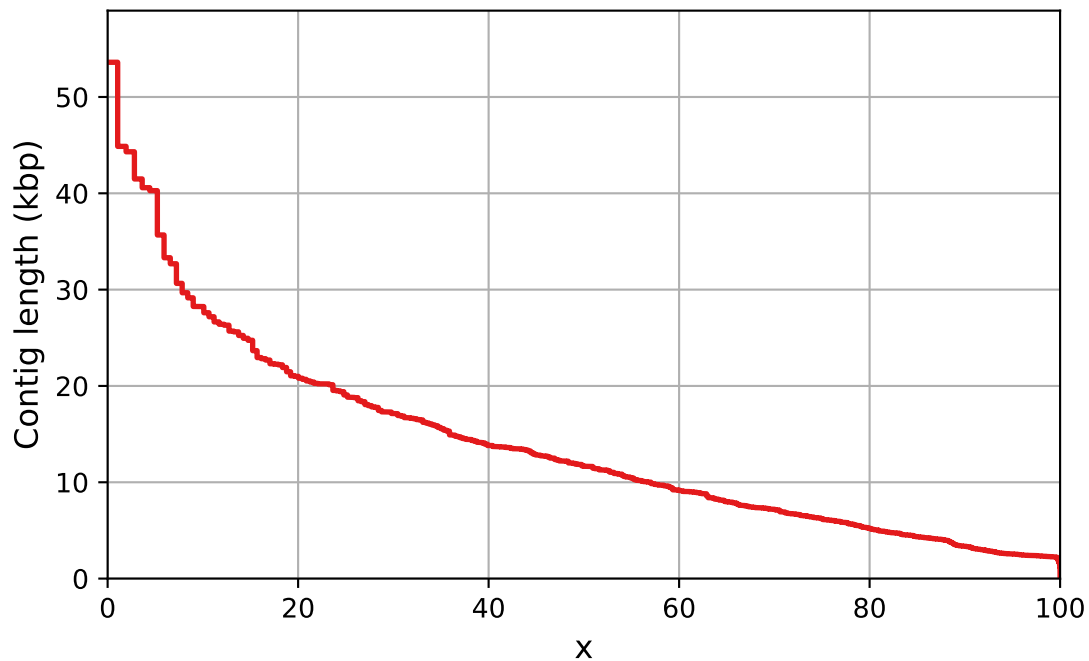


Report

	genome.nextpolish
# contigs (>= 0 bp)	645
# contigs (>= 1000 bp)	644
# contigs (>= 5000 bp)	338
# contigs (>= 10000 bp)	167
# contigs (>= 25000 bp)	22
# contigs (>= 50000 bp)	1
Total length (>= 0 bp)	5077859
Total length (>= 1000 bp)	5077286
Total length (>= 5000 bp)	4111544
Total length (>= 10000 bp)	2896743
Total length (>= 25000 bp)	723468
Total length (>= 50000 bp)	53603
# contigs	645
Largest contig	53603
Total length	5077859
GC (%)	50.75
N50	11696
N90	3356
auN	14174.5
L50	134
L90	447
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	3 + 1 part

