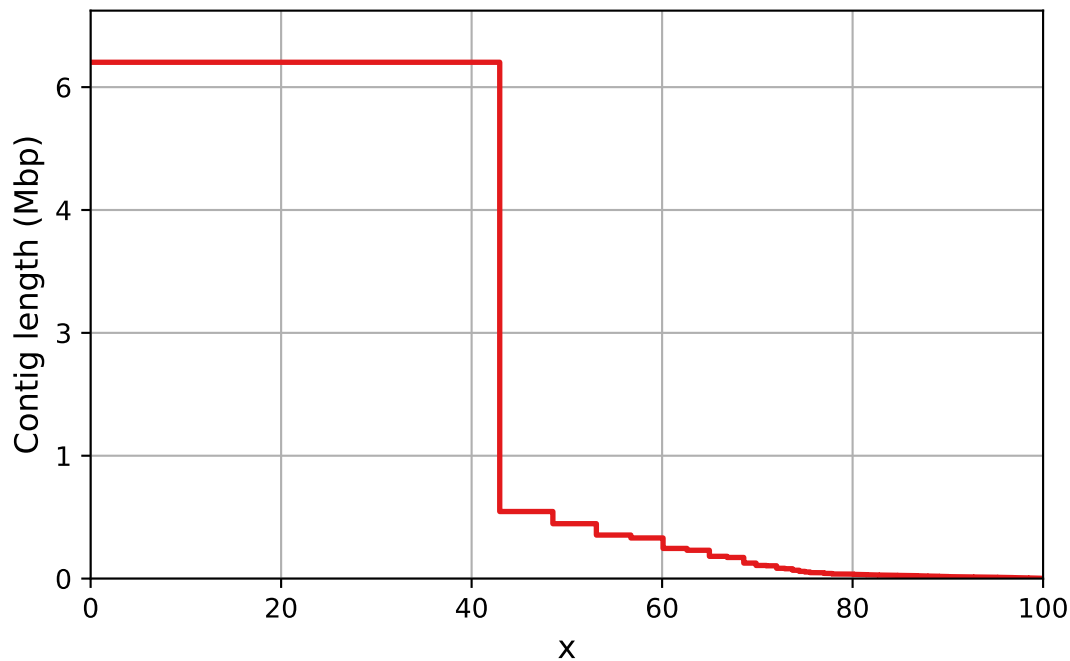


Report

	genome.nextpolish
# contigs (>= 0 bp)	190
# contigs (>= 1000 bp)	190
# contigs (>= 5000 bp)	173
# contigs (>= 10000 bp)	135
# contigs (>= 25000 bp)	67
# contigs (>= 50000 bp)	30
Total length (>= 0 bp)	14675796
Total length (>= 1000 bp)	14675796
Total length (>= 5000 bp)	14608985
Total length (>= 10000 bp)	14336772
Total length (>= 25000 bp)	13199839
Total length (>= 50000 bp)	11860435
# contigs	190
Largest contig	6303579
Total length	14675796
GC (%)	57.95
N50	670165
N90	24606
auN	2864506.5
L50	3
L90	68
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	45 + 1 part

