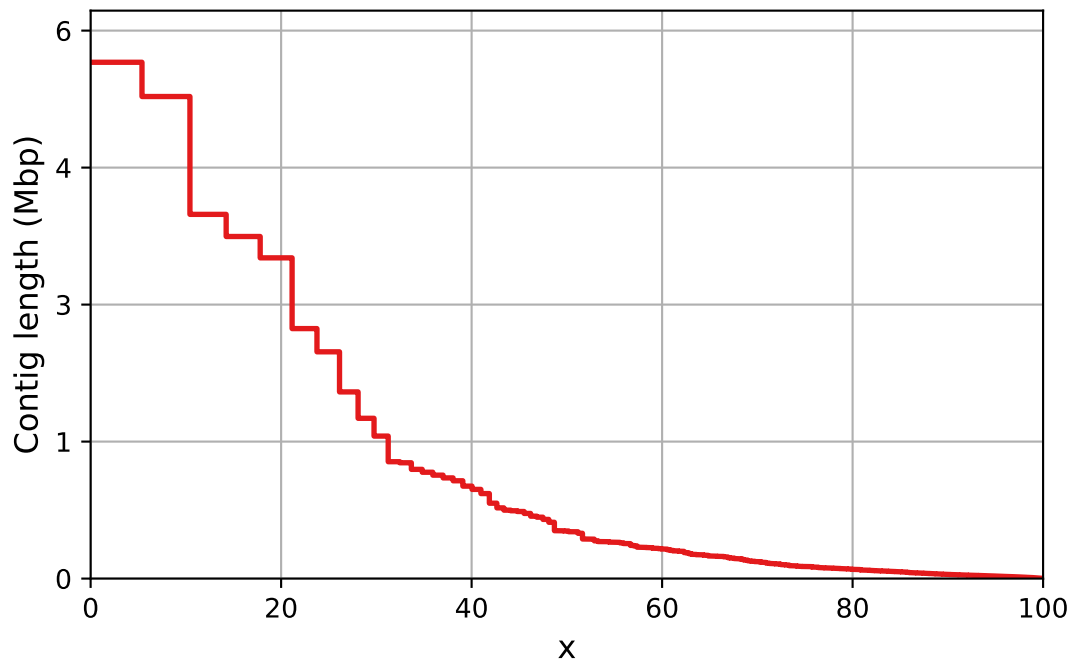


Report

	genome.nextpolish
# contigs (>= 0 bp)	914
# contigs (>= 1000 bp)	907
# contigs (>= 5000 bp)	809
# contigs (>= 10000 bp)	711
# contigs (>= 25000 bp)	505
# contigs (>= 50000 bp)	312
Total length (>= 0 bp)	104843902
Total length (>= 1000 bp)	104839011
Total length (>= 5000 bp)	104503945
Total length (>= 10000 bp)	103806440
Total length (>= 25000 bp)	100284819
Total length (>= 50000 bp)	93382290
# contigs	914
Largest contig	5653315
Total length	104843902
GC (%)	48.83
N50	521022
N90	45343
auN	1461041.3
L50	33
L90	333
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	255 + 14 part

