egg yolk agar plates,  
23 dried and incubated for 20h at 37°C [14-15].
- 24 vii. **Rhamnolipid** 5µl of overnight CAMHB culture were spotted onto M9-based agar plates,  
25 supplemented with 0.2% (v/v) glucose, 2 mM MgSO<sub>4</sub>, 0.05% (v/v) tyrosine, 0.0005% (v/v)

1 methylene blue, 0.02% (v/v) and cetyltrimethylammonium bromide dried and incubated for 48h at  
2 37°C. Colonies were scored as positive by the presence of a violet halo around the colony.

3 viii. **Twitching and swimming motilities** Bacterial patches from 1.5%w/v Luria plates were  
4 scraped with a toothpick and stab inoculated into 10ml (1%w/v Luria-broth) agar. Twitching  
5 motility (light haze of growth at the agar/plate interface) was measured after 24h at 37°C. For  
6 swimming, toothpick scrapings were spotted onto centre of a swim (0.3%w/v Luria-broth) plate and  
7 swim diameters measured after overnight at 30°C.

8 ix. **Swarming motility** Pre-warmed swarm plates (0.8%w/v nutrient broth, 0.5%w/v agar and  
9 0.5%w/v glucose) were inoculated with a colony tooth-picked from a swim plate (above). After 24h  
10 at 37°C, swarming motility was identified as spreading growth from the inoculation site and graded  
11 0-4 where; 0 = no motility, 1= very limited motility. 2 = fair motility, 3 = strongly motile, 4 = very  
12 strongly motile.

13 x. **Biofilm mass**  $10^7$  cells were inoculated into wells of a microtitre plate containing 100µl  
14 CAMHB and plates incubated for 24h at 37°C. Biofilm mass was stained with 1%v/v crystal violet,  
15 ethanol-extracted and absorbance read at  $O_{D595}$ . Readings were then normalised as a percentage of  
16 *P. aeruginosa* PAO1 biofilm grown concurrently.

17 xii. **Colony size** 100µl aliquots of a  $10^{-7}$  dilution of overnight *P. aeruginosa* culture were spread  
18 onto CAMHB plates and incubated at 37°C. Colony diameter was measured in mm after 24h.

19 xiii. **Mucoidy** Cells were scraped from 1.5%w/v Luria-broth patch plate, then patched onto  
20 5%v/v glycerol MacConkey plates (Oxoid) and scored (+/-) for mucoidy after 48h at 37°C.

21

## 22 **Statistical analysis**

23 Patients were followed longitudinally. Each isolate was tested (×3) for each outcome of interest.

24 Nine continuous and three categorical measures of virulence (measured using semi-quantitative

25 scores from 0-to-4) were analysed. These scores were categorised into binary variables to assess the

1 likelihood of having a score of 3 or 4 versus any score below 3. As only four isolates were mucoid,  
2 statistical analysis was not conducted on the fourth categorical measure (mucoidy).

3  
4 The associations between a set of independent factors and each of the virulence measures were  
5 assessed using multivariate regression analysis. The independent factors were gender, exacerbation  
6 at sample collection time, initial vs subsequent strain infection, persistent strain (defined as one  
7 detected continuously or intermittently for at least three-months) and site of sample isolation (BAL  
8 only, BAL+OP, OP-only).

9  
10 We further investigated whether: i) the child's age at acquisition of first infection, ii) the age of the  
11 child at the collection of a strain and iii) time since acquisition of the first strain, had any effect on  
12 the outcomes of interest. The associations between these three time-related variables and each of the  
13 virulence measures were estimated separately using unadjusted regression analysis.

14  
15 Population-averaged panel-data regression models and generalized estimation equations (GEE)  
16 were used to assess the associations of interest. Linear or logistic regression was used depending on  
17 the measure of the outcome of interest. Regression analyses were executed using STATA 12.0  
18 (StataCorp, College Station, TX, USA).

