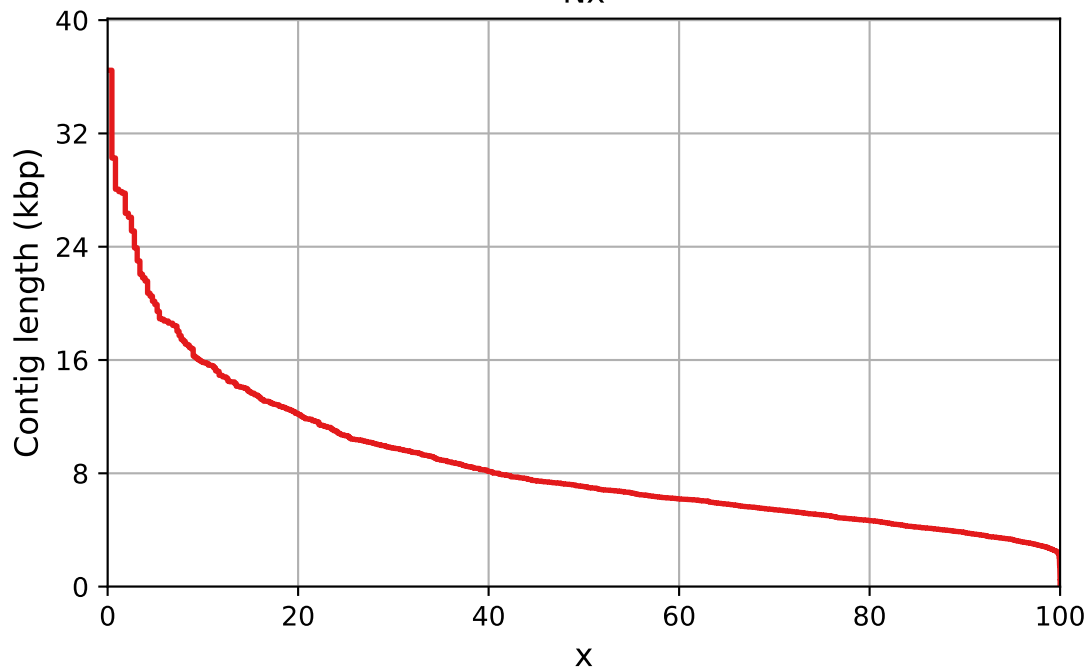


# Report

	genome.nextpolish
# contigs (>= 0 bp)	1246
# contigs (>= 1000 bp)	1243
# contigs (>= 5000 bp)	726
# contigs (>= 10000 bp)	163
# contigs (>= 25000 bp)	8
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	8111426
Total length (>= 1000 bp)	8109742
Total length (>= 5000 bp)	6155167
Total length (>= 10000 bp)	2329739
Total length (>= 25000 bp)	228010
Total length (>= 50000 bp)	0
# contigs	1246
Largest contig	36460
Total length	8111426
GC (%)	43.07
N50	7067
N90	3815
auN	8784.9
L50	372
L90	989
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	14 + 2 part

