

Table S4 Genomes used in this study

Class	Bacterial species	strain	Host	Acc. Number
Betaproteobacteria	<i>Tremblaya princeps</i>	PCVAL	Planococcus citri	CP002918.1
	<i>Tremblaya phenacola</i>	PAVE	Phenacoccus avenae	CP003982.1 and CP003983.1
		PPER	Phenacoccus peruvianus	MKGN00000000*
	<i>Bordetella pertussis</i>	Tohama I	–	BX470248.1
	<i>Burkholderia mallei</i>	ATCC 23344	–	CP000010.1 and CP000011.2
	<i>Neisseria meningitidis</i>	MC58	–	AE002098.2
	<i>Nitrosomonas europaea</i>	ATCC 19718	–	AL954747.1
<i>Ralstonia solanacearum</i>	GMI1000	–	AL646052.1 and AL646053.1	
Gammaproteobacteria	<i>Erwinia amylovora</i>	CFBP1430	–	FN434113.1 and FN434114.1
	<i>Escherichia coli</i>	K-12 substr. MG1655	–	U00096.3
	<i>Klebsiella pneumoniae</i>	HS11286	–	CP003200.1 and CP003223.1 to CP003228.1
	<i>Serratia proteamaculans</i>	568	–	CP000826.1 and CP000827.1
	<i>Sodalis praecaptivus</i>		–	CP006569.1 and CP006570.1
	<i>Doolittlea endobia</i>	MHIR	Maconellicoccus hirsutus	LN999833.1
	<i>Gullanella endobia</i>	FVIR	Ferrisia virgata	LN999832.1
	<i>Mikella endobia</i>	PMAR	Paracoccus marginatus	LN999831.1
<i>Hoaglandella endobia</i>	TPER	Trionymus perrisii	LN999835.1	
Alphaproteobacteria	<i>Bartonella quintana</i>	RM-11	–	CP003784.1
	<i>Rickettsia prowazekii</i>	Madrid E	–	AJ235269.1

\*This Whole Genome Shotgun project has been deposited at DDBJ/ENA/GenBank under the accession MKGN00000000.

The version described in this paper is version MKGN01000000.