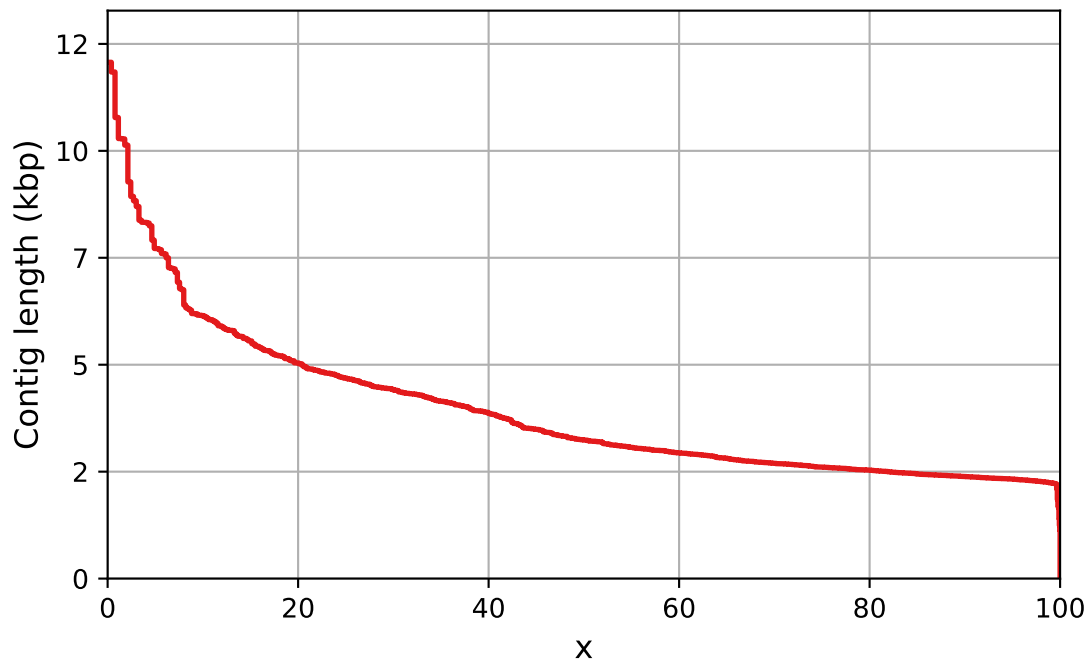


# Report

	genome.nextpolish
# contigs (>= 0 bp)	910
# contigs (>= 1000 bp)	910
# contigs (>= 5000 bp)	97
# contigs (>= 10000 bp)	6
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	3076163
Total length (>= 1000 bp)	3076163
Total length (>= 5000 bp)	627785
Total length (>= 10000 bp)	65392
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	910
Largest contig	12070
Total length	3076163
GC (%)	51.35
N50	3240
N90	2385
auN	3937.1
L50	323
L90	776
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	4 + 0 part