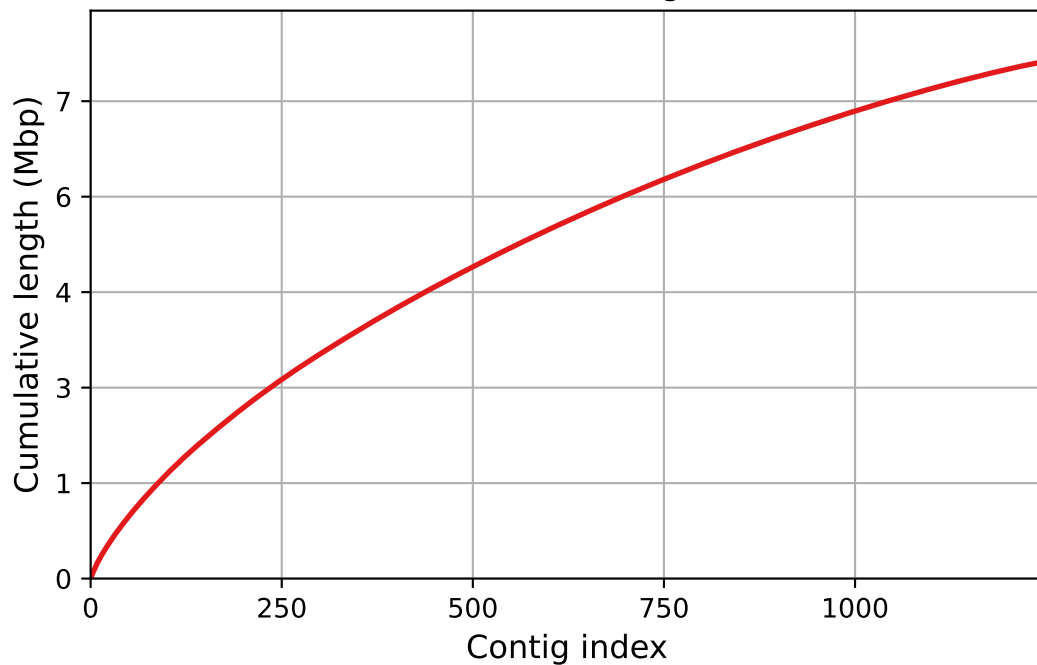
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx



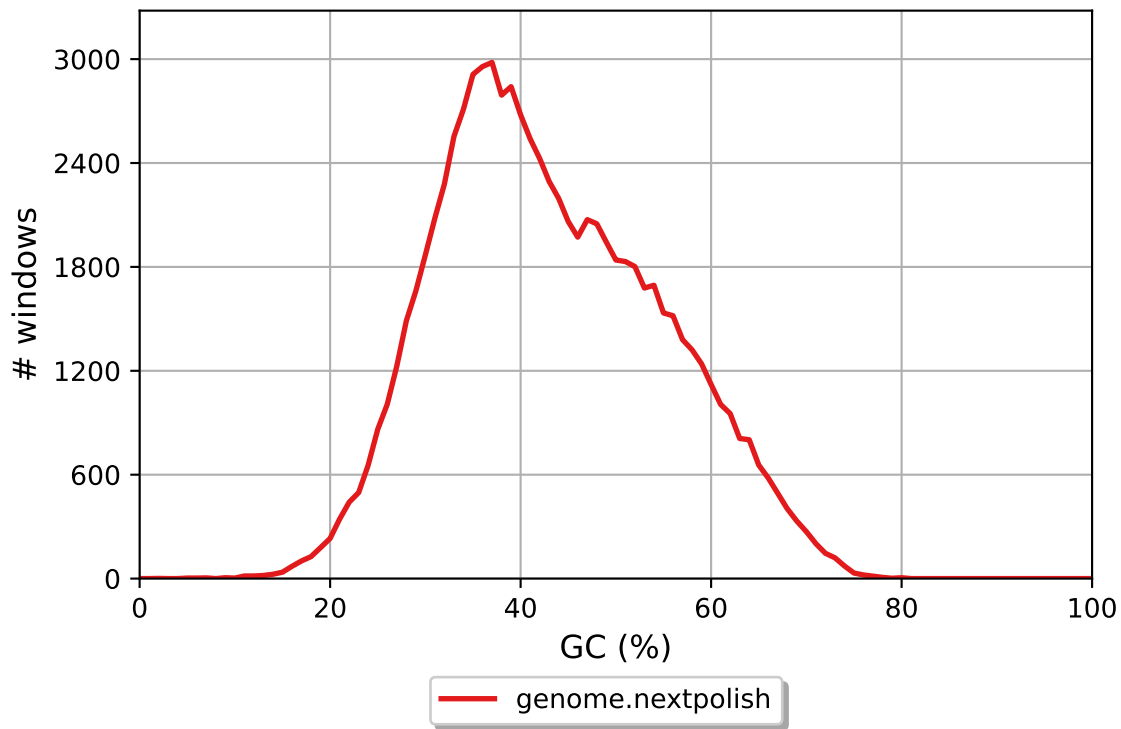
— genome.nextpolish

Cumulative length



— genome.nextpolish

## GC content



genome.nextpolish GC content

