

Supplementary Materials

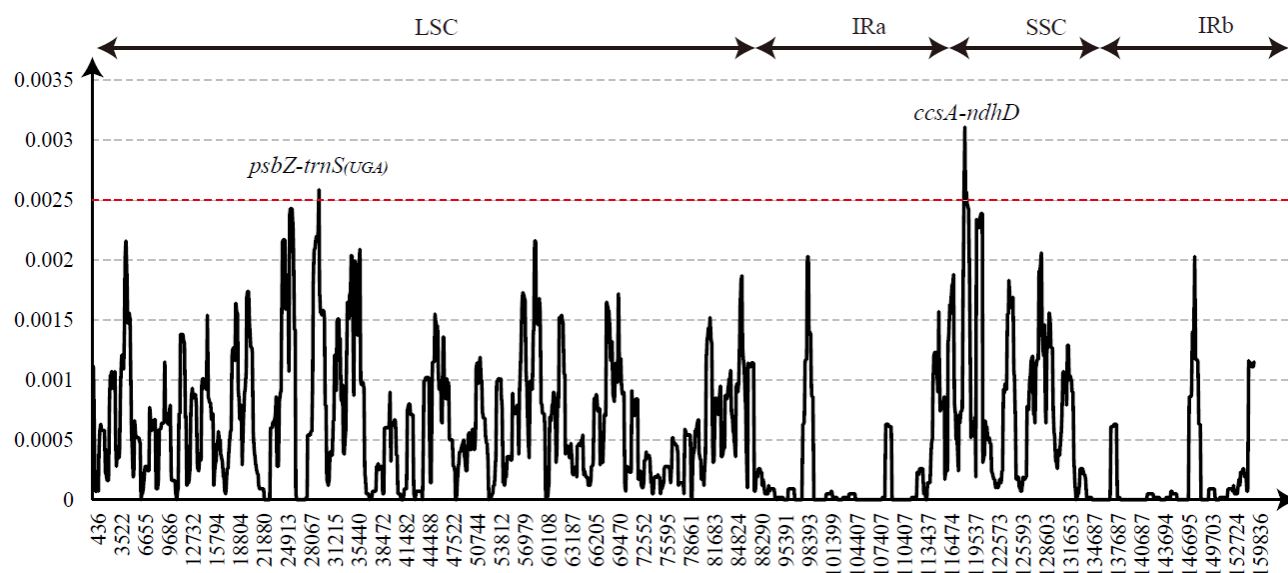


Figure S1. Nucleotide diversity (π) in *P. montana*'s chloroplast genome based on a window size of 800 bp and a step size of 100 bp.

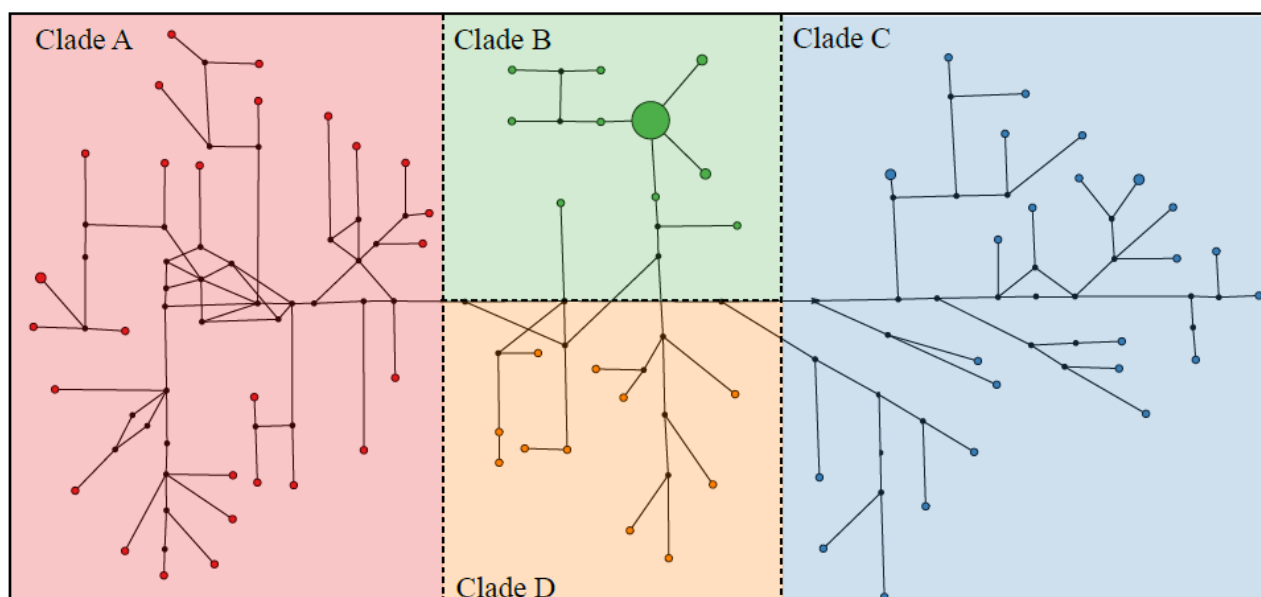


Figure S2. Network of all 104 *P. montana* accessions.

