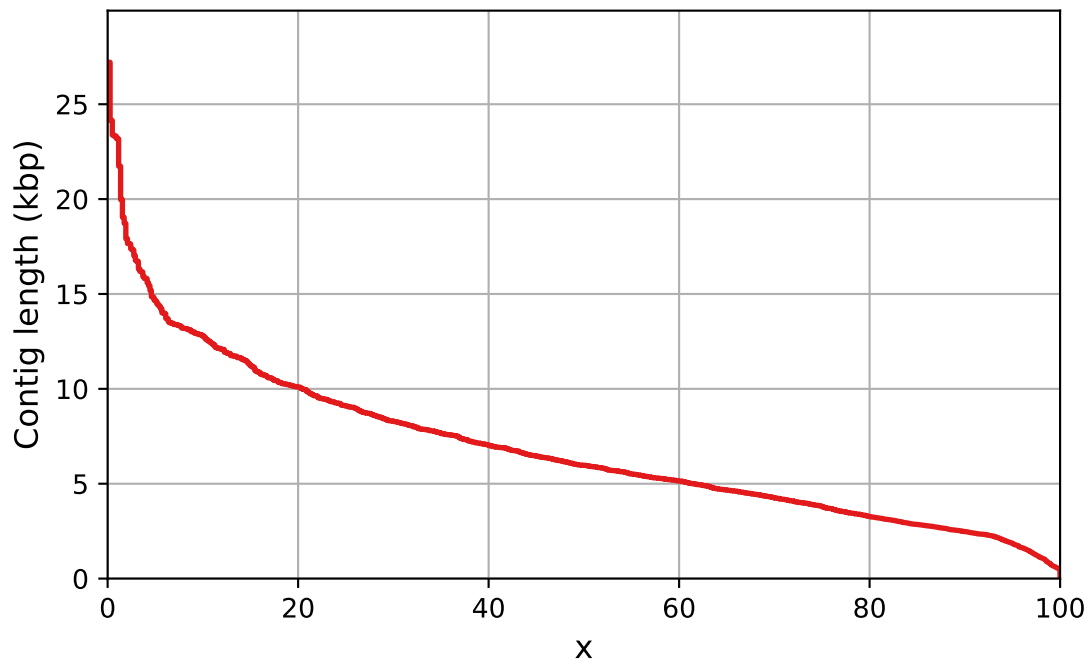


Report

	genome.nextpolish
# contigs (>= 0 bp)	2387
# contigs (>= 1000 bp)	2131
# contigs (>= 5000 bp)	797
# contigs (>= 10000 bp)	167
# contigs (>= 25000 bp)	1
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	10494523
Total length (>= 1000 bp)	10324521
Total length (>= 5000 bp)	6449495
Total length (>= 10000 bp)	2150948
Total length (>= 25000 bp)	27210
Total length (>= 50000 bp)	0
# contigs	2376
Largest contig	27210
Total length	10491360
GC (%)	50.37
N50	5973
N90	2482
auN	6917.8
L50	577
L90	1639
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	12 + 7 part

