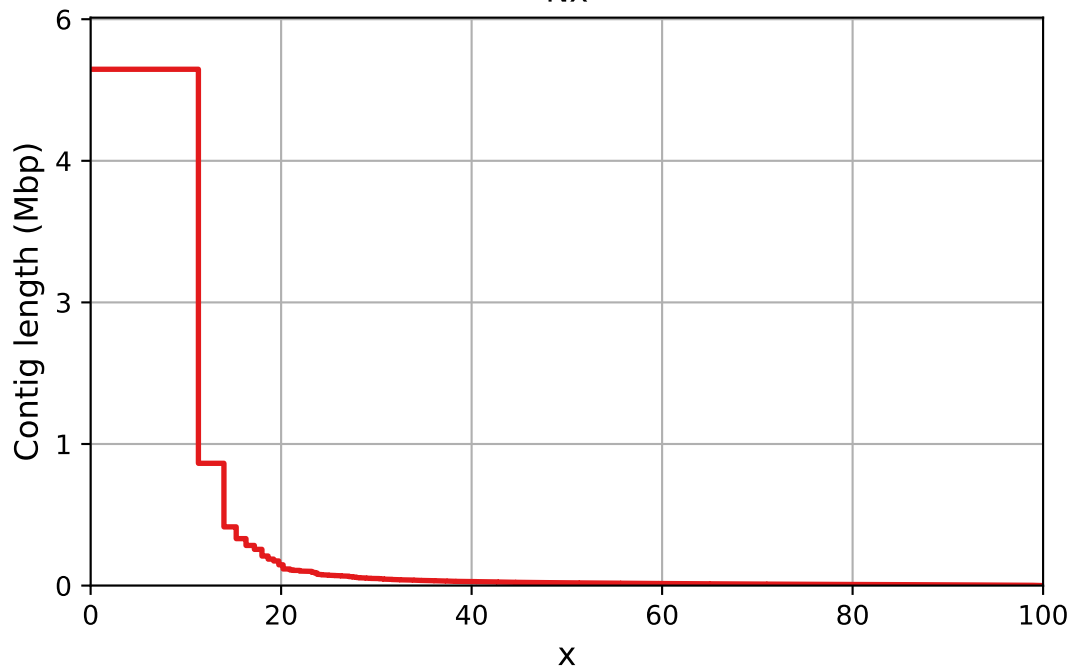


Report

	genome.nextpolish
# contigs (>= 0 bp)	2658
# contigs (>= 1000 bp)	2633
# contigs (>= 5000 bp)	2023
# contigs (>= 10000 bp)	1285
# contigs (>= 25000 bp)	387
# contigs (>= 50000 bp)	105
Total length (>= 0 bp)	48333461
Total length (>= 1000 bp)	48316133
Total length (>= 5000 bp)	46232245
Total length (>= 10000 bp)	40834539
Total length (>= 25000 bp)	26988981
Total length (>= 50000 bp)	17606192
# contigs	2658
Largest contig	5469719
Total length	48333461
GC (%)	55.73
N50	29289
N90	7715
auN	707658.4
L50	283
L90	1587
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	72 + 17 part

