Supplemental Fig. 1. Phylogenetic relationships in *Passiflora* based on nuclear loci (nrITS, ncpGS, and nrITS + ncpGS combined). A–C, strict consensus of trees retained from maximum parsimony analysis. Branches with jackknife support of ≥ 70% in bold. A. nrITS. B. ncpGS. C. nrITS + ncpGS. Number of taxa, number of trees retained, CI, RI, and length reported to the left of each topology.
Supplemental Fig. 2. Phylogenetic relationships in *Passiflora* based on chloroplast loci (*ndhF, trnL-F*, and *ndhF* + *trnL-F* combined). A–C, strict consensus of trees retained from maximum parsimony analysis. Branches with jackknife support of ≥ 70% in bold. A. *ndhF*. B. *trnL-F*. C. *ndhF* + *trnL-F*. Number of taxa, number of trees retained, CI, RI, and length reported to the left of each topology.