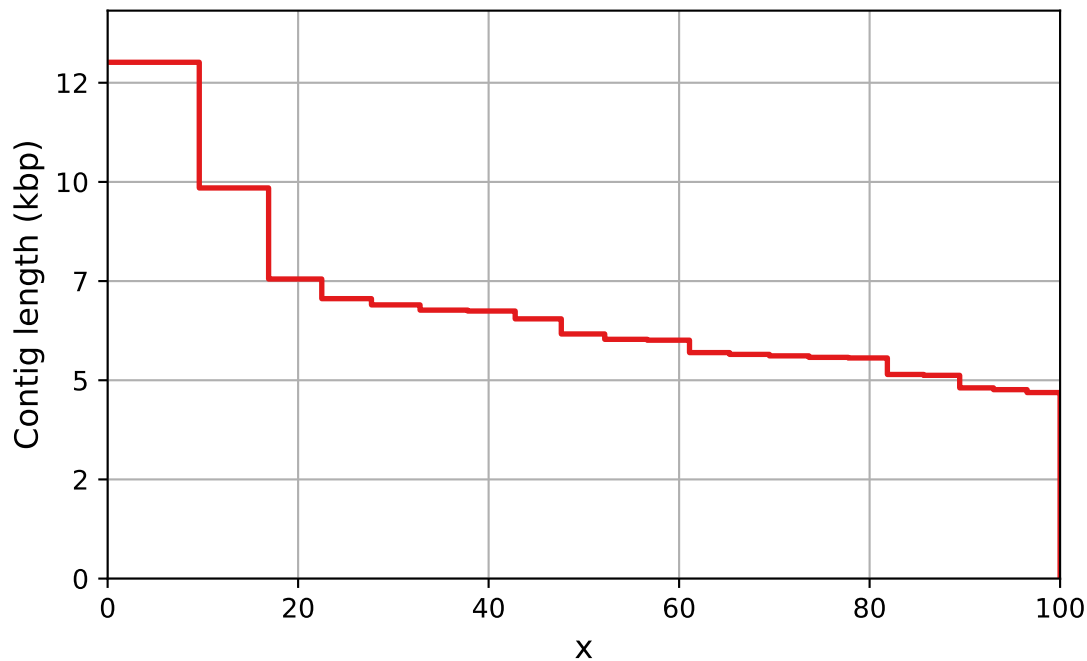


Report

	genome.nextpolish
# contigs (≥ 0 bp)	21
# contigs (≥ 1000 bp)	21
# contigs (≥ 5000 bp)	18
# contigs (≥ 10000 bp)	1
# contigs (≥ 25000 bp)	0
# contigs (≥ 50000 bp)	0
Total length (≥ 0 bp)	135277
Total length (≥ 1000 bp)	135277
Total length (≥ 5000 bp)	121016
Total length (≥ 10000 bp)	13016
Total length (≥ 25000 bp)	0
Total length (≥ 50000 bp)	0
# contigs	21
Largest contig	13016
Total length	135277
GC (%)	52.37
N50	6167
N90	4808
auN	6978.7
L50	9
L90	19
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	0 + 0 part