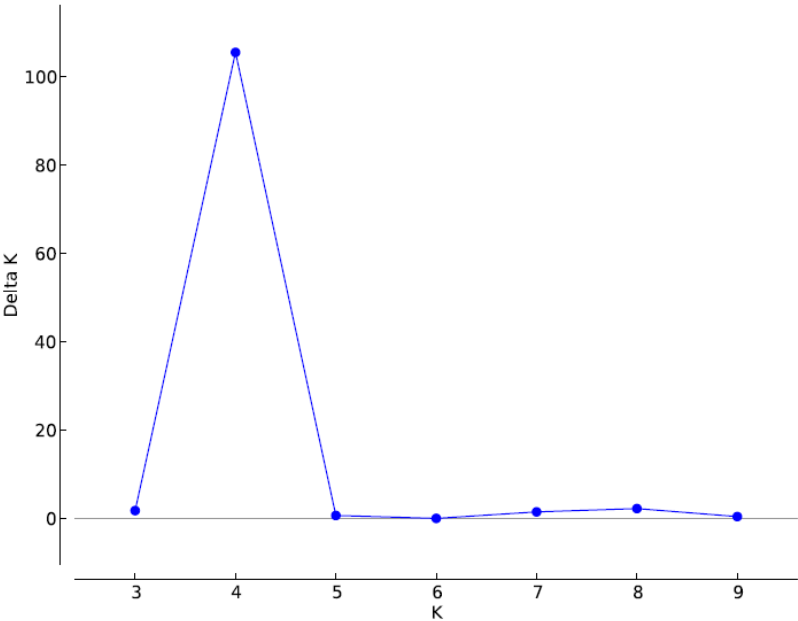


Figure S3. The delta K results of STRUCTURE analysis.

