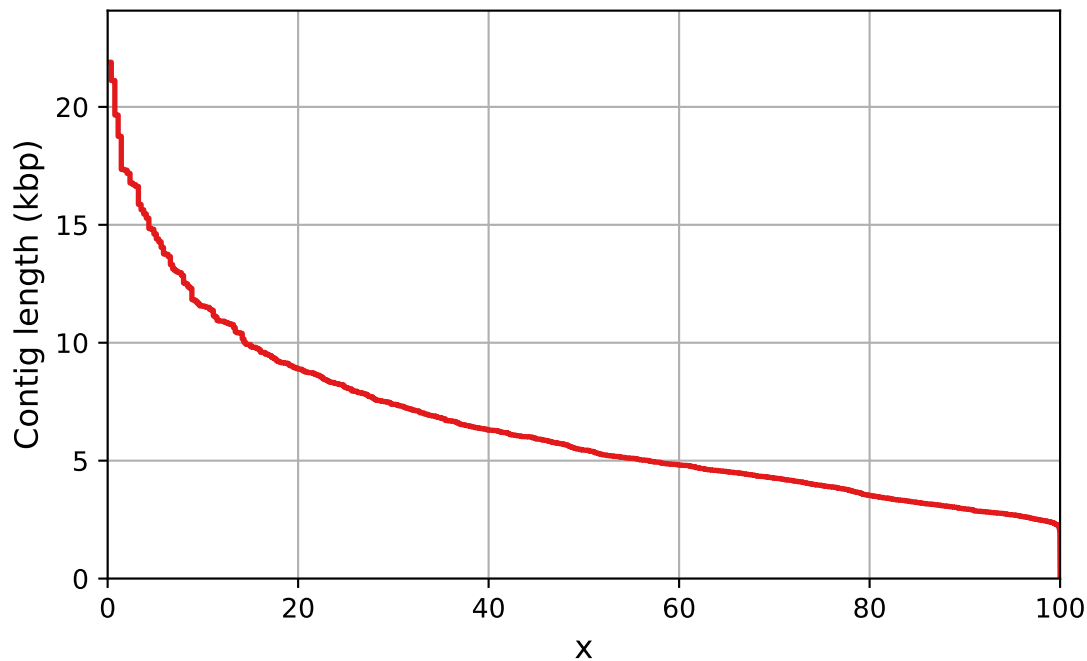


Report

	genome.nextpolish
# contigs (>= 0 bp)	1116
# contigs (>= 1000 bp)	1116
# contigs (>= 5000 bp)	420
# contigs (>= 10000 bp)	62
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	5671627
Total length (>= 1000 bp)	5671627
Total length (>= 5000 bp)	3226378
Total length (>= 10000 bp)	821767
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	1116
Largest contig	21891
Total length	5671627
GC (%)	41.42
N50	5448
N90	2954
auN	6555.4
L50	345
L90	902
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	13 + 2 part

