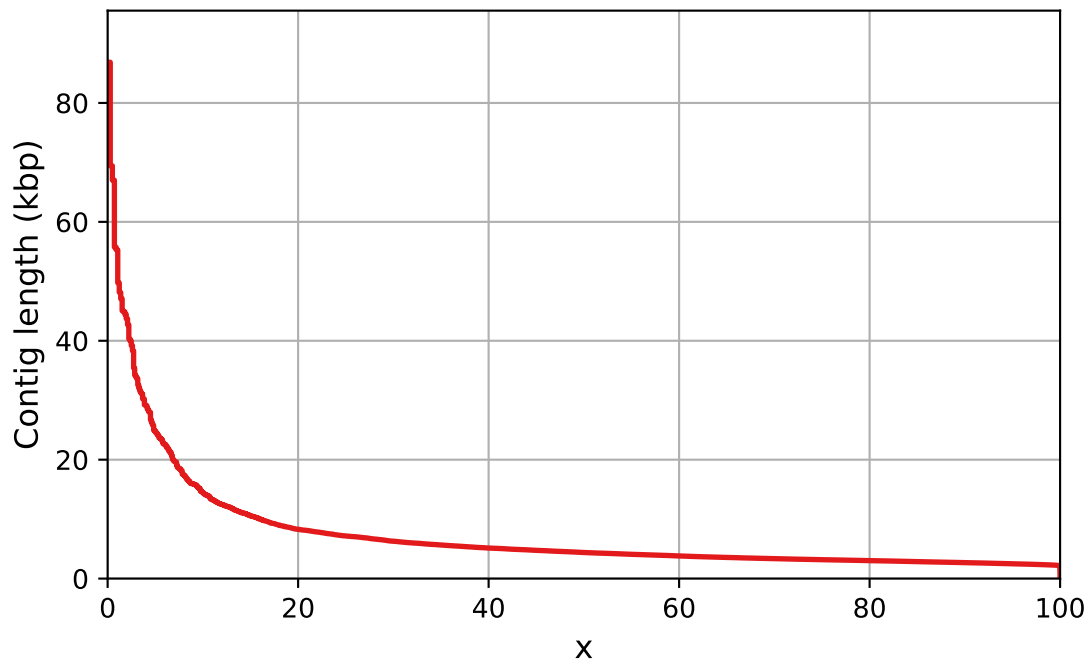


Report

	genome.nextpolish
# contigs (>= 0 bp)	7098
# contigs (>= 1000 bp)	7095
# contigs (>= 5000 bp)	1532
# contigs (>= 10000 bp)	286
# contigs (>= 25000 bp)	39
# contigs (>= 50000 bp)	5
Total length (>= 0 bp)	31439186
Total length (>= 1000 bp)	31436782
Total length (>= 5000 bp)	13151255
Total length (>= 10000 bp)	5035827
Total length (>= 25000 bp)	1519199
Total length (>= 50000 bp)	334348
# contigs	7098
Largest contig	86842
Total length	31439186
GC (%)	39.65
N50	4385
N90	2681
auN	7558.4
L50	2081
L90	5825
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	7 + 5 part

