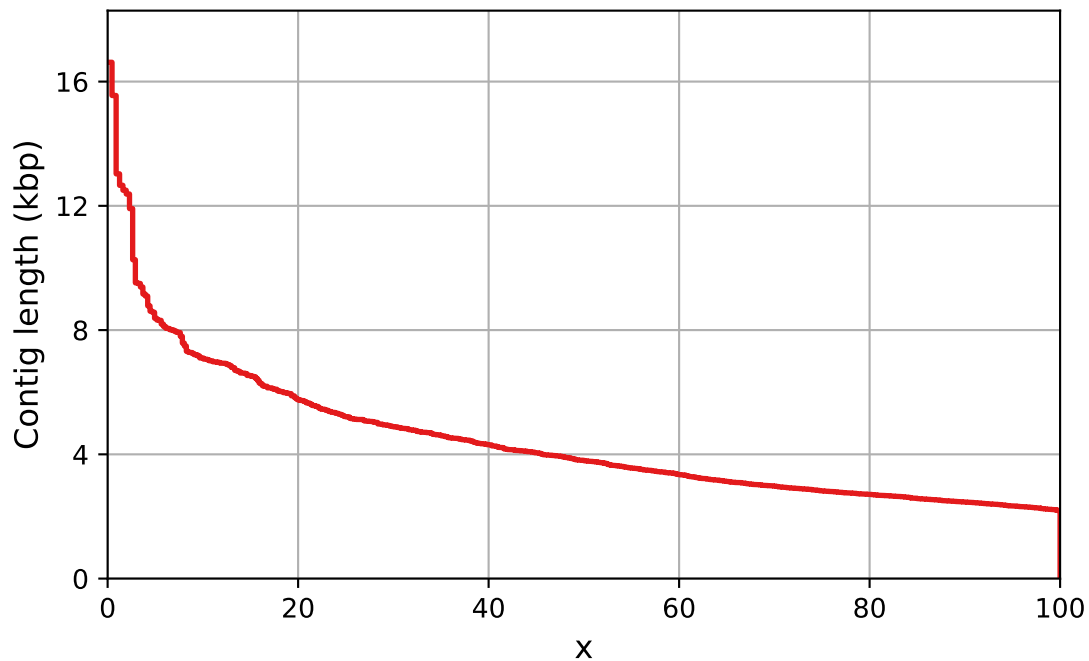


# Report

	genome.nextpolish
# contigs (>= 0 bp)	968
# contigs (>= 1000 bp)	968
# contigs (>= 5000 bp)	152
# contigs (>= 10000 bp)	8
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	3601791
Total length (>= 1000 bp)	3601791
Total length (>= 5000 bp)	1021521
Total length (>= 10000 bp)	104923
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	968
Largest contig	16621
Total length	3601791
GC (%)	41.04
N50	3803
N90	2465
auN	4453.3
L50	332
L90	814
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	9 + 0 part

