S3 Table. Multi-locus $P_{\rm ID}$ and $P_{\rm ID}$ sib. Multi-locus $P_{\rm ID}$ (i.e., the probability that two individuals drawn at random share identical genotypes) and $P_{\rm ID}$ sib (i.e., the probability of identity among siblings) data are provided. Loci are sorted according to the increasing order of their $P_{\rm ID}$ and $P_{\rm ID}$ sib single-locus values (i.e., the locus at the top is the most informative one), and a sequentially multi-loci $P_{\rm ID}$ ($P_{\rm ID}$ sib) is reported for each locus. See Forcina et al. [46] for details on the loci employed in the present study.

Locus	P_{ID}	P _{ID} sib
MCW 252	1.2 x 10 ⁻²	2.9 x 10 ⁻¹
MCW 212	1.3 x 10 ⁻³	1.2 x 10 ⁻¹
MCW 104	2.2 x 10 ⁻⁴	5.5 x 10 ⁻²
MCW 146	4.9 x 10 ⁻⁵	2.8 x 10 ⁻²
MCW 295	1.2 x 10 ⁻⁵	1.4 x 10 ⁻²
Aru 1.23	4.1 x 10 ⁻⁶	8.5 x 10 ⁻³
MCW 127	1.5 x 10 ⁻⁶	5.4 x 10 ⁻³
LEI 30	8.6 x 10 ⁻⁷	4.1 x 10 ⁻³
MCW 280	7.4 x 10 ⁻⁷	3.8 x 10 ⁻³