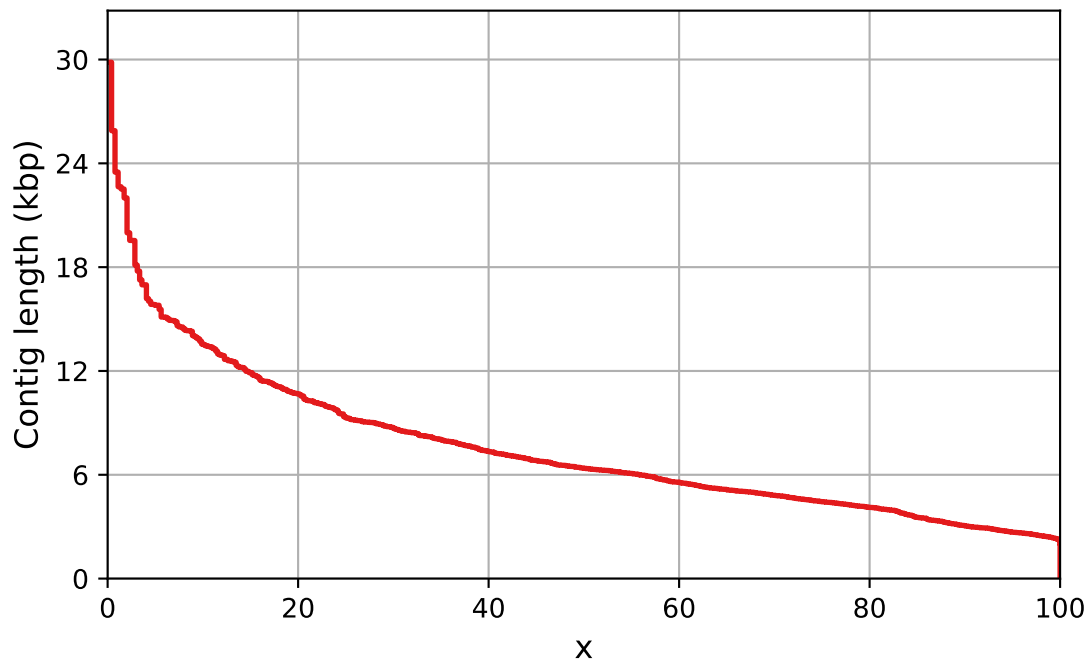


# Report

	genome.nextpolish
# contigs (>= 0 bp)	1264
# contigs (>= 1000 bp)	1264
# contigs (>= 5000 bp)	596
# contigs (>= 10000 bp)	123
# contigs (>= 25000 bp)	2
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	7181450
Total length (>= 1000 bp)	7181450
Total length (>= 5000 bp)	4848704
Total length (>= 10000 bp)	1641876
Total length (>= 25000 bp)	55729
Total length (>= 50000 bp)	0
# contigs	1264
Largest contig	29841
Total length	7181450
GC (%)	54.43
N50	6377
N90	3053
auN	7587.1
L50	374
L90	996
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	13 + 1 part

