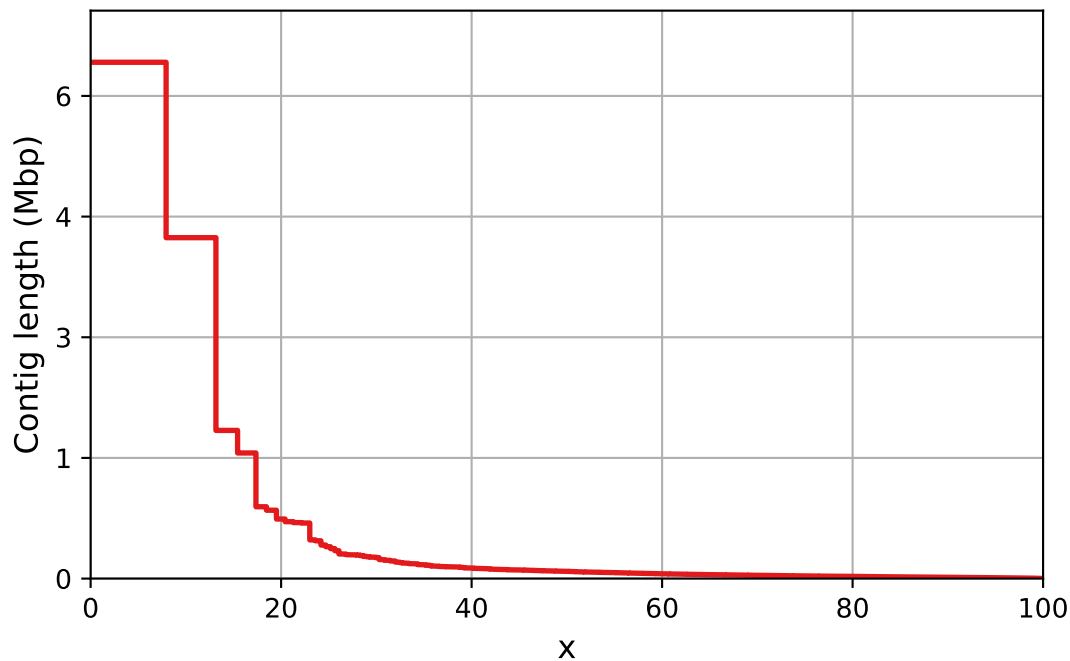


# Report

	genome.nextpolish
# contigs (>= 0 bp)	2039
# contigs (>= 1000 bp)	2025
# contigs (>= 5000 bp)	1783
# contigs (>= 10000 bp)	1392
# contigs (>= 25000 bp)	731
# contigs (>= 50000 bp)	331
Total length (>= 0 bp)	81000029
Total length (>= 1000 bp)	80988863
Total length (>= 5000 bp)	80150660
Total length (>= 10000 bp)	77191485
Total length (>= 25000 bp)	66307236
Total length (>= 50000 bp)	52339199
# contigs	2039
Largest contig	6418430
Total length	81000029
GC (%)	53.36
N50	89778
N90	16221
auN	917303.5
L50	151
L90	1060
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	137 + 8 part

