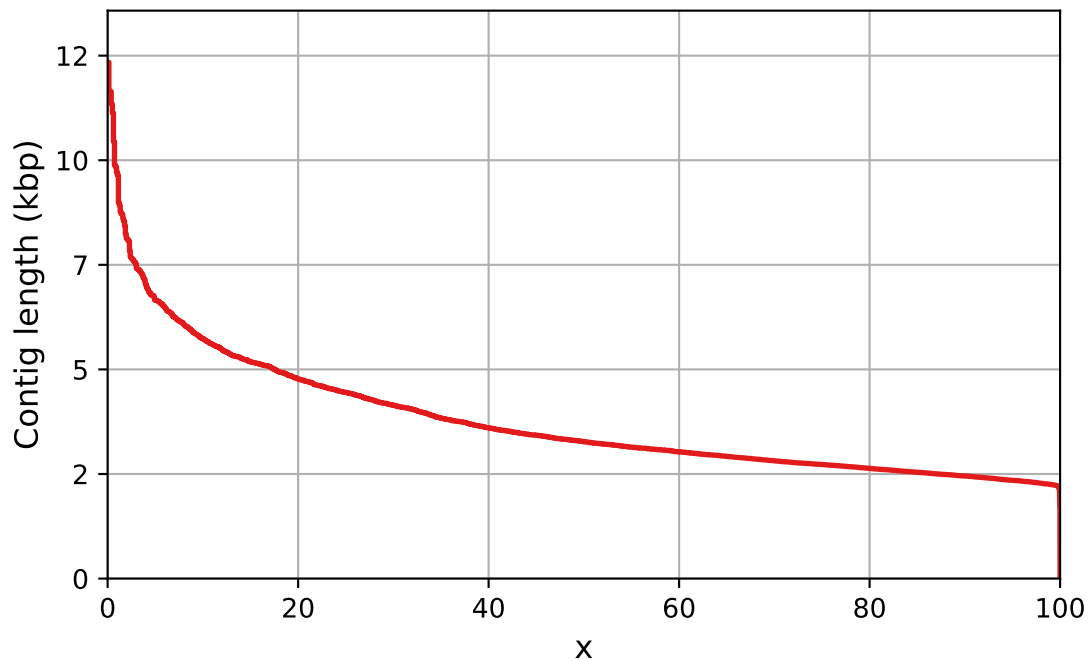


Report

	genome.nextpolish
# contigs (>= 0 bp)	2815
# contigs (>= 1000 bp)	2813
# contigs (>= 5000 bp)	271
# contigs (>= 10000 bp)	6
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	9506172
Total length (>= 1000 bp)	9504749
Total length (>= 5000 bp)	1669081
Total length (>= 10000 bp)	68561
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	2815
Largest contig	12341
Total length	9506172
GC (%)	36.74
N50	3272
N90	2453
auN	3788.6
L50	1057
L90	2408
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	8 + 0 part

