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Please, contact us if you have suggestions, find errors, inconsistencies, or any other problems with the files.

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### 1. COI\_tree.nex

The dated phylogenetic gene tree based on COI sequences taken from (Damgaard *et al.* 2008). The methods used to generate the tree are described in the section "Genetic diversity & cross-species amplification" in the Materials and Methods section of the manuscript.

### 2. PrimerNote\_FSTAT.dat

The genotypes for *Karoophasma biedouwense* from Clanwilliam and Biedouw Valley used to examine the properties of the newly developed microsatellite loci (in FSTAT format).

### 3. Microsatellite Sequences\_KB1\_KB50.fas

The microsatellite containing sequences from which primers were designed for the current study (fasta format).

## 4. Dool\_SupplTable 1\_2\_4\_5\_JHeredity.pdf

Supplementary files which accompany the manuscript:

Supplementary Table 1: details of 24 *Karoophasma biedouwense* sent for Illumina sequencing at Plön. These sequences were used for microsatellite discovery and primer design.

Supplementary Table 2: unlabelled microsatellite loci designed for *Karoophasma biedouwense* which were not selected for further optimisation or multiplexing.

Supplementary Table 4: Amplification success of 23 microsatellite loci designed for *Karoophasma biedouwense* across the order Mantophasmatodea.

Supplementary Table 5: Genbank accession numbers and melting temperatures (forward primer, reverse primer) for each microsatellite locus (see Table 1).

# 5. Supp\_Table3\_Genotypes.xlsx

Genotypes of *Karoophasma biedouwense* from Clanwilliam and Biedouw Valley and all other species of Mantophasmatodea which were used in the study.

### References

Damgaard J., Klass K.-D., Picker M.D. & Buder G. (2008) Phylogeny of the Heelwalkers (Insecta: Mantophasmatodea) based on mtDNA sequences, with evidence for additional taxa in South Africa. *Molecular phylogenetics and evolution* **47**, 443-62.