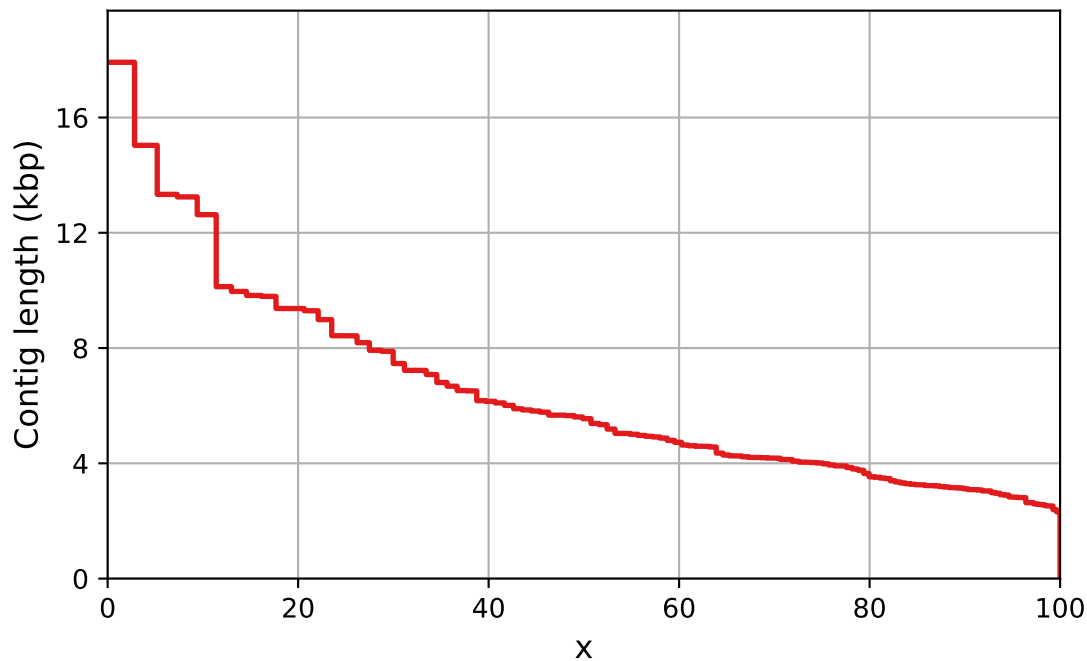


Report

	genome.nextpolish
# contigs (>= 0 bp)	123
# contigs (>= 1000 bp)	123
# contigs (>= 5000 bp)	45
# contigs (>= 10000 bp)	6
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	632876
Total length (>= 1000 bp)	632876
Total length (>= 5000 bp)	352178
Total length (>= 10000 bp)	82288
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	123
Largest contig	17917
Total length	632876
GC (%)	37.70
N50	5549
N90	3115
auN	6589.9
L50	39
L90	101
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	1 + 0 part

