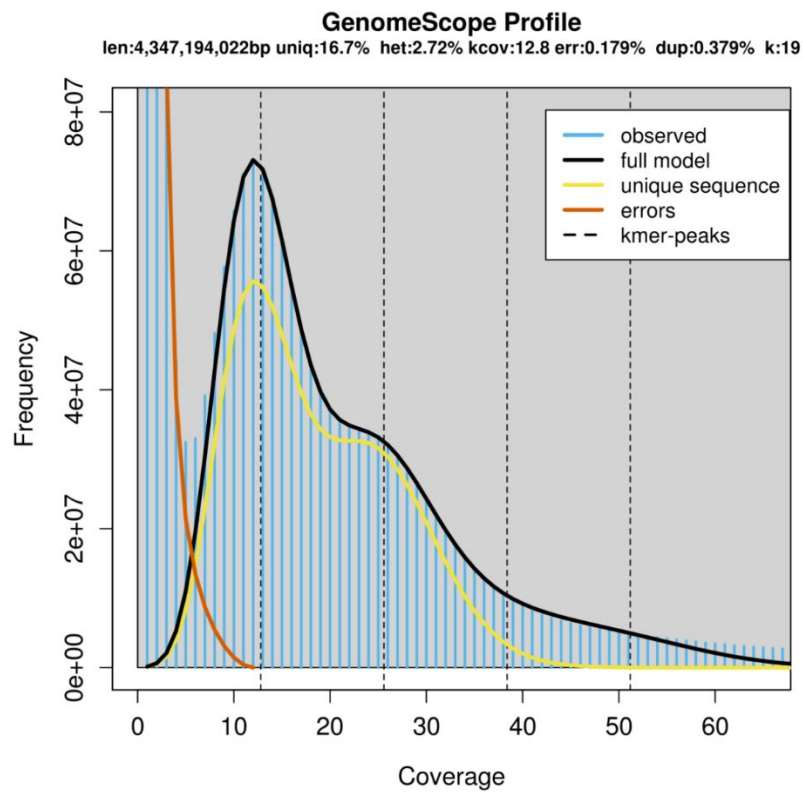


Supplementary Figures



Supplementary Figure 1. Genome size and heterozygosity estimation using 19 *K*-mer distribution.