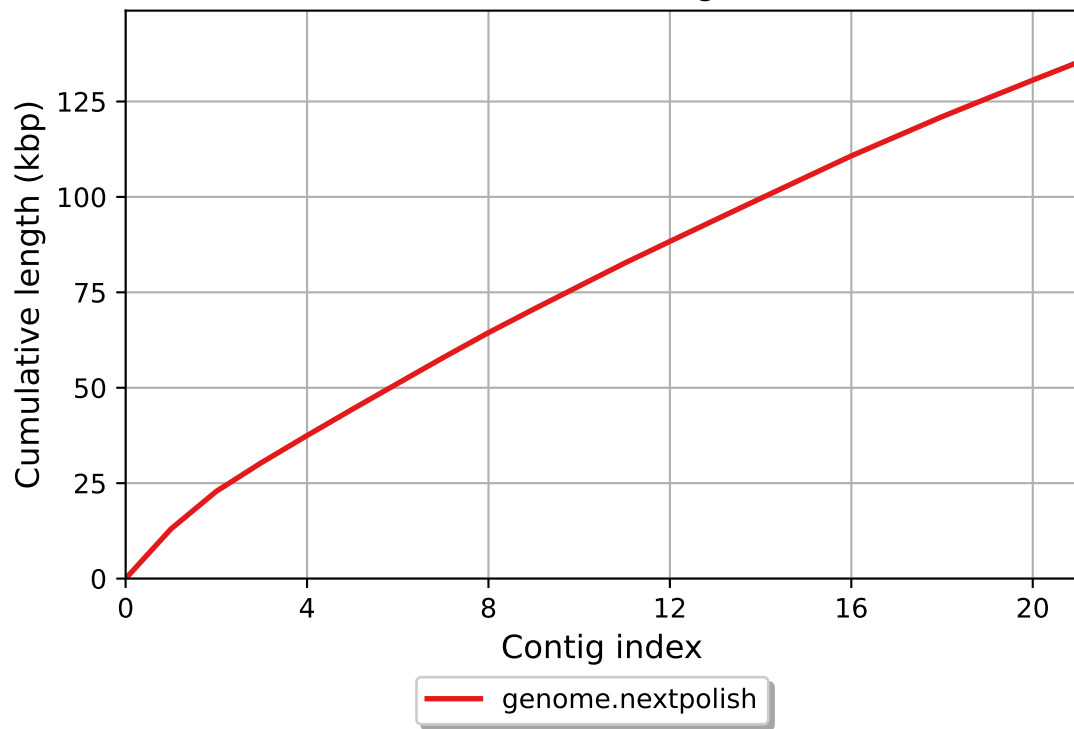
All statistics are based on contigs of size ≥ 500 bp, unless otherwise noted (e.g., "# contigs (≥ 0 bp)" and "Total length (≥ 0 bp)" include all contigs).

Nx

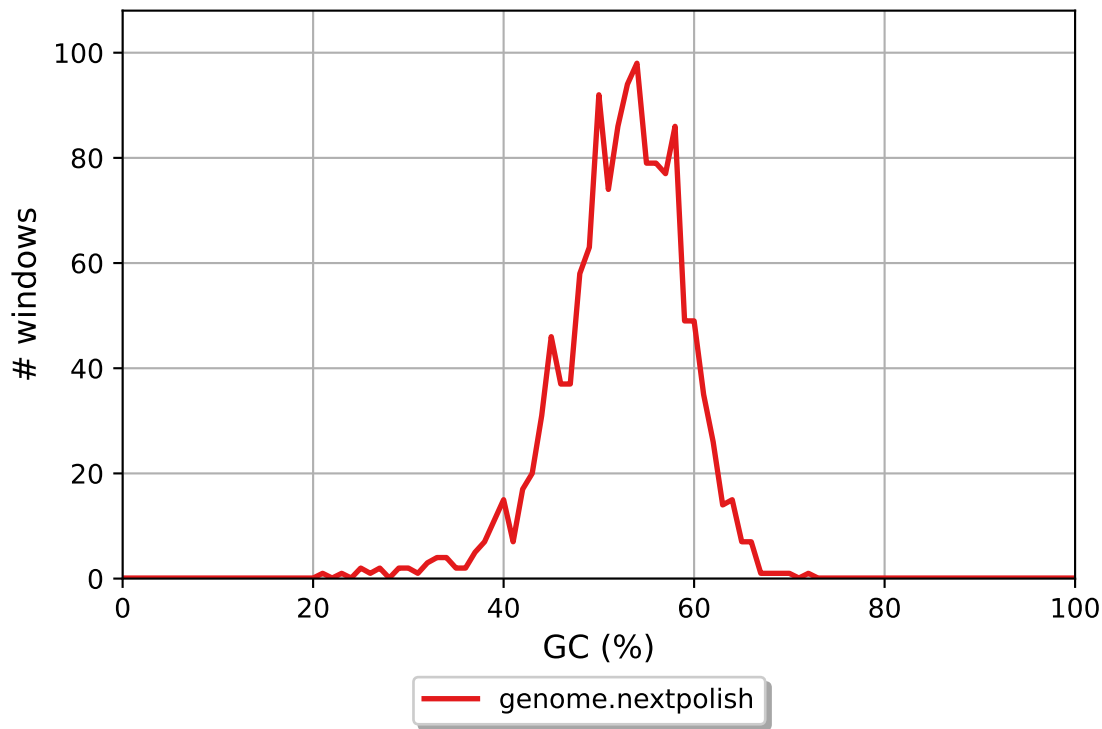


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

