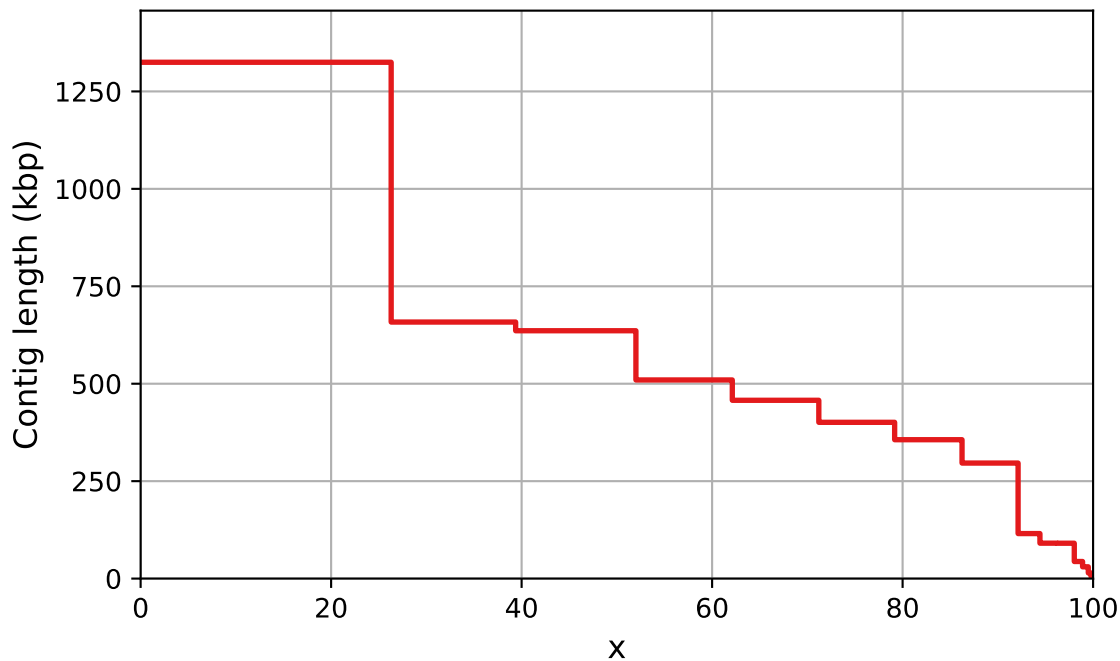


Report

	genome.nextpolish
# contigs (>= 0 bp)	17
# contigs (>= 1000 bp)	17
# contigs (>= 5000 bp)	15
# contigs (>= 10000 bp)	14
# contigs (>= 25000 bp)	13
# contigs (>= 50000 bp)	11
Total length (>= 0 bp)	5036423
Total length (>= 1000 bp)	5036423
Total length (>= 5000 bp)	5030919
Total length (>= 10000 bp)	5025473
Total length (>= 25000 bp)	5011003
Total length (>= 50000 bp)	4936913
# contigs	17
Largest contig	1324795
Total length	5036423
GC (%)	55.18
N50	635941
N90	296409
auN	689049.6
L50	3
L90	8
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	9 + 5 part

