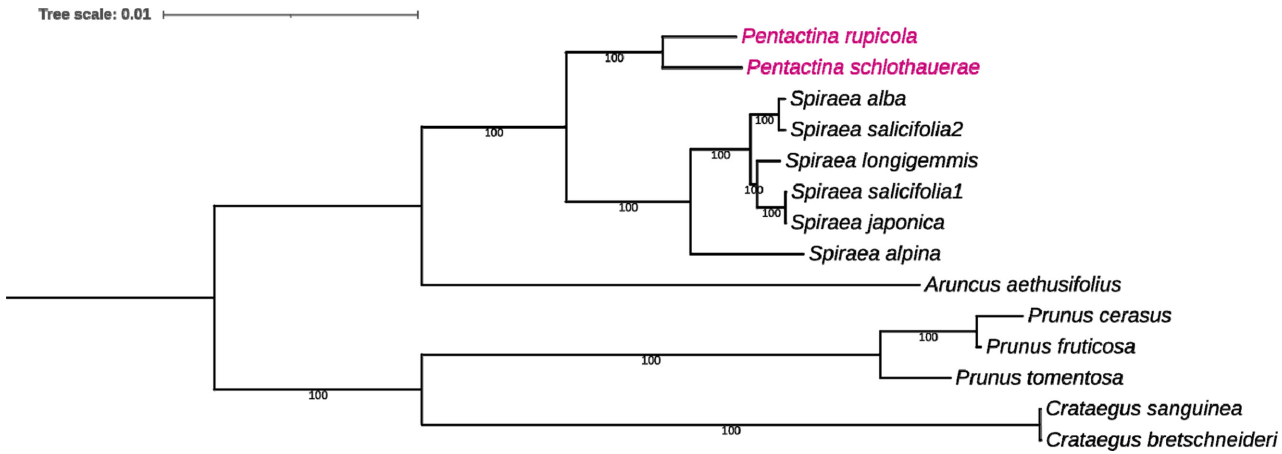

On-line Supplemental Data Fig. S3. Maximum-likelihood tree inferred from complete plastome sequences of *Pentactina* and its related genera. Numbers below the branches represent bootstrap support values. Sequences were obtained from NCBI: *Aruncus aethusifolius* (MZ882398), *Crataegus bretschneideri* (OP964009), *Crataegus sanguinea* (OP964003), *Prunus cerasus* (MW477432), *Prunus fruticosa* (MT916286), *Prunus tomentosa* (MF624726), *Spiraea alba* (OX465087), *Spiraea alpina* (OR601530), *Spiraea japonica* (OR513066), *Spiraea longigemmis* (OR513059), *Spiraea salicifolia*1 (OR513063), *Spiraea salicifolia*2 (ON159999), *Pentactina rupicola* (NC_016921), and *Pentactina schlothauerae* (PP189928).