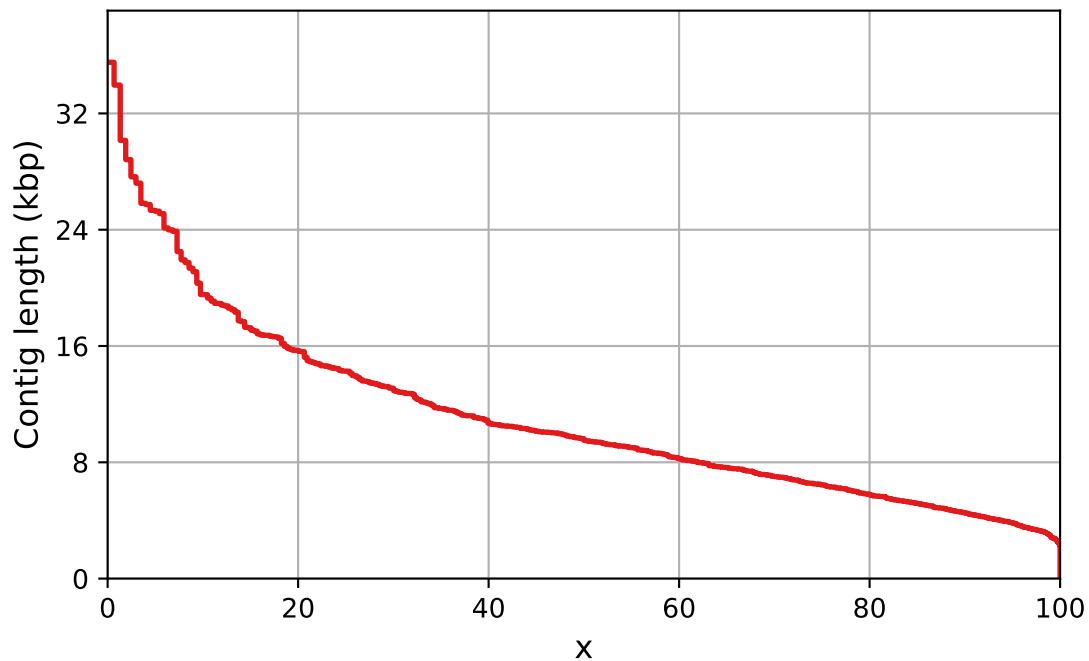


# Report

	genome.nextpolish
# contigs (>= 0 bp)	640
# contigs (>= 1000 bp)	640
# contigs (>= 5000 bp)	457
# contigs (>= 10000 bp)	169
# contigs (>= 25000 bp)	11
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	5247891
Total length (>= 1000 bp)	5247891
Total length (>= 5000 bp)	4539346
Total length (>= 10000 bp)	2476836
Total length (>= 25000 bp)	310546
Total length (>= 50000 bp)	0
# contigs	640
Largest contig	35519
Total length	5247891
GC (%)	39.02
N50	9516
N90	4531
auN	11161.6
L50	185
L90	496
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	0 + 0 part

