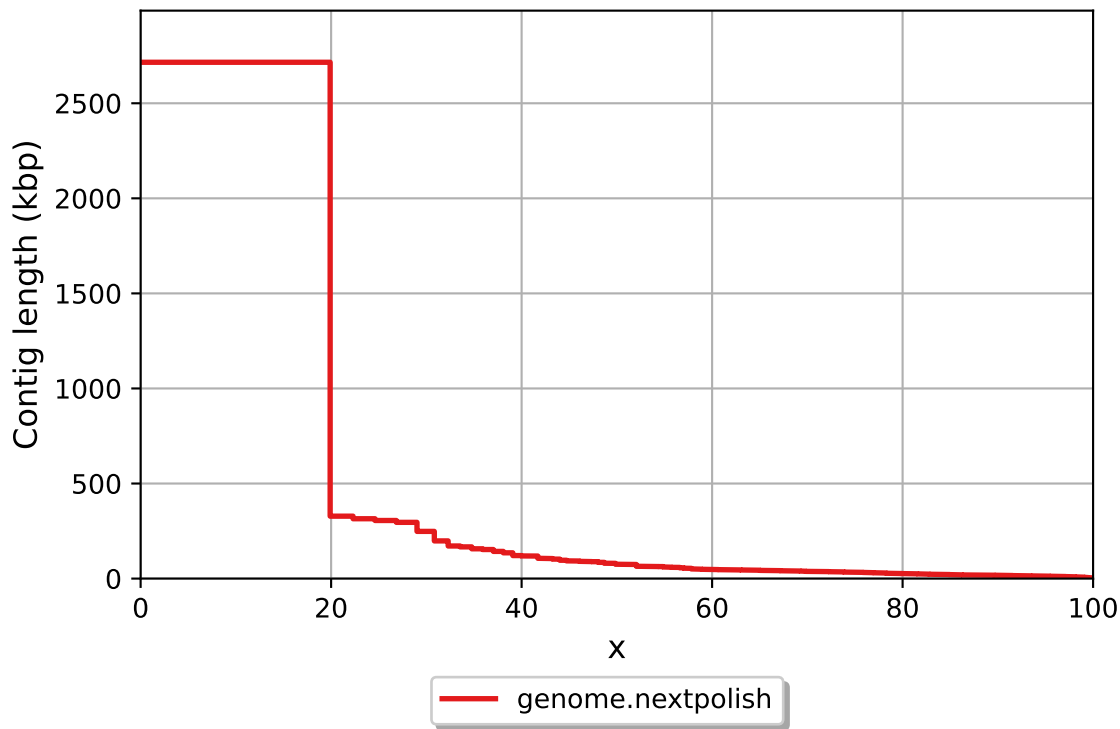


Report

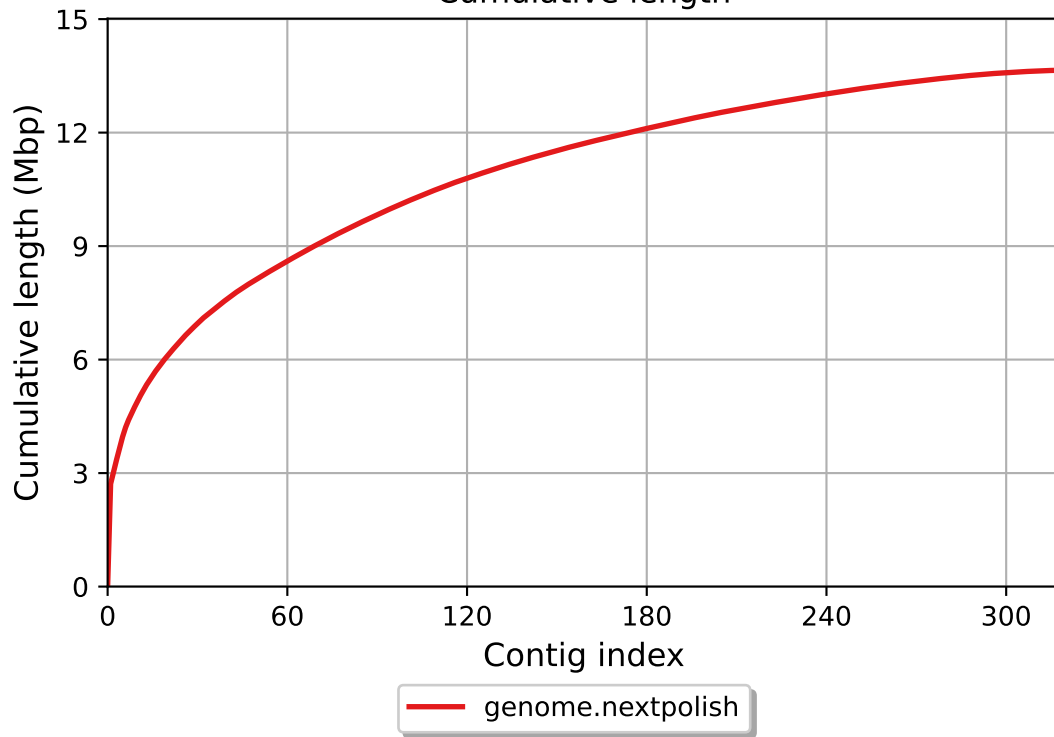
	genome.nextpolish
# contigs (>= 0 bp)	318
# contigs (>= 1000 bp)	318
# contigs (>= 5000 bp)	301
# contigs (>= 10000 bp)	264
# contigs (>= 25000 bp)	132
# contigs (>= 50000 bp)	46
Total length (>= 0 bp)	13647776
Total length (>= 1000 bp)	13647776
Total length (>= 5000 bp)	13589876
Total length (>= 10000 bp)	13298163
Total length (>= 25000 bp)	11108464
Total length (>= 50000 bp)	7941456
# contigs	318
Largest contig	2715631
Total length	13647776
GC (%)	47.04
N50	75468
N90	17578
auN	614675.1
L50	29
L90	190
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	32 + 1 part

