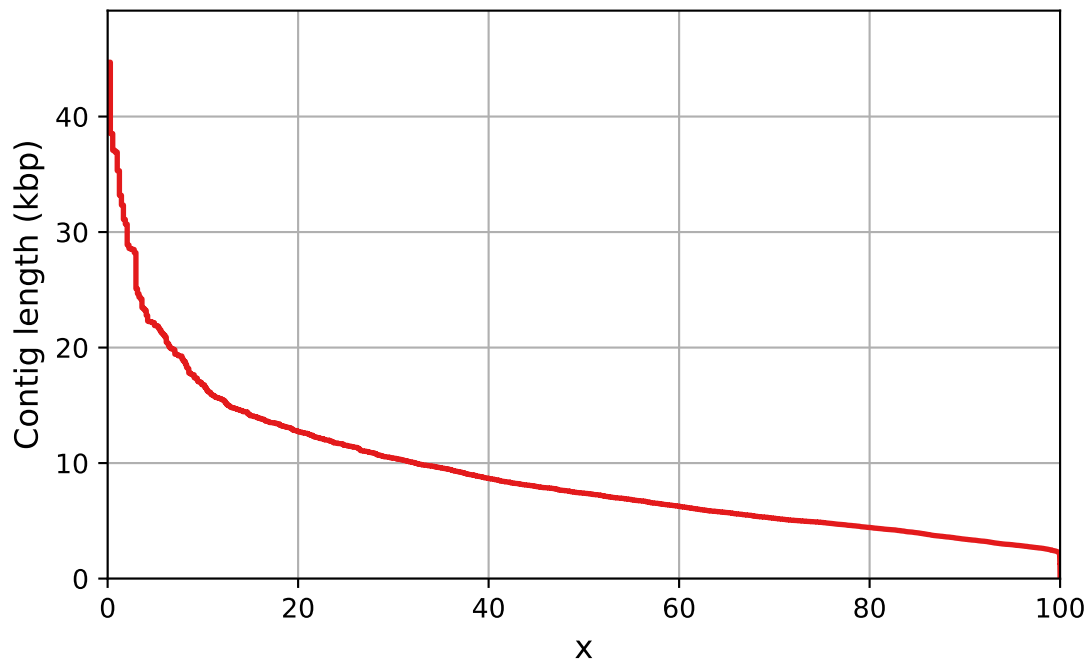


Report

	genome.nextpolish
# contigs (>= 0 bp)	2437
# contigs (>= 1000 bp)	2433
# contigs (>= 5000 bp)	1258
# contigs (>= 10000 bp)	348
# contigs (>= 25000 bp)	15
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	15563232
Total length (>= 1000 bp)	15560364
Total length (>= 5000 bp)	11329227
Total length (>= 10000 bp)	5030728
Total length (>= 25000 bp)	487615
Total length (>= 50000 bp)	0
# contigs	2437
Largest contig	44699
Total length	15563232
GC (%)	41.97
N50	7386
N90	3417
auN	9202.6
L50	672
L90	1893
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	24 + 2 part

