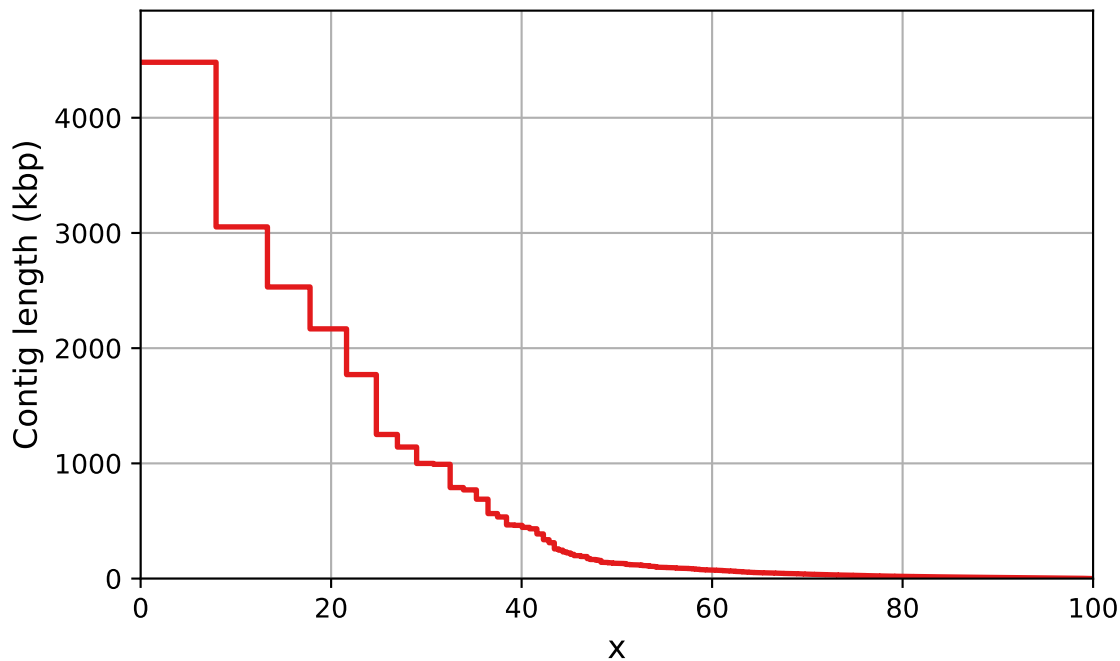


Report

	genome.nextpolish
# contigs (>= 0 bp)	2341
# contigs (>= 1000 bp)	2156
# contigs (>= 5000 bp)	1304
# contigs (>= 10000 bp)	803
# contigs (>= 25000 bp)	328
# contigs (>= 50000 bp)	152
Total length (>= 0 bp)	56591269
Total length (>= 1000 bp)	56465847
Total length (>= 5000 bp)	54165192
Total length (>= 10000 bp)	50586110
Total length (>= 25000 bp)	43306951
Total length (>= 50000 bp)	37093826
# contigs	2316
Largest contig	4481583
Total length	56582290
GC (%)	49.21
N50	132671
N90	9432
auN	950849.4
L50	42
L90	838
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	112 + 2 part

