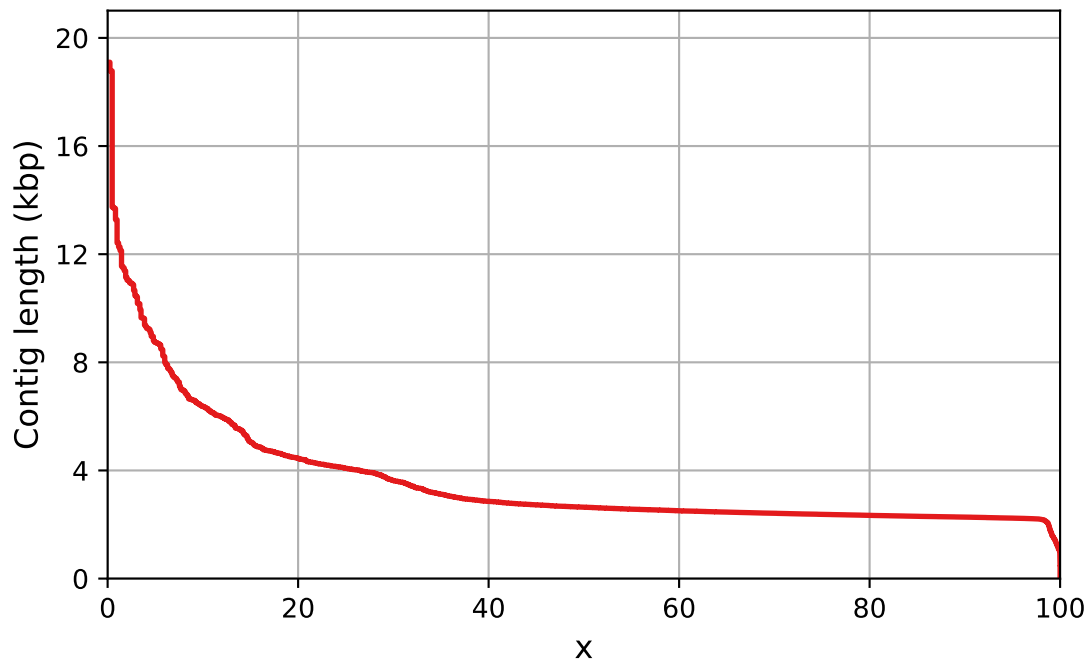


Report

	genome.nextpolish
# contigs (>= 0 bp)	2665
# contigs (>= 1000 bp)	2662
# contigs (>= 5000 bp)	162
# contigs (>= 10000 bp)	22
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	7911433
Total length (>= 1000 bp)	7909585
Total length (>= 5000 bp)	1204248
Total length (>= 10000 bp)	267636
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	2665
Largest contig	19099
Total length	7911433
GC (%)	36.83
N50	2640
N90	2273
auN	3652.2
L50	979
L90	2284
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	8 + 3 part

