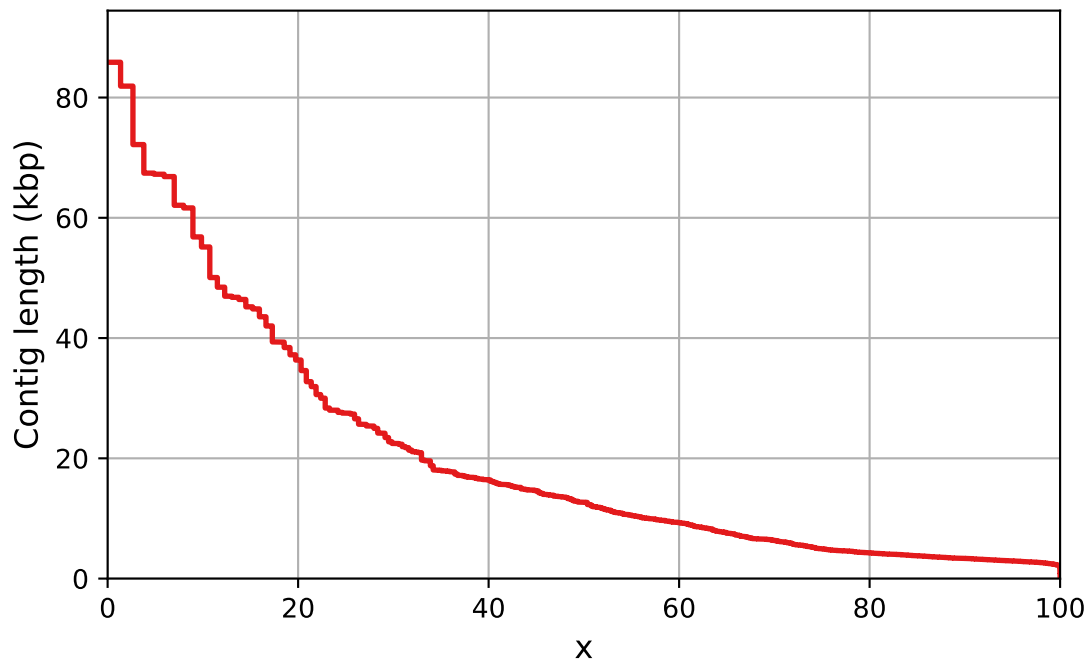


# Report

	genome.nextpolish
# contigs (>= 0 bp)	781
# contigs (>= 1000 bp)	780
# contigs (>= 5000 bp)	318
# contigs (>= 10000 bp)	163
# contigs (>= 25000 bp)	41
# contigs (>= 50000 bp)	11
Total length (>= 0 bp)	6312080
Total length (>= 1000 bp)	6311110
Total length (>= 5000 bp)	4705972
Total length (>= 10000 bp)	3586238
Total length (>= 25000 bp)	1765006
Total length (>= 50000 bp)	727224
# contigs	781
Largest contig	85865
Total length	6312080
GC (%)	43.61
N50	12683
N90	3356
auN	20607.6
L50	125
L90	560
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	10 + 1 part

