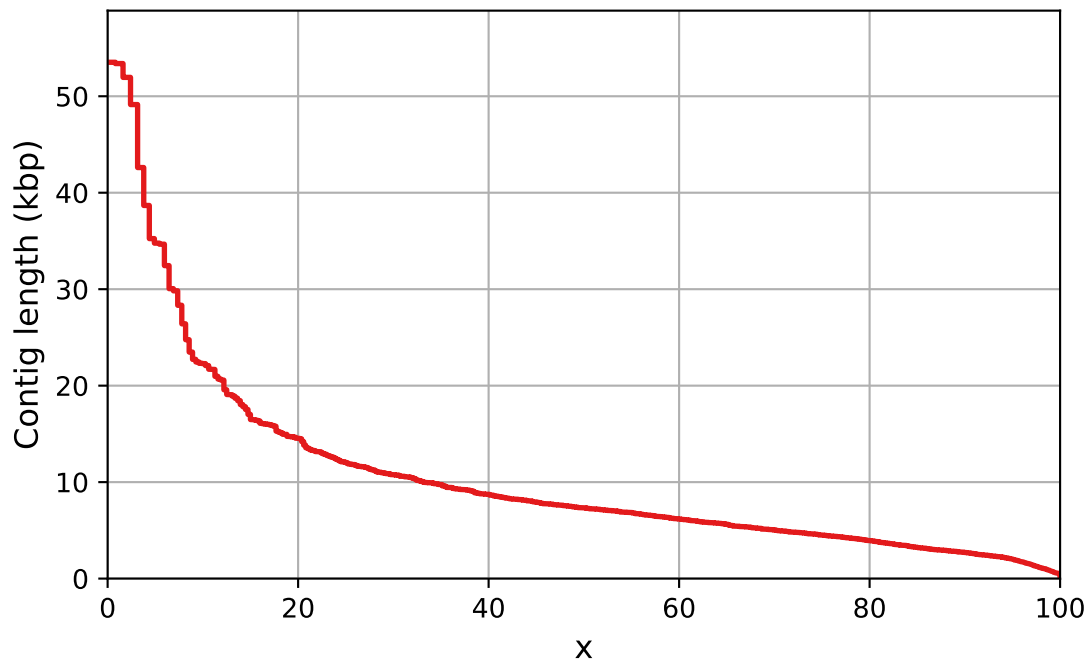


# Report

	genome.nextpolish
# contigs (>= 0 bp)	1283
# contigs (>= 1000 bp)	1131
# contigs (>= 5000 bp)	483
# contigs (>= 10000 bp)	132
# contigs (>= 25000 bp)	14
# contigs (>= 50000 bp)	3
Total length (>= 0 bp)	6612711
Total length (>= 1000 bp)	6510093
Total length (>= 5000 bp)	4643024
Total length (>= 10000 bp)	2193835
Total length (>= 25000 bp)	541045
Total length (>= 50000 bp)	158867
# contigs	1275
Largest contig	53522
Total length	6609203
GC (%)	60.08
N50	7335
N90	2713
auN	10721.0
L50	264
L90	832
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	6 + 1 part

