Supplementary files

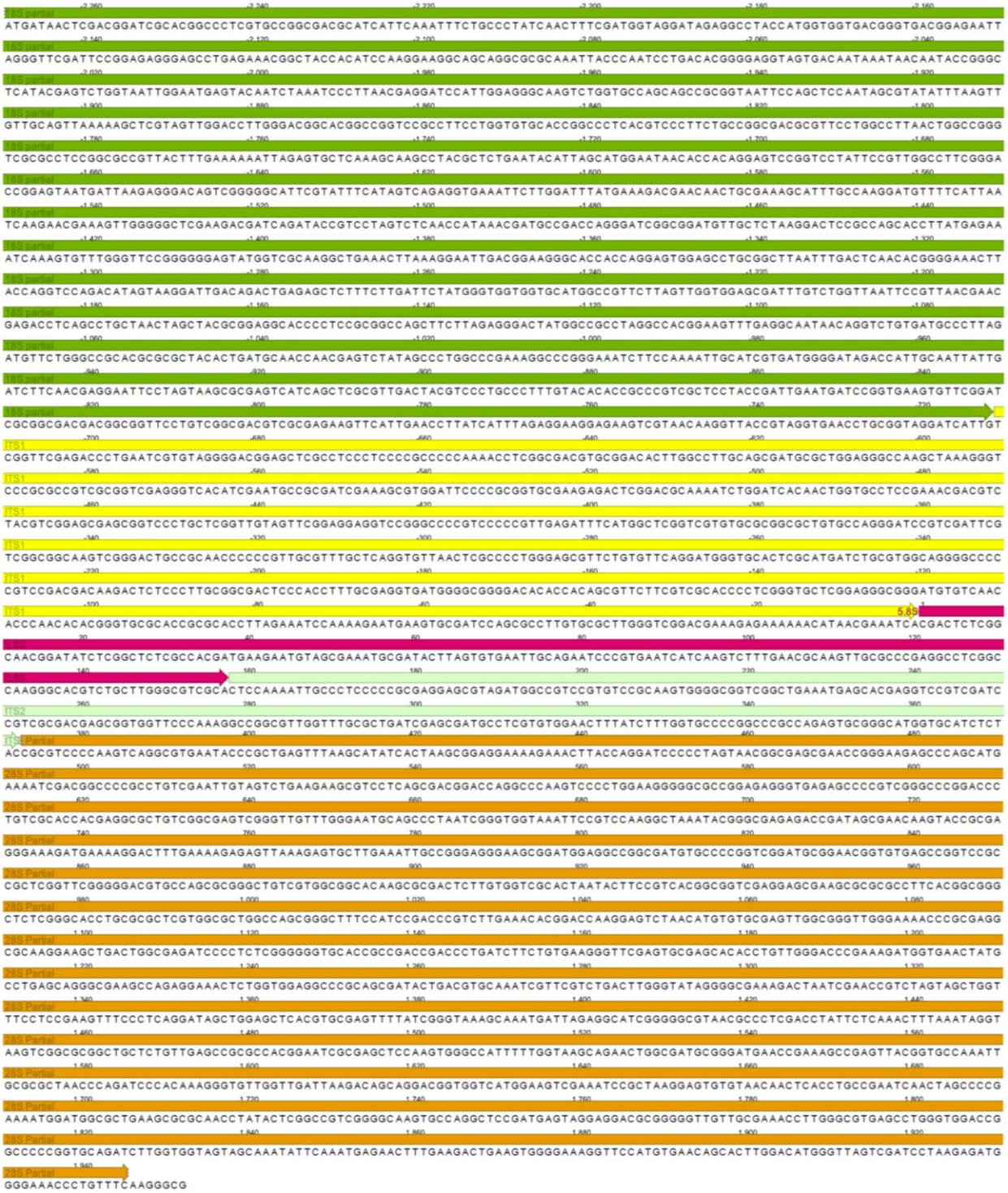


Figure Supplementary S1. Complete annotated ITS-rDNA reference sequence of *Juniperus* spp. The sequence is 4,231 bp long and includes partial 18S rRNA (positions –2276 to –719), ITS1 (positions –718

to –1), 5.8S rRNA (positions 1 to 157), ITS2 (positions 158 to 375), and partial 28S rRNA (positions 376 to 1948). Nucleotide positions are numbered relative to the first base of the 5.8S region, designated as +1, with negative values indicating upstream regions. We assembled the reference sequence from 570 ITS records retrieved from NCBI. Because it was assembled, no NCBI accession number was assigned. The complete sequence is provided here in FASTA format for researchers to use directly.

