



**Figure Appendix S6:** Phylogram for 152 species of the alpine flora of the Sawtooth National Forest (98% of species collected). Relationships estimated under maximum likelihood from concatenated gene regions (ITS, *atpB*, *matK*, *ndhF*, *rbcL*, and *trnTLF*) of the total dataset (combining molecular sequence data from GenBank and targeted high-throughput MiSeq). Node values show bootstrap support (BS).