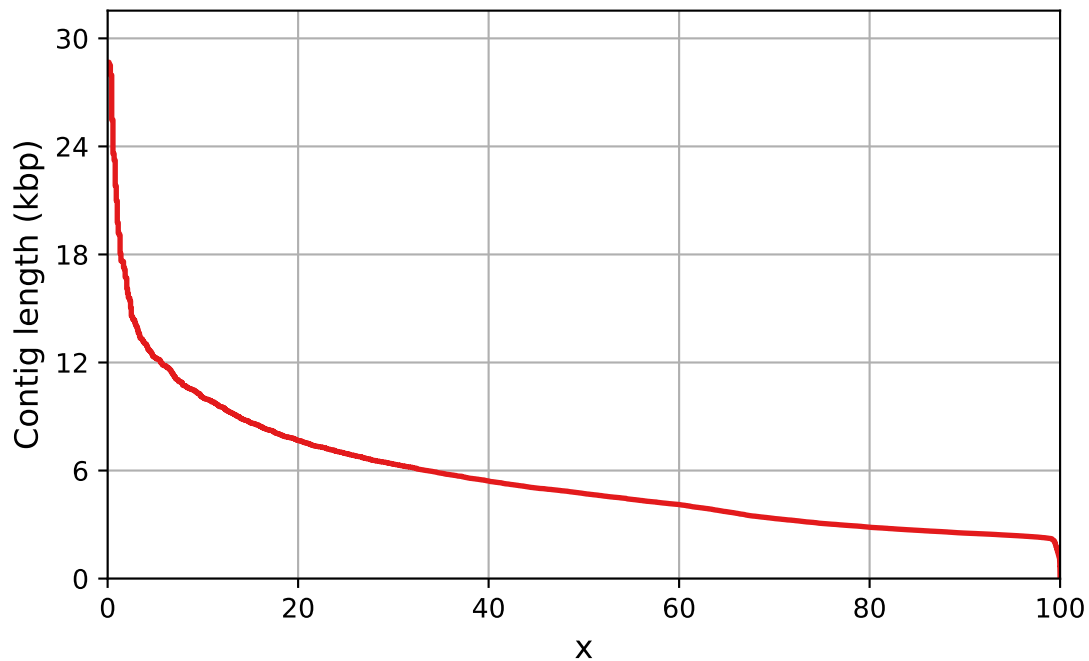


Report

	genome.nextpolish
# contigs (>= 0 bp)	4615
# contigs (>= 1000 bp)	4605
# contigs (>= 5000 bp)	1210
# contigs (>= 10000 bp)	155
# contigs (>= 25000 bp)	4
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	19752649
Total length (>= 1000 bp)	19746574
Total length (>= 5000 bp)	8989840
Total length (>= 10000 bp)	2007640
Total length (>= 25000 bp)	110626
Total length (>= 50000 bp)	0
# contigs	4615
Largest contig	28672
Total length	19752649
GC (%)	43.31
N50	4724
N90	2525
auN	5686.3
L50	1392
L90	3755
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	14 + 5 part