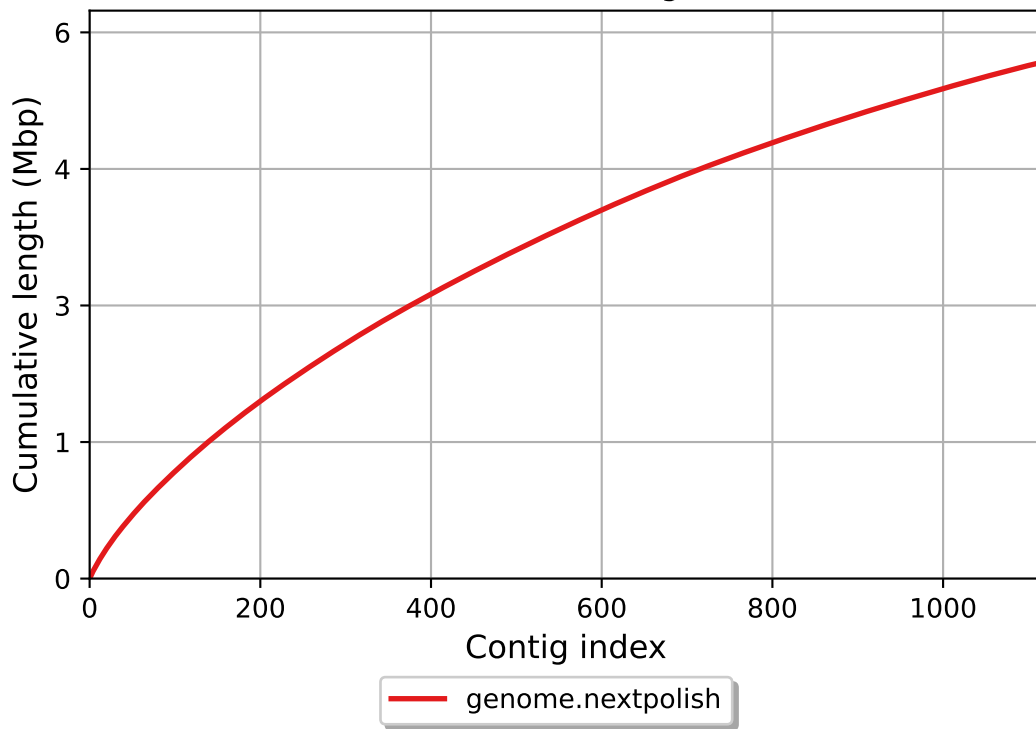
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

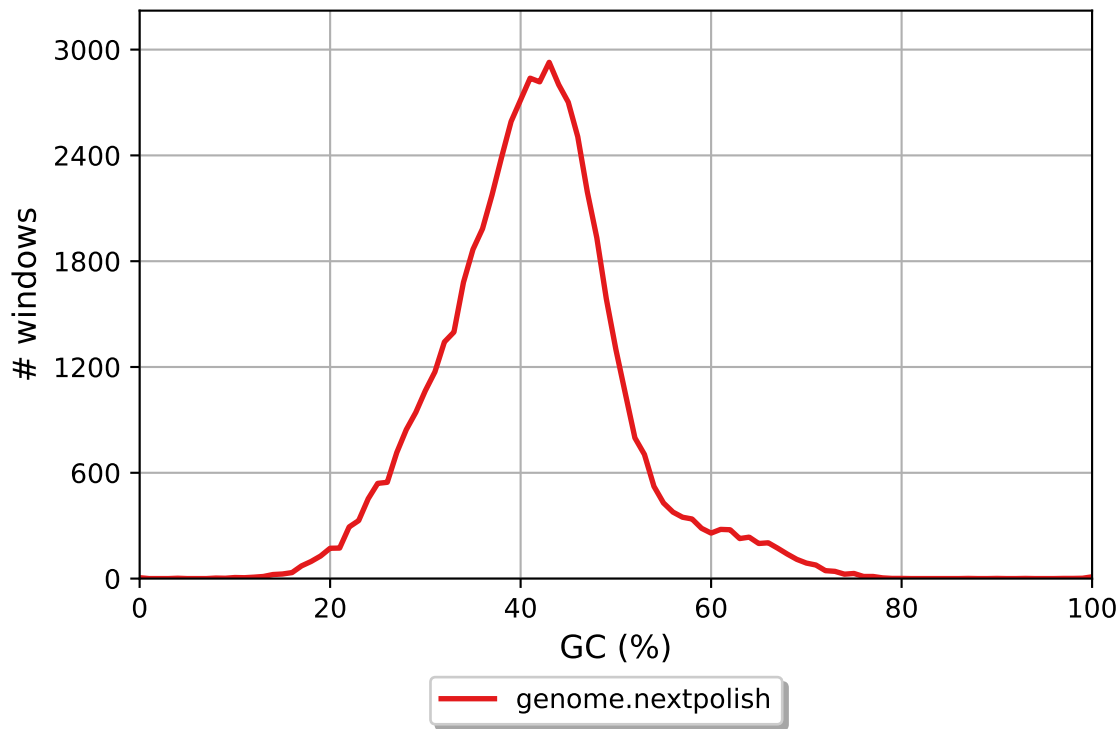


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

