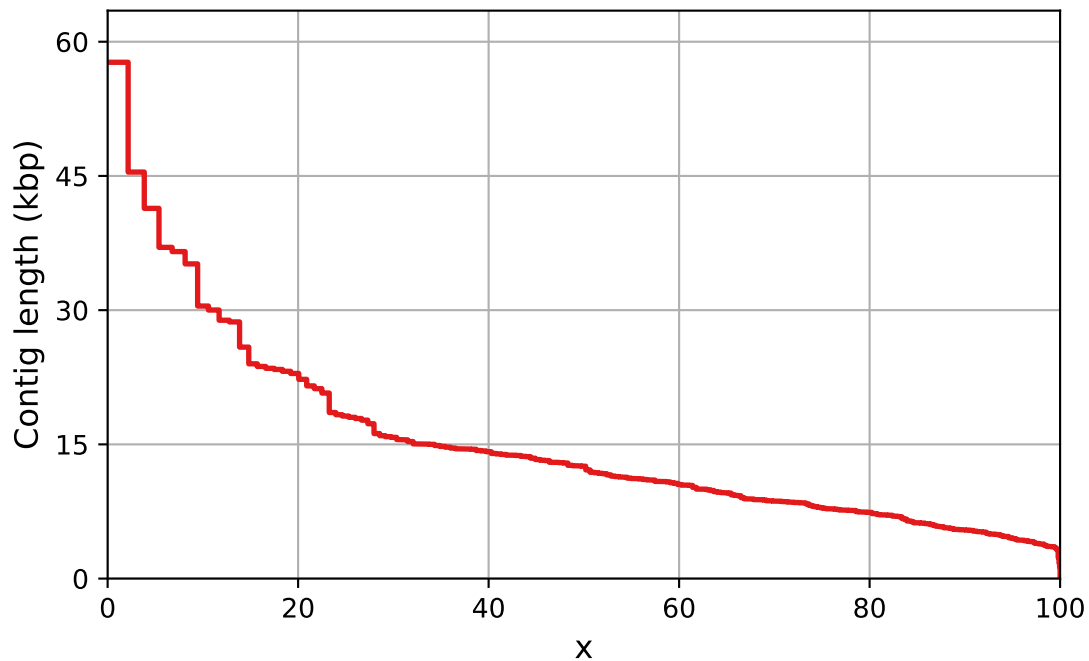


# Report

	genome.nextpolish
# contigs (>= 0 bp)	260
# contigs (>= 1000 bp)	260
# contigs (>= 5000 bp)	211
# contigs (>= 10000 bp)	99
# contigs (>= 25000 bp)	11
# contigs (>= 50000 bp)	1
Total length (>= 0 bp)	2680090
Total length (>= 1000 bp)	2680090
Total length (>= 5000 bp)	2482186
Total length (>= 10000 bp)	1666365
Total length (>= 25000 bp)	397152
Total length (>= 50000 bp)	57704
# contigs	260
Largest contig	57704
Total length	2680090
GC (%)	48.81
N50	12557
N90	5441
auN	15545.7
L50	70
L90	198
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	10 + 0 part