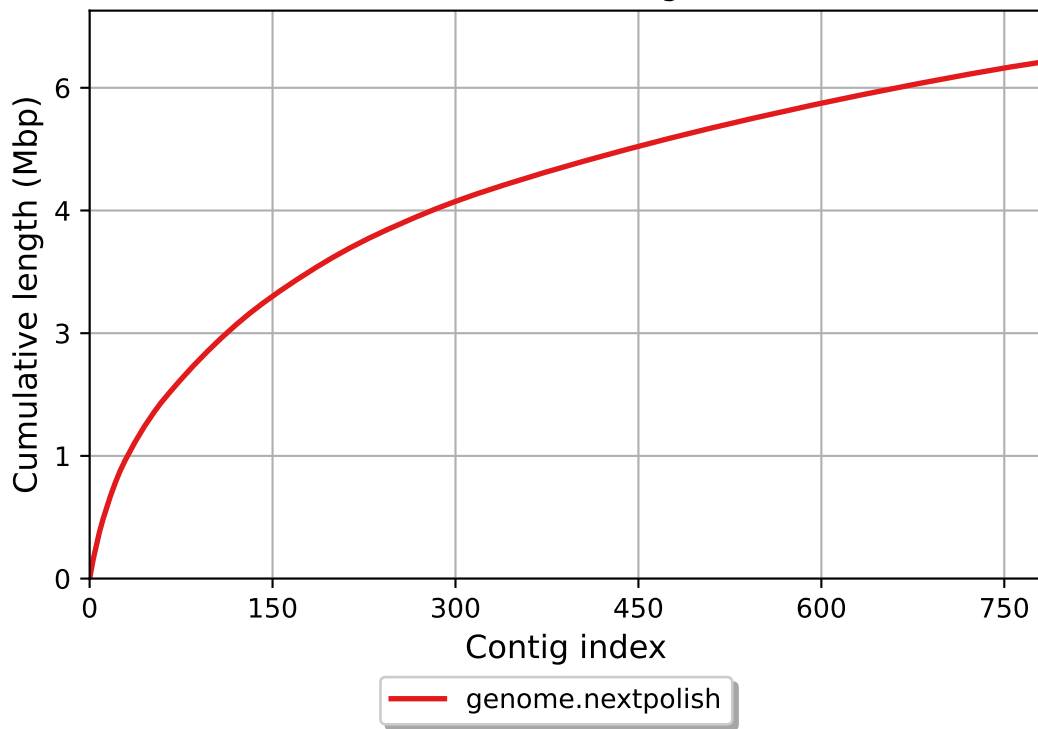
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

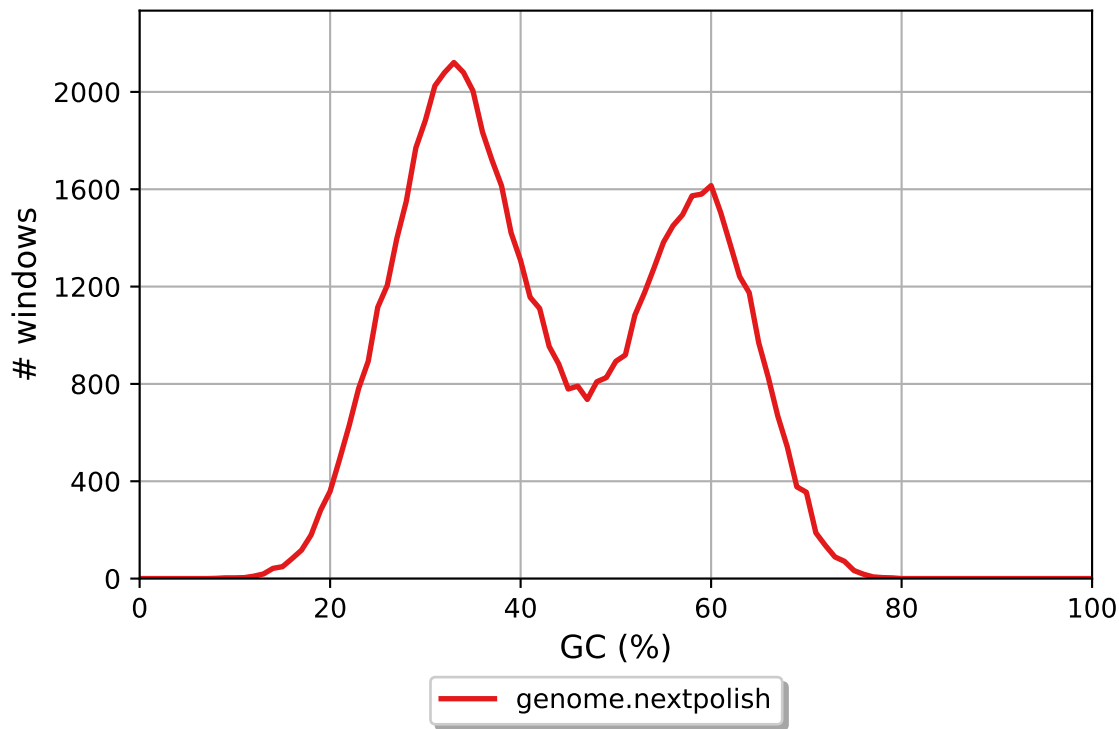


— genome.nextpolish

Cumulative length



## GC content



genome.nextpolish GC content