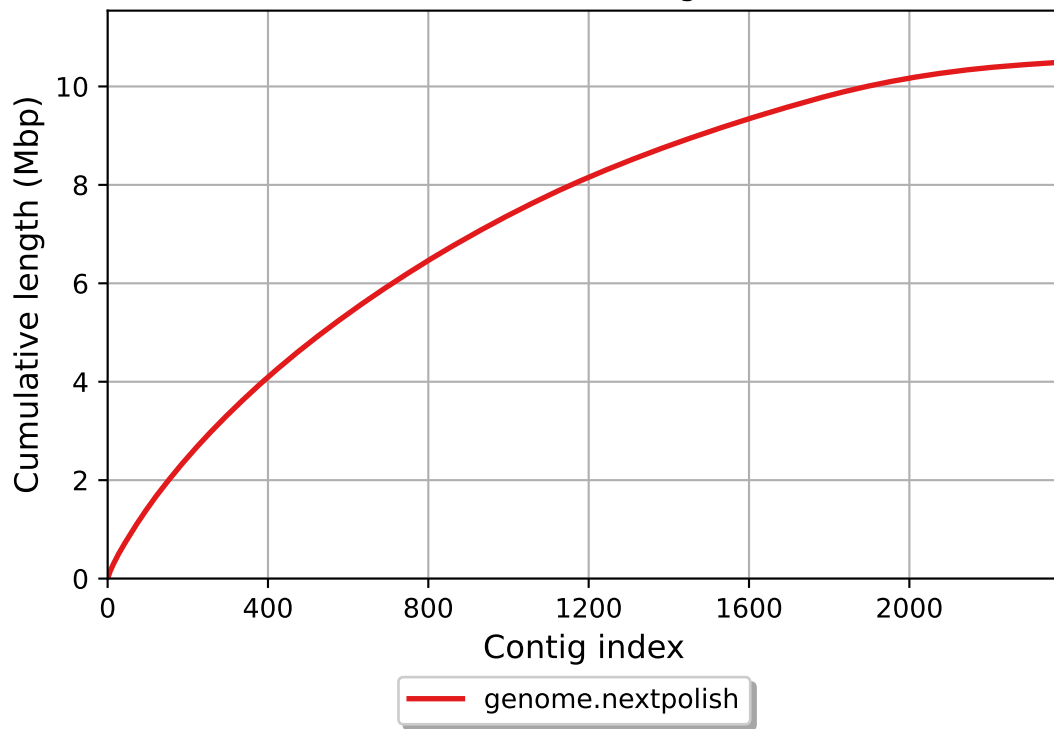
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

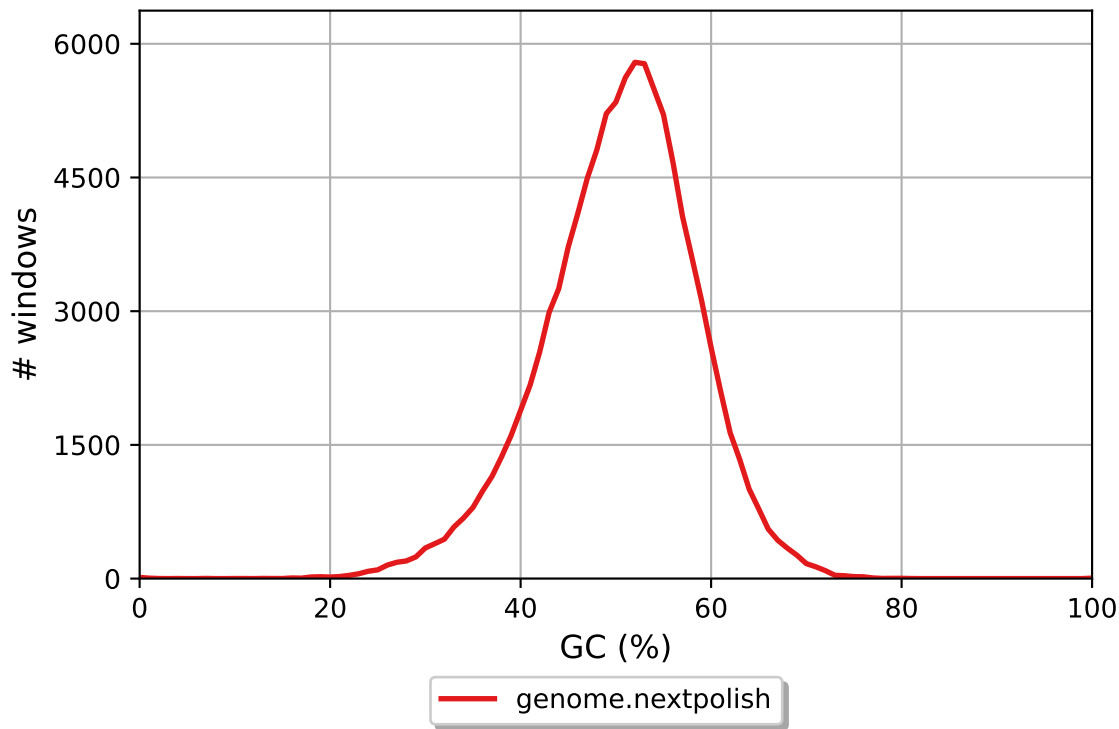


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