1 **RESULTS**

2 The 58 children fell into three categories. Twenty-eight had only one strain detected on a single  
3 occasion; 10 had one strain detected on multiple occasions and the remaining 20 had multiple  
4 strains detected on multiple occasions.

5 **Genotype**

6 Ninety-two strains were identified by PFGE. Of these, 90 strains had unique pulsotypes, while the  
7 remaining two strains had macrorestriction patterns indistinguishable from Aust-01 and Aust-02,  
8 which are highly-prevalent strains found frequently in older, chronically infected patients attending  
9 Australian CF clinics [16]. Only one patient had Aust-01, and another two had the Aust-02 strain  
10 identified. Since transmissibility and other properties of these clonal strains may have biased  
11 results, all analyses were repeated without them and results remained comparable (data not shown).

12  
13 Of the 92 strains, 52 (56.5%) were initial infecting strains detected in children at an average age of  
14 26.8 (range 3-63, standard deviation 14.6) months. Seventy-three strains (79.3%) came from an  
15 exacerbation, 22 strains (23.9%) were persistent (14 were persistent from the initial infection and  
16 eight from a subsequent infection), 59 (64.1%) cleared, while the persistence of 11 (12.0%) strains  
17 was unknown as genotyping was limited to the available sample set. Thirty-five strains were from  
18 the upper airways (OP) only, 31 from the lower airways collected by BAL and 18 were found in  
19 both sites (eight strains from follow-up sputa were excluded from analyses for site, as the quality of  
20 these specimens was unknown and they may have contained both upper and lower airway isolates).

21 **Phenotype**

22 Pyocyanin was scored at 3 or 4 in 20.5% of the tests, pyoverdine was scored at 3 or 4 in 31.8% of  
23 the tests and swarming was scored at 3 or 4 in 12.6% of the tests. The distributions of all continuous  
24 measures of interest are presented in Table 1.

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1 *Persistent strains versus cleared strains*

2 Multivariate analysis (Table 2) showed that the levels of six virulence factors: pyoverdine,  
3 rhamnolipid, haemolysin, total protease and swimming and twitching motilities were significantly  
4 lower in persistent strains compared to cleared strains, while another six factors; pyocyanin,  
5 elastase, phospholipase C, biofilm mass, colony size and swarming motility showed non-significant  
6 reductions. No factor was significantly higher in persistent strains compared to cleared strains.

7  
8 *Effects of gender, timing of isolation, site of infection and exacerbation on phenotype*

9 With respect to gender, pyoverdine expression and swimming motility were significantly lower in  
10 strains from males compared to females (Table 2). In comparisons of initial and subsequent  
11 (genotypically-different) strains at their first isolation, initial strains had higher expression of all  
12 virulence factors, except pyocyanin, and this reached statistical significance for phospholipase C.  
13 All virulence factors showed an increase in strains from both OP and BAL compared to BAL-only,  
14 while swimming motility was significantly higher. Only smaller colony size was associated with  
15 exacerbation.

16  
17 *Influence of age of child and time of isolation on phenotype*

18 Table 3 shows changes in virulence factor expression and Figure 1 shows the changes for the eight  
19 clearance-zone based measurements, with respect to: age at initial strain isolation, age at isolation of  
20 a strain and time since initial strain isolation.

21  
22 *Age at initial strain isolation:* Phospholipase C, haemolysin and total protease levels were  
23 significantly lower in initial strains acquired at age 12-23 months compared with  $\leq 12$ -months, but  
24 were not different at age  $>36$ -months compared with  $\leq 12$ -months (Table 3, part 1). Pyoverdine and  
25 swarming motility were lower in initial strains acquired at 24-35 months compared with  $\leq 12$ -  
26 months, but were the same at  $\geq 36$  months as at  $\leq 12$  months. Biofilm mass increased with age of

1 initial strain isolation, becoming significant at 24-35 months. Of the eight virulence factors  
2 measured by area of clearance zone, all except biofilm mass produced a non-linear, non-monotonic  
3 U-shaped curve (Figure 1A).