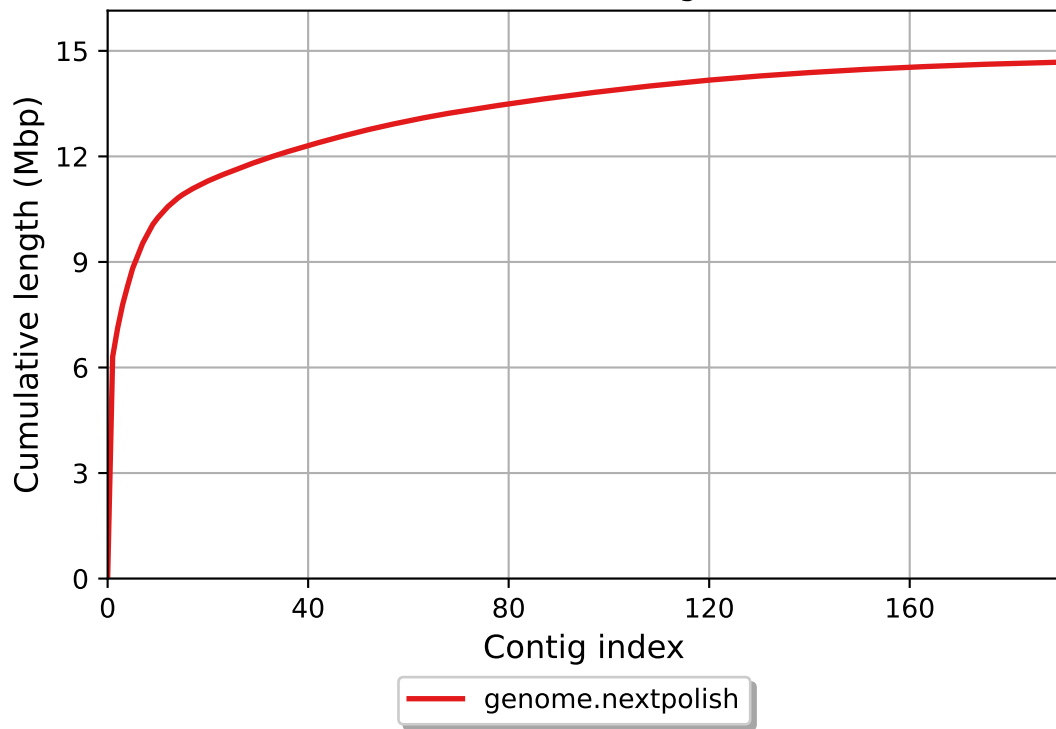
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

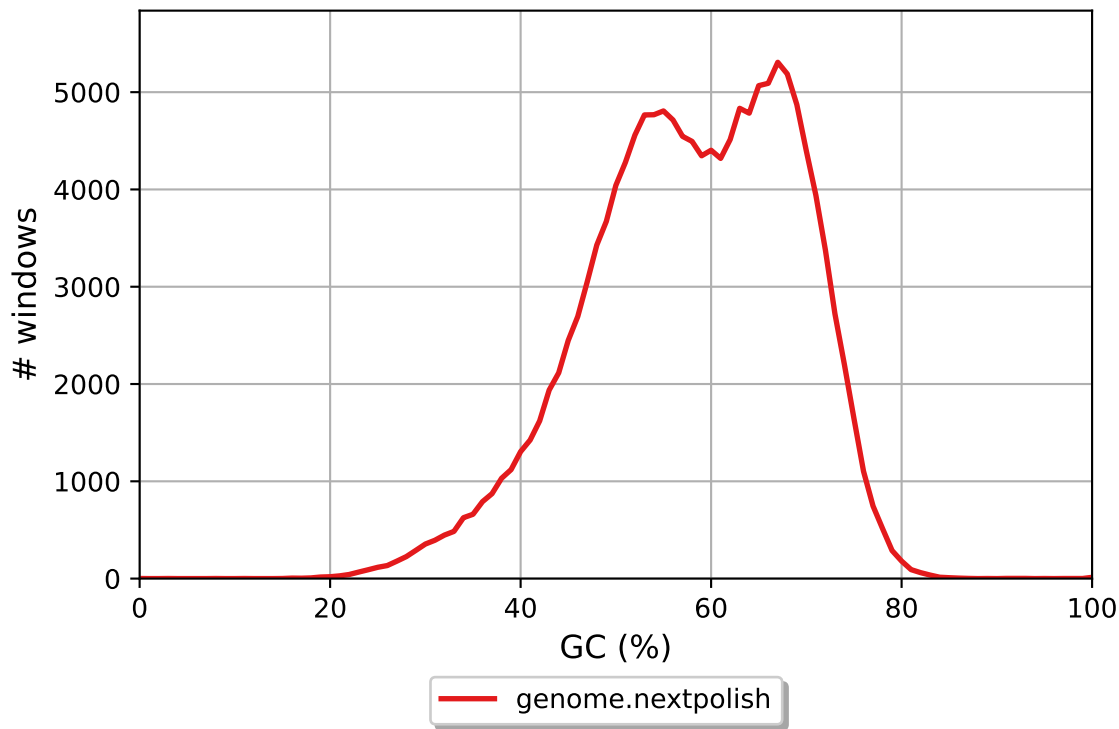


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