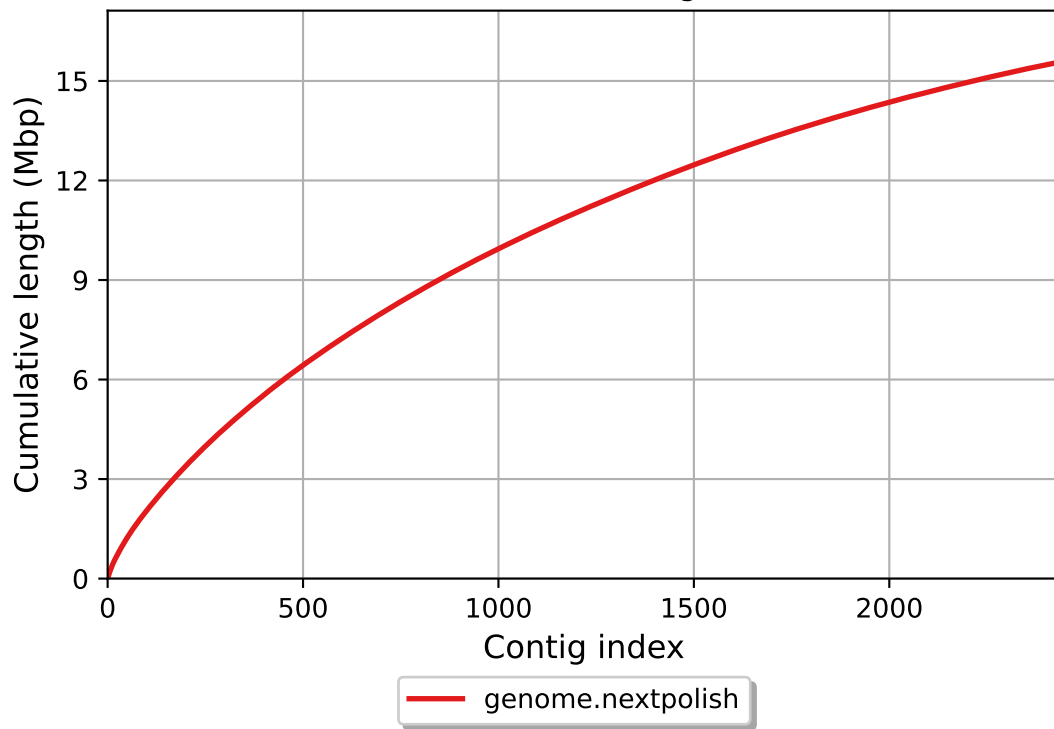
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

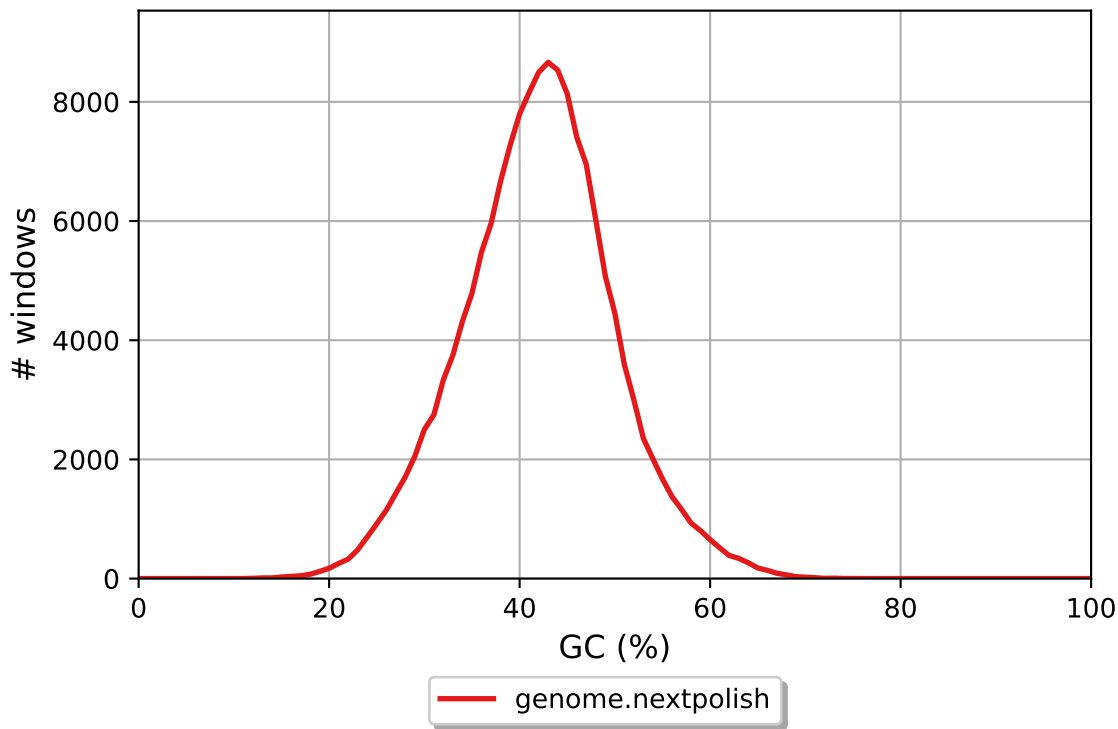


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