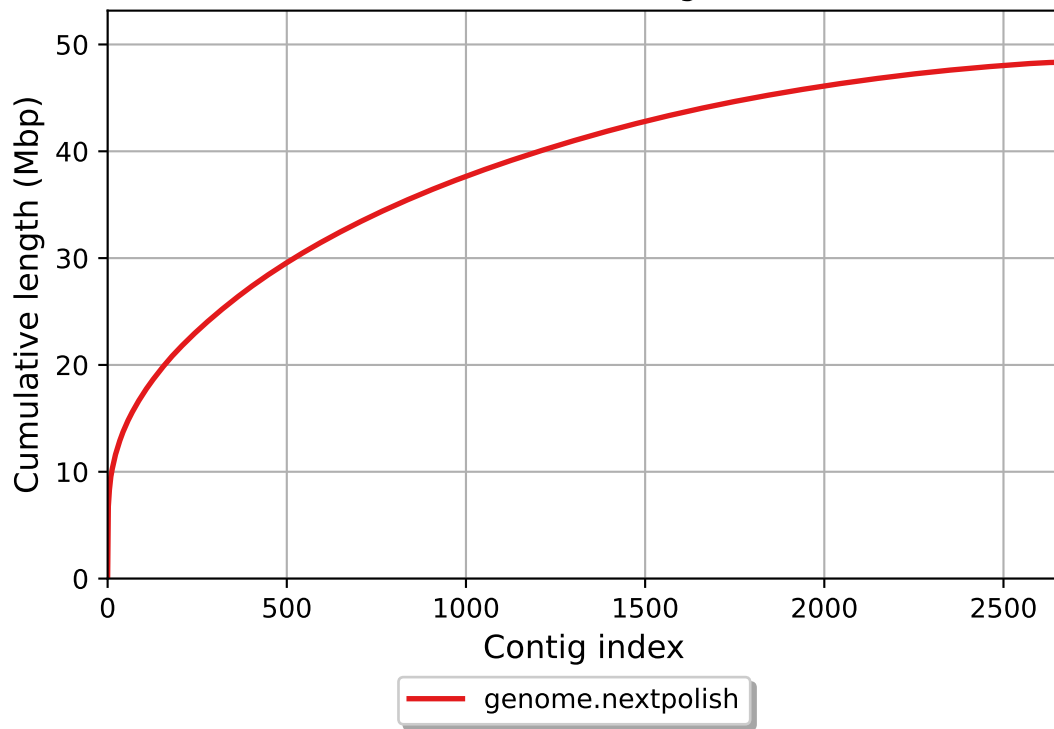
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

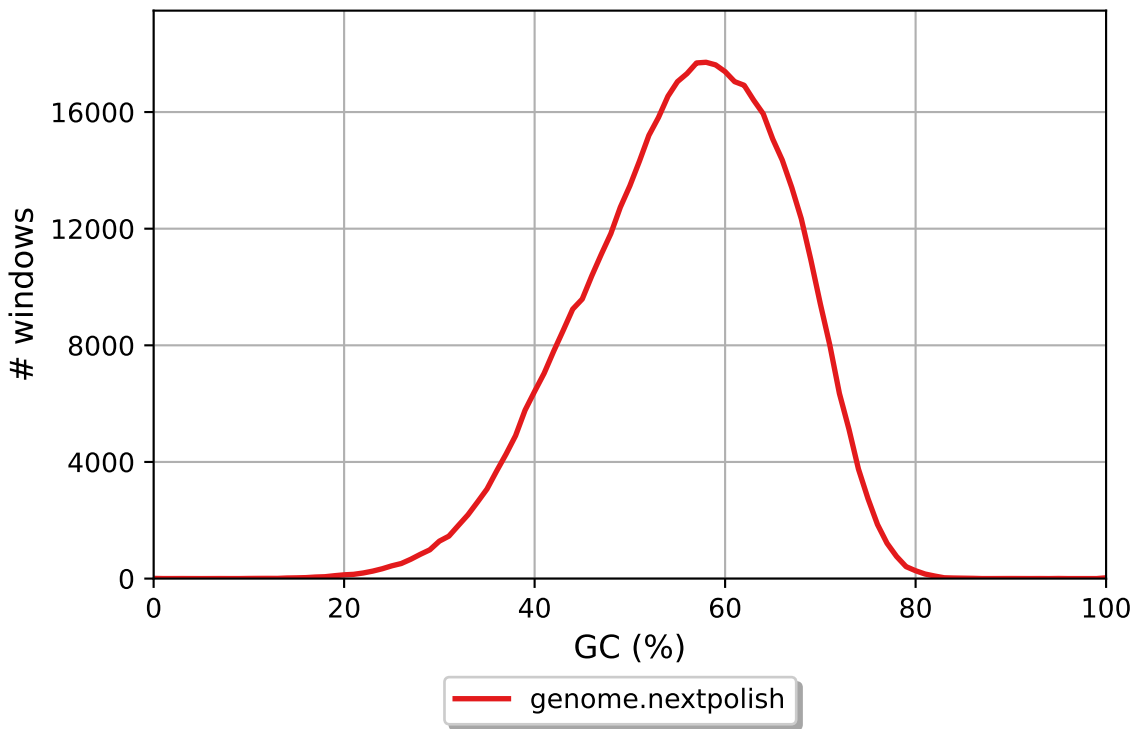


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

