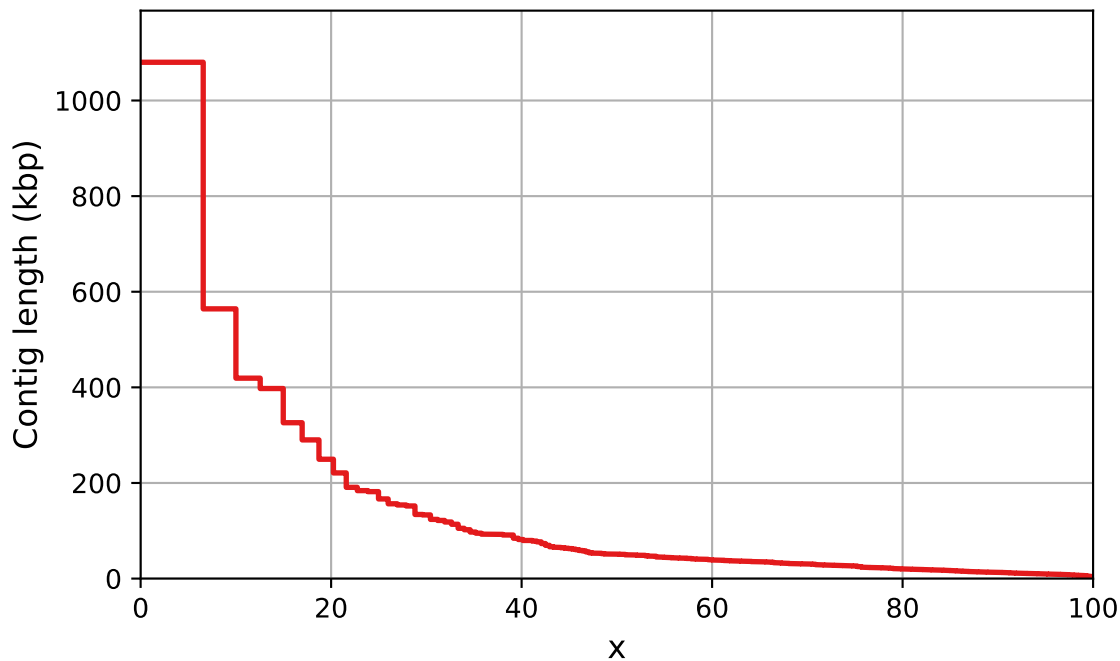


Report

	genome.nextpolish
# contigs (>= 0 bp)	498
# contigs (>= 1000 bp)	498
# contigs (>= 5000 bp)	481
# contigs (>= 10000 bp)	375
# contigs (>= 25000 bp)	176
# contigs (>= 50000 bp)	62
Total length (>= 0 bp)	16433979
Total length (>= 1000 bp)	16433979
Total length (>= 5000 bp)	16362214
Total length (>= 10000 bp)	15525387
Total length (>= 25000 bp)	12413123
Total length (>= 50000 bp)	8360750
# contigs	498
Largest contig	1079983
Total length	16433979
GC (%)	49.14
N50	51062
N90	12911
auN	172720.4
L50	60
L90	310
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	46 + 4 part

