Table S3. Haplotype diversity assessed by bootstrap resampling with 1000 replicates

Gene	Country	Haplotype diversity	95% Confidence intervals	P-value
P1	China	0.988	0.970 - 0.998	0.990^{ns}
	Japan	0.989	0.972 - 0.998	0.098^{ns}
CP	China	0.967	0.924 - 0.989	1.000^{ns}
	Japan	0.972	0.936 - 0.991	0.050^{*}

^{ns}, not significant; *, $0.01 < P\text{-}value < 0.05$; **, $0.001 < P\text{-}value < 0.01$; ***, $P\text{-}value < 0.001$