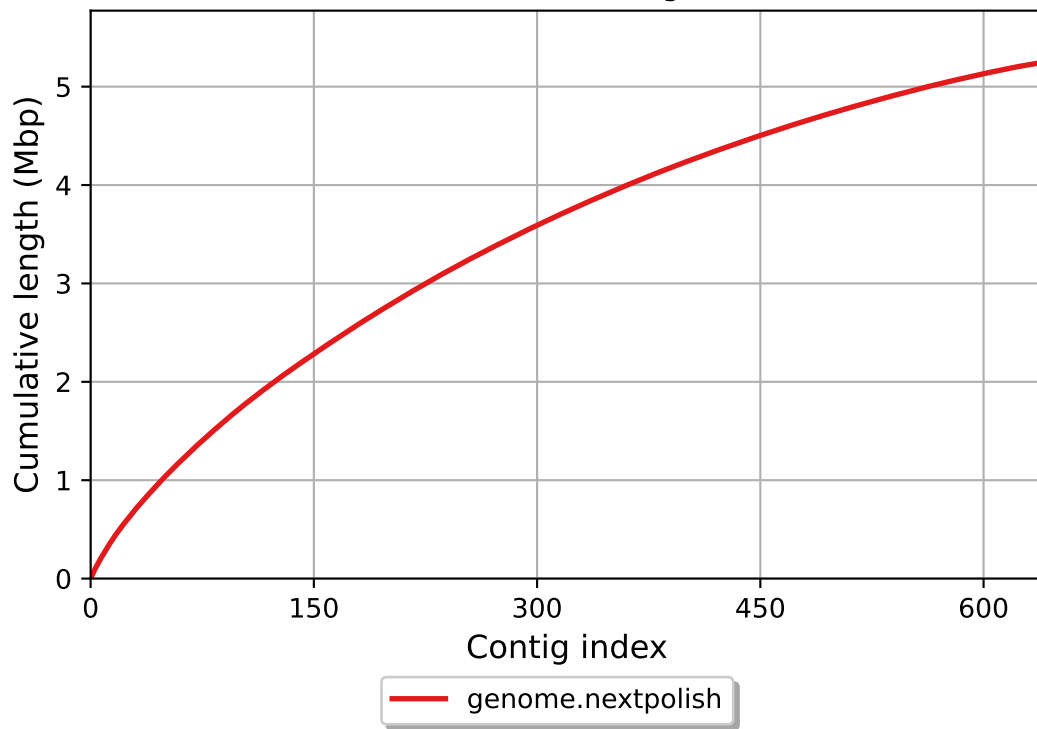
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

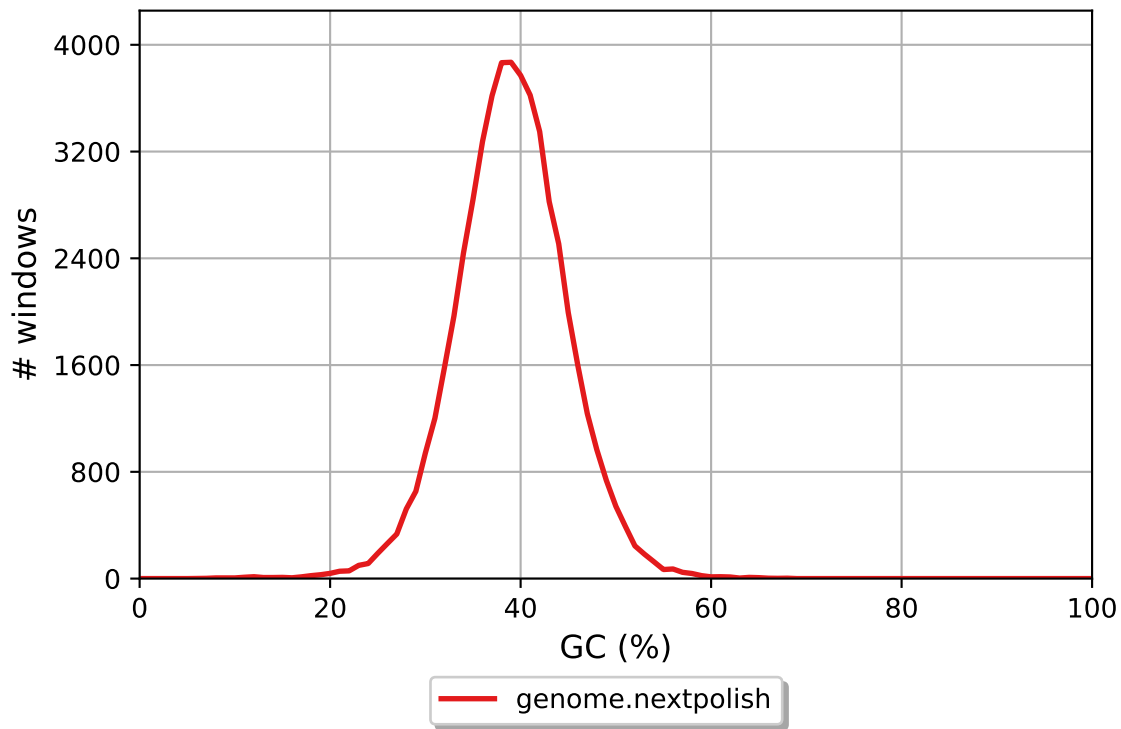


— genome.nextpolish

Cumulative length



## GC content



genome.nextpolish GC content

