

Bacterial Phylum/Class	Related Species	Accession #	Similarity (%)	Clones	
Alphaproteobacteria	<i>Agrobacterium</i> sp. Mei-Qs6	EF090606	99	1	
	<i>Hyphomicrobium vulgare</i>	AB543807	95-98	7	
	<i>Hyphomicrobium</i> sp. TW4	AY934492	96	3	
	<i>Nordella</i> sp. P-63	AM411927	96	3	
	<i>Rubellimicrobium aerolatum</i>	EU338486	95	1	
	<i>Rhizobiaceae</i> bacterium KVD-unk-43	DQ490336	95-96	2	
	<i>Rhizobium</i> sp. 3-4	FJ598330	98	1	
	<i>Pleomorphomonas oryzae</i>	AB159685	99	1	
	<i>Agrobacterium tumefaciens</i>	DQ993282	96	1	
	<i>Mesorhizobium</i> sp. ORS1080	AJ295082	91	1	
	<i>Holospira obtusa</i>	X58198	92	1	
	Alpha proteobacterium CRIB-02	DQ123619	98	2	
	Gammaproteobacteria	<i>Acidithiobacillus ferrooxidans</i>	AJ459800	85	1
		<i>Legionella busanensis</i>	NR_025196	91	6
<i>Legionella taurinensis</i>		DQ667196	84-86	2	
<i>Legionella longbeachae</i>		FN650140	91	1	
<i>Legionella impletisoli</i>		AB233209	87	1	
<i>Legionella</i> sp. HB09011		GU319886	85	1	
<i>Legionella yabuuchiae</i>		AB233212	91	1	
<i>Legionella lytica</i>		NR_026334	95	1	
Clostridia	<i>Clostridium</i> sp. SL206	EU816420	94	1	
	<i>Clostridium intestinale</i>	AY781385	99	4	
	<i>Clostridium chromoreductans</i>	AY228334	99	1	
	<i>Clostridium</i> sp. MK11	AB275140	97	2	
	<i>Clostridium</i> sp. CYP5	DQ479415	98	2	
	<i>Clostridium saccharolyticum</i>	FJ957875	97	2	
	<i>Clostridium</i> sp. FRC C11	EU331374	99	1	
	<i>Clostridium</i> sp. BL-26	DQ196630	99	1	
Flavobacteria	<i>Flavobacterium</i> sp. TISTR	AB465580	95-98	4	
Planctomycetacia	<i>Planctomyces</i> sp. (strain 599)	AJ231189	90-95	3	
	<i>Pirellula</i> sp. (Schlesner 678)	X81947	85-86	2	
	<i>Planctomyces</i> sp. (Schlesner 269)	X81953	89	1	
	<i>Planctomycete</i> str. 292	AJ231182	91	1	
Bacilli	<i>Geobacillus pallidus</i>	FJ808721	99	1	
TM7	TM7 phylum sp. oral taxon 355 clone	GU410617	87	2	
Total				67	