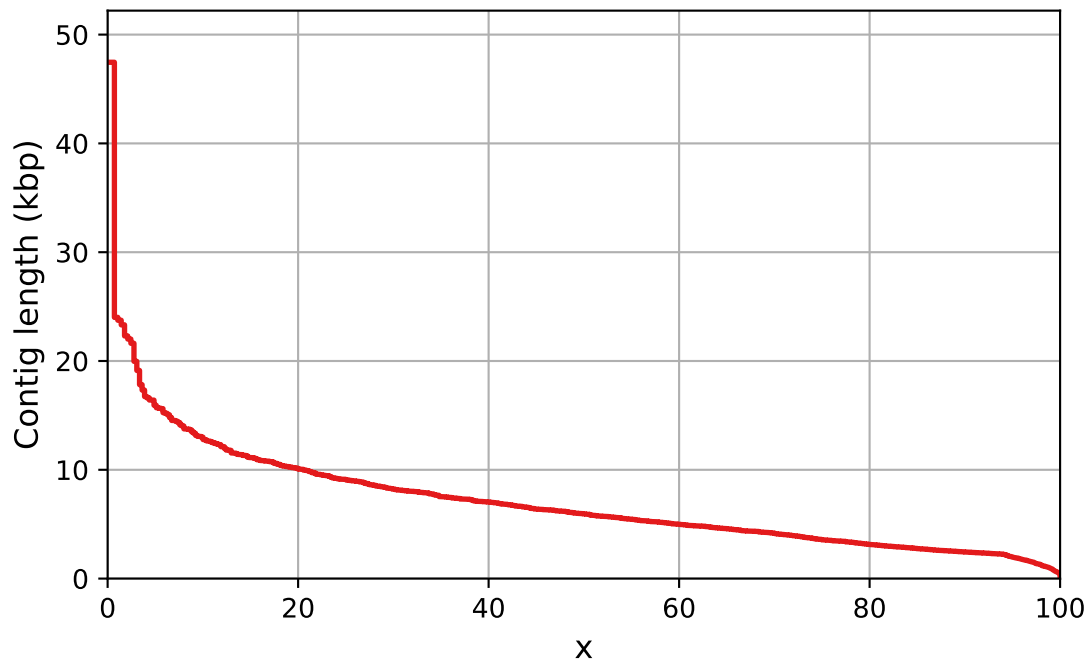


# Report

	genome.nextpolish
# contigs (>= 0 bp)	1493
# contigs (>= 1000 bp)	1380
# contigs (>= 5000 bp)	484
# contigs (>= 10000 bp)	102
# contigs (>= 25000 bp)	1
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	6672503
Total length (>= 1000 bp)	6596062
Total length (>= 5000 bp)	3994284
Total length (>= 10000 bp)	1370611
Total length (>= 25000 bp)	47457
Total length (>= 50000 bp)	0
# contigs	1488
Largest contig	47457
Total length	6670924
GC (%)	49.34
N50	5950
N90	2445
auN	7188.6
L50	364
L90	1059
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	7 + 10 part