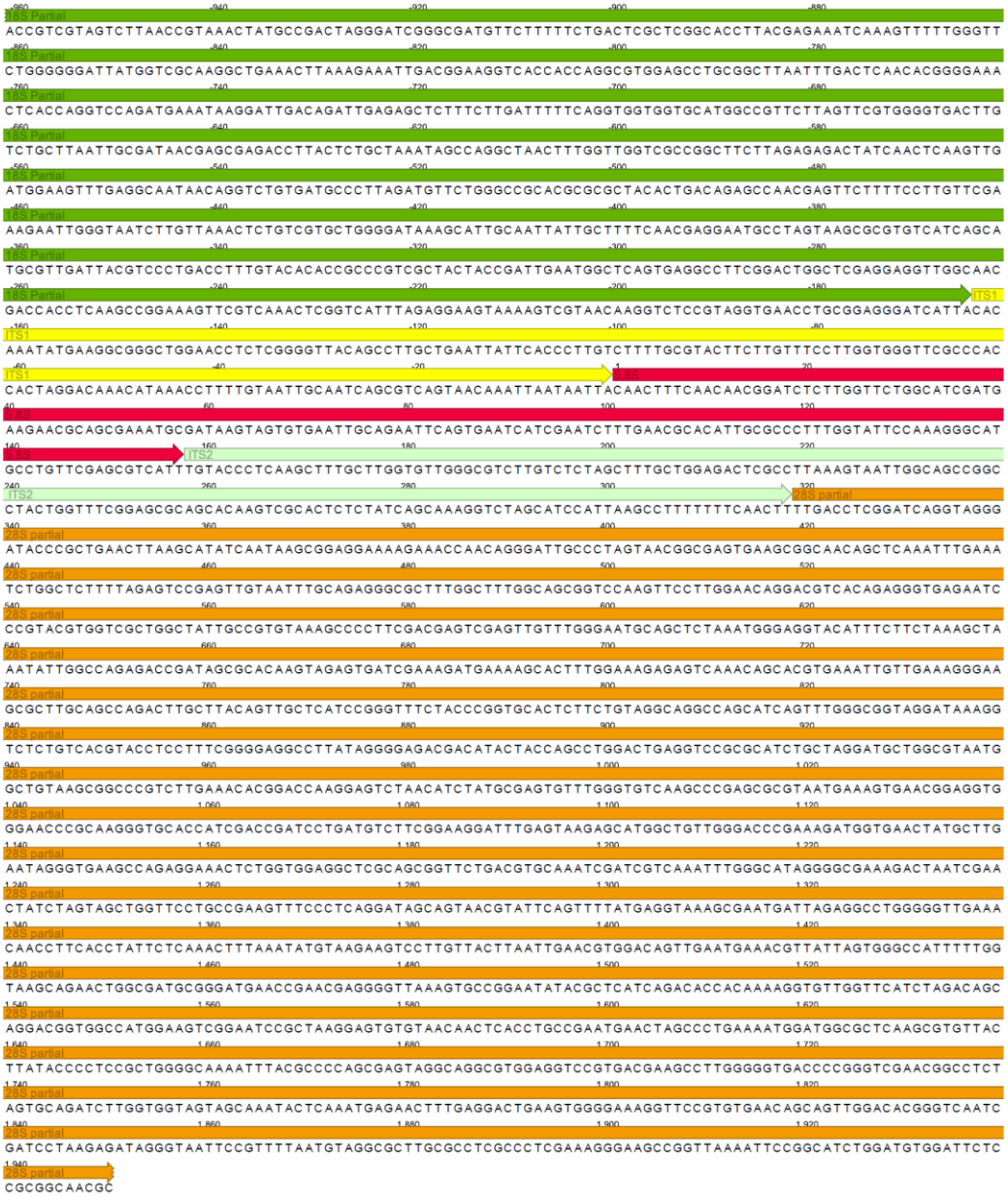


Figure Supplementary S2. Complete annotated ITS-rDNA reference sequence of *Alternaria alternata* (submission ID SUB16216726). The sequence is 2,911 bp long and includes partial 18S rRNA (positions –958 to –165), ITS1 (positions –164 to –1), 5.8S rRNA (positions 1 to 157), ITS2 (positions 158 to 318),

and partial 28S rRNA (positions 319 to 1950). Nucleotide positions are numbered relative to the first base of the 5.8S region, designated as +1, with negative values indicating upstream regions. We assembled the reference sequence from 8,050 ITS records retrieved from NCBI. Because NCBI does not assign accession numbers to assembled reference sequences, we provide the complete sequence here in FASTA format; it is also available from the corresponding author upon request. This article may be cited as the source of the reference sequence.