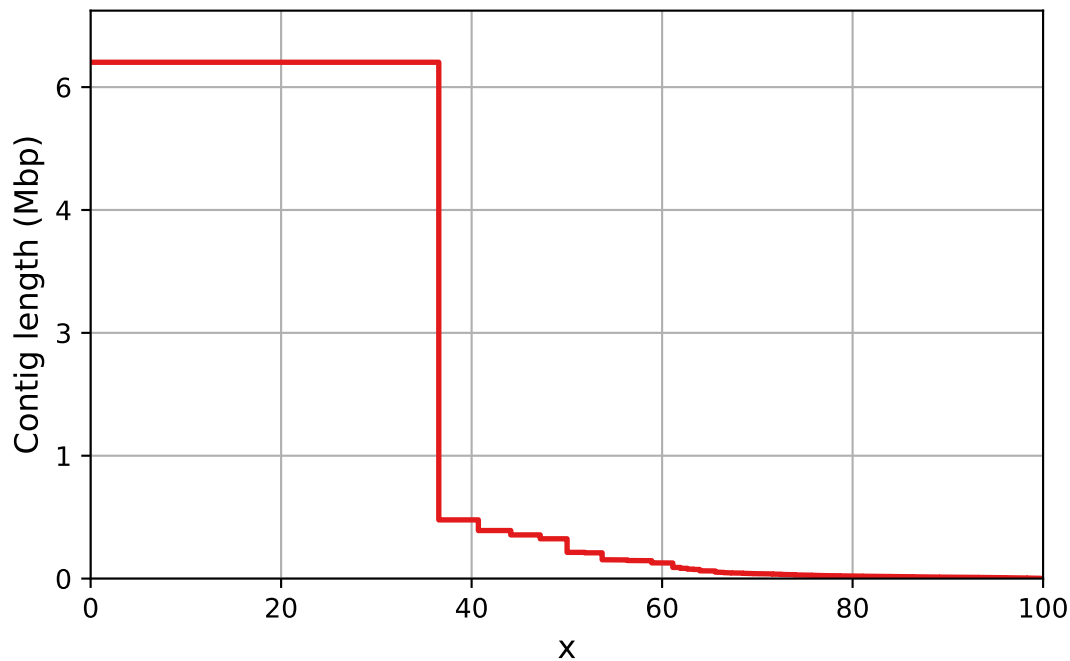


Report

	genome.nextpolish
# contigs (>= 0 bp)	332
# contigs (>= 1000 bp)	332
# contigs (>= 5000 bp)	298
# contigs (>= 10000 bp)	227
# contigs (>= 25000 bp)	100
# contigs (>= 50000 bp)	40
Total length (>= 0 bp)	17244094
Total length (>= 1000 bp)	17244094
Total length (>= 5000 bp)	17133384
Total length (>= 10000 bp)	16612540
Total length (>= 25000 bp)	14556476
Total length (>= 50000 bp)	12556788
# contigs	332
Largest contig	6303389
Total length	17244094
GC (%)	57.53
N50	485996
N90	17075
auN	2427565.2
L50	5
L90	148
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	42 + 2 part