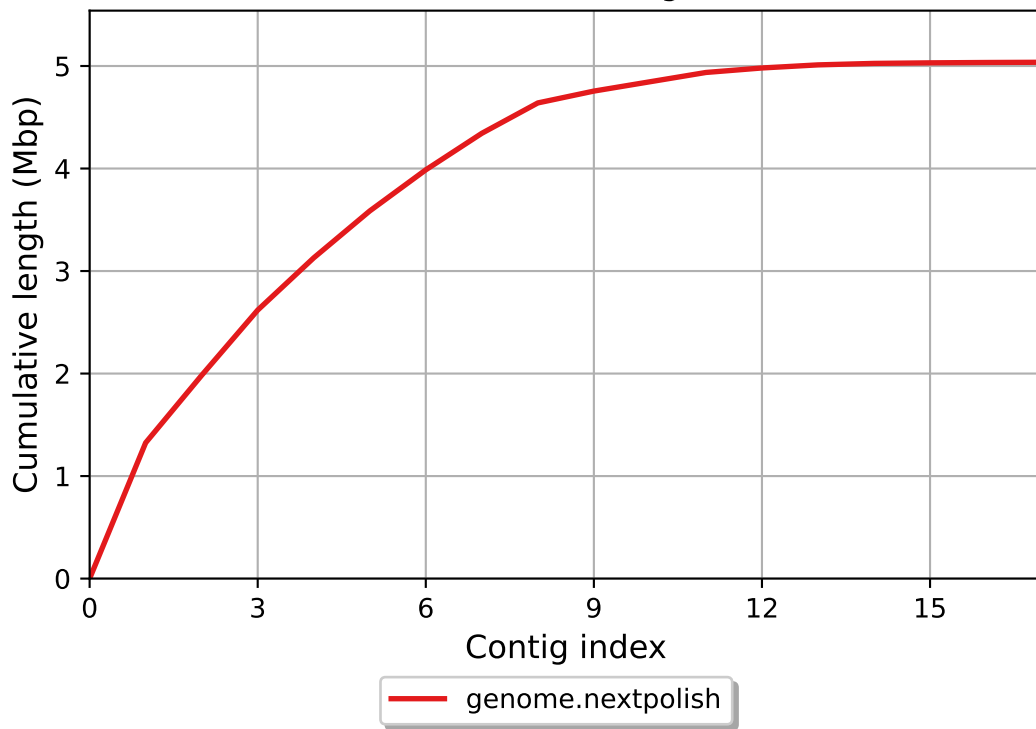
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

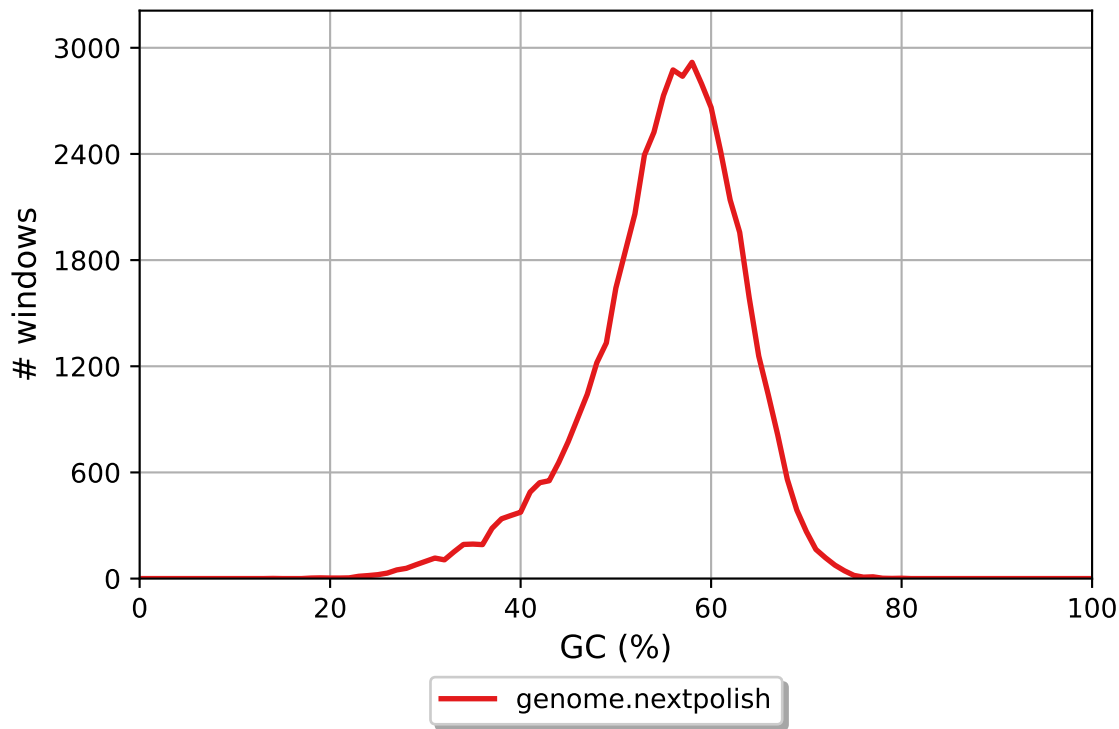


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

