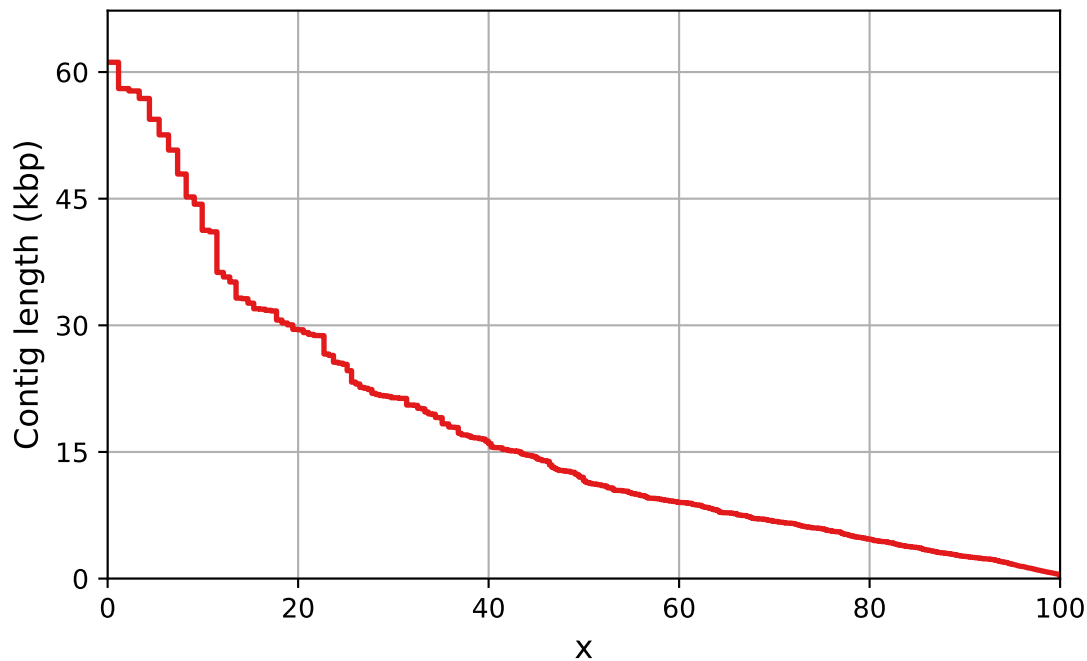


Report

	genome.nextpolish
# contigs (>= 0 bp)	890
# contigs (>= 1000 bp)	710
# contigs (>= 5000 bp)	309
# contigs (>= 10000 bp)	141
# contigs (>= 25000 bp)	36
# contigs (>= 50000 bp)	7
Total length (>= 0 bp)	5330925
Total length (>= 1000 bp)	5207838
Total length (>= 5000 bp)	4171637
Total length (>= 10000 bp)	2952011
Total length (>= 25000 bp)	1339946
Total length (>= 50000 bp)	391549
# contigs	881
Largest contig	61158
Total length	5328182
GC (%)	40.36
N50	11605
N90	2639
auN	17431.9
L50	115
L90	479
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	11 + 1 part