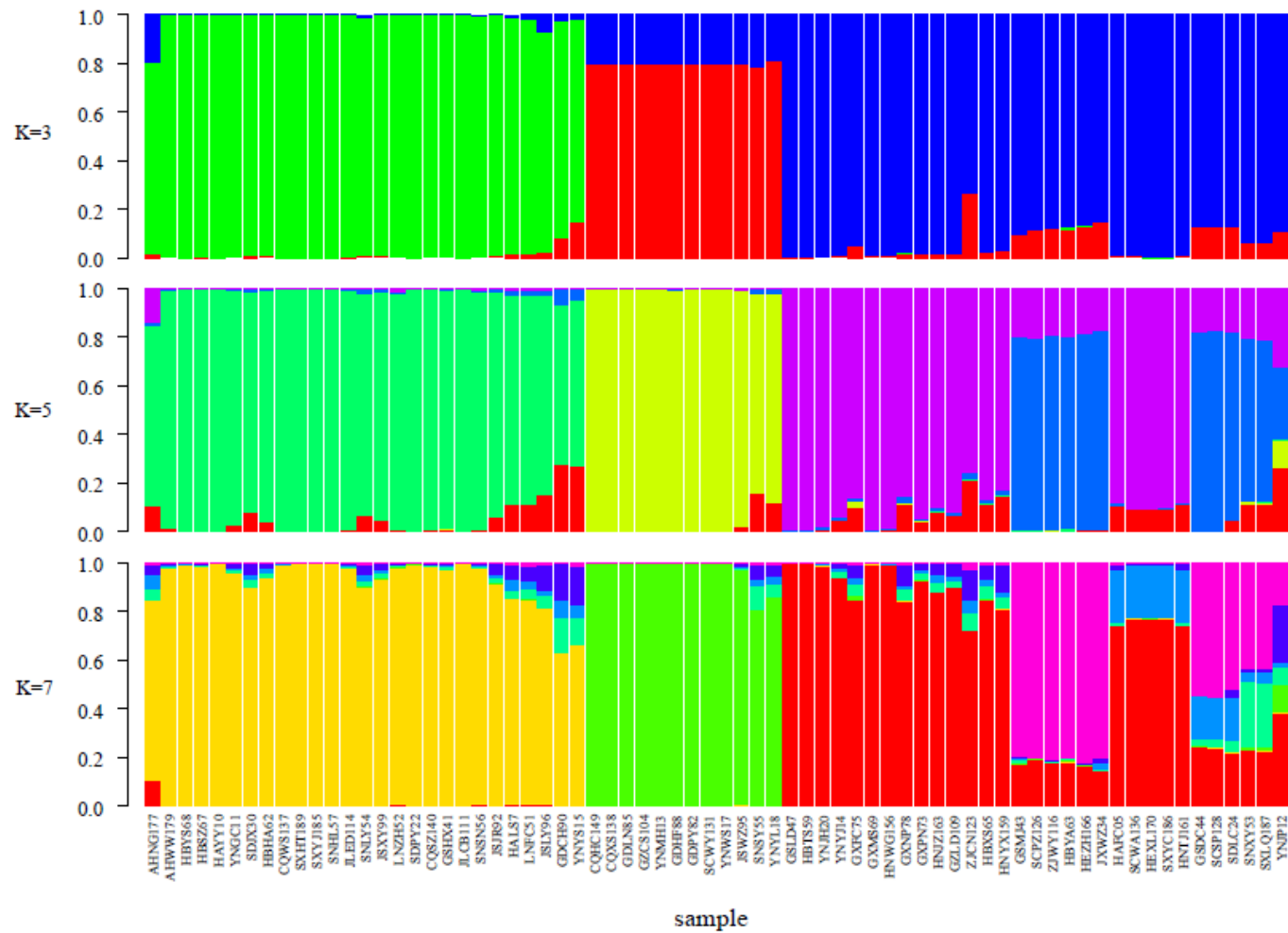


Figure S4. Population structure of all 104 accessions with K = 3, 5, and 6.

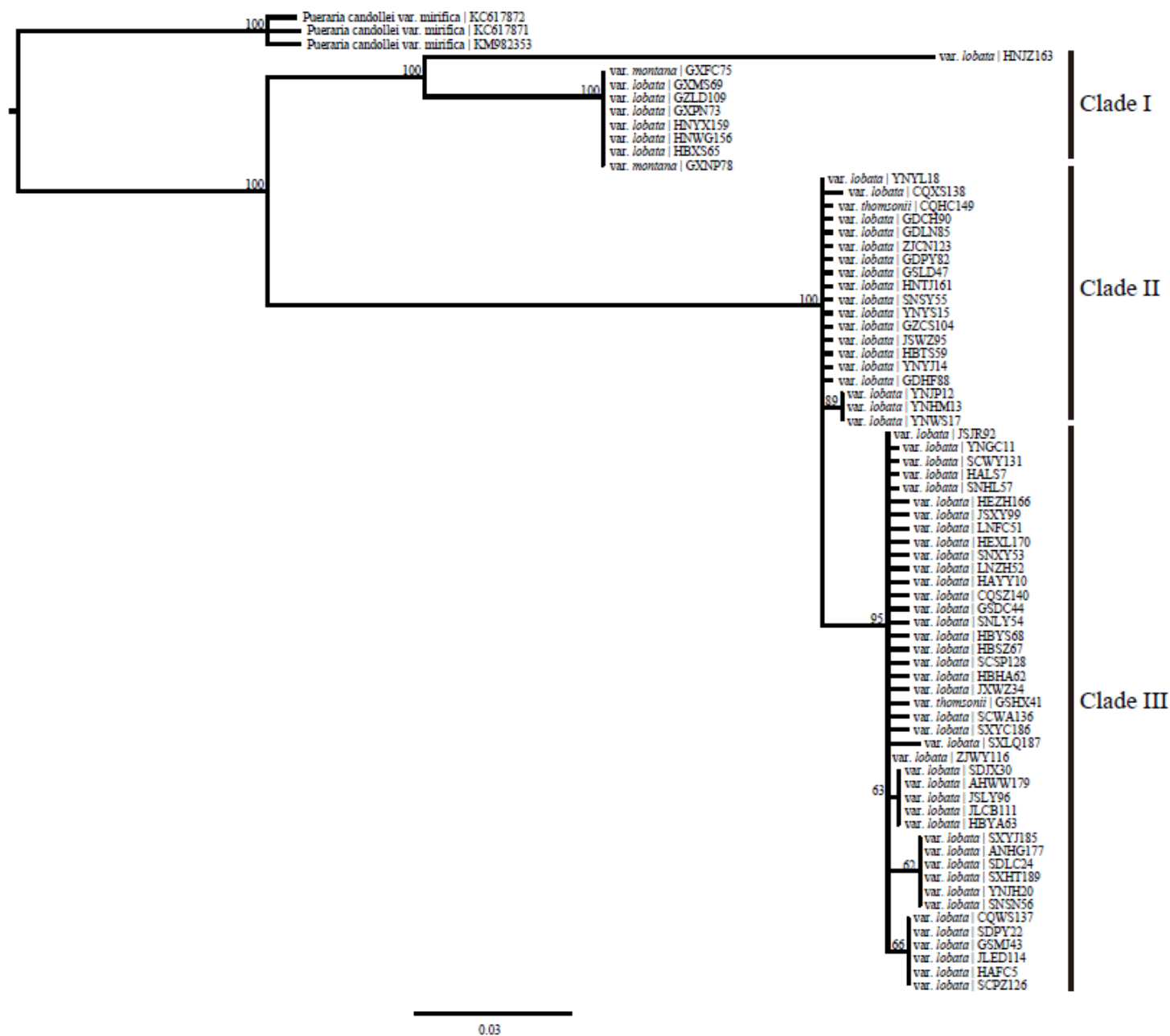


Figure S5. The ITS phylogenetic tree of 70 *P. montana* collected accessions.