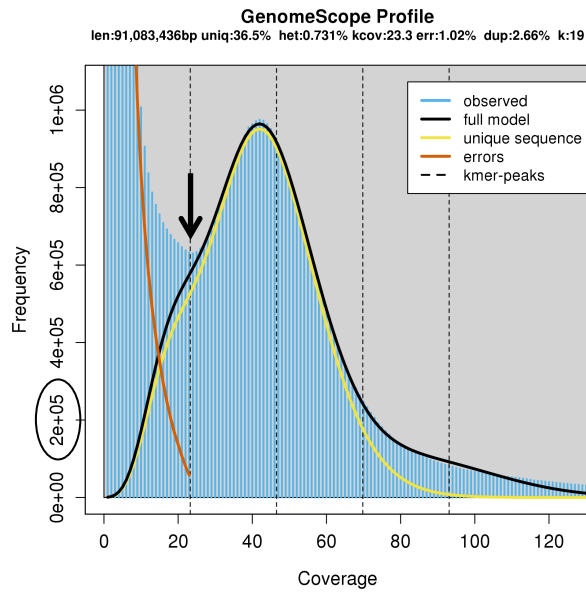
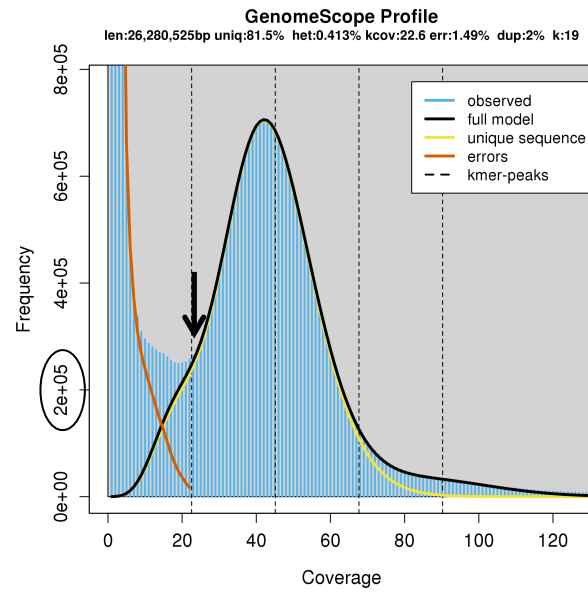


