Metadata associated with Yu et al. Molecular Ecology DRYAD.xls

Sheet 1: Genotypic data of *Valasia javana*

Nuclear genotypes (9 loci) for 568 individuals of *Valisia javana*. Population refers to collection sites, with GPS coordinates provided in the main text. The sampling spanned a distance of over 4100 km. For each nuclear locus, alleles names are arbitrary (i.e., do not indicate size or sequence characteristics), and “0” indicates missing data.

Sheet 2: Genotypic data of *Ficus hirta*

Nuclear genotypes (7 loci) for 449 individuals of *Ficus hirta*. Population refers to collection sites, with GPS coordinates provided in the main text. The sampling spanned a distance of over 4100 km. For each nuclear locus, alleles names are arbitrary (i.e., do not indicate size or sequence characteristics), and “0” indicates missing data.

Notebook 1: Mitochondrial DNA sequence (COI) of *Valasia javana*

The aligned partial mtDNA COI sequences of *Valisia javana* with 330 individuals and the GPS coordinates for the sampling locations were included. Local population designations follow Sheet 1 and the main text.

Notebook 2: nuclear DNA sequence (ITS2) of *Valasia javana*

The aligned partial nrDNA ITS2 sequences of *Valisia javana* with species and haplotypes. Species and haplotype designations follow Table S5 in supplementary data and the main text.

Notebook 3: Chloraplast DNA sequence (trnLF+trnSG) of *Ficus hirta*

The aligned partial cpDNA (trnLF+trnSG) sequences of *Ficus hirta* with 309 individuals and the GPS coordinates for the sampling locations were included. Local population designations follow Sheet 2 and the main text.