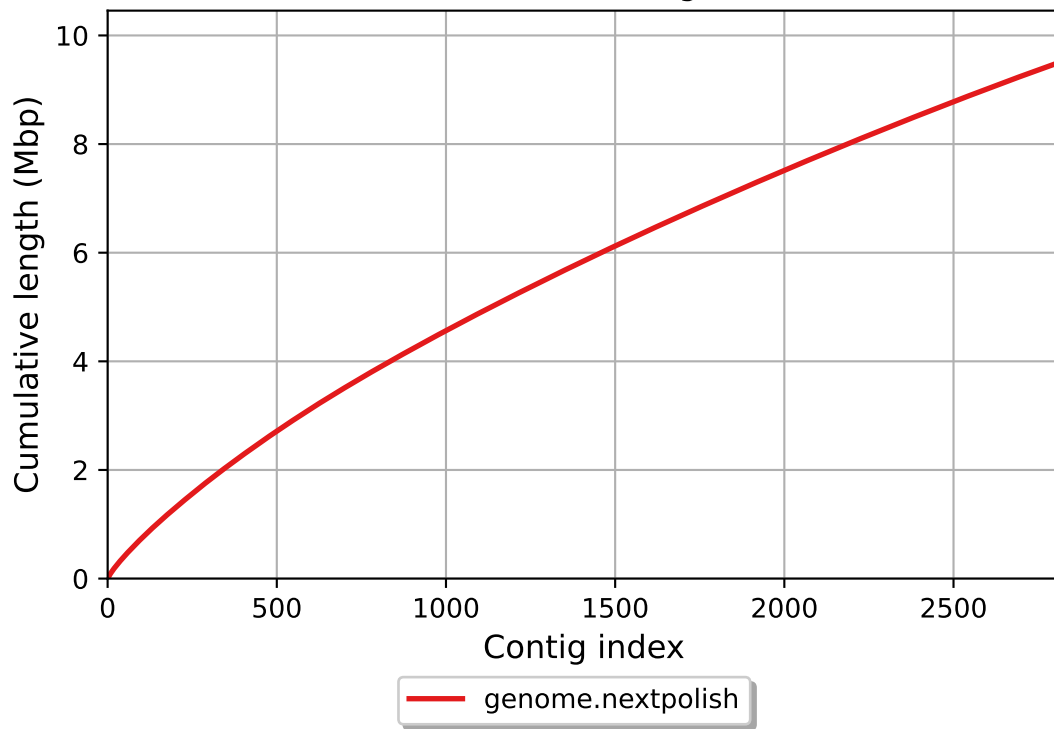
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

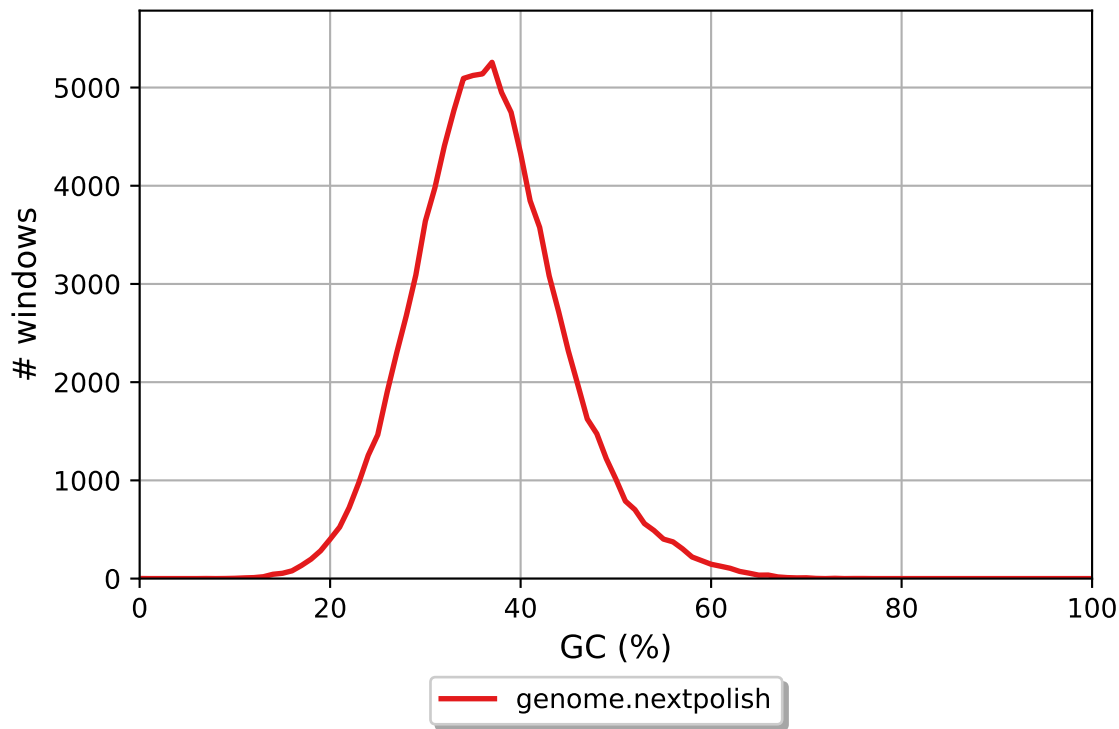


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

