

Table S1. 28S gene sequence identities of mealybugs in this study and other mealybug species.

Species	GenBank Accession nos.	Sequences identity (%) to mealybugs in this study
<i>Dysmicoccus neobrevipes</i>	KX015048	99.68%
<i>Dysmicoccus neobrevipes</i>	KX015042	99.68%
<i>Dysmicoccus neobrevipes</i>	KX015055	99.68%
<i>Dysmicoccus neobrevipes</i>	KX015040	99.68%
<i>Planococcus citri</i>	GU134660	99.68%
<i>Dysmicoccus brevipes</i>	GU134658	94.43%
<i>Pseudococcus viburni</i>	GU134653	94.39%
<i>Pseudococcus viburni</i>	GU134652	94.08%
<i>Pseudococcus nr maritimus</i>	GU134655	93.17%
<i>Pseudococcus calceolariae</i>	GU168802	90.31%
<i>Pseudococcus longispinus</i>	GU134657	90.06%
<i>Pseudococcus nr microadonidum</i>	GU134656	90.06%
<i>Pseudococcus comstocki</i>	GU134654	89.81%
<i>Vryburgia rimariae</i>	GU134659	86.05%
<i>Planococcus minor</i>	GU134661	85.14%
<i>Planococcus ficus</i>	GU134662	84.88%
<i>Phenacoccus parvus</i>	GU134663	77.84%