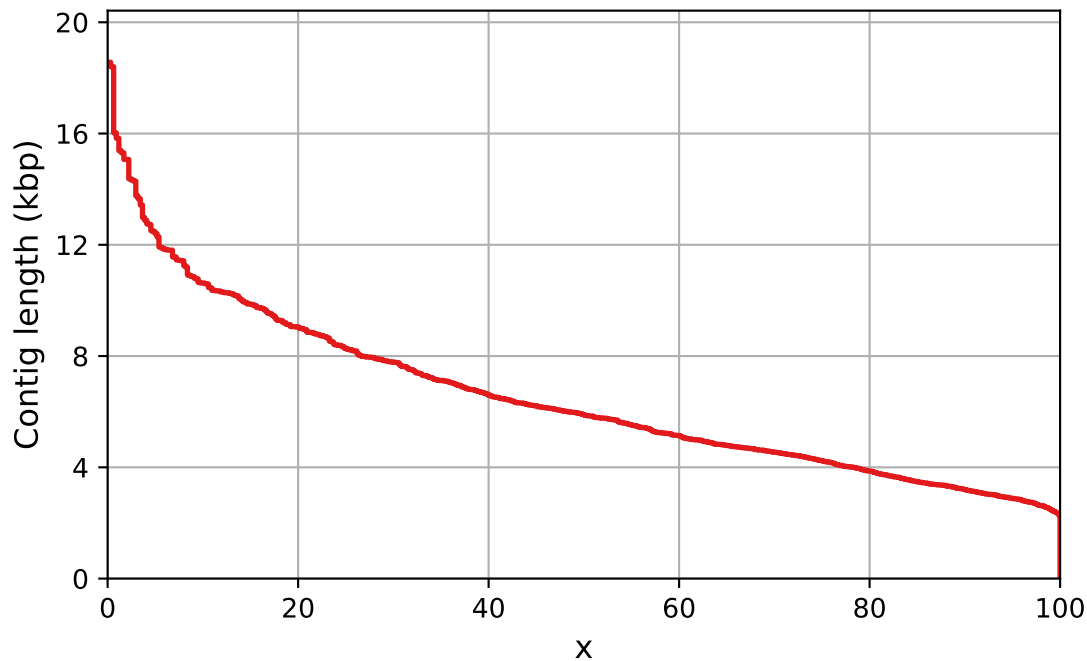


Report

	genome.nextpolish
# contigs (>= 0 bp)	1090
# contigs (>= 1000 bp)	1090
# contigs (>= 5000 bp)	473
# contigs (>= 10000 bp)	69
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	5833810
Total length (>= 1000 bp)	5833810
Total length (>= 5000 bp)	3581838
Total length (>= 10000 bp)	823037
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	1090
Largest contig	18561
Total length	5833810
GC (%)	41.25
N50	5901
N90	3197
auN	6549.5
L50	351
L90	884
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	9 + 6 part

