

Supplementary tables

Table S1: Primers used for Multi Locus Sequences Typing of *Wolbachia*

Genes	Designation	Primer sequences 5'-3'	References
gatB	gatB_F1	GAKTTAAAYCGYGCAGGBGTT	[6]
	gatB_R1	TGGYAAAYTCRGGYAAAGATGA	
coxA	coxA_F1	TTGGRGCRATYAACTTTATAG	[6]
	coxA_R1	CTAAAGACTTTKACRCCAGT	
hcpA	hcpA_F1	GAAATARCAGTTGCTGCAAA	[6]
	hcpA_R1	GAAAGTYRAGCAAGYTCTG	
ftsZ	ftsZ_F1	ATYATGGARCATATAAARGATAG	[6]
	ftsZ_R1	TCRAGYAATGGATTRGATAT	
fbpA	fbpA_F1	GCTGCTCCRCTTGGYWTGAT	[6]
	fbpA_R1	CCRCCAGARAAAAYYACTATTC	

Table S2: Blast results of fbpA sequences with those of *Wolbachia* strains in the database

<i>Wolbachia</i> species	Size of fragments (bp)	Query coverage	E-value	Identity	Accession number
<i>Wolbachia</i> sp. <i>wRi</i> . Complete genome	490 498	100% 99%	0.0 0.0	90% 89%	CP001391.1
<i>Wolbachia</i> endosymbiont of <i>Drosophila simulans wHa</i> . Complete genome	490 498	100% 99%	0.0 0.0	90% 89%	CP003884.1
<i>Wolbachia</i> endosymbiont of <i>Hyla</i> <i>variegatus</i> isolate <i>HVa</i> fructose- bisphosphate aldolase (fbpA) gene	490 498	96% 94%	0.0 7e- 179	91% 90%	KP183278.1
<i>Wolbachia pipientis</i> voucher <i>wol_sc_02621</i> fructose-bisphosphate aldolase (fbpA) gene	490 498	93% 91%	0.0 3e- 177	91% 90%	KX843420.1