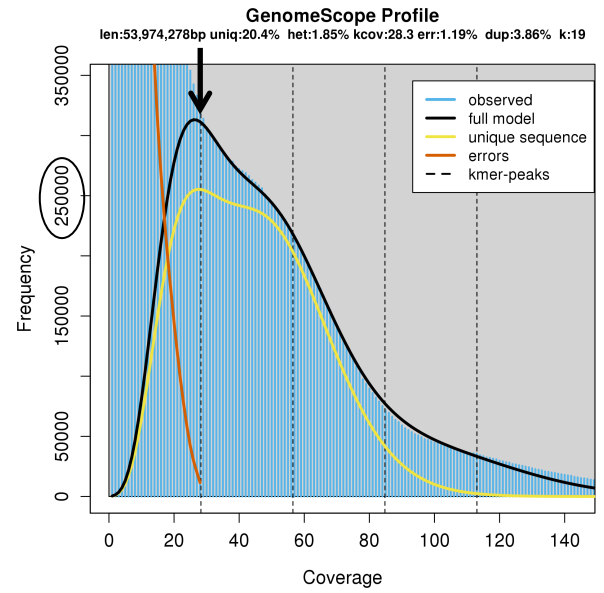
(a) All data



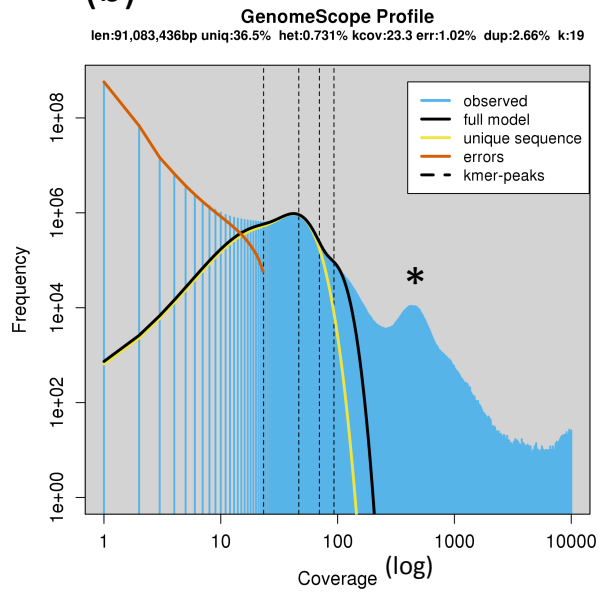
(c) Depleted of bacterial reads



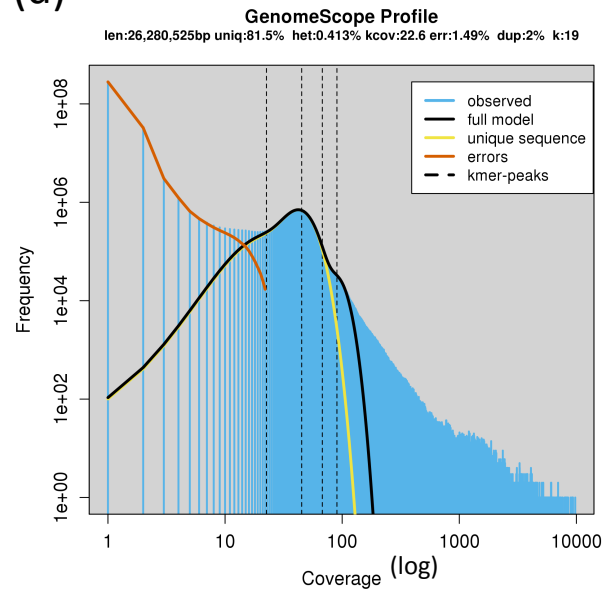
(e) Depleted of nuclear reads



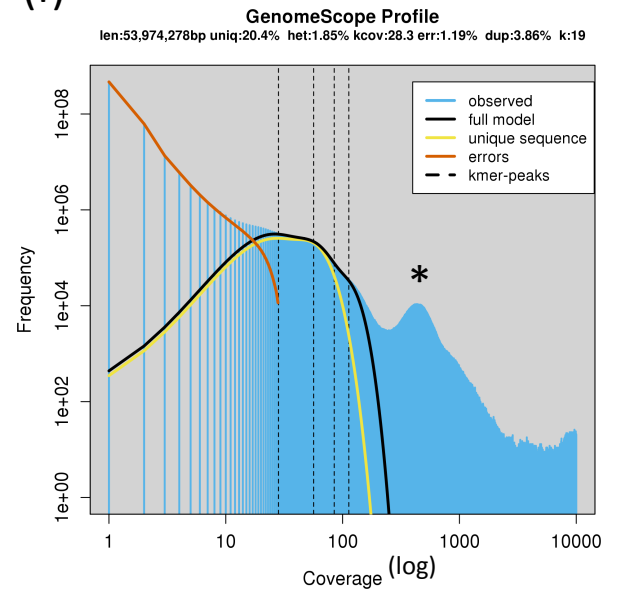
(b)



(d)



(f)



GenomeScope version 1.0

k = 19

property	min	max
Heterozygosity	0.409485%	0.416116%
Genome Haploid Length	26,268,957 bp	26,280,525 bp
Genome Repeat Length	4,867,943 bp	4,870,086 bp
Genome Unique Length	21,401,014 bp	21,410,439 bp
Model Fit	97.5292%	99.2466%
Read Error Rate	1.49446%	1.49446%