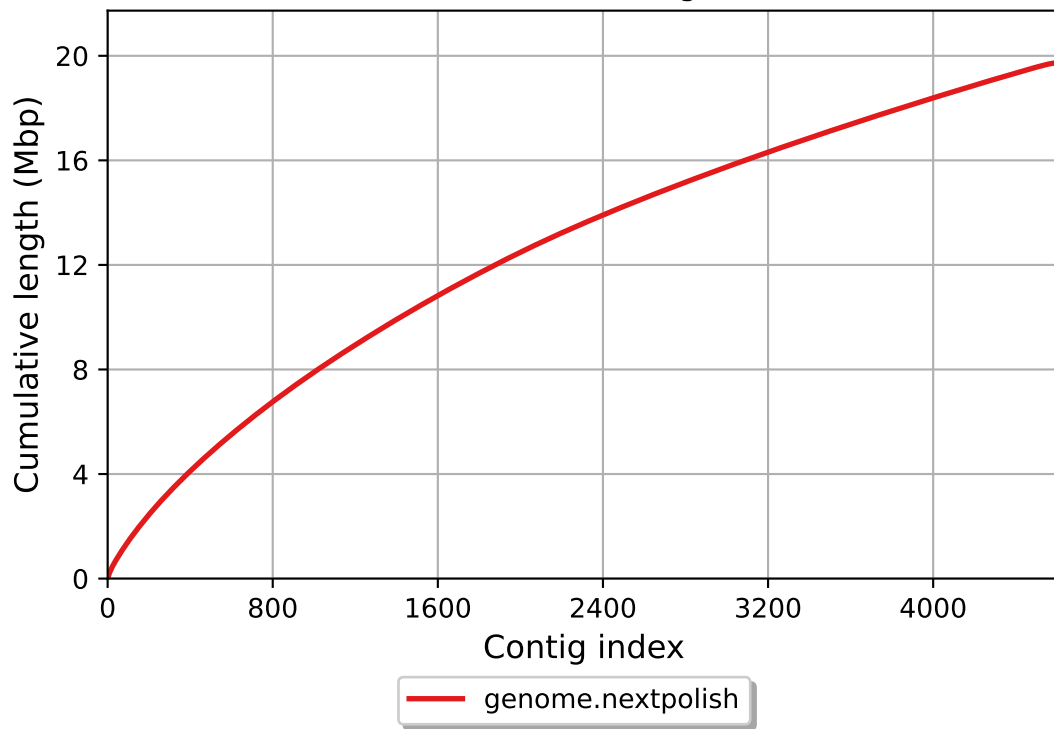
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

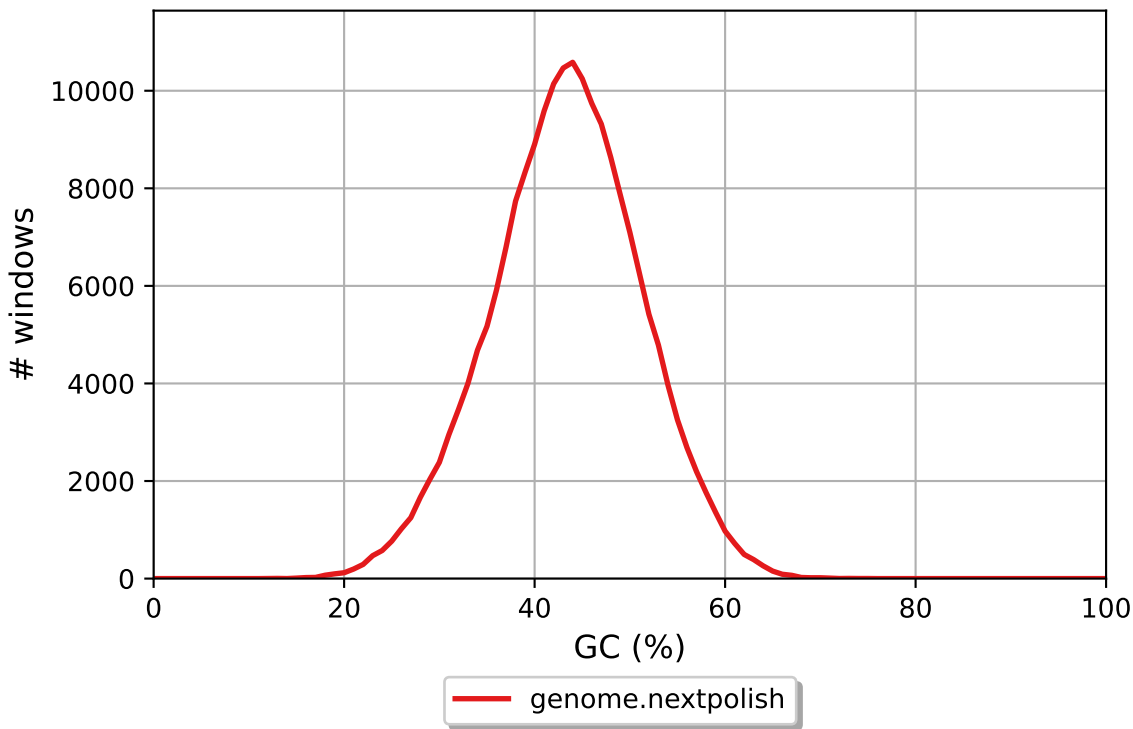


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

