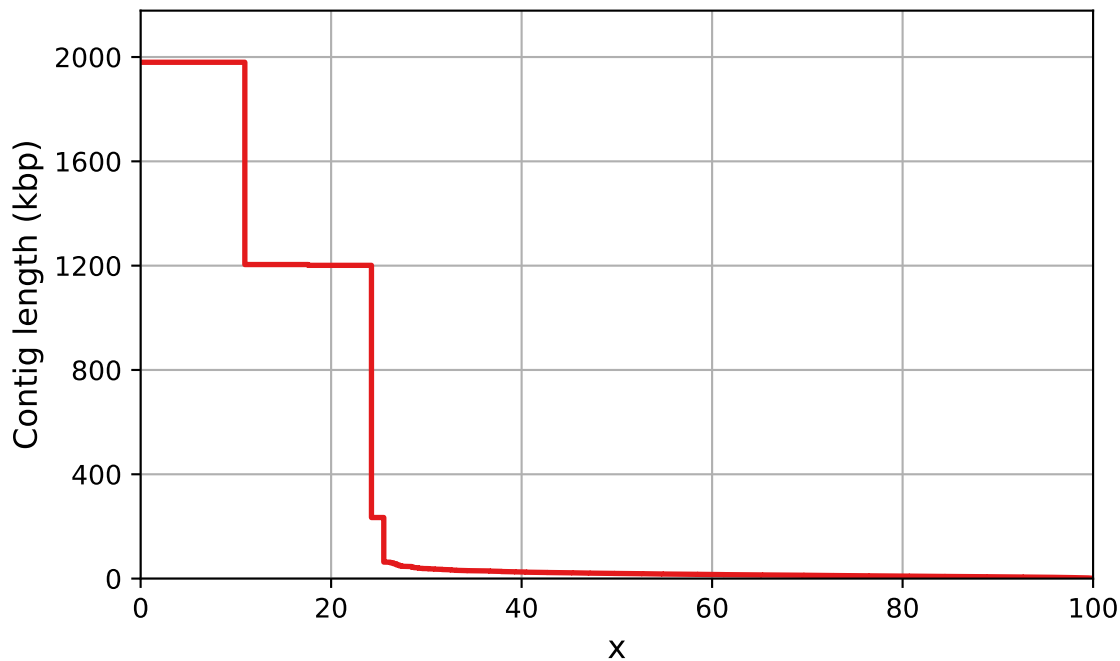


Report

	genome.nextpolish
# contigs (>= 0 bp)	1211
# contigs (>= 1000 bp)	1190
# contigs (>= 5000 bp)	950
# contigs (>= 10000 bp)	523
# contigs (>= 25000 bp)	84
# contigs (>= 50000 bp)	9
Total length (>= 0 bp)	18094815
Total length (>= 1000 bp)	18080415
Total length (>= 5000 bp)	17245081
Total length (>= 10000 bp)	14106712
Total length (>= 25000 bp)	7320826
Total length (>= 50000 bp)	4915333
# contigs	1211
Largest contig	1979896
Total length	18094815
GC (%)	47.60
N50	19659
N90	6744
auN	392602.8
L50	163
L90	787
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	53 + 7 part

