

Table S1. Accession numbers for DNA sequences used in this study

	<i>recA</i>	<i>mdh</i>	<i>katA</i>	<i>luxR-I</i> intergenic	<i>glpA-fdhA</i> intergenic
CG101	HQ595306	EU907966.1	EU907990.1	HQ436471	HQ535993
CG103	HQ595307	HQ595322	HQ595331	HQ436472	HQ535994
EM17	HQ595308	EU907967.1	EU907991.1	HQ436484	HQ535997
ES114	VF_0535 ^a	VF_0276 ^a	VF_A0009 ^a	HQ436469	CP000021.2 (bp 279820- 279517)
ES12	HQ595309	HQ595323	HQ595332	HQ436480	HQ535986
ES213	HQ595310	EU907971.1	EU907995.1	HQ436481	HQ535984
ES401	HQ595311	HQ595324	HQ595333	HQ436483	HQ535991
ET101	HQ595312	HQ595325	HQ595334	HQ436475	HQ535999
ET401	HQ595313	HQ595326	HQ595335	HQ436476	HQ535990
H905	HQ595314	EU907972.1	EU907996.1	HQ436478	HQ535998
MJ1S	HQ595315	HQ595327	HQ595336	HQ436470	HQ535987
MJ11	VFMJ11_0538 ^a	VFMJ11_0264 ^a	VFMJ11_A0023 ^a	HQ436479	CP001133.1 (bp 321512- 321209)
MJ1	HQ595316	HQ595328	HQ595337	HQ436485	HQ535996
PP3	HQ595317	HQ595329	HQ595338	HQ436468	HQ535988
SA1	HQ595318	EU907986.1	EU908010.1	HQ436473	HQ535992
SR5	HQ595319	EU907987.1	EU908011.1	HQ436482	HQ535985
VLS2	HQ595320	HQ595330	HQ595339	HQ436477	HQ535995
WH1	HQ595321	EU907988.1	EU908012.1	HQ436474	HQ535989
LF11238	VSAL_I0634 ^a	VSAL_I0359 ^a	VSAL_II0215 ^a	NA ^b	FM178380.1 (bp 445307- 445005)

^a Locus tag of full gene given in table; partial sequence used for alignment.

^b NA indicates “not applicable” – this sequence was too divergent to provide a reasonable alignment and was not included in phylogenetic reconstructions