4  
5 *Age at isolation of a strain:* Expression of pyoverdine and haemolysin, and colony size were  
6 significantly lower in strains from children aged 24-35-months compared to children aged  
7  $\leq 12$  months. Total protease expression was significantly lower in both 12-23 and 24-35-month-old  
8 children compared with children aged  $\leq 12$  months (Table 3, part 2). Of the eight virulence factors  
9 measured by area of clearance zone (Figure 1B) all except swimming and twitching motility gave a  
10 U-shaped effect.

11  
12 *Time since isolation of initial strain;* The longer the time, the lower the level of most virulence  
13 factors (Table 3, part 3). Significant decreases occurred for biofilm mass and swimming motility by  
14 age 11-months, for twitching motility and colony size by 23-months and for elastase, phospholipase  
15 C, haemolysin and total protease by  $>36$ -months. Figure 1C, shows an initial fall, followed by a  
16 stable period, with a sharp decline at  $>36$ -months, for all factors except rhamnolipid. The presence  
17 of subsequent infections does not explain the sharp decline in virulence observed in the  $\geq 36$   
18 month's group. Subsequent strains were isolated from patients between 2 and 97 months after initial  
19 strains (mean=22.8 months, median=15 months), meaning that most subsequent infections occurred  
20 before 36 months.

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## 1 **DISCUSSION**

2 *P. aeruginosa* virulence factors are usually associated with acute infection within the host. The  
3 complex relationships between virulence factor expression and pathogenesis of airway infection in  
4 CF have been investigated mainly in chronically infected adults [17-19]. Our study is the first to  
5 comprehensively examine *P. aeruginosa* virulence factor expression profiles in CF infants and  
6 young children.

### 7 *Virulence versus persistence, initial/subsequent strain, gender, exacerbation and site of infection*

8 This study provides the first evidence that virulence factor expression profiles of *P aeruginosa*  
9 strains at their first isolation in the CF airway of young children may help predict the ability of the  
10 strain to persist in the face of the immune response and antibiotic therapy. Expressions of  
11 pyoverdine, rhamnolipid, haemolysin, total protease and swimming and twitching motilities were  
12 all significantly lower in persister than clearer strains and this finding was independent of potential  
13 confounders such as site of isolation (upper or lower airway) and whether the airway was *P.*  
14 *aeruginosa* naive or potentially damaged by a previous *P. aeruginosa* strain. Since these persister  
15 strains were not clonally related to each other or to frequent clones circulating in Australian clinics  
16 they are assumed to have been environmentally acquired. *P. aeruginosa* is known to down-regulate  
17 these virulence factors as a survival mechanism in the CF lung [17, 20]. However it seems unlikely  
18 that such adaptations begin so early in the infection process. Perhaps instead some environmental  
19 strains first colonise the sinuses where adaptation to the local microenvironment takes place before  
20 aspiration into the lower airways occurs. Studies show pyoverdine-negative mutants increase with  
21 lung colonisation time [21], which would indicate that persisters do not have a greater requirement  
22 for pyoverdine. Rhamnolipid is a biosurfactant involved in protecting cells against oxidative stress  
23 [22] and in increased swarming [23]. It is also cytotoxic to eukaryotic cells, decreases liquid surface  
24 tension and facilitates access to nutrients within biofilms [24-25]. Since these functions are  
25 important in long-term bacterial survival, the down regulation of rhamnolipid is noteworthy.  
26 Decreased haemolysin expression has been correlated with increased mucoidy [26], however all

1 persistent strains in our study were non-mucoid. Some bacterial proteases promote mucoidy [27], so  
2 strains may begin to display mucoidy as persistence continues.