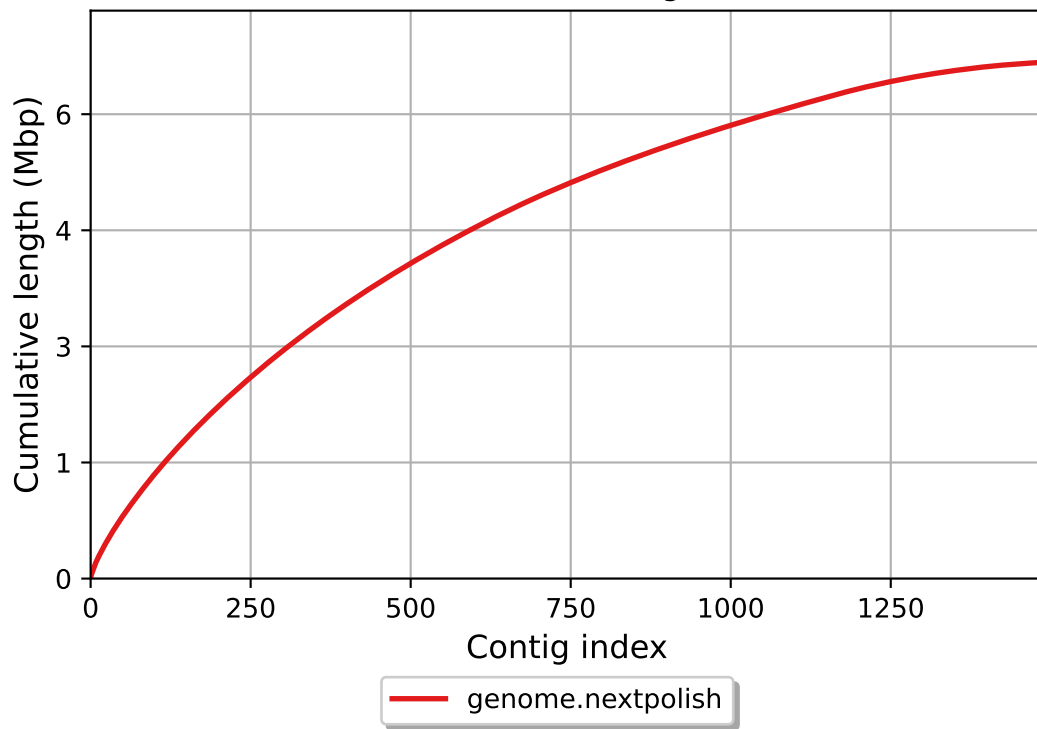
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

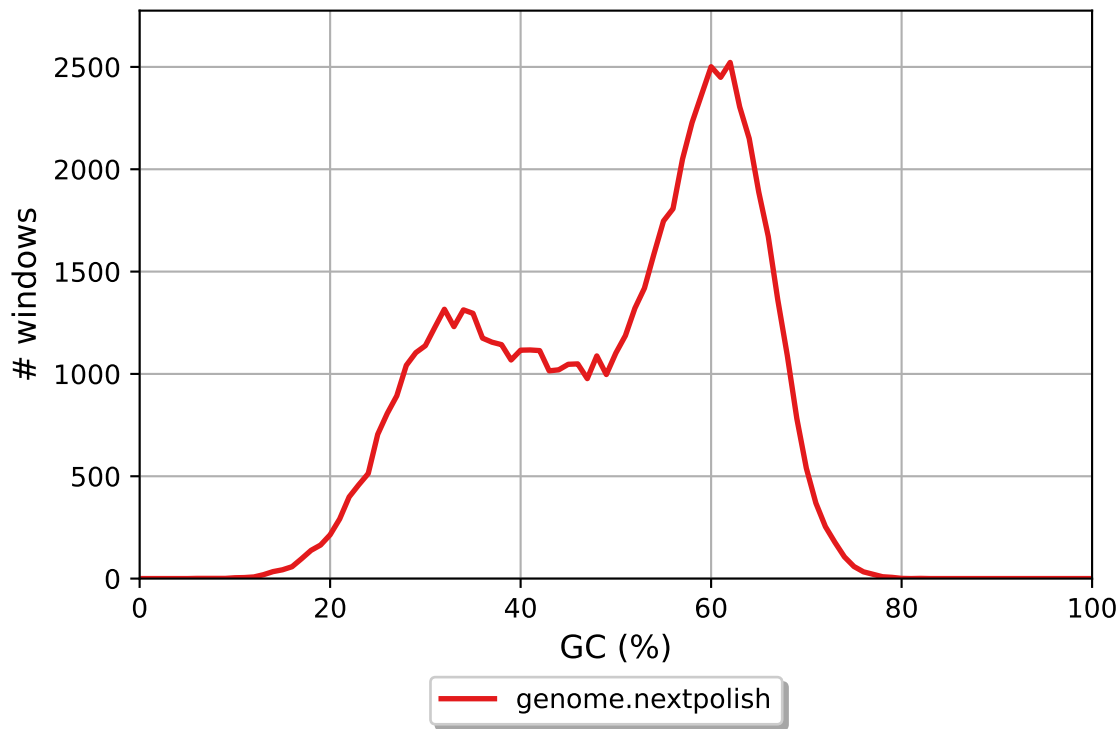


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

