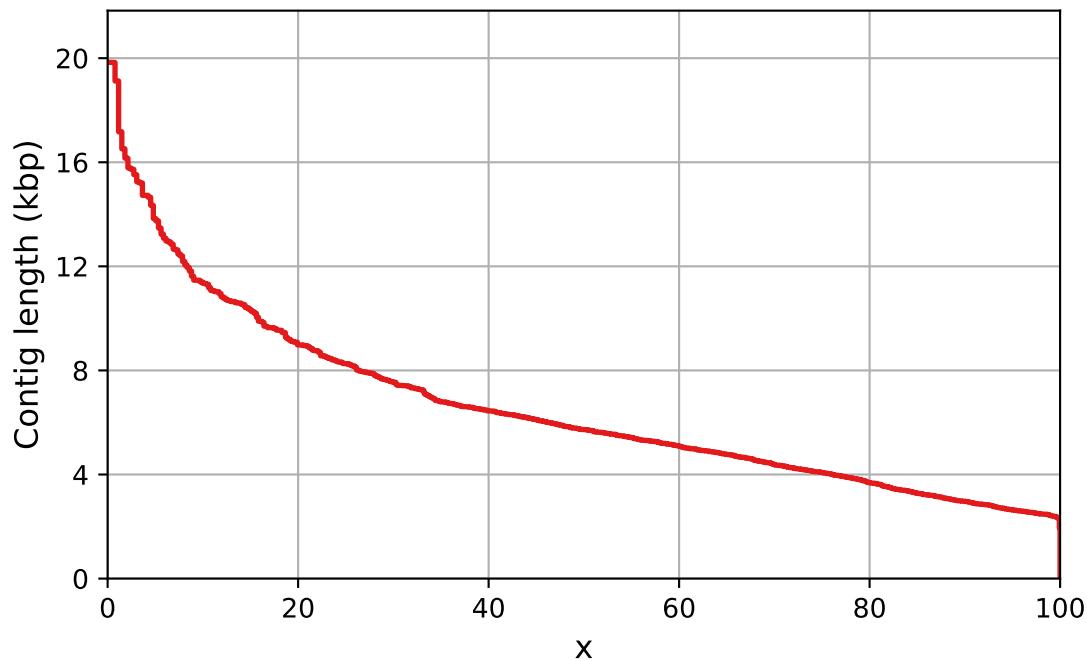


Report

	genome.nextpolish
# contigs (>= 0 bp)	981
# contigs (>= 1000 bp)	981
# contigs (>= 5000 bp)	412
# contigs (>= 10000 bp)	64
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	5095958
Total length (>= 1000 bp)	5095958
Total length (>= 5000 bp)	3117722
Total length (>= 10000 bp)	806448
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	981
Largest contig	19842
Total length	5095958
GC (%)	43.23
N50	5732
N90	2968
auN	6609.5
L50	306
L90	788
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	17 + 5 part

