

**Table S3** (a) Genetic diversity parameters estimated for the CP gene of OrMV. (b) Summary of test statistics for population differentiation

Phylogenetic Group	Sample size	Haplotypes	Haplotype diversity		Nucleotide diversity		
(a)							
Group 1	23	23	1.000 ( $\pm 0.013$ )		0.106 ( $\pm 0.014$ )		
Group 2	13	13	1.000 ( $\pm 0.030$ )		0.117 ( $\pm 0.015$ )		
Total	36	30	1.000 ( $\pm 0.007$ )		0.156 ( $\pm 0.010$ )		
Phylogenetic Group	$K_{ST}$	$K_S$	<i>P-value</i>	$S_{nn}$	<i>P-value</i>	$F_{ST}$	<i>P-value</i>
(b)							
Group 1 vs. Group 2	0.296	83.545	< 0.001 <sup>***</sup>	1.00	< 0.001 <sup>***</sup>	0.470	< 0.001 <sup>***</sup>

Significance thresholds: \*,  $0.01 < P\text{-value} < 0.05$ ; \*\*,  $0.001 < P\text{-value} < 0.01$ ; \*\*\*,  $P\text{-value} < 0.001$