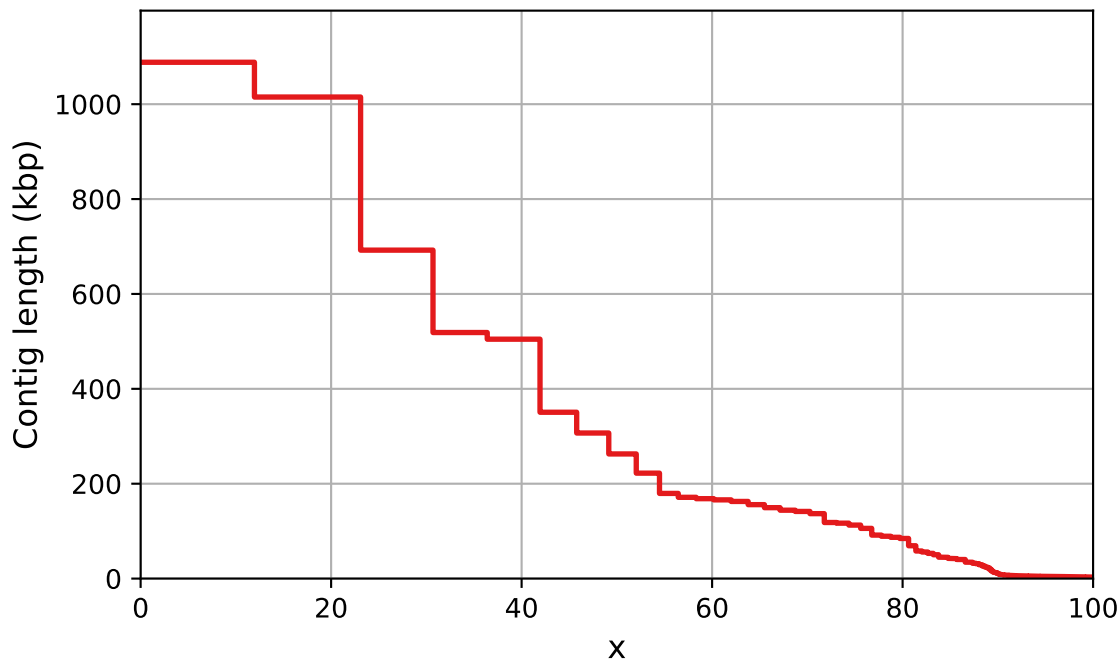


# Report

	genome.nextpolish
# contigs (>= 0 bp)	243
# contigs (>= 1000 bp)	243
# contigs (>= 5000 bp)	113
# contigs (>= 10000 bp)	52
# contigs (>= 25000 bp)	44
# contigs (>= 50000 bp)	32
Total length (>= 0 bp)	9107706
Total length (>= 1000 bp)	9107706
Total length (>= 5000 bp)	8580690
Total length (>= 10000 bp)	8202778
Total length (>= 25000 bp)	8073760
Total length (>= 50000 bp)	7630232
# contigs	243
Largest contig	1088350
Total length	9107706
GC (%)	55.50
N50	262742
N90	10803
auN	431113.6
L50	8
L90	52
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	31 + 4 part