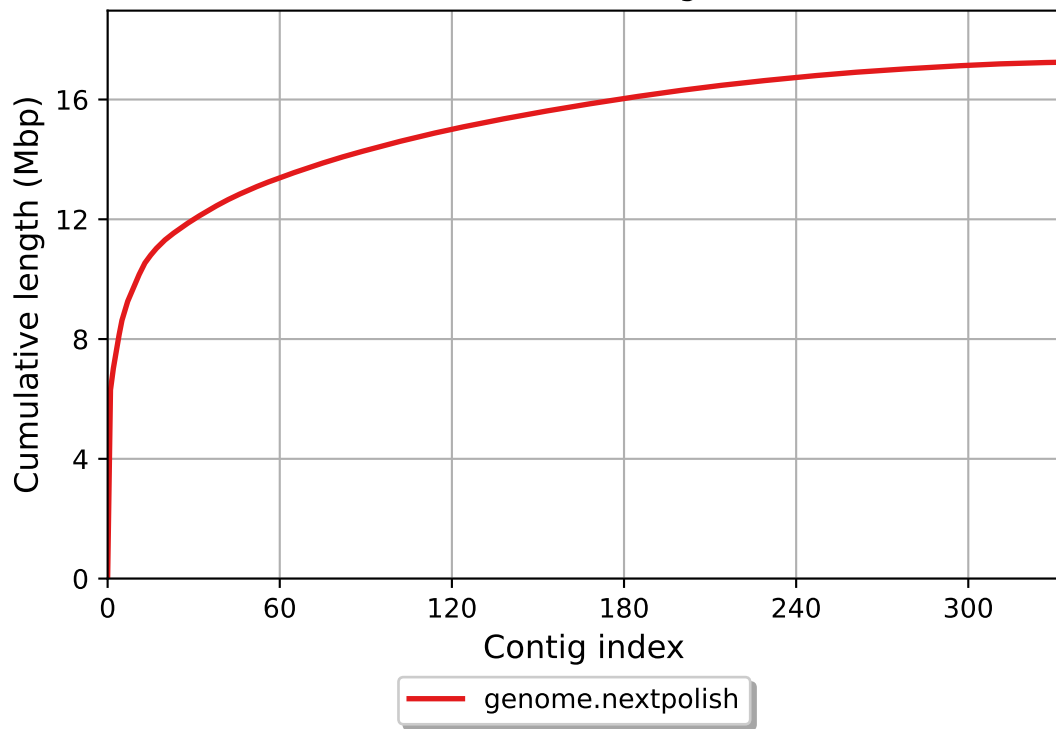
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

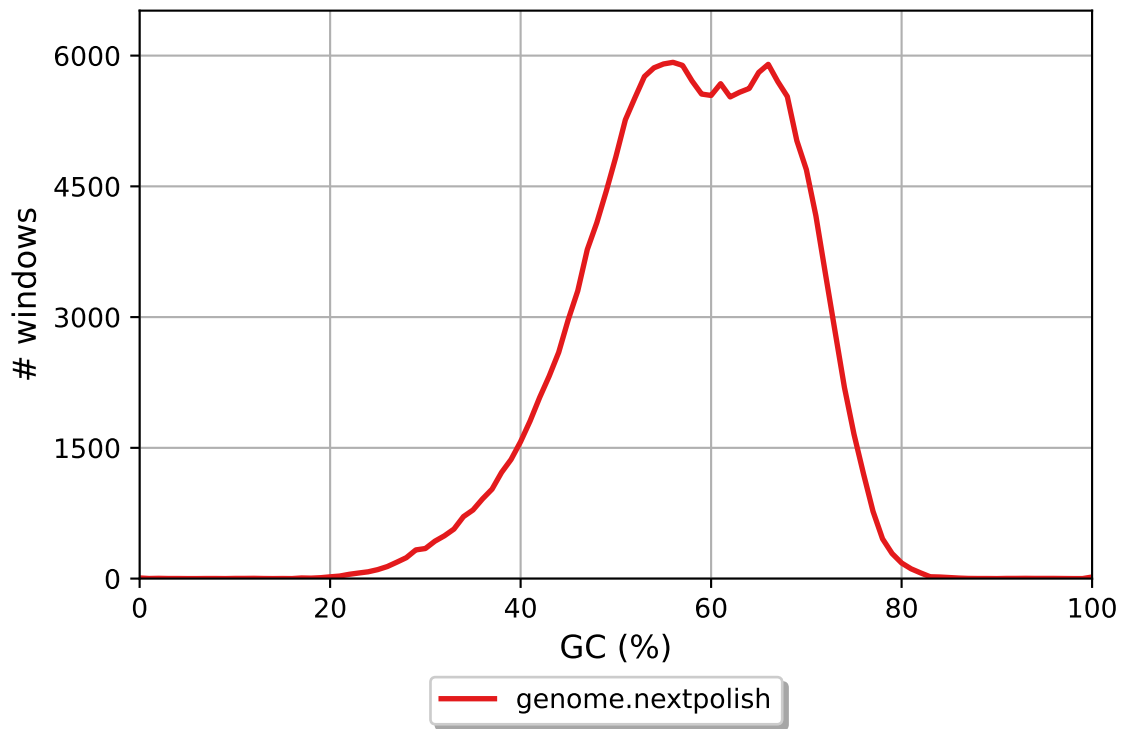


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