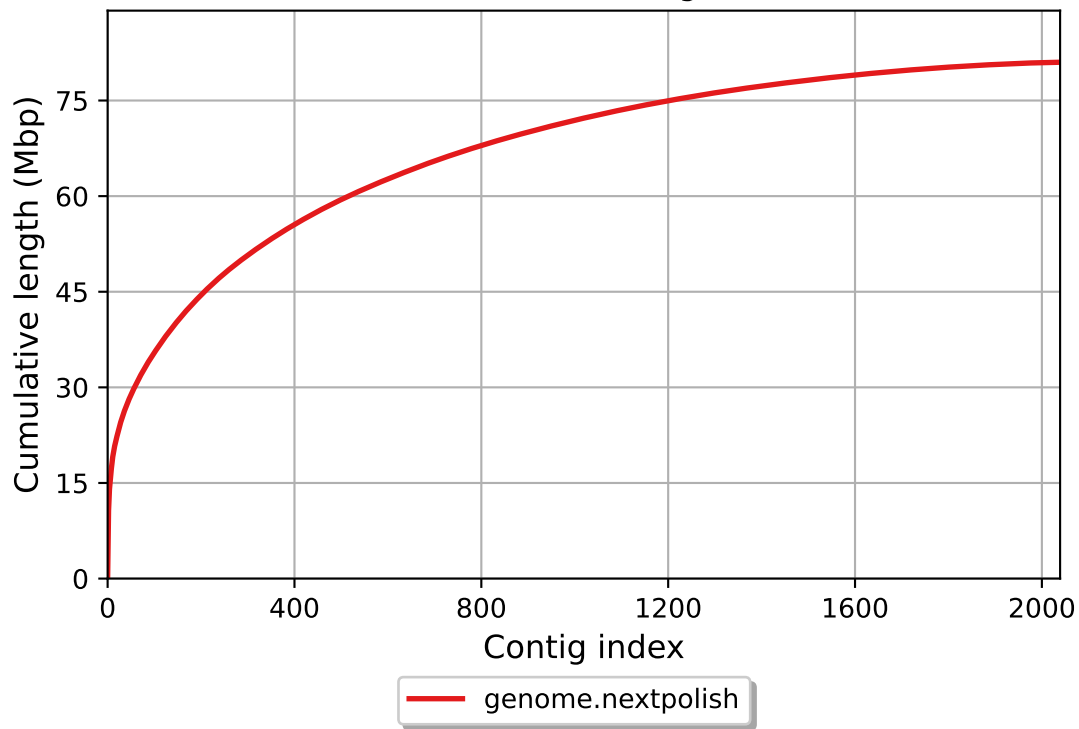
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

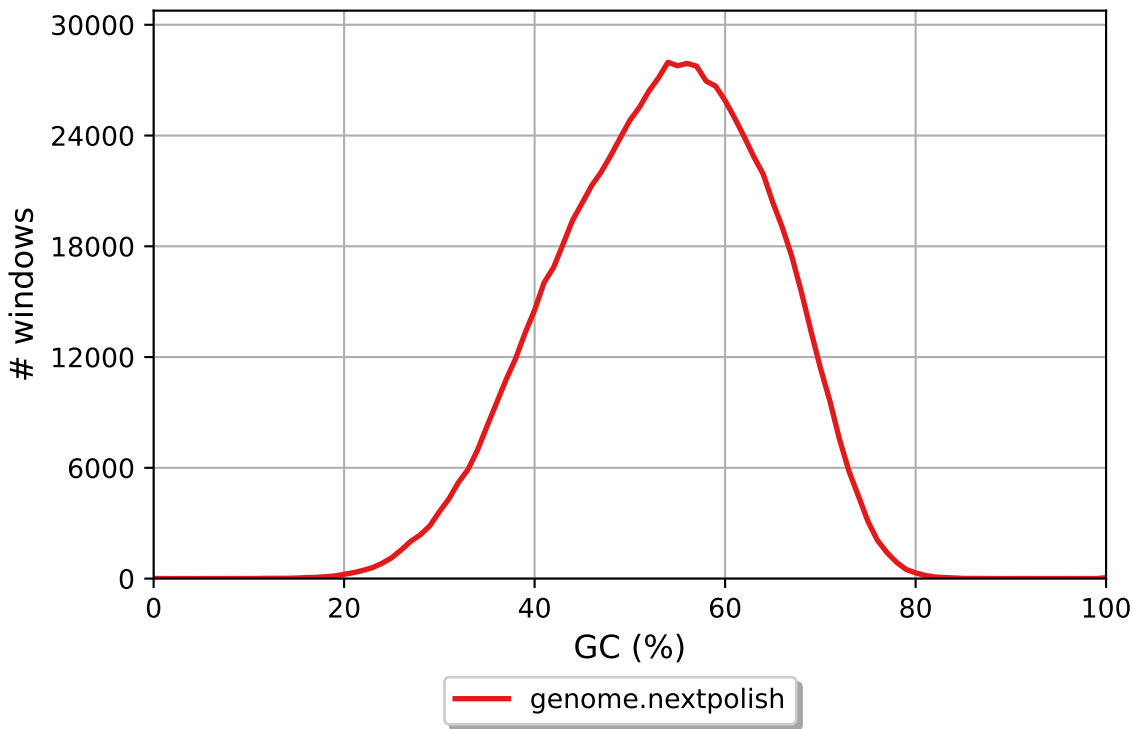


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