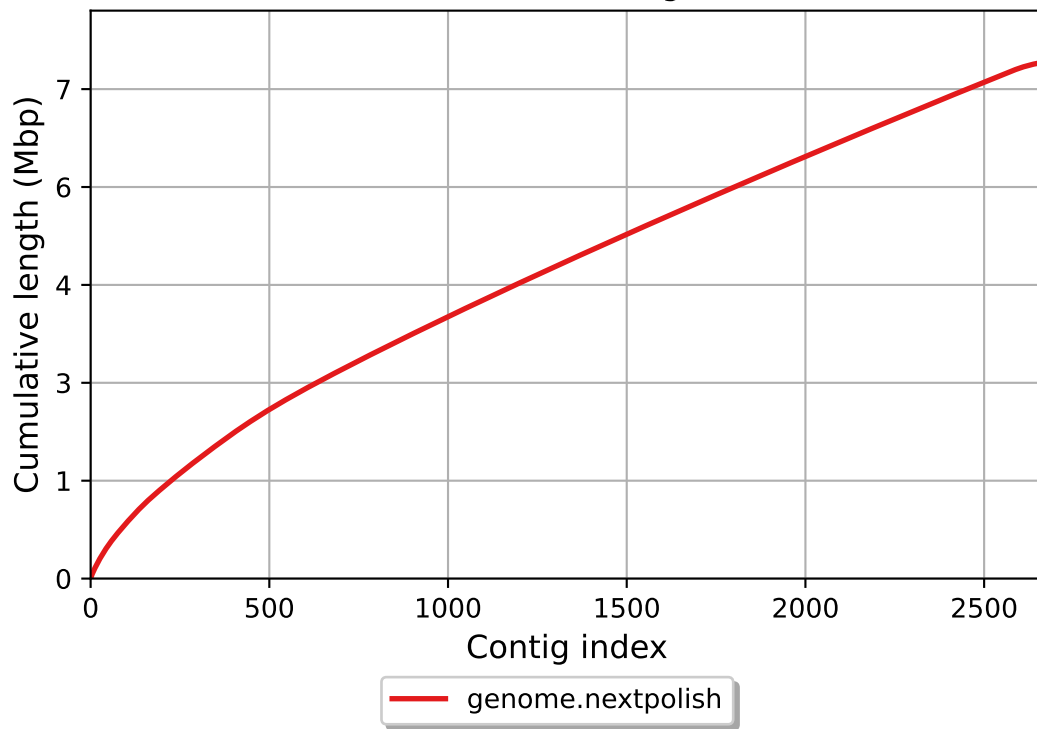
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

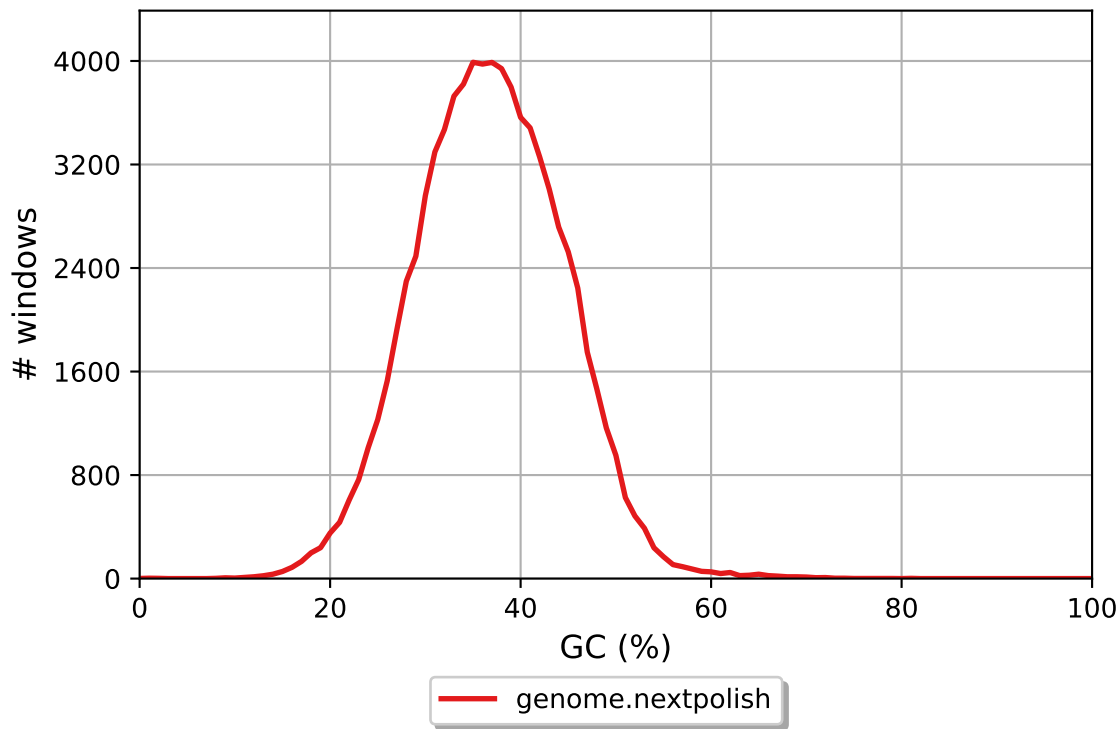


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

