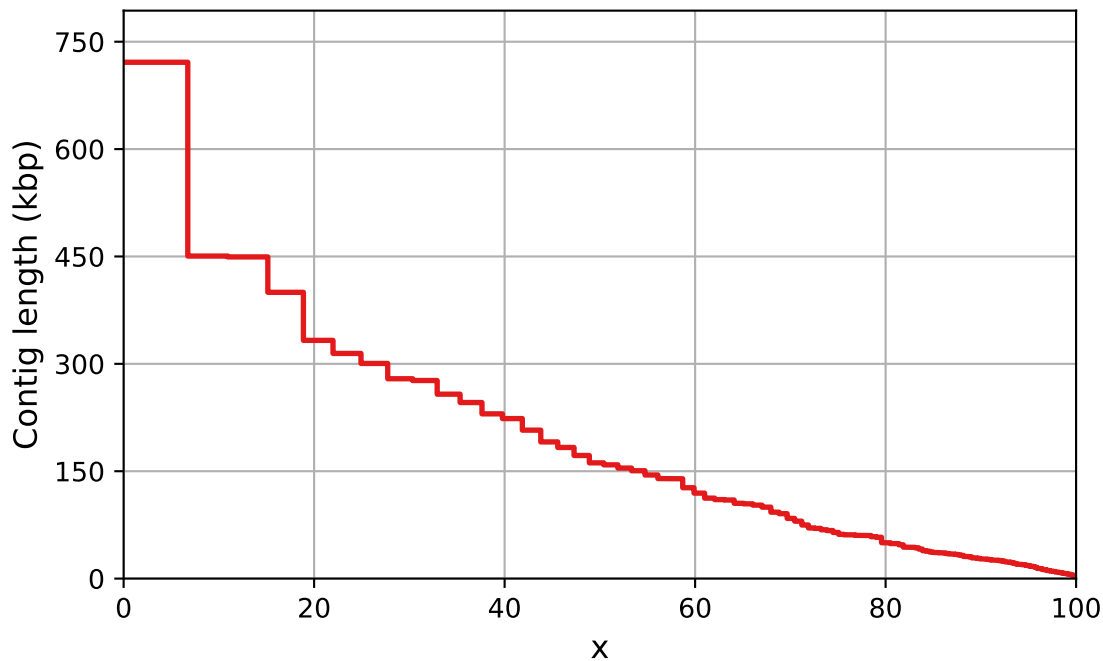


Report

	genome.nextpolish
# contigs (>= 0 bp)	163
# contigs (>= 1000 bp)	163
# contigs (>= 5000 bp)	154
# contigs (>= 10000 bp)	125
# contigs (>= 25000 bp)	89
# contigs (>= 50000 bp)	52
Total length (>= 0 bp)	10707471
Total length (>= 1000 bp)	10707471
Total length (>= 5000 bp)	10668423
Total length (>= 10000 bp)	10455876
Total length (>= 25000 bp)	9858793
Total length (>= 50000 bp)	8570135
# contigs	163
Largest contig	721288
Total length	10707471
GC (%)	58.07
N50	161684
N90	27609
auN	217549.3
L50	18
L90	81
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	39 + 0 part

