

Table S3. The proportion of CDSs shared among genomes in the *Pseudomonas fluorescens* group^a

Strains	Pf-5	30-84	O6	Pf0-1	Q8r1-96	Q2-87	BG33R	SBW25	A506	SS101
Pf-5										
30-84	0.74									
O6	0.72	0.90								
Pf0-1	0.70	0.72	0.73							
Q8r1-96	0.67	0.69	0.69	0.69						
Q2-87	0.66	0.69	0.70	0.71	0.83					
BG33R	0.69	0.70	0.71	0.68	0.64	0.64				
SBW25	0.67	0.67	0.67	0.68	0.68	0.68	0.77			
A506	0.72	0.72	0.72	0.70	0.68	0.67	0.84	0.78		
SS101	0.72	0.72	0.72	0.70	0.67	0.67	0.83	0.78	0.87	

^a CDSs shared between each pair of genomes in the *P. fluorescens* group as a proportion of the total number of CDSs in the strain with the smallest genome. Shading designates comparisons between strains within a specific sub-clade: pink, Sub-clade 1; blue, Sub-clade 2; green, Sub-clade 3.