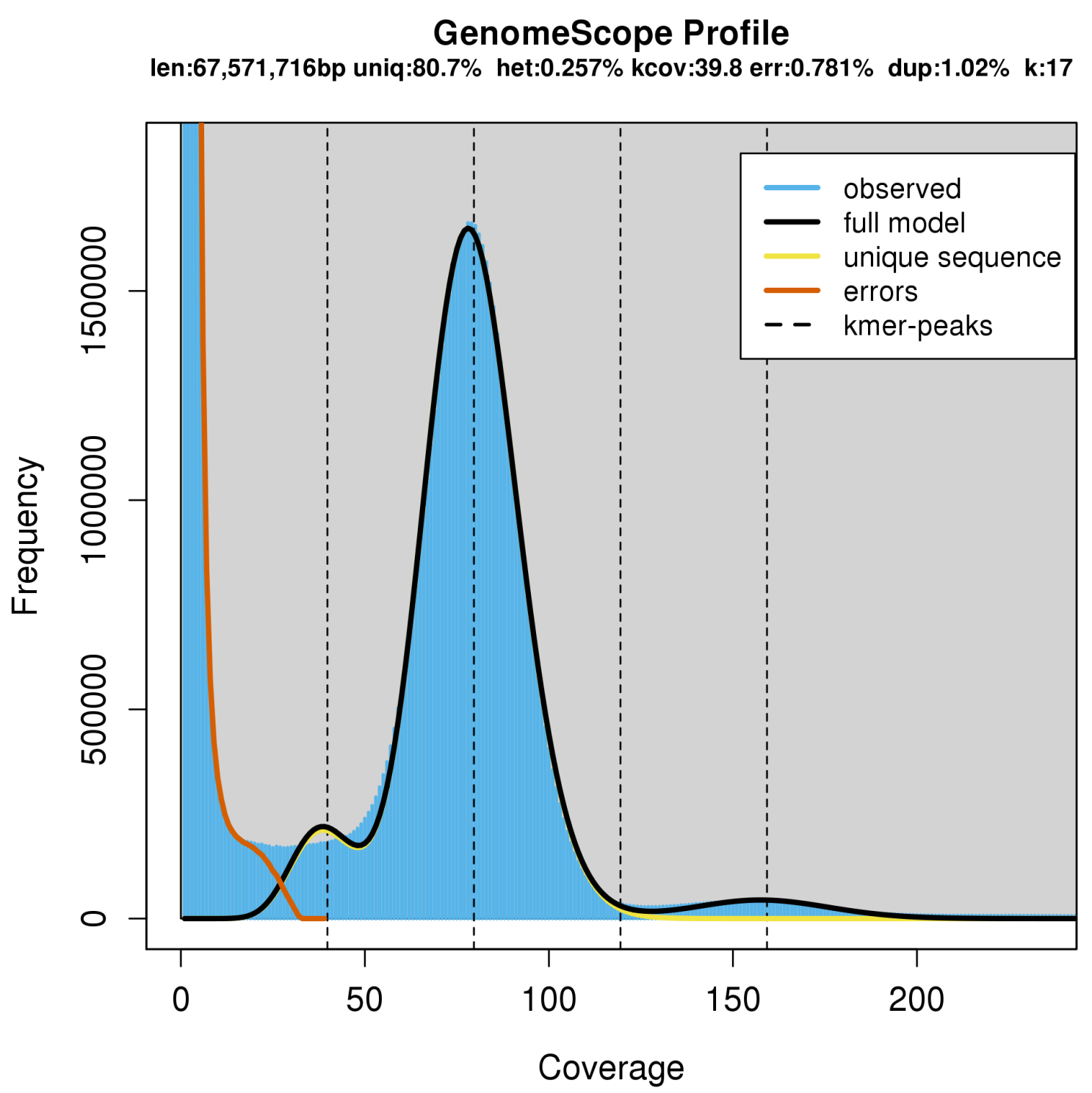
Figure S1


GenomeScope Profile
len:67,571,716bp uniq:80.7% het:0.257% kcov:39.8 err:0.781% dup:1.02% k:17
observed
full model
unique sequence
errors
kmer-peaks
Frequency
0
500000
1000000
1500000
0
50
100
150
200
Coverage