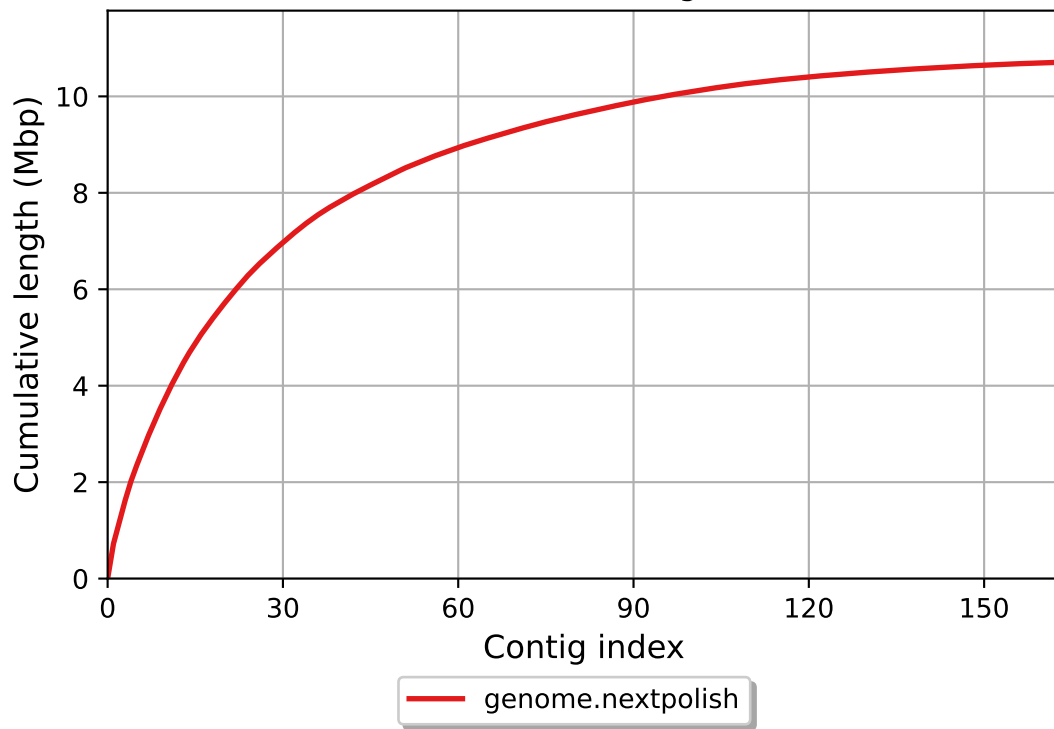
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

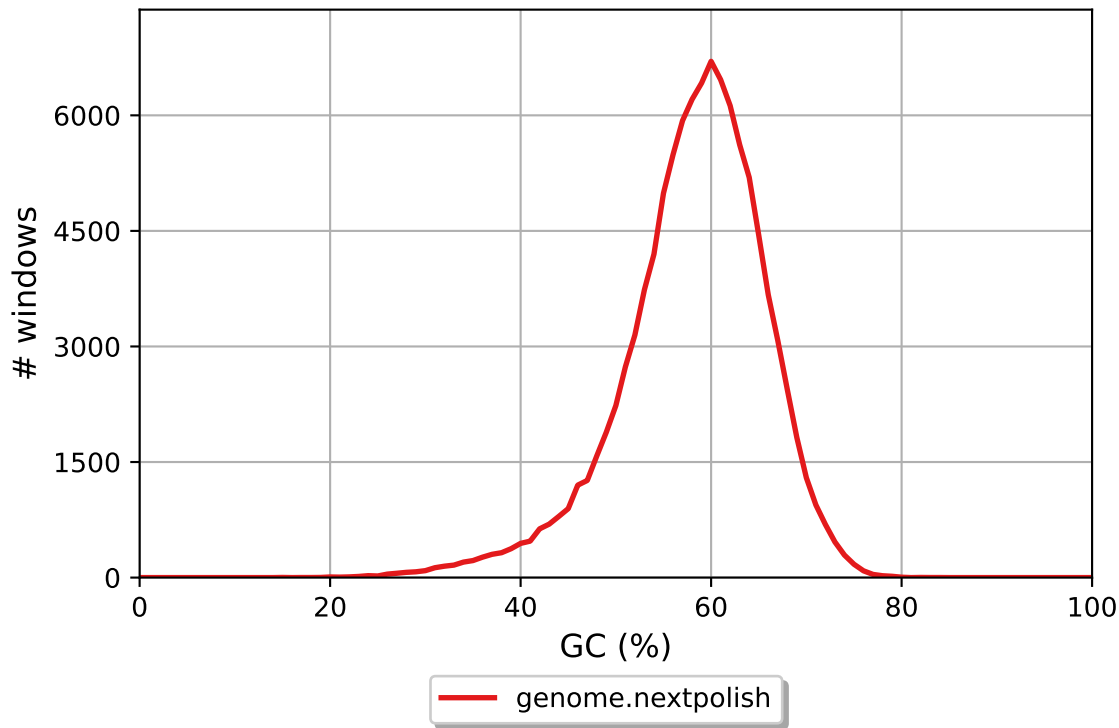


genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