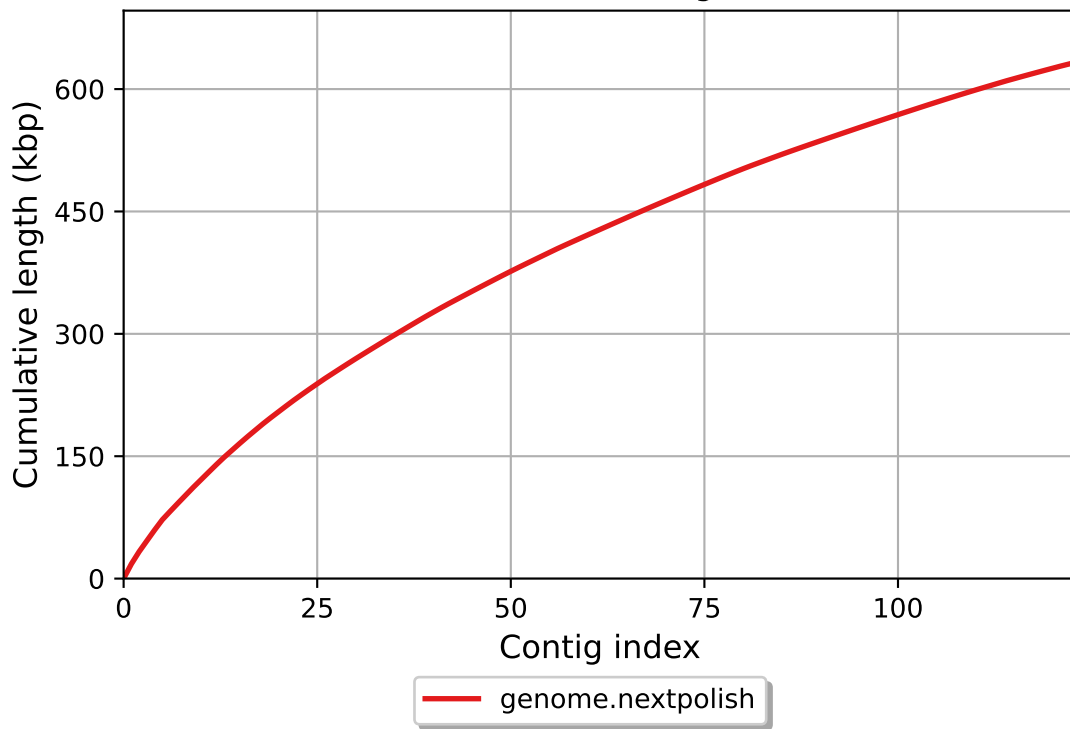
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

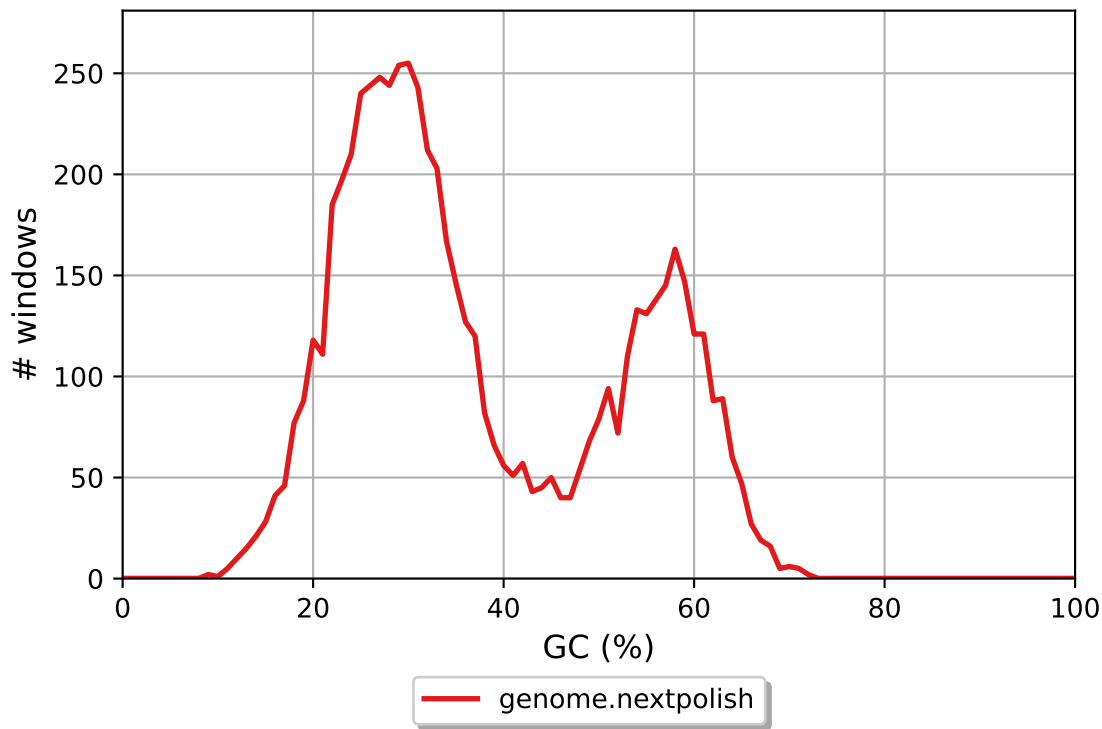


genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

