

Table S2. Genes shared by and unique to ten fully-sequenced strains in the *Pseudomonas fluorescens* group^a

Annotated function	Pf-5	30-84	O6	Pf0-1	Q8r1-96	Q2-87	BG33R	SBW25	A506	SS101
Conserved hypothetical protein	PFL_0092	Pchl3084_0090	PchlO6_0094	PfI01_0045	PfIQ8_0052	PfIQ2_0066	PseBG33_0108	PFLU0089	PfIA506_0088	PfISS101_0088
Biofilm PGA synthesis protein PgaA	PFL_0161	Pchl3084_0182	PchlO6_0181	PfI01_0177	PfIQ8_0168	PfIQ2_5565	PseBG33_0177	PFLU0143	PfIA506_0154	PfISS101_0155
Biofilm PGA synthesis lipoprotein PgaB	PFL_0162	Pchl3084_0183	PchlO6_0182	PfI01_0178	PfIQ8_0169	PfIQ2_5564	PseBG33_0178	PFLU0144	PfIA506_0155	PfISS101_0156
Biofilm PGA synthesis N-glycosyltransferase PgaC	PFL_0163	Pchl3084_0184	PchlO6_0183	PfI01_0179	PfIQ8_0170	PfIQ2_5563	PseBG33_0179	PFLU0145	PfIA506_0156	PfISS101_0157
Putative biofilm PGA synthesis protein PgaD	PFL_0164	Pchl3084_0185	PchlO6_0184	PfI01_0180	PfIQ8_0171	PfIQ2_5562	PseBG33_0180	PFLU0146	PfIA506_0157	PfISS101_0158
Conserved hypothetical protein	PFL_1293	Pchl3084_1294	PchlO6_1369	PfI01_1241	PfIQ8_1215	PfIQ2_4257	PseBG33_1466	PFLU1339	PfIA506_1296	PfISS101_4100
Conserved hypothetical protein	PFL_2043	Pchl3084_1970	PchlO6_2191	PfI01_1872	PfIQ8_3930	PfIQ2_1714	PseBG33_3750	PFLU4227	PfIA506_3549	PfISS101_3585
Transcriptional regulator, DeoR family	PFL_2429	Pchl3084_2237	PchlO6_2468	PfI01_2141	PfIQ8_2635	PfIQ2_2551	PseBG33_3183	PFLU3612	PfIA506_2356	PfISS101_3062
Response regulator	PFL_2454	Pchl3084_2272	PchlO6_2498	PfI01_3529	PfIQ8_2054	PfIQ2_2059	PseBG33_4589	PFLU5118	PfIA506_4409	PfISS101_4477
Putative cytochrome P450 oxidoreductase	PFL_2514	Pchl3084_2315	PchlO6_2537	PfI01_3472	PfIQ8_3443	PfIQ2_3353	PseBG33_2095	PFLU3256	PfIA506_2140	PfISS101_2021
Conserved hypothetical protein	PFL_2570	Pchl3084_2373	PchlO6_2593	PfI01_3422	PfIQ8_3419	PfIQ2_2108	PseBG33_2088	PFLU2168	PfIA506_2134	PfISS101_2014
DNA-binding response regulator	PFL_3269	Pchl3084_3225	PchlO6_3454	PfI01_2892	PfIQ8_3453	PfIQ2_3362	PseBG33_3035	PFLU2722	PfIA506_2968	PfISS101_2399
Transcriptional regulator, LysR family	PFL_3291	Pchl3084_2626	PchlO6_2861	PfI01_2190	PfIQ8_3335	PfIQ2_2187	PseBG33_3106	PFLU2649	PfIA506_2457	PfISS101_2240
GDP-mannose pyrophosphatase NudK / conserved domain protein	PFL_3331	Pchl3084_2574	PchlO6_2807	PfI01_3567	PfIQ8_5806	PfIQ2_5681	PseBG33_3179	PFLU3608	PfIA506_2359	PfISS101_3059
Conserved hypothetical protein	PFL_3866	Pchl3084_3665	PchlO6_3958	PfI01_3573	PfIQ8_3620	PfIQ2_1921	PseBG33_3124	PFLU3541	PfIA506_2422	PfISS101_2998
Conserved hypothetical protein	PFL_4007	Pchl3084_3794	PchlO6_4089	PfI01_3715	PfIQ8_3935	PfIQ2_1712	PseBG33_2051	PFLU2131	PfIA506_2096	PfISS101_1977
Transporter, CPA2 family	PFL_4229	Pchl3084_4091	PchlO6_4345	PfI01_3967	PfIQ8_1711	PfIQ2_3768	PseBG33_1826	PFLU1748	PfIA506_1790	PfISS101_1749
Aminotransferase, class V	PFL_4607	Pchl3084_4478	PchlO6_4719	PfI01_4359	PfIQ8_1258	PfIQ2_4212	PseBG33_4327	PFLU4835	PfIA506_4147	PfISS101_4223
Putative lipoprotein	PFL_5371	Pchl3084_5148	PchlO6_5415	PfI01_4896	PfIQ8_5035	PfIQ2_4771	PseBG33_4824	PFLU5351	PfIA506_4646	PfISS101_4700
Conserved hypothetical protein	PFL_6244	Pchl3084_2038	PchlO6_2252	PfI01_1927	PfIQ8_3792	PfIQ2_1840	PseBG33_3664	PFLU4152	PfIA506_3461	PfISS101_3478

^a Genes are present in genomes of the *P. fluorescens* group strains, but are not present in the genomes of other *Pseudomonas* spp. in Figure 1.