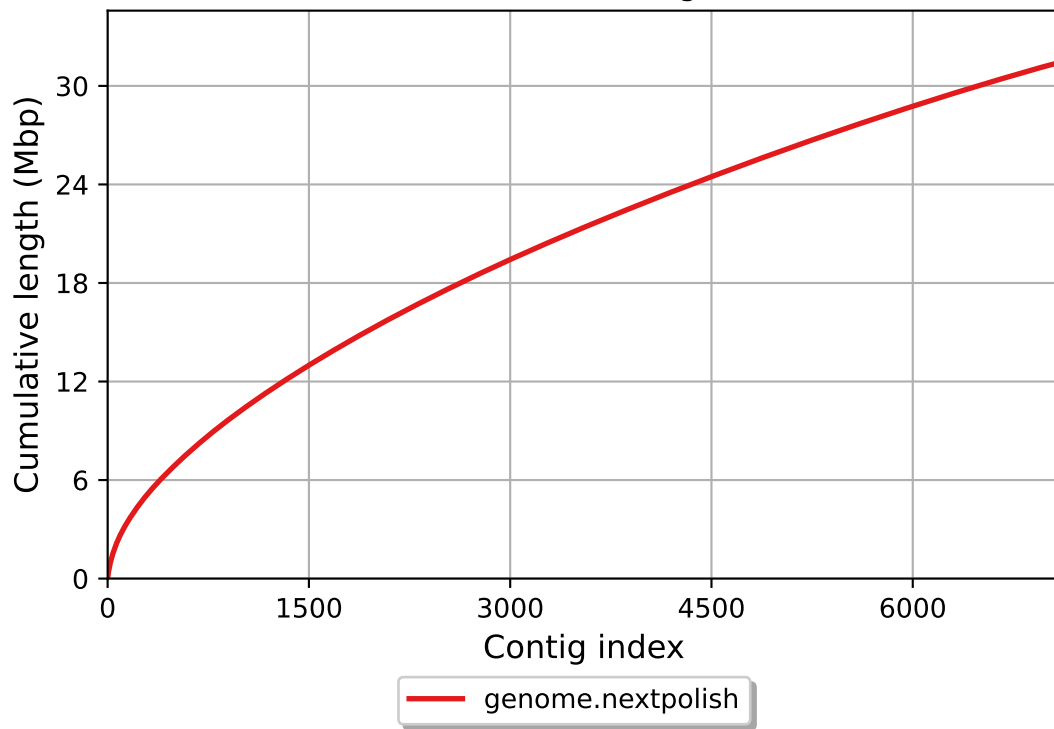
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

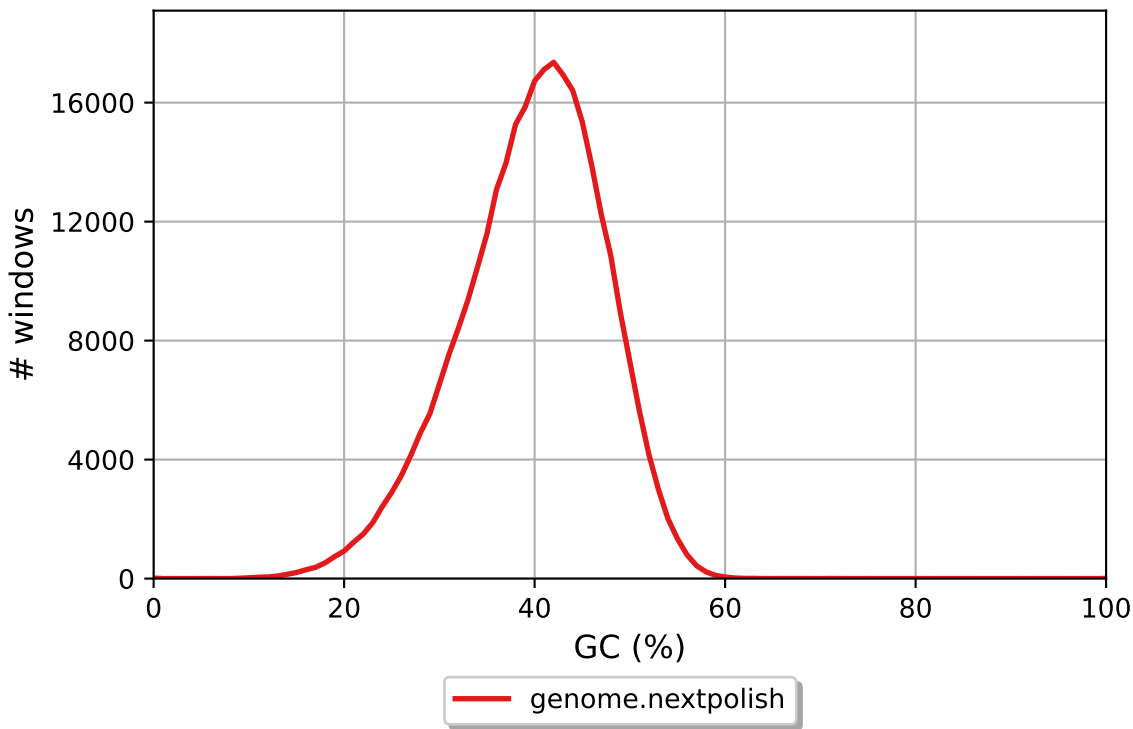


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

