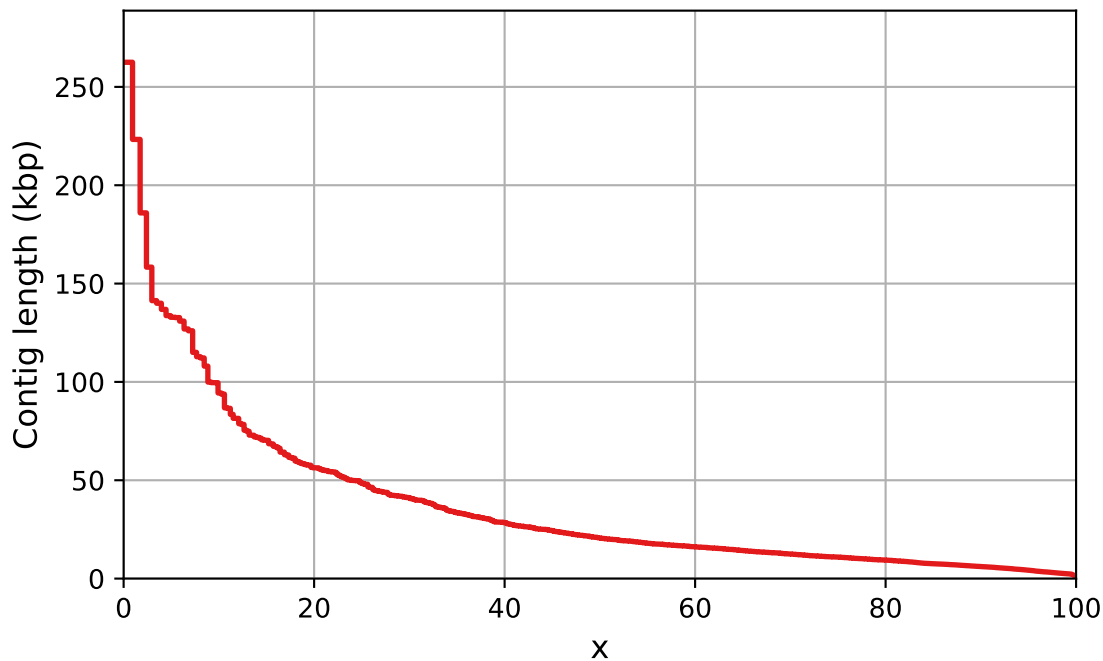


# Report

	genome.nextpolish
# contigs (>= 0 bp)	2065
# contigs (>= 1000 bp)	2055
# contigs (>= 5000 bp)	1466
# contigs (>= 10000 bp)	868
# contigs (>= 25000 bp)	247
# contigs (>= 50000 bp)	81
Total length (>= 0 bp)	28030433
Total length (>= 1000 bp)	28023143
Total length (>= 5000 bp)	26149630
Total length (>= 10000 bp)	21824977
Total length (>= 25000 bp)	12393293
Total length (>= 50000 bp)	6694551
# contigs	2065
Largest contig	262519
Total length	28030433
GC (%)	45.50
N50	20656
N90	6132
auN	38313.6
L50	319
L90	1301
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	51 + 5 part