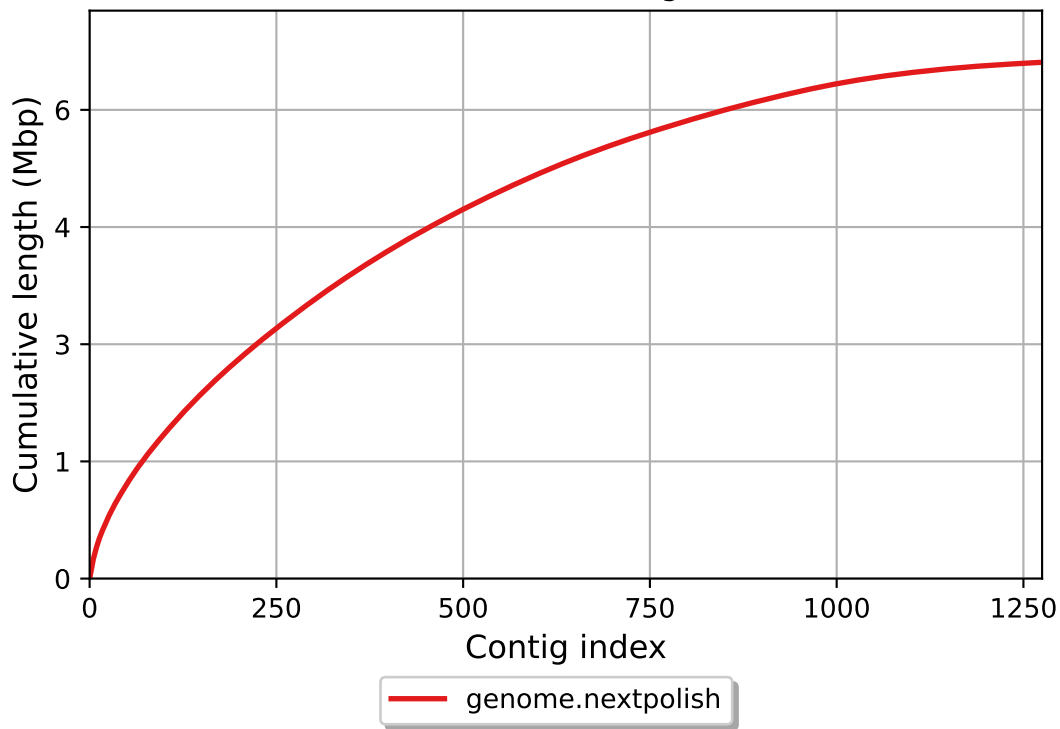
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

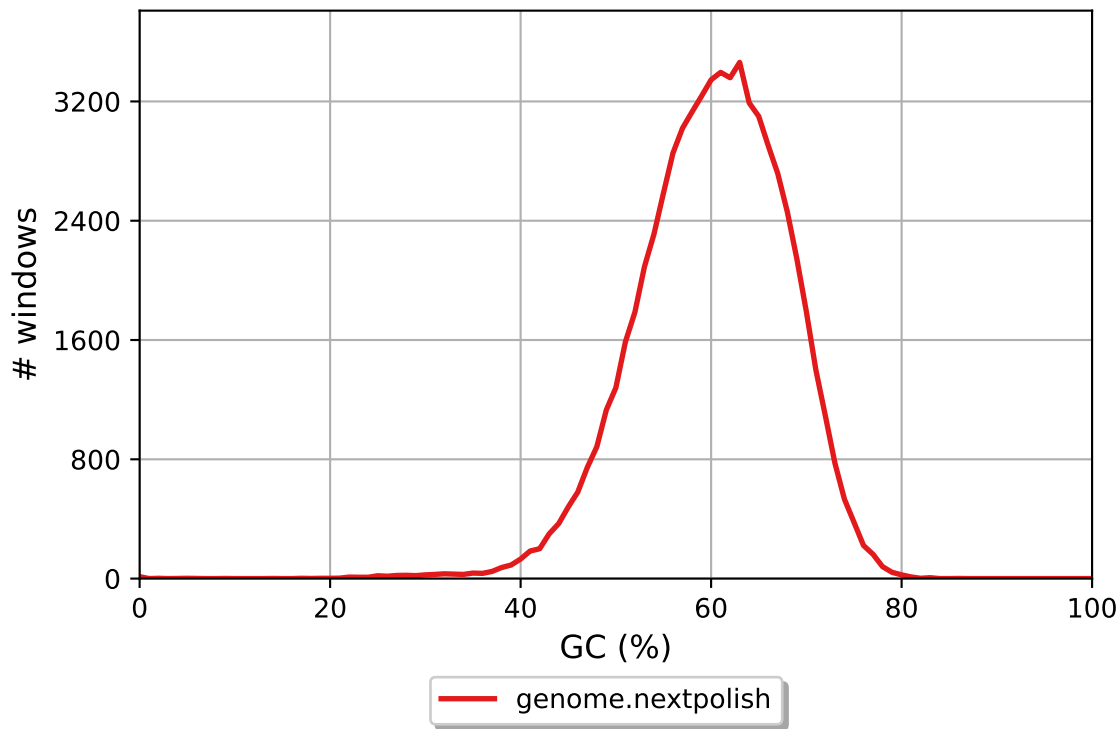


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

