

APPENDIX S1. Bayesian inference analysis obtained from the plastid *trnS-trnG* intergenic spacer including species from *Cereus*, *Cipocereus*, and *Praecereus*. Posterior probabilities (>0.90) are shown below the node branches. The subgenera of *Cereus* are indicated by symbols according to the legend (Franco et al., unpublished data).

