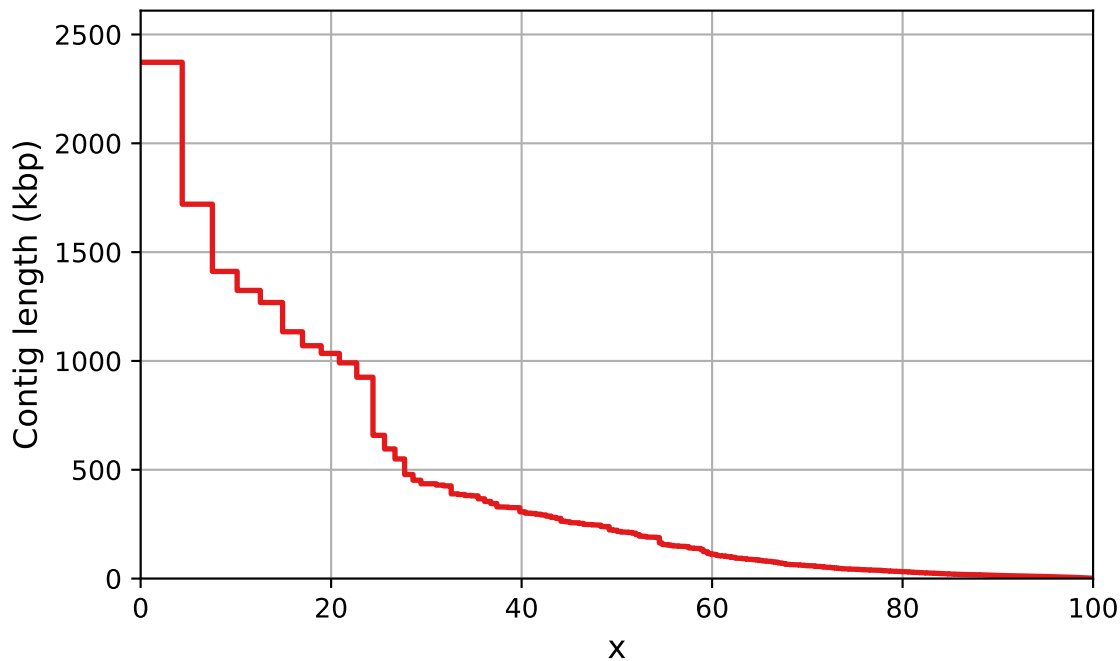


Report

	genome.nextpolish
# contigs (>= 0 bp)	1364
# contigs (>= 1000 bp)	1321
# contigs (>= 5000 bp)	1025
# contigs (>= 10000 bp)	738
# contigs (>= 25000 bp)	336
# contigs (>= 50000 bp)	178
Total length (>= 0 bp)	54320395
Total length (>= 1000 bp)	54292854
Total length (>= 5000 bp)	53401073
Total length (>= 10000 bp)	51306196
Total length (>= 25000 bp)	45067745
Total length (>= 50000 bp)	39475600
# contigs	1361
Largest contig	2372343
Total length	54319561
GC (%)	45.95
N50	221988
N90	14554
auN	482400.3
L50	51
L90	539
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	102 + 0 part

