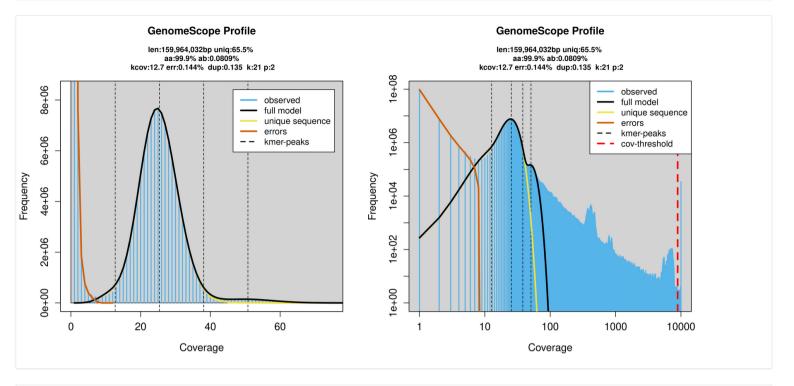
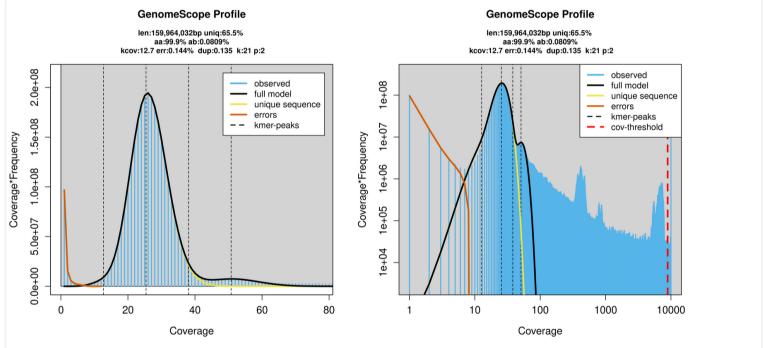
GenomeScope Home Info Examples → My results →

my sample





Results

GenomeScope version 2.0
input file = user_uploads/wCgJJUbU3okwQLADSZx4
output directory = user_data/wCgJJUbU3okwQLADSZx4
p = 2
p = 2
h = -21

Homozygous (aa) Heterozygous (ab) Genome Haploid Length Genome Repeat Length Genome Unique Length Model Fit 99.8909% 0.0527612% 159,746,355 b 55,164,242 bp 104,582,113 b 67.8311% 0.144188% max 99.9472% 0.109119% 159,964,032 bp 55,239,411 bp 104,724,621 bp 98.6673%

View analysis later

Return to view your results at any time:

Progress

starting

round 0 trimming to 7 trying 2p peak model... converged. score: 723793932593932 round 1 trimming to 12 trying 2p peak model... converged. score: 723815890395177

round 2 trimming to 17 trying 2p peak model... converged. score: 734465441027511

round 3 trimming to 22 trying 2p peak model... converged. score: 697778731018068 done