3  
4 The overall virulence profiles of initial strains were higher than those of subsequent strains at their  
5 first isolation. In particular, phospholipase C was significantly higher in initial strains.  
6 Phospholipase C is a heat-labile lecithin-degrading haemolysin. The phospholipase C/sphingo-  
7 myelinase gene-pair acts as an inhibitor of lung pulmonary surfactant [28], thus phospholipase C  
8 may be required to establish the initial, but not subsequent infection.

9  
10 Associations with gender proved interesting. Previous reports had indicated high virulence factor  
11 expression during early CF lung disease played a role in the worse prognosis of females [29],  
12 however there was no evidence of this in our analysis. The significance of the lower pyoverdine  
13 expression and lower swimming motility in strains from males is unknown. There was a lack of  
14 association between virulence factor expression and exacerbation other than for colony size, and  
15 this was at odds with our early study of CF where higher extracellular enzyme levels were found in  
16 hospitalised than non-hospitalised CF adolescent patients with chronic *P. aeruginosa* infection [30].  
17 We found no significant associations between expression and site of infection, except for swimming  
18 motility being greater in strains from both BAL and OP than those from BAL-alone. These  
19 conflicting results could reflect differences in age, disease stage and the small number of non-  
20 exacerbation samples.

21  
22 Flagella and pili-driven twitching motility are required for biofilm formation [31]. Alternatively,  
23 swarming motility is inversely proportional to biofilm formation [32]. Our finding that swarming  
24 motility and biofilm mass were not significantly different, while twitching and swimming motility  
25 were significantly reduced in persisters highlights differences between early persisters in young  
26 children and chronic infection in CF adults where biofilm formation is a distinctive feature [33].

1 *Influence of age of child and time of isolation on phenotype*

2 Particularly noteworthy were associations between age and virulence factor expression (Table 3 and  
3 Figures 1A 1B 1C). The initial reduction in virulence factor production with increasing age of initial  
4 infection (Figure 1A) could be an adaptation to the host immune response. However, the increase in  
5 almost all factors in initial strains in the older age groups was unexpected. This was particularly  
6 marked for haemolysin, which appears independent of age of acquiring the initial strain. A U-  
7 shaped curve was also evident in the analyses of virulence by age at isolation of a strain (Figure  
8 1B), with strains from children >36-months showing a modest rise after an initial fall for all factors  
9 except swimming and twitching motilities, and haemolysin again showing a significant fall at 24-35  
10 months followed by a rise at  $\geq 36$  months. While we are unsure of the reasons for this U-shaped  
11 effect, the fall in virulence factor expression seen in new *P. aeruginosa* isolates from older children,  
12 could offer a partial explanation. As the sinuses develop and grow in CF children, they may act as a  
13 reservoir for initial and newly acquired *P. aeruginosa* strains, where adaptation to the lung  
14 environment may already begin prior to them infecting lower airways [34-35].

15  
16 In contrast, results of the analyses examining time since isolation of initial strain (Figure 1C), which  
17 showed significant decreased expression for eight of the twelve virulence factors by the  $\geq 36$  month  
18 time period. The significant decreases start from the earliest time point (1-11 months) in the case of  
19 biofilm mass and swimming. These results are comparable to those found by a study of *P.*  
20 *aeruginosa* isolates from CF children of undisclosed ages [36], except that this trend was evident at  
21 24 months compared with over >36-months in our study. Interestingly, four virulence factors,  
22 (pyocyanin, pyoverdine, swarming and rhamnolipid) remained virtually unchanged over the entire  
23 period.

24  
25 This study provides important new information on factors influencing virulence factor expression of  
26 *P. aeruginosa* infecting the CF airway in the first years of life. While it is generally accepted that *P.*

1 *aeruginosa* loses its virulence in the transition to chronic infection in CF, our results which show a  
2 a diverse range of up and downregulation as well as maintenance of expression amongst the twelve  
3 virulence factors over time, challenge this view. It seems likely that there is considerable diversity  
4 within microbial population evolution, with some strains retaining or enhancing virulence and  
5 others losing virulence over time. Our data suggest that this process begins within 36 or so months  
6 after initial infection.