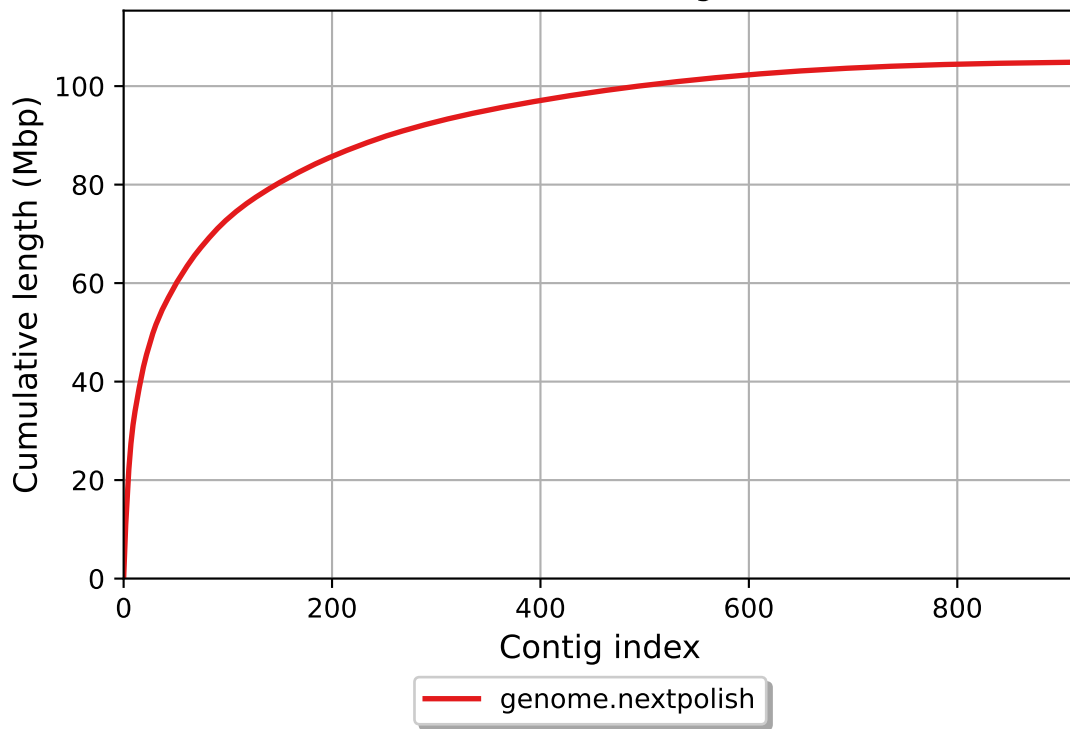
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

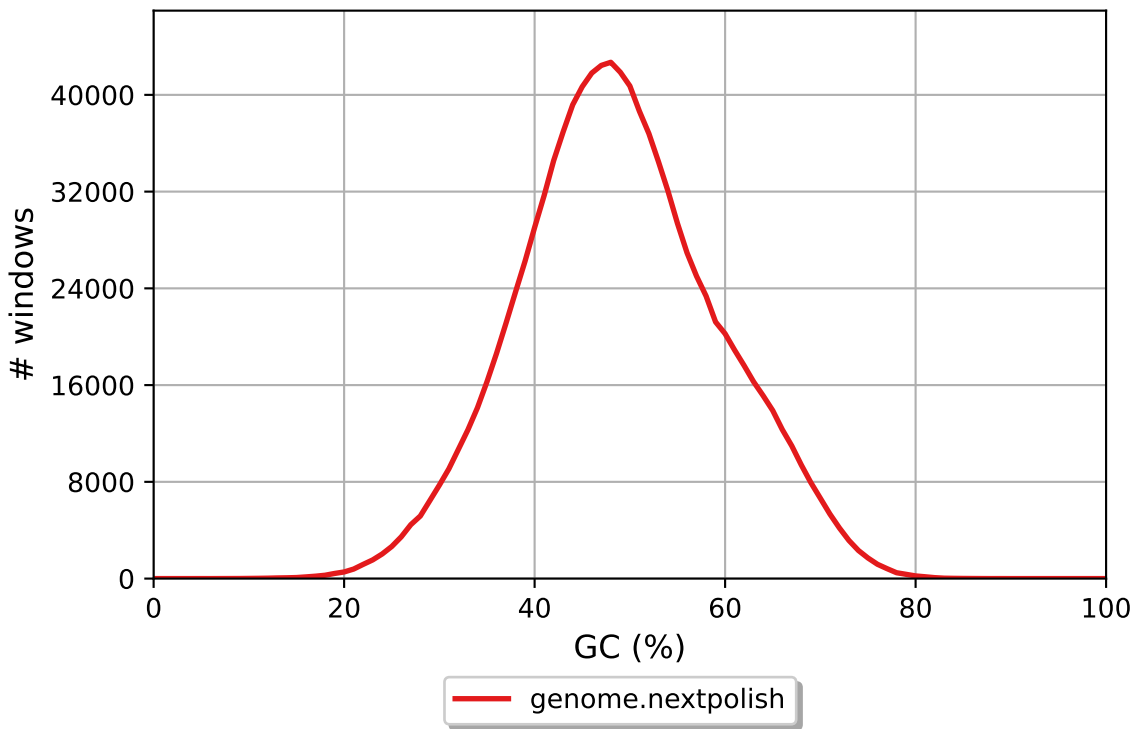


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