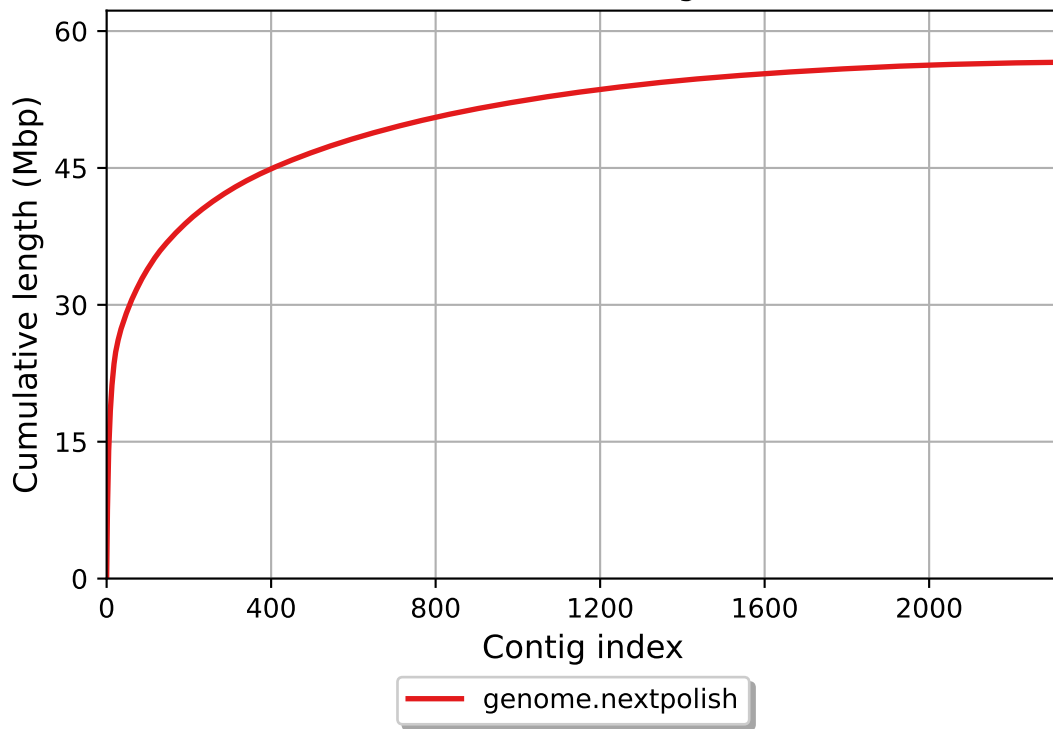
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

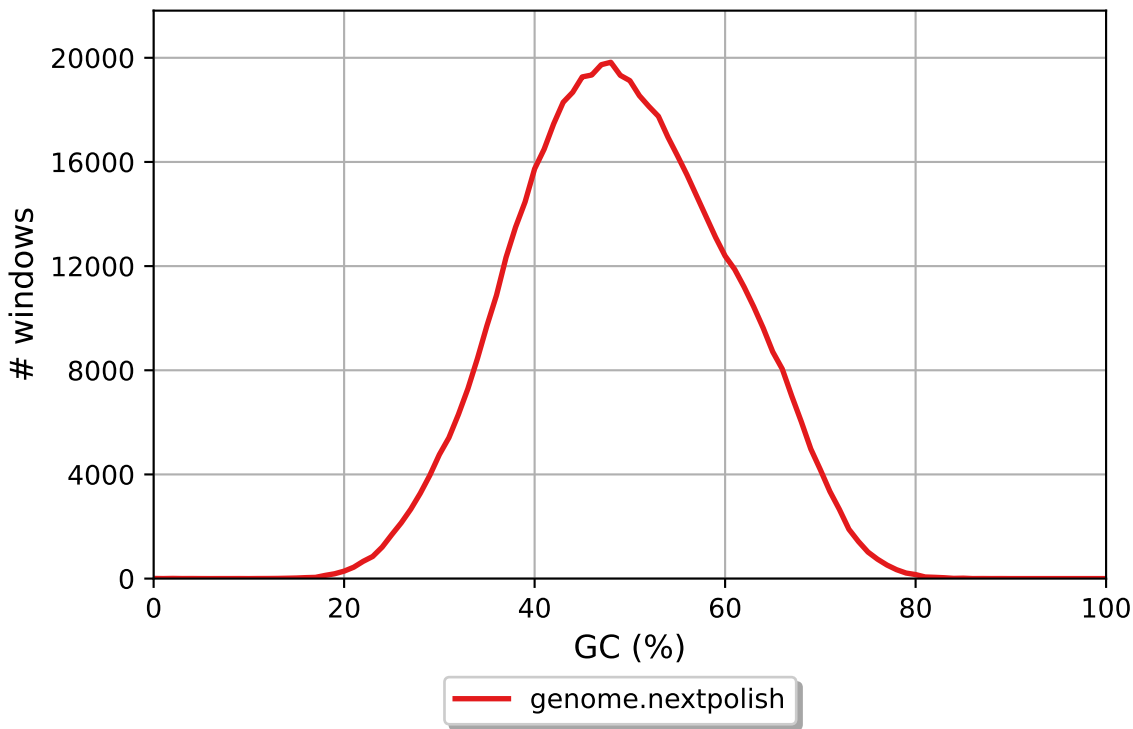


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

