

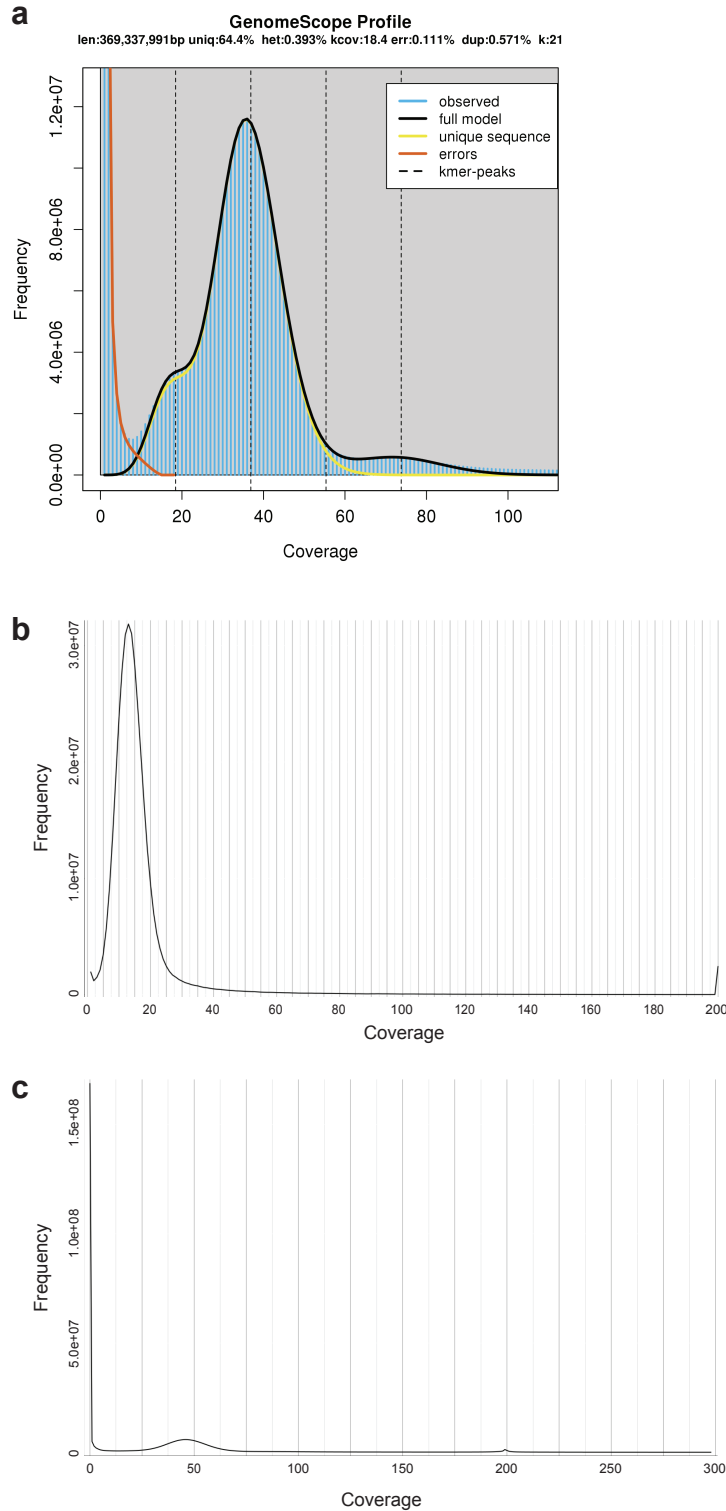


Figure S1. Estimated heterozygosity plots for *L. erythrorhizon*. a) GenomeScope k-mer profile plot of the *L. erythrorhizon* illumina gDNA reads. The blue bars indicate the observed kmer frequencies, and the black and yellow lines indicate expected distribution of kmers using the GenomeScope model for diploid genomes. b) Purge-haplotigs read-depth histogram of the *L. erythrorhizon* genome displaying coverage of nanopore reads. c) Purge-haplotigs read-depth histogram of the *L. erythrorhizon* genome displaying coverage of illumina gDNA reads.