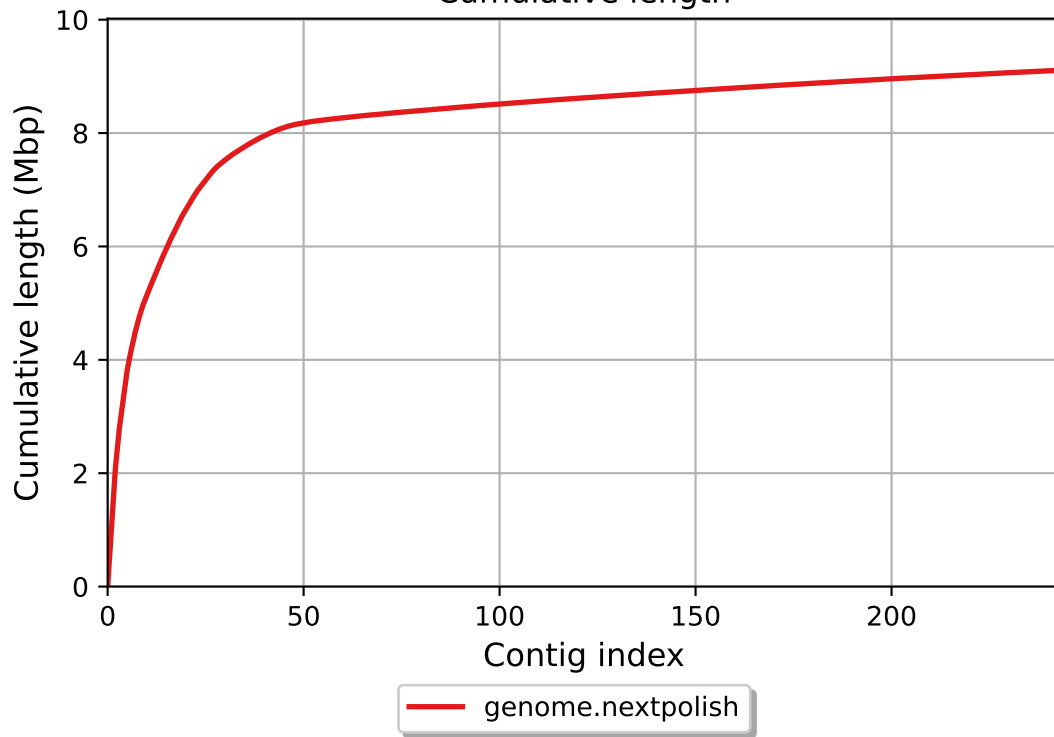
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

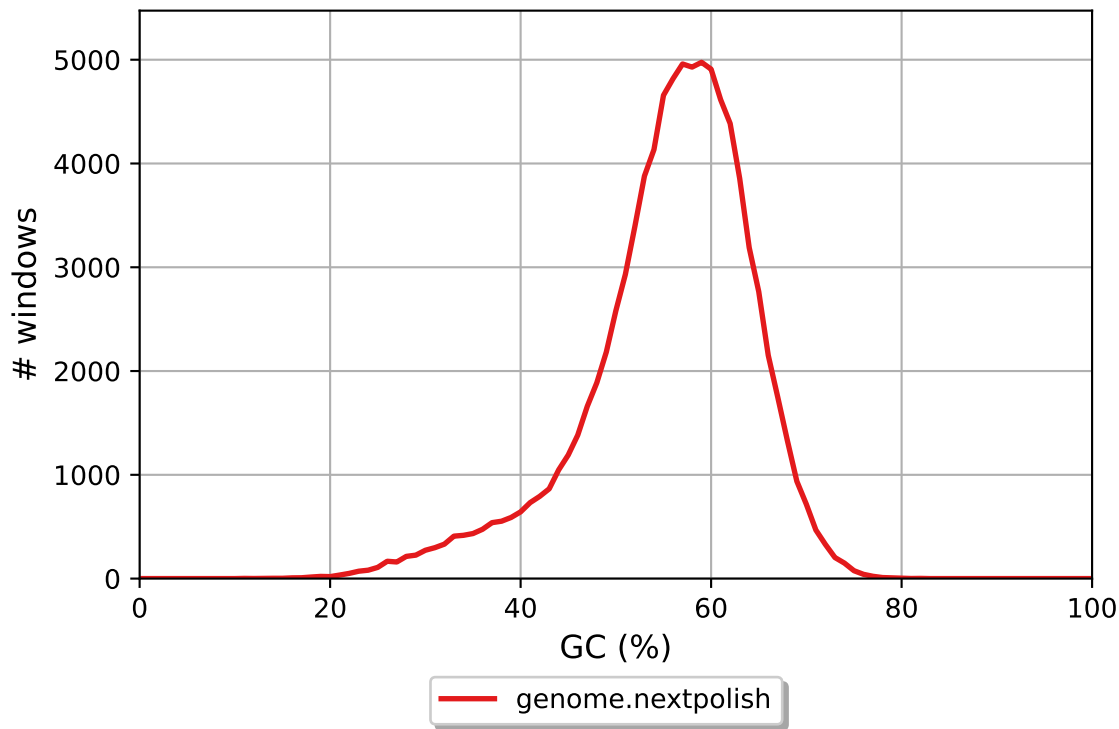


— genome.nextpolish

Cumulative length



## GC content



genome.nextpolish GC content