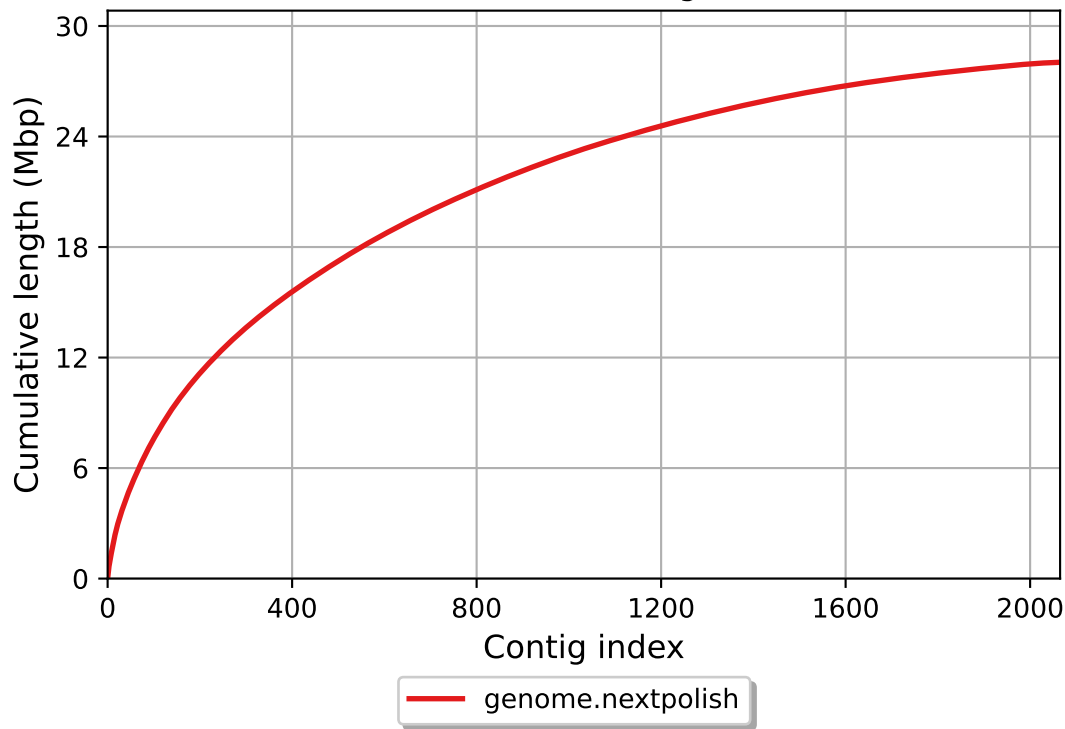
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

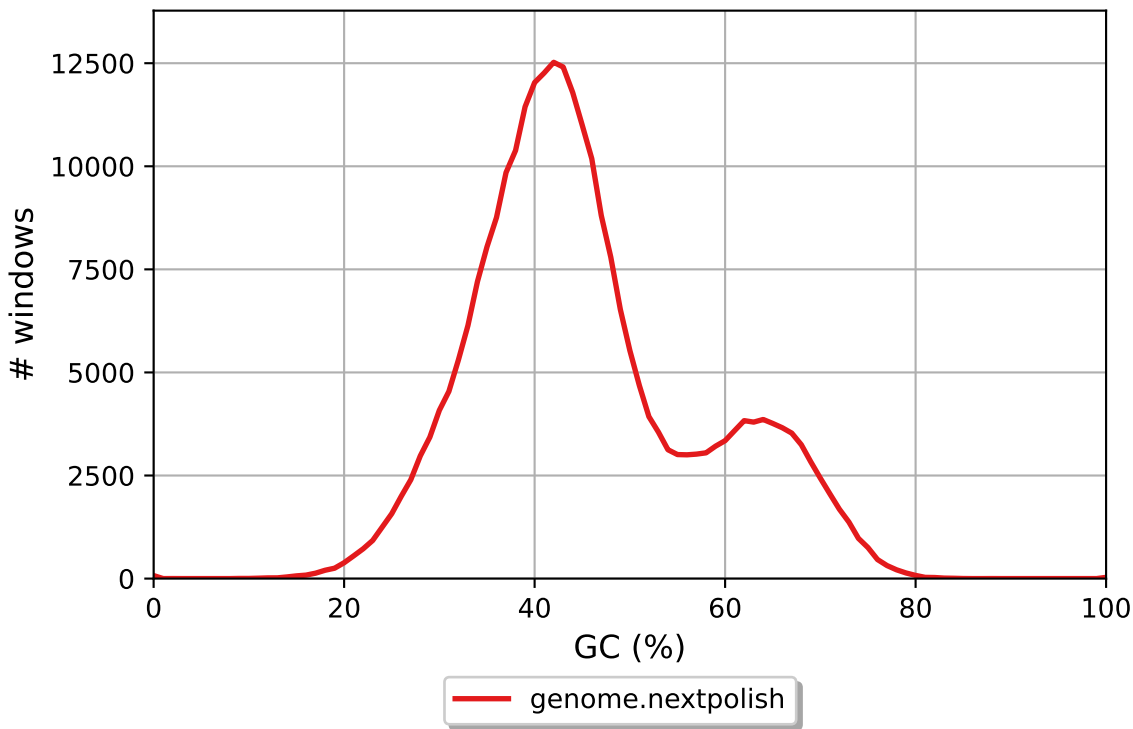


— genome.nextpolish

Cumulative length



## GC content



genome.nextpolish GC content

