**Title page**

**Testing mate recognition through reciprocal crosses of two native populations of the whitefly *Bemisia tabaci* (Gennadius) in Australia**

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**Supplementary Figure 1**: Bayesian phylogenetic tree using MrBayes (GTR+I+G model model) based on the 3’ end of the mitochondrial cytochrome oxidase I (mtCOI) gene (655 bp) of parents and progeny involved in the cross mating test AUSI-M x AUSII-F. The outgroup is *Bemisia atriplex*.