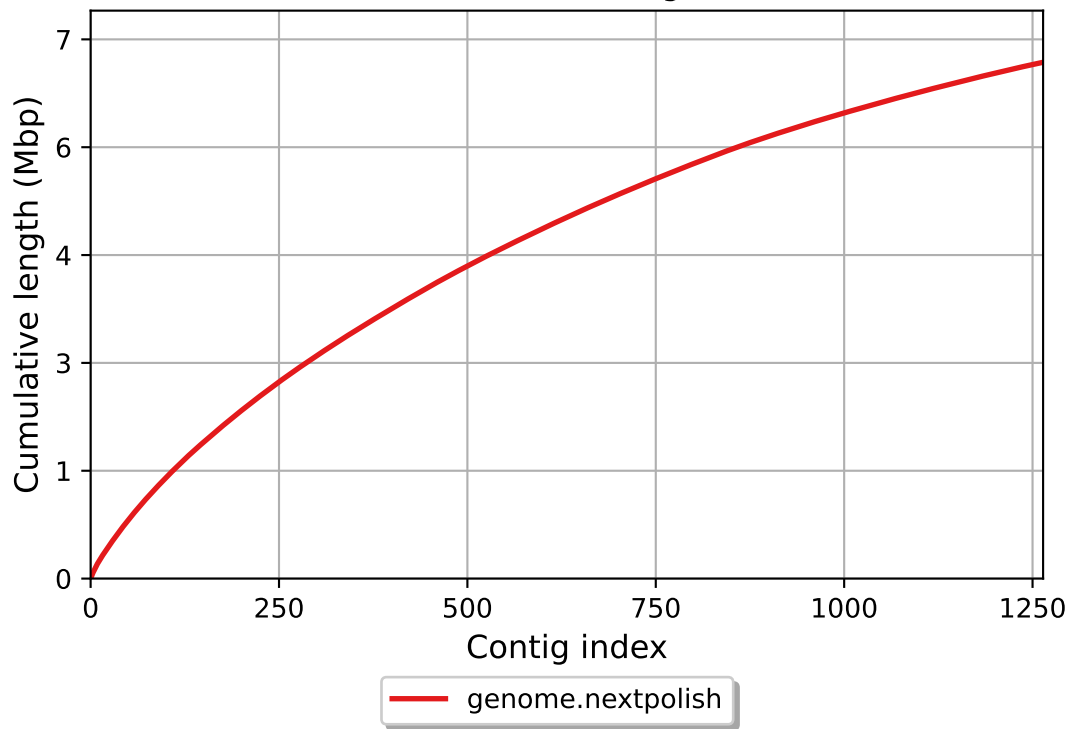
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

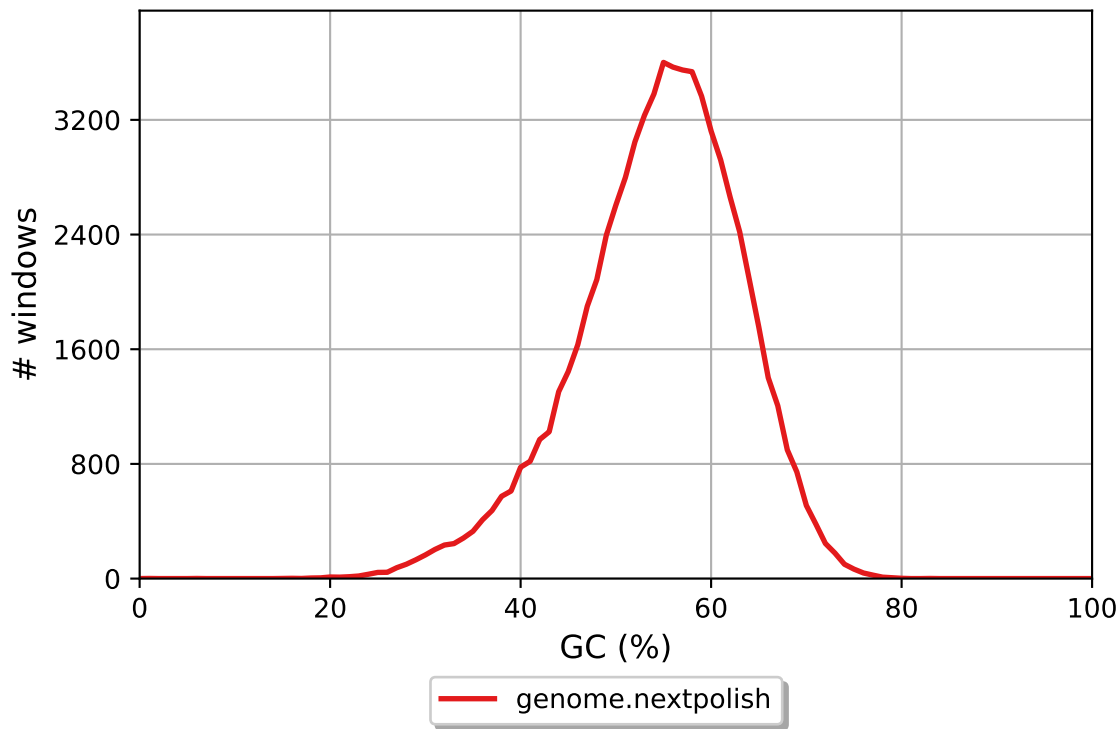


— genome.nextpolish

Cumulative length



## GC content



genome.nextpolish GC content