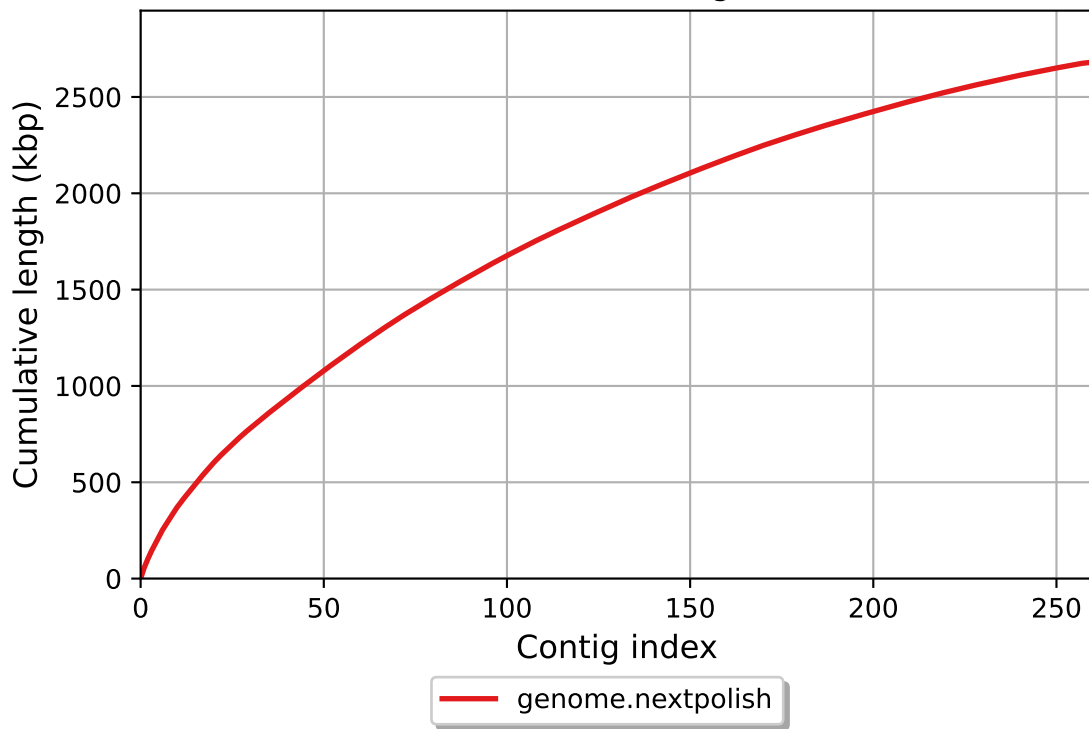
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

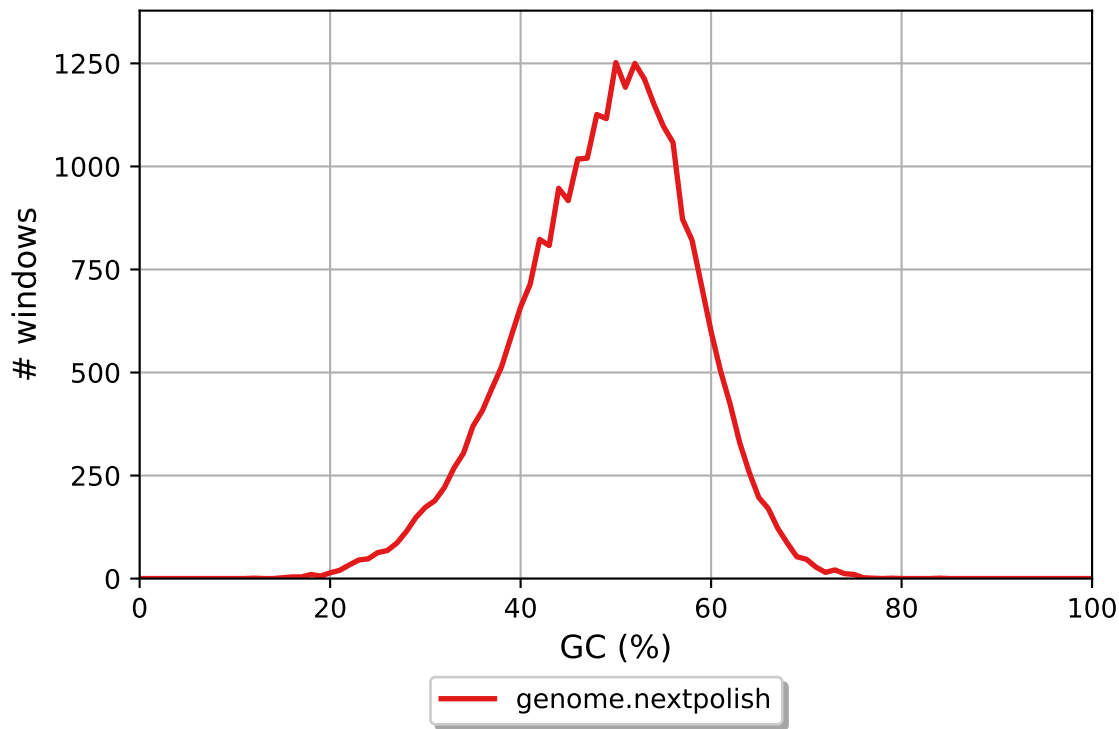


— genome.nextpolish

Cumulative length



## GC content



genome.nextpolish GC content