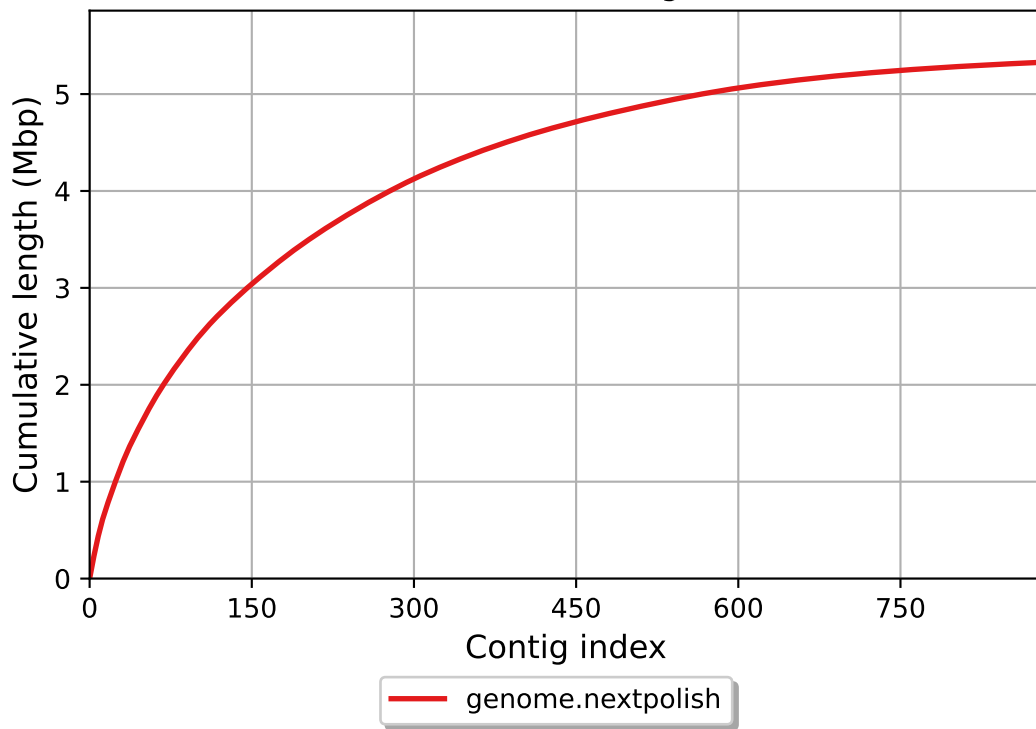
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

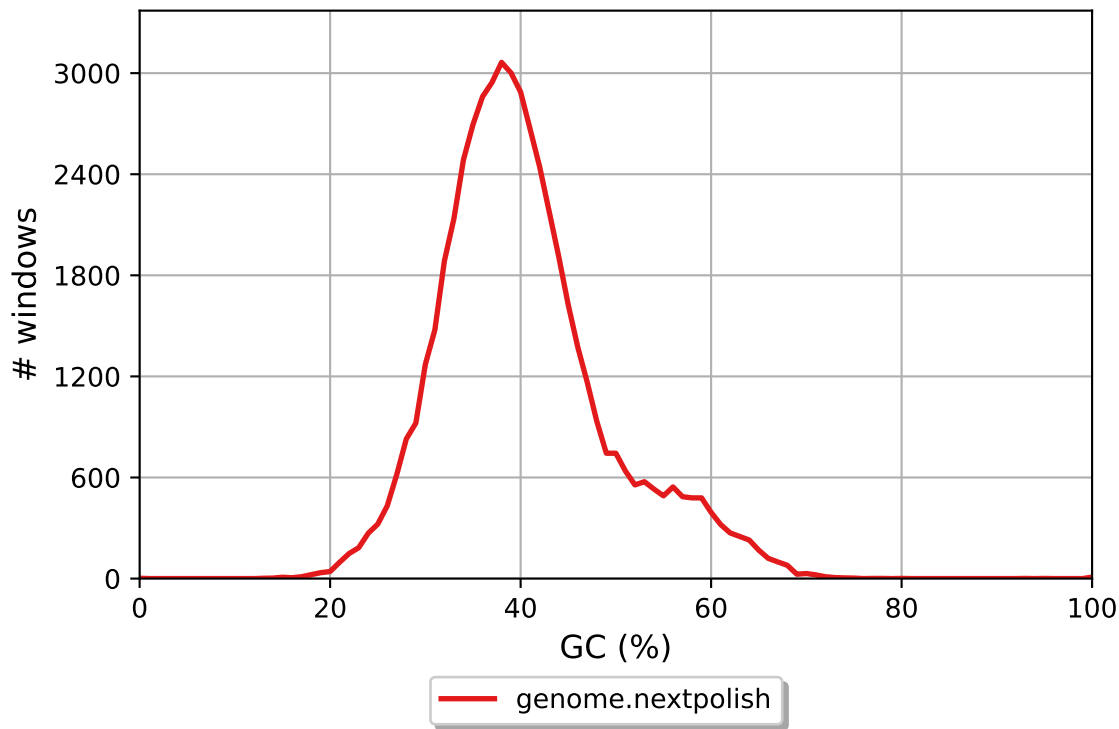


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