7  
8 Our findings may help in understanding which virulence factors are important in establishing  
9 infection in CF infants where, despite aggressive antibiotic therapy, impaired airway clearance  
10 mechanisms exist. This in turn will be important in developing new more effective strategies to  
11 reduce the impact of *P. aeruginosa* on morbidity and mortality in CF.

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3 Trust. Cynthia B. Whitchurch was supported by a NHMRC Career Development Award and a  
4 NHMRC Senior Research Fellowship.

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6

1 **CONFLICT OF INTEREST**

2 All authors declare that they have no conflict of interest.

3

4

1 **TABLES AND FIGURES**

2 **Table 1: Statistical distributions of all continuous measures of interest**

3

<b>Variable</b>	<b>Mean</b>	<b>Std. Dev.</b>	<b>Min</b>	<b>Max</b>
Elastase (mm <sup>2</sup> )	82.8	40.5	0	144.0
Rhamnolipid (mm <sup>2</sup> )	60.7	28.0	0	125.3
Haemolysin (mm <sup>2</sup> )	84.8	63.2	0	233.1
Total Protease (mm <sup>2</sup> )	87.1	30.2	0	134.4
Phospholipase C (mm <sup>2</sup> )	90.4	39.0	0	155.0
Biofilm mass (% of PAO1 biomass)	88.6	49.0	11.3	358.3
Swimming (mm <sup>2</sup> )	69.6	35.5	0	198.8
Twitching (mm <sup>2</sup> )	60.6	43.2	0	136.5
Colony Size (diameter in mm)	2.6	1.3	0.5	6.0

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1 **Table 2: Multivariate analysis of associations between a set of independent variables and individual virulence factors**

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<b>Independent variables</b>	<b>Pyoverdine OR<sup>a</sup> (95% CI)</b>	<b>Pyocyanin OR<sup>a</sup> (95% CI)</b>	<b>Swarming OR<sup>a</sup> (95% CI)</b>	<b>Elastase AEC<sup>b</sup> (95% CI)</b>
<b>Gender:</b> Male vs Female	<b>0.48 (0.23-0.99)*</b>	0.87 (0.36-2.11)	0.50 (0.18-1.41)	-5.55 (-19.6-8.85)
<b>Exacerbation:</b> Yes vs No	0.53 (0.20-1.42)	1.48 (0.42-5.13)	0.65 (0.21-2.02)	-8.9 (-26.58-8.79)
<b>Strain:</b> Initial vs Subsequent	1.64 (0.55-4.89)	0.55 (0.16-1.88)	1.62 (0.28-9.34)	17.84 (-3.84-39.52)
<b>Strain:</b> Persister vs clearer	<b>0.21 (0.08-0.56)**</b>	0.77 (0.27-2.17)	0.50 (0.16-1.55)	-11.03 (-28.40-6.34)
<b>Sample:</b> OP vs BAL	1.34 (0.43-4.16)	1.53 (0.39-5.99)	0.40 (0.07-2.20)	3.7 (-17.88-25.27)
<b>Sample:</b> (BAL+OP) vs BAL	3.28 (0.83-12.88)	3.27 (0.05-12.9)	1.48 (0.31-7.05)	8.0 (-16.89-32.90)
<b>Independent variables</b>	<b>Total Protease AEC<sup>b</sup> (95% CI)</b>	<b>Rhamnolipid AEC<sup>b</sup> (95% CI)</b>	<b>PLC<sup>c</sup> AEC<sup>b</sup> (95% CI)</b>	<b>Haemolysin AEC<sup>b</sup> (95% CI)</b>
<b>Gender:</b> Male vs Female	2.75 (-7.23-12.7)	-6.99 (-16.40-2.42)	7.0 (-6.12-20.12)	-2.32 (-23.2-18.56)
<b>Exacerbation:</b> Yes vs No	-3.49 (-16.1-9.11)	-1.6 (-13.48-10.28)	-8.05 (-24.61-8.51)	-12.13 (-38.48-14.22)
<b>Strain:</b> Initial vs Subsequent	11.37 (-4.08-26.82)	9.25 (-5.31-23.83)	<b>26.08 (5.78-46.39)*</b>	15.99 (-16.31-48.3)
<b>Strain:</b> Persister vs clearer	<b>-12.92 (-25.30--0.54)*</b>	<b>-12.32 (-23.99--0.64)*</b>	-9.72 (-25.99-6.55)	<b>-32.93 (-58.82--7.04)*</b>
<b>Sample:</b> OP vs BAL	3.03 (-12.34-18.40)	-4.87 (-19.37-9.63)	-7.8 (28.1-12.40)	-10.1 (-42.24-22.06)
<b>Sample:</b> (BAL+OP) vs BAL	10.63 (-7.10-28.37)	8.42 (-8.31-25.15)	2.79 (-20.52-26.10)	19.54 (-17.55-56.63)
<b>Independent variables</b>	<b>Swimming AEC<sup>b</sup> (95% CI)</b>	<b>Twitching AEC<sup>b</sup> (95% CI)</b>	<b>Biofilm Mass AEC<sup>b</sup> (95% CI)</b>	<b>Colony Size AEC<sup>b</sup> (95% CI)</b>
<b>Gender:</b> Male vs Female	<b>-13.95 (-25.55--2.36)*</b>	-2.71 (-17.14-11.71)	1.19 (-16.34-18.72)	0.23 (-0.04-0.5)
<b>Exacerbation:</b> Yes vs No	0.3 (-14.33-14.94)	2.98 (-15.22-21.19)	-14.32 (-37.86-9.21)	<b>-0.4 (-0.74--0.07)*</b>
<b>Strain:</b> Initial vs Subsequent	16.90 (-1.04-34.84)	21.61 (-0.7-43.9)	14.76 (-11.7-41.22)	0.11 (-0.3-0.53)

