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Transcriptional regulator, GntR familyPfl01_4144PflQ8_4104PflQ2_1539Tat (twin-arginine translocation) pathway signal sequence domain proteinPfl01_4469PflQ8_4593PflQ2_1056Membrane protein, putativePfl01_4512PflQ8_4475PflQ2_1171Autoinducer-binding transcriptional regulator, LuxRValueValueValue		Pfl01_3959	PfIQ8_1719	PfIQ2_3760
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Autoinducer-binding transcriptional regulator, LuxR		Pfl01_4469	PfIQ8_4593	PfIQ2_1056
	Membrane protein, putative	Pfl01_4512	PflQ8_4475	PflQ2_1171
family PfI01_4889 PfIQ8_5023 PfIQ2_4761	Autoinducer-binding transcriptional regulator, LuxR			
	family	Pfl01_4889	PflQ8_5023	PfIQ2_4761
	Conserved hypothetical protein	Pfl01_5217		PfIQ2_0504
		Pfl01_5218	PflQ8_5381	PfIQ2_0503
Oxidoreductase, GMC family PfI01_5219 PfIQ8_5382 PfIQ2_0502			PflQ8_5382	PfIQ2_0502

Table S7. Genes shared by and unique to Sub-clade 2^a

^a Genes are present in genomes of the strains found in Sub-clade 2, but are not present in the genomes of other *Pseudomonas* spp. in Figure 1.