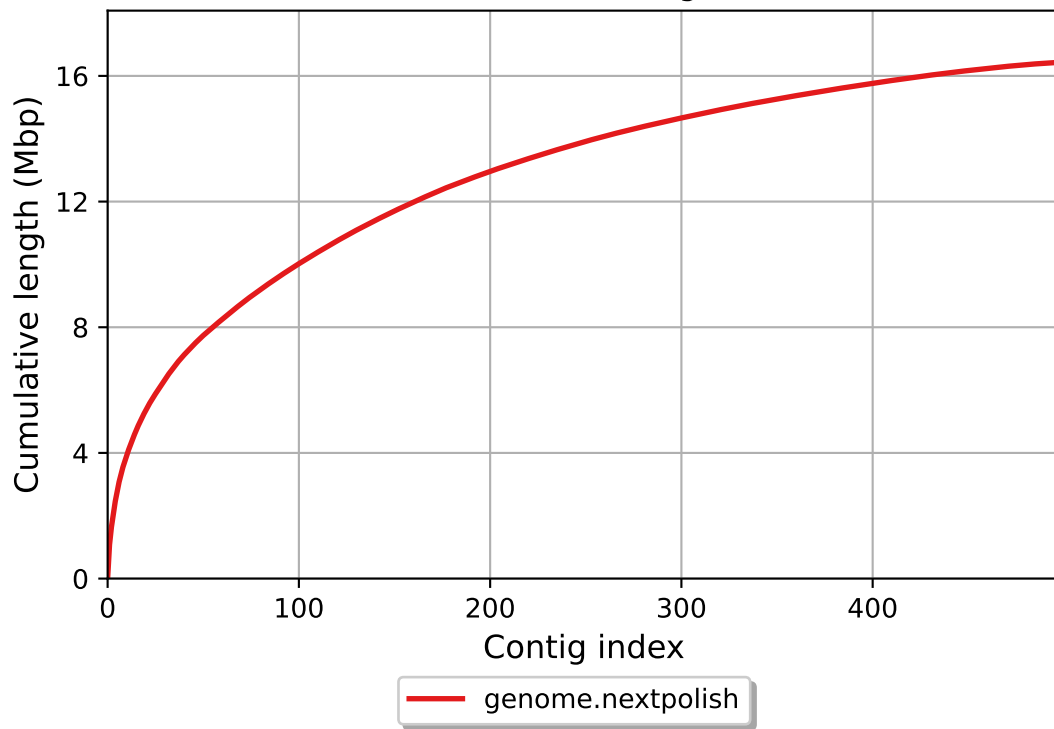
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

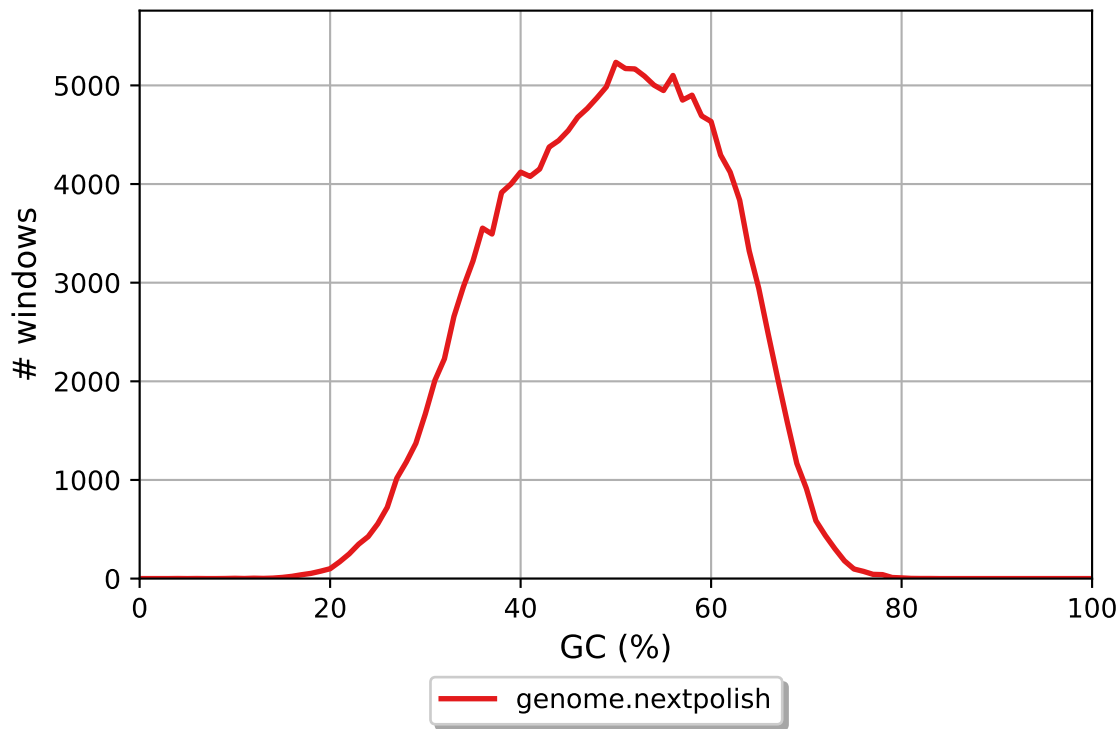


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

