

APPENDIX S2. Principal coordinate analyses (PCoA) based on microsatellite data at 13 loci and across 81 *Fosterella* accessions, including 29 accessions of *F. christophii*, 31 of *F. micrantha*, and 21 of *F. villosula*. Each species is represented by a unique symbol. Seven identical multilocus genotypes were identified. The first three axes together represent 53.5% of the total genetic variation (Dim-1 = 22.7%, Dim-2 = 18.7%, Dim-3 = 11.8%). Three distinct groups (indicated by dotted lines) could be discerned, illustrating the potential of the newly developed microsatellite markers for species delimitation.