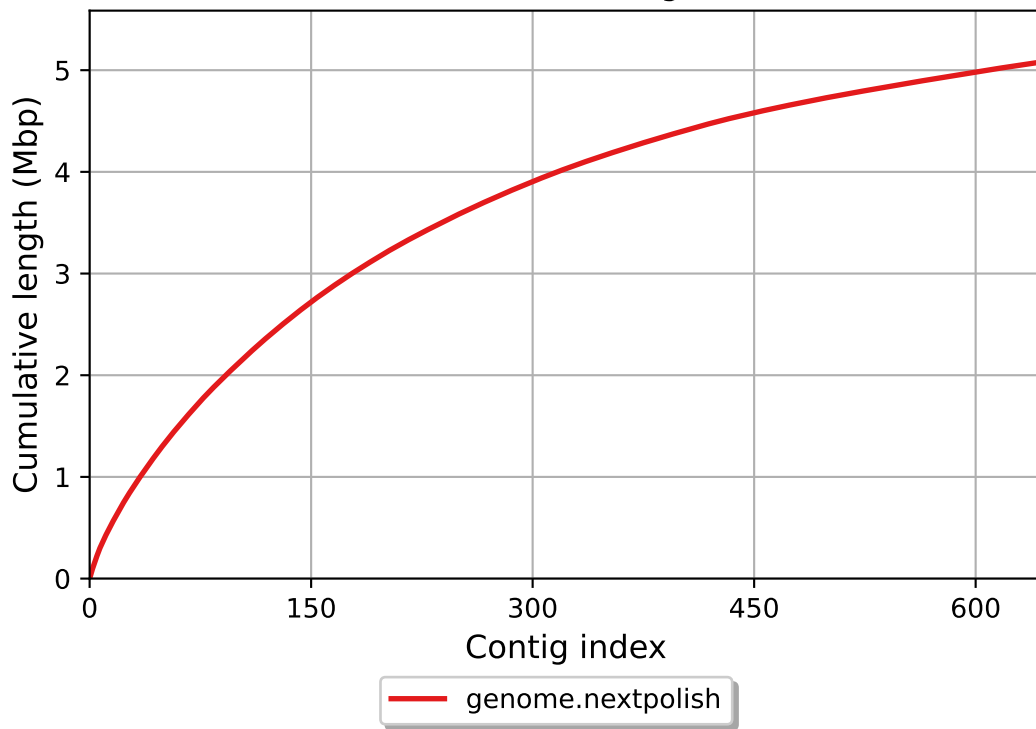
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

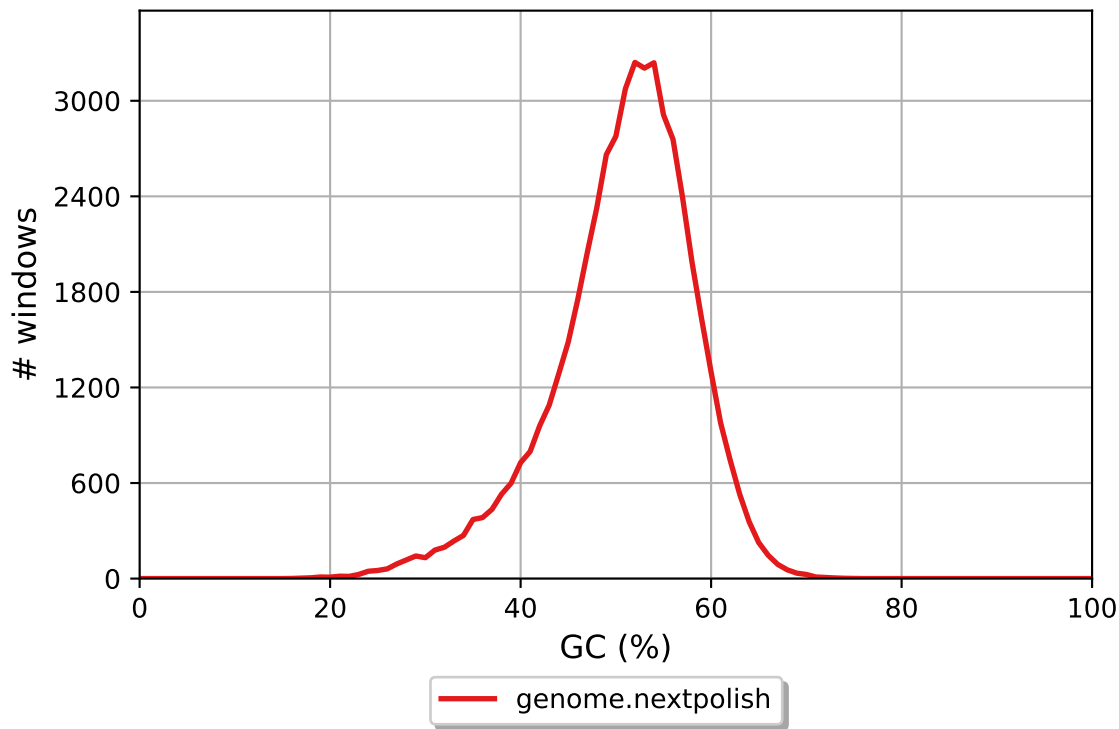


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

