

Appendix S1: Sequence alignments for 372 base pairs of *Mycetophylax simplex* wingless gene from 51 specimens (GenBank accession numbers: KP939178-KP939228). The 51 individuals sequenced here are the same individuals sequenced for COI gene. The alignment included none variable sites and none parsimony informative sites. When phased in the DnaSP program the data show only eight segregating sites and none number of polymorphic segregating sites with more than two variants.

	1	10	20	30	40	50	60																																																							
mx001	G	T	T	T	C	G	A	C	G	G	G	G	C	T	T	C	T	C	T	C	G	G	G	T	A	A	T	G	G	T	C	A	G	T	A	A	C	T	C	G	G	A	A	C	G	A	G	C	G	C	G	T	T	A	C	C	A	A	C	A		
mx087	G	T	T	T	C	G	A	C	G	G	G	G	C	T	T	C	T	C	T	C	G	G	G	T	A	A	T	G	G	T	C	A	G	T	A	A	C	T	C	G	G	A	A	C	G	A	G	C	G	C	G	C	G	T	T	A	C	C	A	A	C	A
mx035	G	T	T	T	C	G	A	C	G	G	G	G	C	T	T	C	T	C	T	C	G	G	G	T	A	A	T	G	G	T	C	A	G	T	A	A	C	T	C	G	G	W	R	C	G	W	G	C	G	C	G	C	G	T	T	A	C	C	A	A	C	A
mx004	G	T	T	T	C	G	A	C	G	G	G	G	C	T	T	C	T	C	T	C	G	G	G	T	A	A	T	G	G	T	C	A	G	T	A	A	C	T	C	G	G	W	R	C	G	W	G	C	G	C	G	C	G	T	T	A	C	C	A	A	C	A
mx008	G	T	T	T	C	G	A	C	G	G	G	G	C	T	T	C	T	C	T	C	G	G	G	T	A	A	T	G	G	T	C	A	G	T	A	A	C	T	C	G	G	A	A	C	G	A	G	C	G	C	G	C	G	T	T	A	C	C	A	A	C	A
mx042	G	T	T	T	C	G	A	C	G	G	G	G	C	T	T	C	T	C	T	C	G	G	G	T	A	A	T	G	G	T	C	A	G	T	A	A	C	T	C	G	G	A	A	C	G	A	G	C	G	C	G	C	G	T	T	A	C	C	A	A	C	A
mx002	G	T	T	T	C	G	A	C	G	G	G	G	C	T	T	C	T	C	T	C	G	G	G	T	A	A	T	G	G	T	C	A	G	T	A	A	C	T	C	G	G	A	A	C	G	A	G	C	G	C	G	C	G	T	T	A	C	C	A	A	C	A
mx053	G	T	T	T	C	G	A	C	G	G	G	G	C	T	T	C	T	C	T	C	G	G	G	T	A	A	T	G	G	T	C	A	G	T	A	A	C	T	C	G	G	W	R	C	G	W	G	C	G	C	G	C	G	T	T	A	C	C	A	A	C	A
mx109	G	T	T	T	C	G	A	C	G	G	G	G	C	T	T	C	T	C	T	C	G	G	G	T	A	A	T	G	G	T	C	A	G	T	A	A	C	T	C	G	G	A	A	C	G	A	G	C	G	C	G	C	G	T	T	A	C	C	A	A	C	A
mx110	G	T	T	T	C	G	A	C	G	G	G	G	C	T	T	C	T	C	T	C	G	G	G	T	A	A	T	G	G	T	C	A	G	T	A	A	C	T	C	G	G	A	A	C	G	A	G	C	G	C	G	C	G	T	T	A	C	C	A	A	C	A