1 **Table 3: Change in virulence factor expression over time**  
 2 **Part 1: Age at initial strain isolation**

Virulence factor	Units	<12mths	12-23mths	pvalue <sup>a</sup>	24-35mths	pvalue <sup>a</sup>	36+mths	pvalue <sup>a</sup>
Pyocyanin	%	23.08	26.79	0.720	12.12	0.271	4.35	0.069
Pyoverdine	%	42.31	28.57	0.207	20.20	0.057†	46.38	0.767
Swarming	%	12.8	14.3	0.850	2.0	<b>0.027*</b>	17.4	0.643
Elastase	mm <sup>2</sup>	85.27	72.63	0.158	89.45	0.672	102.44	0.112
Rhamnolipid	mm <sup>2</sup>	65.23	59.85	0.412	59.22	0.407	68.50	0.679
PLC <sup>b</sup>	mm <sup>2</sup>	97.26	79.52	<b>0.044*</b>	92.92	0.656	114.04	0.115
Haemolysin	mm <sup>2</sup>	103.78	71.50	<b>0.017*</b>	69.53	<b>0.008**</b>	132.45	0.080
Total Protease	mm <sup>2</sup>	94.52	80.15	<b>0.026*</b>	85.66	0.213	102.28	0.317
Biofilm Mass	% <sup>c</sup>	73.47	83.05	0.402	110.14	<b>0.004**</b>	90.11	0.265
Swimming	mm <sup>2</sup>	70.40	72.12	0.836	66.33	0.658	74.20	0.705
Twitching	mm <sup>2</sup>	67.82	53.53	0.167	63.76	0.723	66.69	0.928
Colony Size	mm	2.5	2.83	0.077	2.49	0.941	2.74	0.288
<b>Part 2: Age at isolation of a strain</b>								
Virulence factor	Units	<12mths	12-23mths	pvalue <sup>a</sup>	24-35mths	pvalue <sup>a</sup>	36+mths	pvalue <sup>a</sup>
Pyocyanin	%	28.57	33.33	0.752	17.39	0.372	14.75	0.238
Pyoverdine	%	54.76	36.67	0.245	18.12	<b>0.008**</b>	34.43	0.152
Swarming	%	14.3	13.3	0.932	8.0	0.477	15.3	0.924
Elastase	mm <sup>2</sup>	94.33	75.31	0.147	80.82	0.275	85.28	0.451
Rhamnolipid	mm <sup>2</sup>	63.86	58.86	0.574	55.33	0.309	64.84	0.904
PLC <sup>b</sup>	mm <sup>2</sup>	101.29	85.21	0.200	85.01	0.169	94.43	0.550
Haemolysin	mm <sup>2</sup>	115.72	87.01	0.137	56.49	<b>0.001**</b>	98.03	0.317
Total Protease	mm <sup>2</sup>	100.41	81.45	<b>0.047*</b>	81.51	<b>0.036*</b>	91.07	0.285
Biofilm Mass	% <sup>c</sup>	85.28	83.14	0.894	88.46	0.832	92.29	0.634
Swimming	mm <sup>2</sup>	74.37	72.51	0.871	70.12	0.696	66.72	0.469
Twitching	mm <sup>2</sup>	81.43	64.42	0.221	60.70	0.114	53.09	<b>0.031*</b>
Colony Size	mm	3.07	2.95	0.626	2.54	<b>0.025*</b>	2.44	<b>0.006**</b>