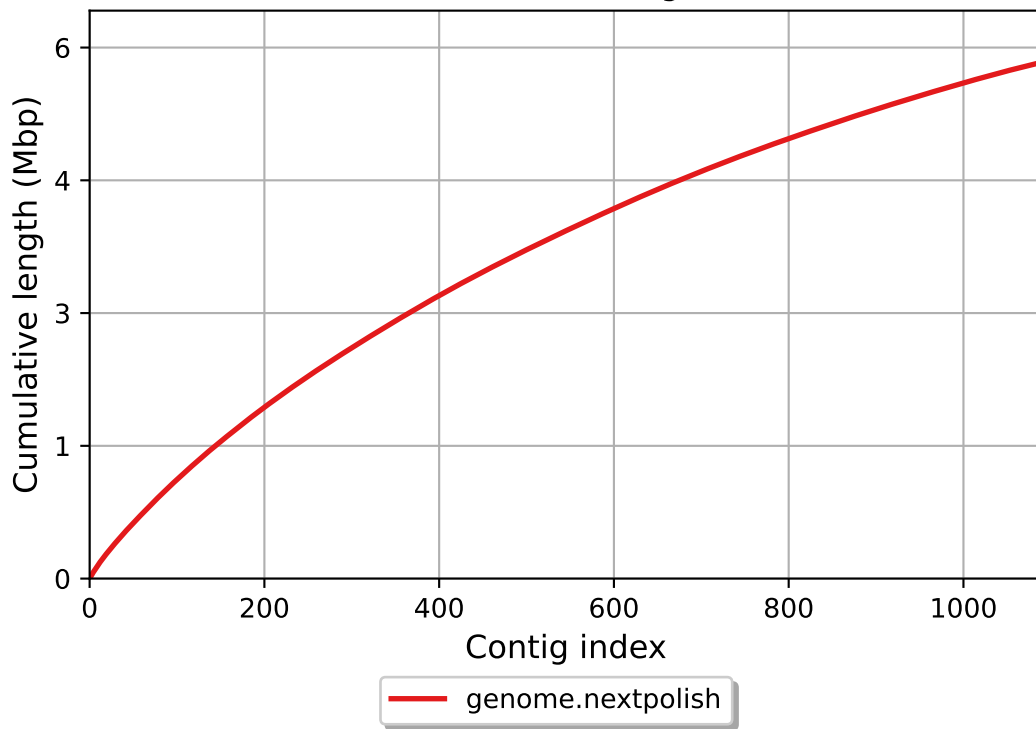
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

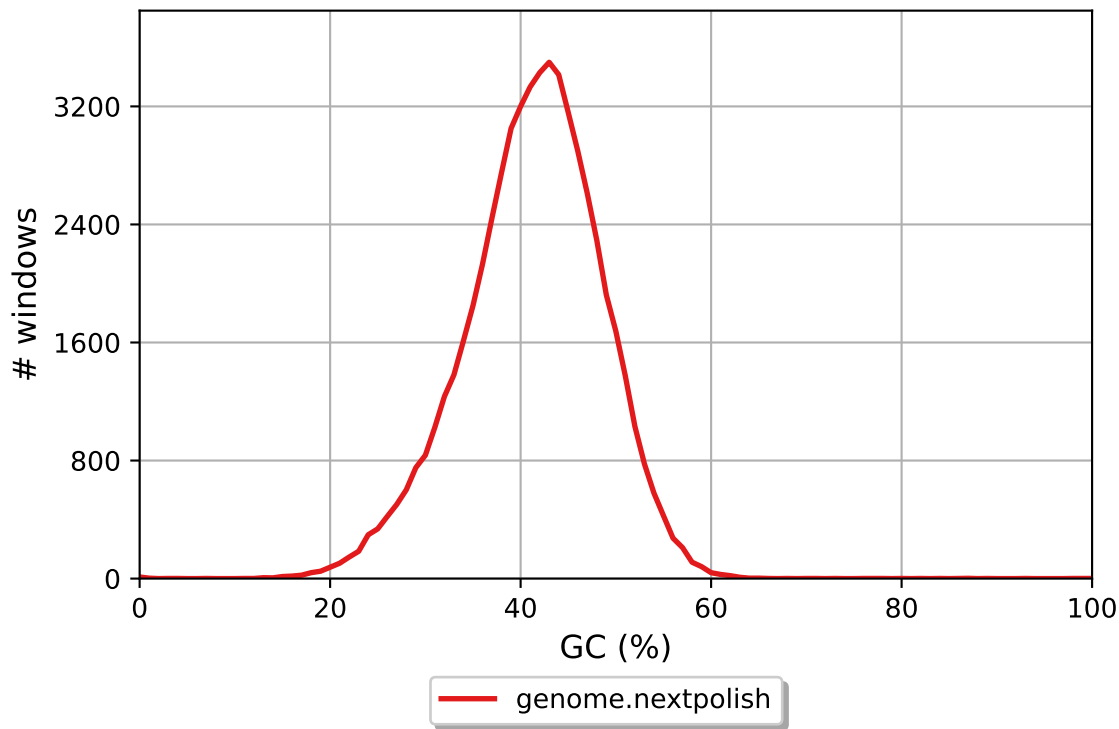


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