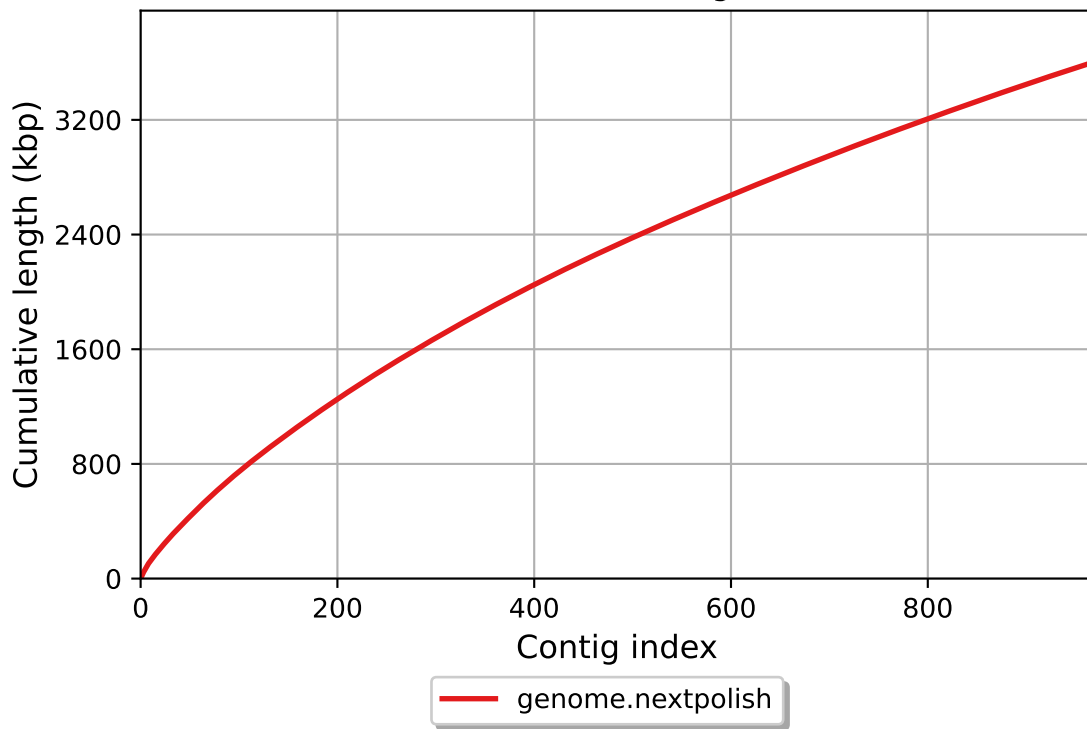
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

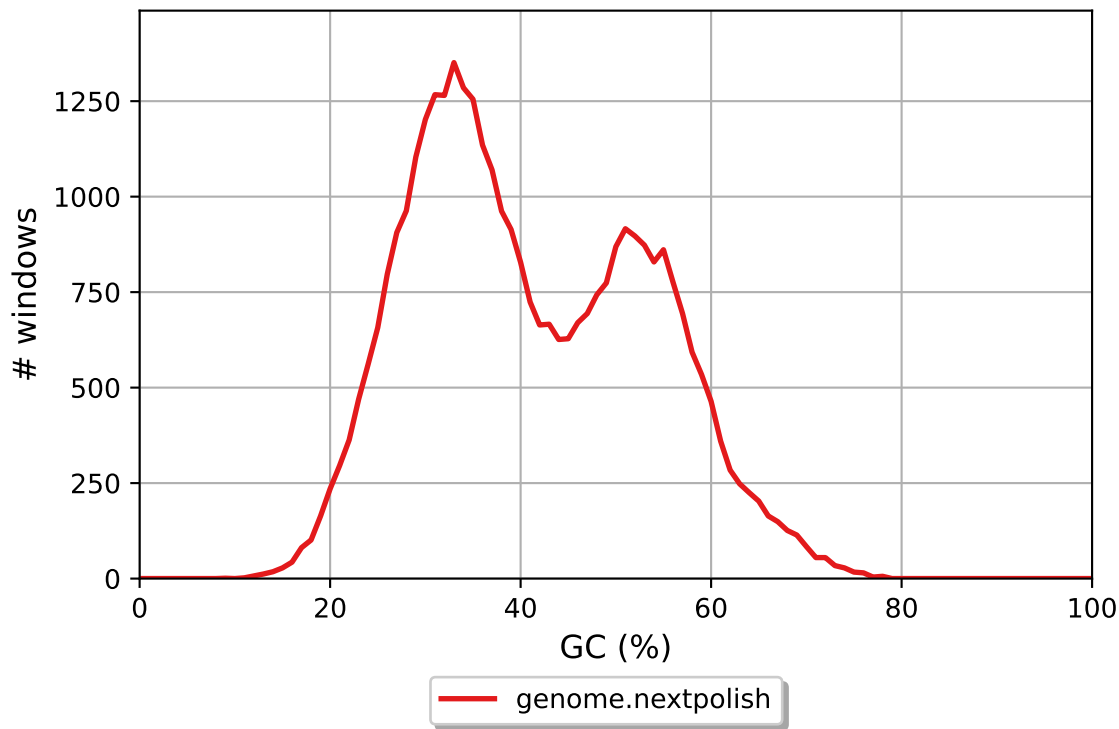


— genome.nextpolish

Cumulative length



## GC content



genome.nextpolish GC content