**Part 3: Time since isolation of initial strain**

Virulence factor	Units	0mths	1-11mths	pvalue <sup>a</sup>	12-23mths	pvalue <sup>a</sup>	24-35mths	pvalue <sup>a</sup>	+36mths	pvalue <sup>a</sup>
Pyocyanin	%	18.03	20.00	0.822	19.05	0.918	0.00	0.061	33.33	0.202
Pyoverdine	%	40.98	26.67	0.180	25.40	0.171	18.18	0.166	26.67	0.309
Swarming	%	13.7	11.1	0.725	4.8	0.292	18.2	0.695	8.9	0.552
Elastase	mm <sup>2</sup>	89.88	79.36	0.203	89.20	0.942	100.84	0.367	49.75	<b>0.000***</b>
Rhamnolipid	mm <sup>2</sup>	62.07	59.60	0.690	60.23	0.794	66.77	0.607	66.93	0.545
PLC <sup>b</sup>	mm <sup>2</sup>	97.72	88.68	0.287	95.62	0.827	96.20	0.903	65.56	<b>0.003**</b>
Haemolysin	mm <sup>2</sup>	96.88	76.96	0.147	83.34	0.385	90.92	0.767	59.88	<b>0.037*</b>
Total Protease	mm <sup>2</sup>	91.50	84.34	0.249	88.99	0.722	94.35	0.755	73.82	<b>0.028*</b>
Biofilm Mass	% <sup>c</sup>	95.07	70.61	<b>0.033*</b>	92.71	0.850	100.82	0.721	85.02	0.480
Swimming	mm <sup>2</sup>	78.01	61.71	<b>0.034*</b>	68.68	0.284	75.34	0.812	58.90	0.054†
Twitching	mm <sup>2</sup>	70.96	62.13	0.354	45.34	<b>0.018*</b>	55.89	0.282	42.67	0.054†
Colony Size	mm	2.86	2.83	0.874	2.43	<b>0.027*</b>	2.64	0.375	1.93	<b>0.000***</b>

1 <sup>a</sup> Wald statistic; p values obtained from unadjusted regression analysis <sup>b</sup> Phospholipase C <sup>c</sup> % of *P. aeruginosa* PAO1 biofilm grown concurrently  
2 \* p<0.05; \*\* p<0.01; \*\*\*p<0.001; †borderline p<0.05