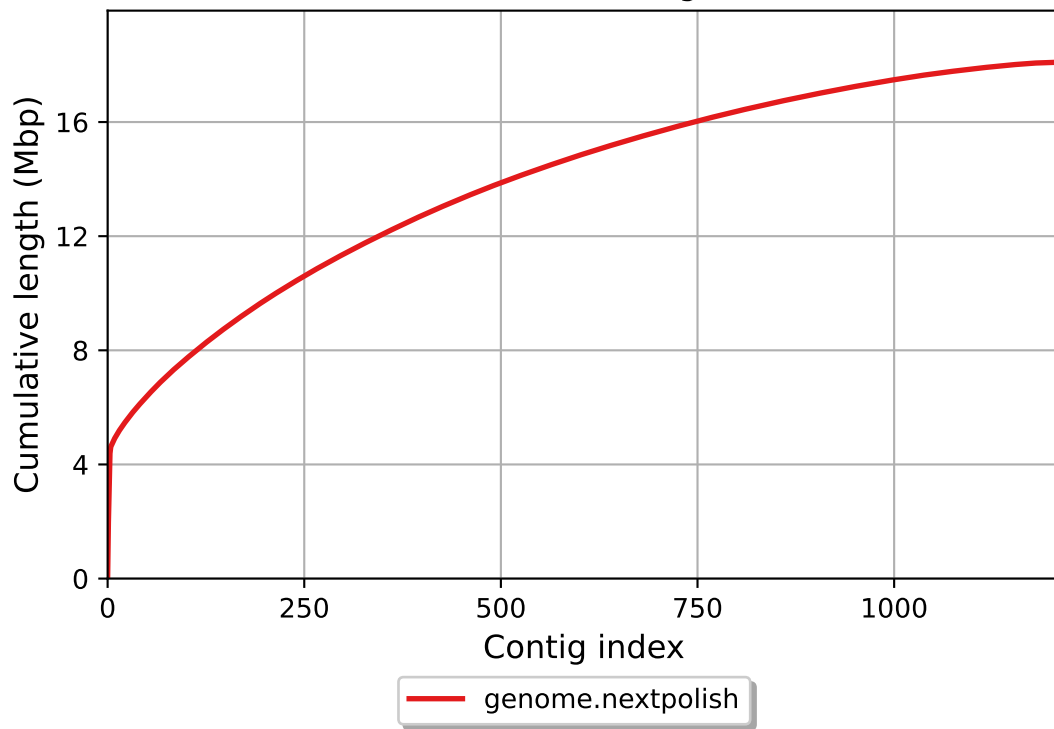
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

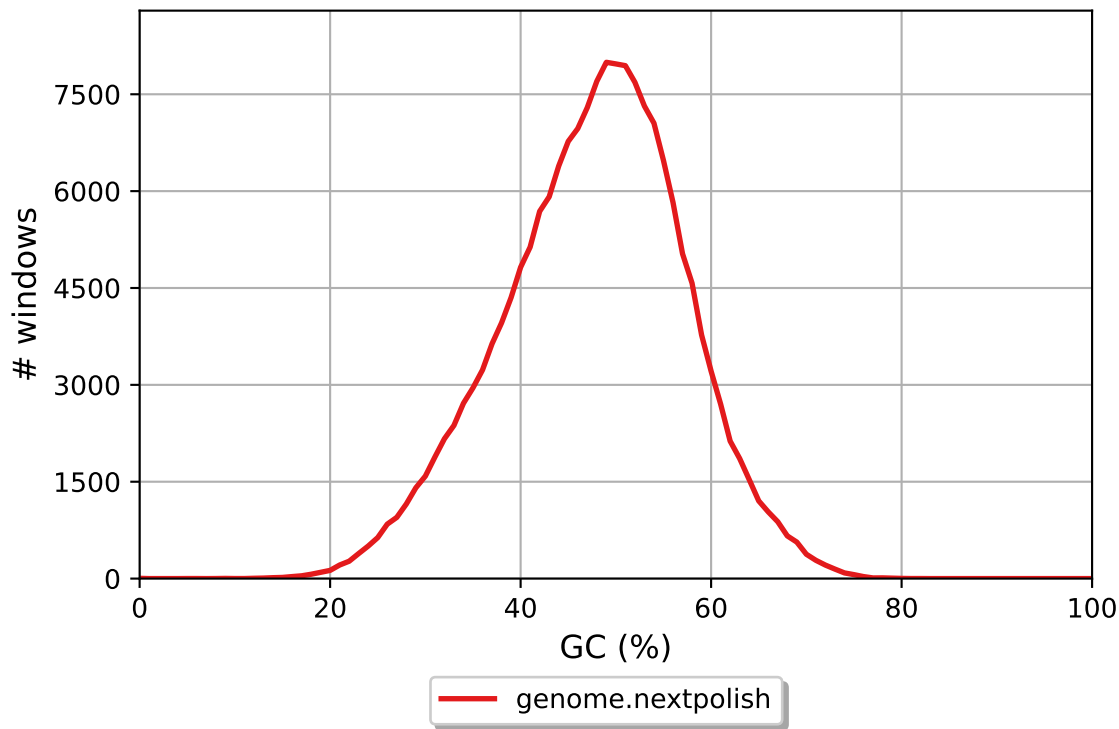


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