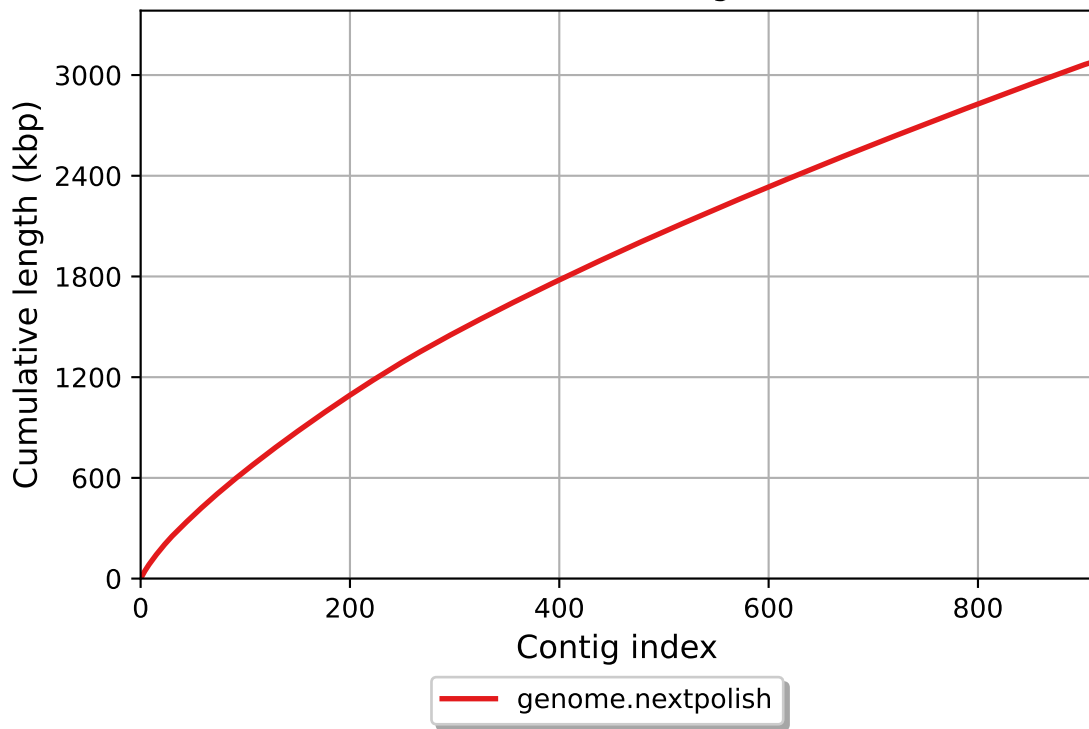
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

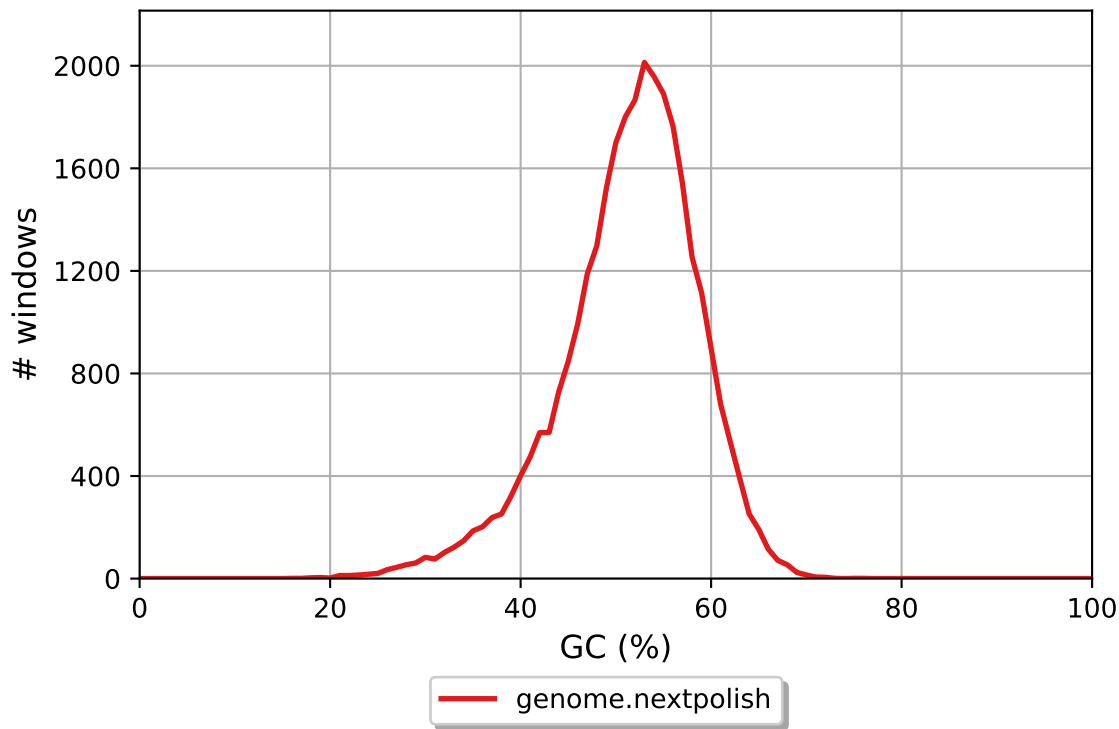


— genome.nextpolish

Cumulative length



## GC content



genome.nextpolish GC content