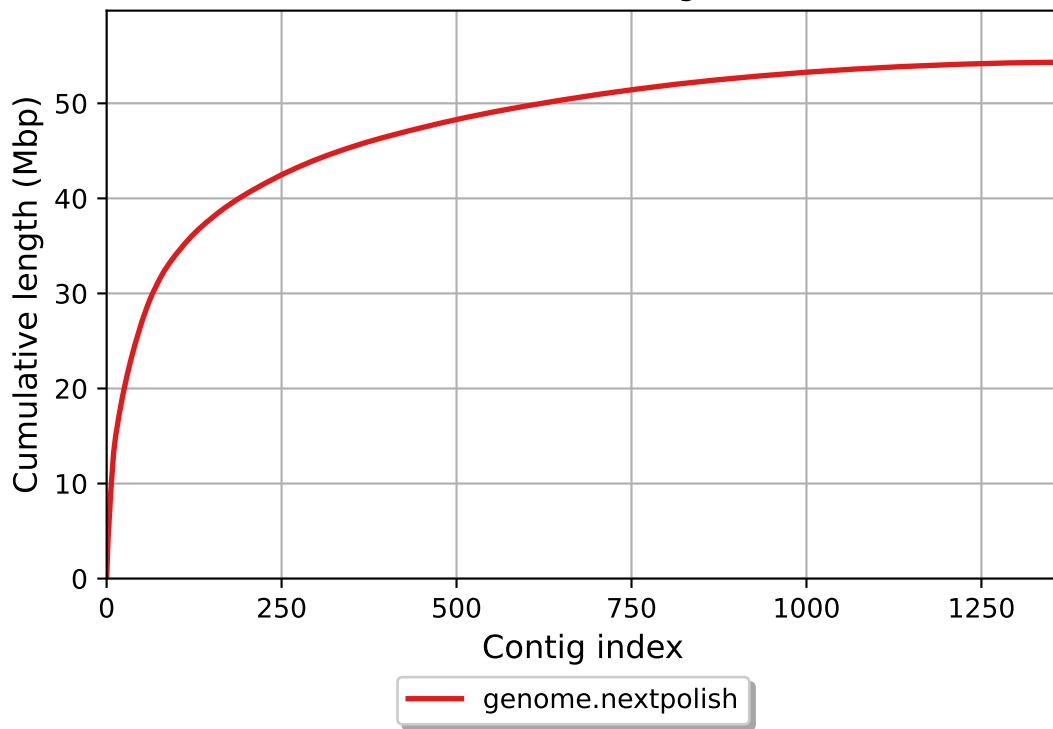
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

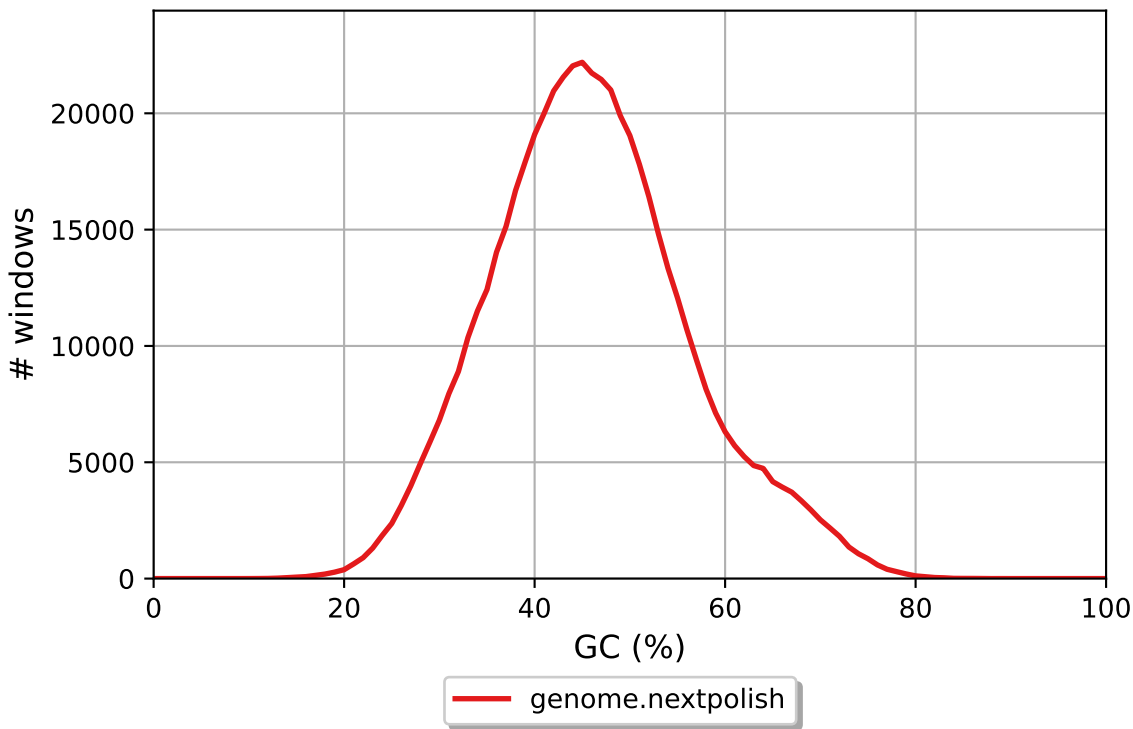


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