mx013	G	T	T	T	C	G	A	C	G	G	G	G	C	T	T	C	T	C	T	C	G	G	G	T	A	A	T	G	G	T	C	A	G	T	A	A	C	T	C	G	G	A	A	C	G	A	G	C	G	C	G	C	G	T	T	A	C	C	A	A	C	A
mx034	G	T	T	T	C	G	A	C	G	G	G	G	C	T	T	C	T	C	T	C	G	G	G	T	A	A	T	G	G	T	C	A	G	T	A	A	C	T	C	G	G	A	A	C	G	A	G	C	G	C	G	C	G	T	T	A	C	C	A	A	C	A
mx020	G	T	T	T	C	G	A	C	G	G	G	G	C	T	T	C	T	C	T	C	G	G	G	T	A	A	T	G	G	T	C	A	G	T	A	A	C	T	C	G	G	A	A	C	G	A	G	C	G	C	G	C	G	T	T	A	C	C	A	A	C	A
mx048	G	T	T	T	C	G	A	C	G	G	G	G	C	T	T	C	T	C	T	C	G	G	G	T	A	A	T	G	G	T	C	A	G	T	A	A	C	T	C	G	G	A	A	C	G	A	G	C	G	C	G	C	G	T	T	A	C	C	A	A	C	A
mx045	G	T	T	T	C	G	A	C	G	G	G	G	C	T	T	C	T	C	T	C	G	G	G	T	A	A	T	G	G	T	C	A	G	T	A	A	C	T	C	G	G	A	A	C	G	A	G	C	G	C	G	C	G	T	T	A	C	C	A	A	C	A
mx025	G	T	T	T	C	G	A	C	G	G	G	G	C	T	T	C	T	C	T	C	G	G	G	T	A	A	T	G	G	T	C	A	G	T	A	A	C	T	C	G	G	A	A	C	G	A	G	C	G	C	G	C	G	T	T	A	C	C	A	A	C	A
mx024	G	T	T	T	C	G	A	C	G	G	G	G	C	T	T	C	T	C	T	C	G	G	G	T	A	A	T	G	G	T	C	A	G	T	A	A	C	T	C	G	G	A	A	C	G	A	G	C	G	C	G	C	G	T	T	A	C	C	A	A	C	A
mx044	G	T	T	T	C	G	A	C	G	G	G	G	C	T	T	C	T	C	T	C	G	G	G	T	A	A	T	G	G	T	C	A	G	T	A	A	C	T	C	G	G	A	A	C	G	A	G	C	G	C	G	C	G	T	T	A	C	C	A	A	C	A
mx049	G	T	T	T	C	G	A	C	G	G	G	G	C	T	T	C	T	C	T	C	G	G	G	T	A	A	T	G	G	T	C	A	G	T	A	A	C	T	C	G	G	A	A	C	G	A	G	C	G	C	G	C	G	T	T	A	C	C	A	A	C	A
mx050	G	T	T	T	C	G	A	C	G	G	G	G	C	T	T	C	T	C	T	C	G	G	G	T	A	A	T	G	G	T	C	A	G	T	A	A	C	T	C	G	G	A	A	C	G	A	G	C	G	C	G	C	G	T	T	A	C	C	A	A	C	A
mx051	G	T	T	T	C	G	A	C	G	G	G	G	C	T	T	C	T	C	T	C	G	G	G	T	A	A	T	G	G	T	C	A	G	T	A	A	C	T	C	G	G	A	A	C	G	A	G	C	G	C	G	C	G	T	T	A	C	C	A	A	C	A
mx055	G	T	T	T	C	G	A	C	G	G	G	G	C	T	T	C	T	C	T	C	G	G	G	T	A	A	T	G	G	T	C	A	G	T	A	A	C	T	C	G	G	A	A	C	G	A	G	C	G	C	G	C	G	T	T	A	C	C	A	A	C	A
mx111	G	T	T	T	C	G	A	C	G	G	G	G	C	T	T	C	T	C	T	C	G	G	G	T	A	A	T	G	G	T	C	A	G	T	A	A	C	T	C	G	G	A	A	C	G	A	G	C	G	C	G	C	G	T	T	A	C	C	A	A	C	A
mx069	G	T	T	T	C	G	A	C	G	G	G	G	C	T	T	C	T	C	T	C	G	G	G	T	A	A	T	G	G	T	C	A	G	T	A	A	C	T	C	G	G	W	R	C	G	W	G	C	G	C	G	C	G	T	T	A	C	C	A	A	C	A
