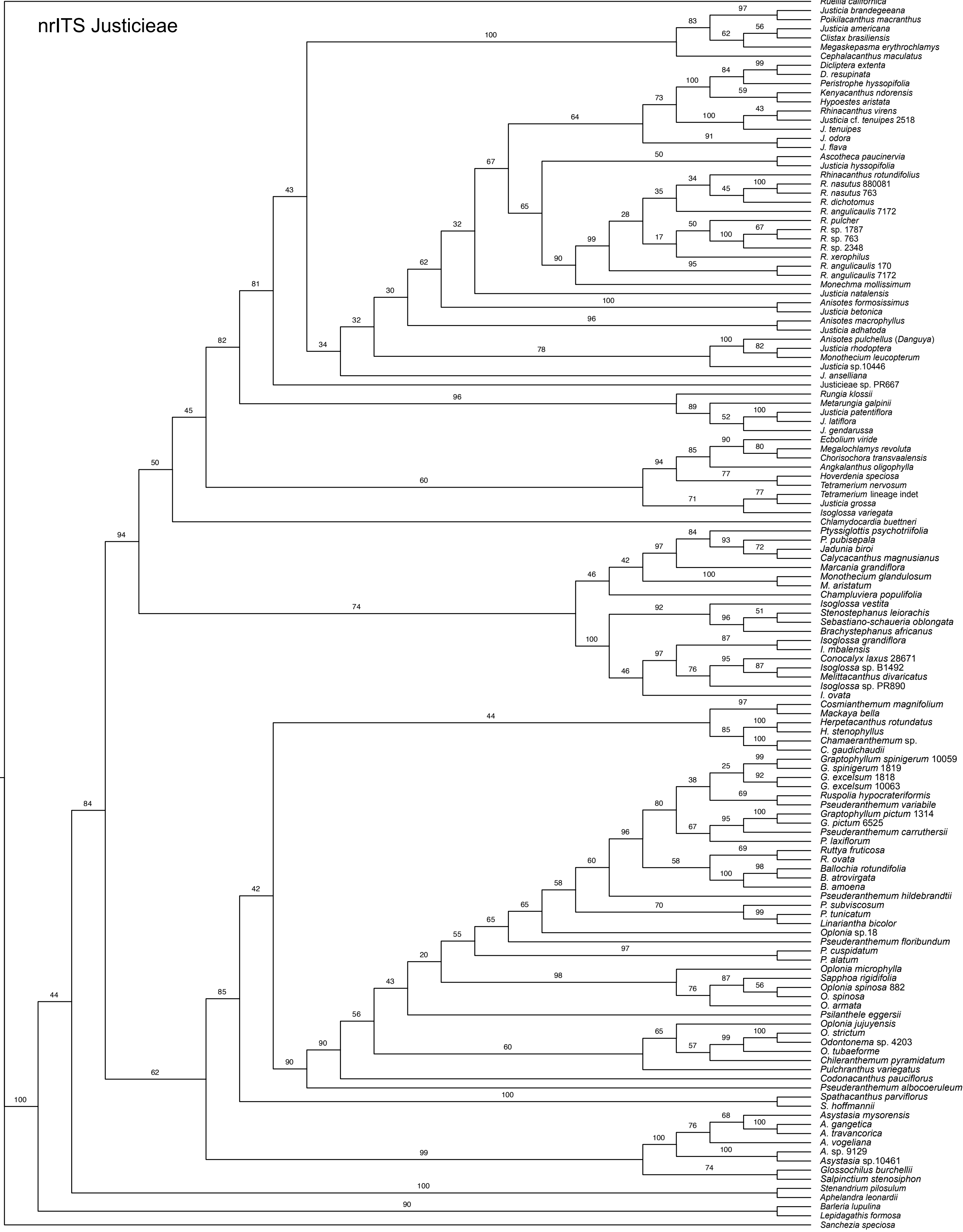


nrITS Justiceae



S2. The most likely phylogenetic hypothesis for relationships among 148 taxa dataset of the Justiceae lineage generated from nrITS data. Maximum likelihood bootstrap support corresponds to values above the branches.