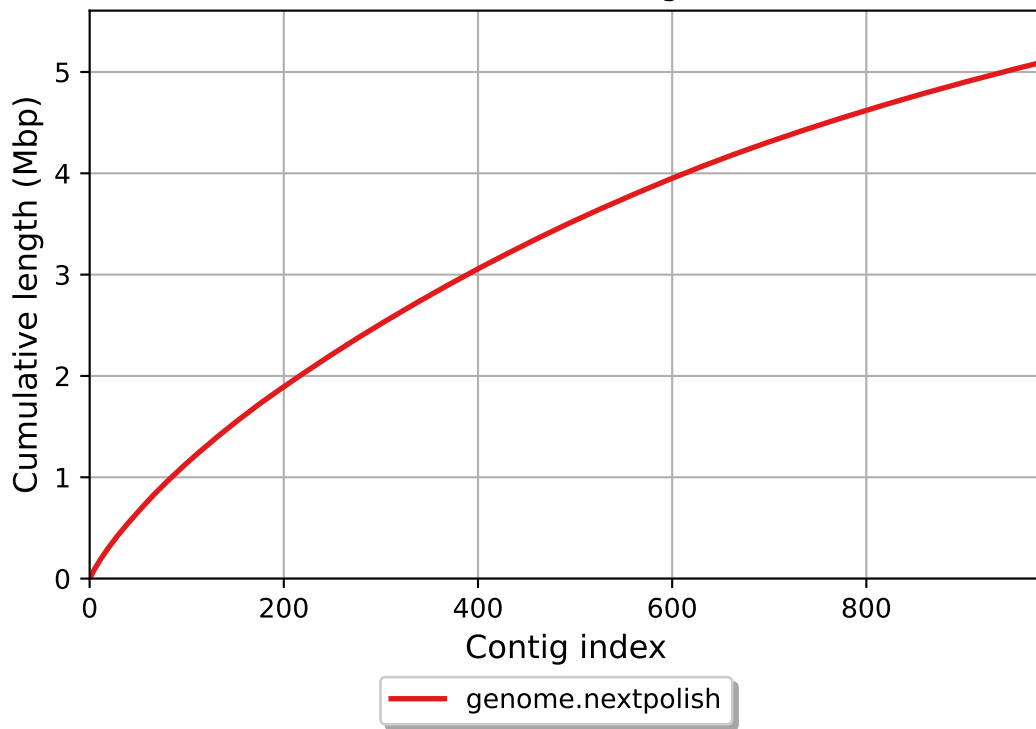
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

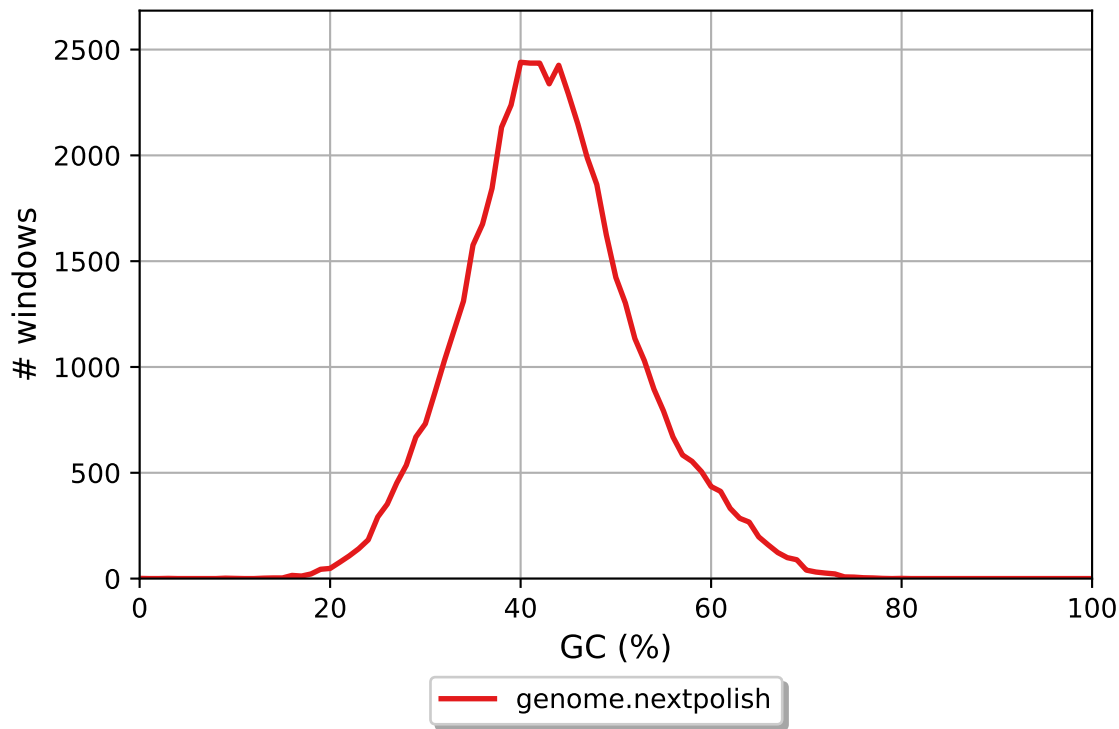


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