mx062	G	T	T	T	C	G	A	C	G	G	G	G	C	T	T	C	T	C	T	C	G	G	G	T	A	A	T	G	G	T	C	A	G	T	A	A	C	T	C	G	G	W	R	C	G	W	G	C	G	C	G	C	G	T	T	A	C	C	A	A	C	A
mx112	G	T	T	T	C	G	A	C	G	G	G	G	C	T	T	C	T	C	T	C	G	G	G	T	A	A	T	G	G	T	C	A	G	T	A	A	C	T	C	G	G	A	A	C	G	A	G	C	G	C	G	C	G	T	T	A	C	C	A	A	C	A
mx065	G	T	T	T	C	G	A	C	G	G	G	G	C	T	T	C	T	C	T	C	G	G	G	T	A	A	T	G	G	T	C	A	G	T	A	A	C	T	C	G	G	W	R	C	G	W	G	C	G	C	G	C	G	T	T	A	C	C	A	A	C	A
mx074	G	T	T	T	C	G	A	C	G	G	G	G	C	T	T	C	T	C	T	C	G	G	G	T	A	A	T	G	G	T	C	A	G	T	A	A	C	T	C	G	G	A	A	C	G	A	G	C	G	C	G	C	G	T	T	A	C	C	A	A	C	A
mx084	G	T	T	T	C	G	A	C	G	G	G	G	C	T	T	C	T	C	T	C	G	G	G	T	A	A	T	G	G	T	C	A	G	T	A	A	C	T	C	G	G	W	R	C	G	W	G	C	G	C	G	C	G	T	T	A	C	C	A	A	C	A
mx066	G	T	T	T	C	G	A	C	G	G	G	G	C	T	T	C	T	C	T	C	G	G	G	T	A	A	T	G	G	T	C	A	G	T	A	A	C	T	C	G	G	W	R	C	G	W	G	C	G	C	G	C	G	T	T	A	C	C	A	A	C	A
mx072	G	T	T	T	C	G	A	C	G	G	G	G	C	T	T	C	T	C	T	C	G	G	G	T	A	A	T	G	G	T	C	A	G	T	A	A	C	T	C	G	G	W	R	C	G	W	G	C	G	C	G	C	G	T	T	A	C	C	A	A	C	A
mx086	G	T	T	T	C	G	A	C	G	G	G	G	C	T	T	C	T	C	T	C	G	G	G	T	A	A	T	G	G	T	C	A	G	T	A	A	C	T	C	G	G	W	R	C	G	W	G	C	G	C	G	C	G	T	T	A	C	C	A	A	C	A
mx091	G	T	T	T	C	G	A	C	G	G	G	G	C	T	T	C	T	C	T	C	G	G	G	T	A	A	T	G	G	T	C	A	G	T	A	A	C	T	C	G	G	A	A	C	G	A	G	C	G	C	G	C	G	T	T	A	C	C	A	A	C	A
mx113	G	T	T	T	C	G	A	C	G	G	G	G	C	T	T	C	T	C	T	C	G	G	G	T	A	A	T	G	G	T	C	A	G	T	A	A	C	T	C	G	G	W	R	C	G	W	G	C	G	C	G	C	G	T	T	A	C	C	A	A	C	A
mx067	G	T	T	T	C	G	A	C	G	G	G	G	C	T	T	C	T	C	T	C	G	G	G	T	A	A	T	G	G	T	C	A	G	T	A	A	C	T	C	G	G	A	A	C	G	A	G	C	G	C	G	C	G	T	T	A	C	C	A	A	C	A
mx014	G	T	T	T	C	G	A	C	G	G	G	G	C	T	T	C	T	C	T	C	G	G	G	T	A	A	T	G	G	T	C	A	G	T	A	A	C	T	C	G	G	A	A	C	G	A	G	C	G	C	G	C	G	T	T	A	C	C	A	A	C	A
mx007	G	T	T	T	C	G	A	C	G	G	G	G	C	T	T	C	T	C	T	C	G	G	G	T	A	A	T	G	G	T	C	A	G	T	A	A	C	T	C	G	G	A	A	C	G	A	G	C	G	C	G	C	G	T	T	A	C	C	A	A	C	A
mx017	G	T	T	T	C	G	A	C	G	G	G	G	C	T	T	C	T	C	T	C																																										