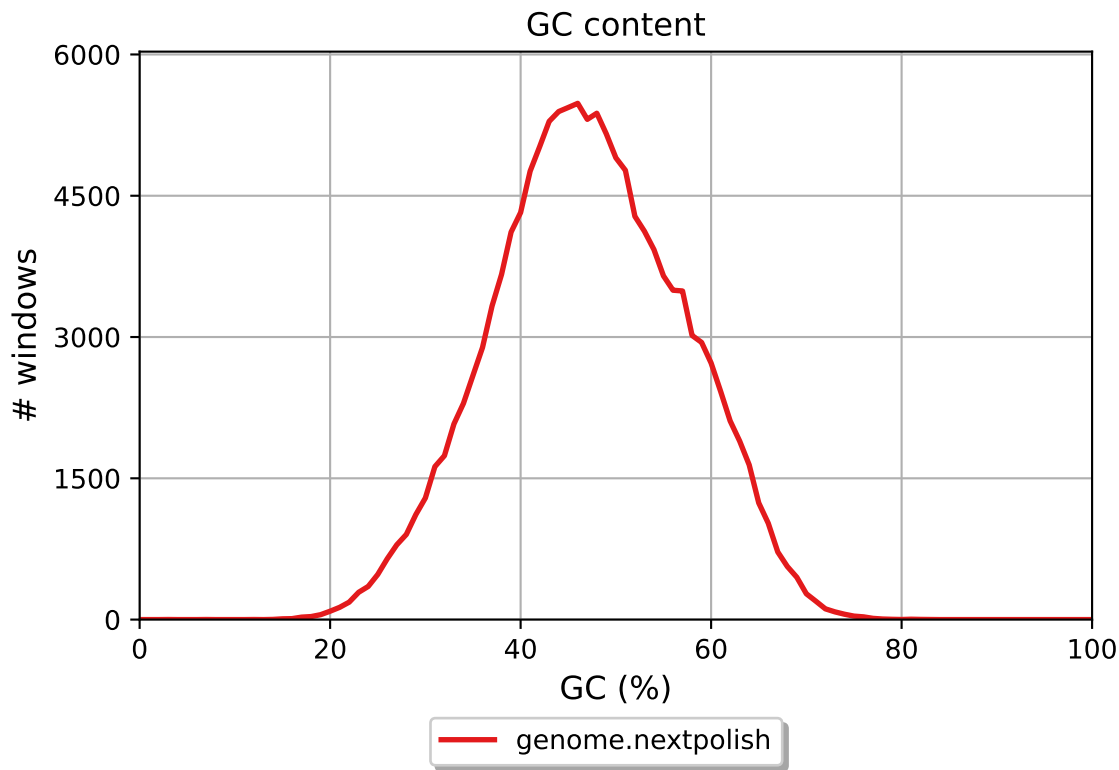
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx



Cumulative length





genome.nextpolish GC content

