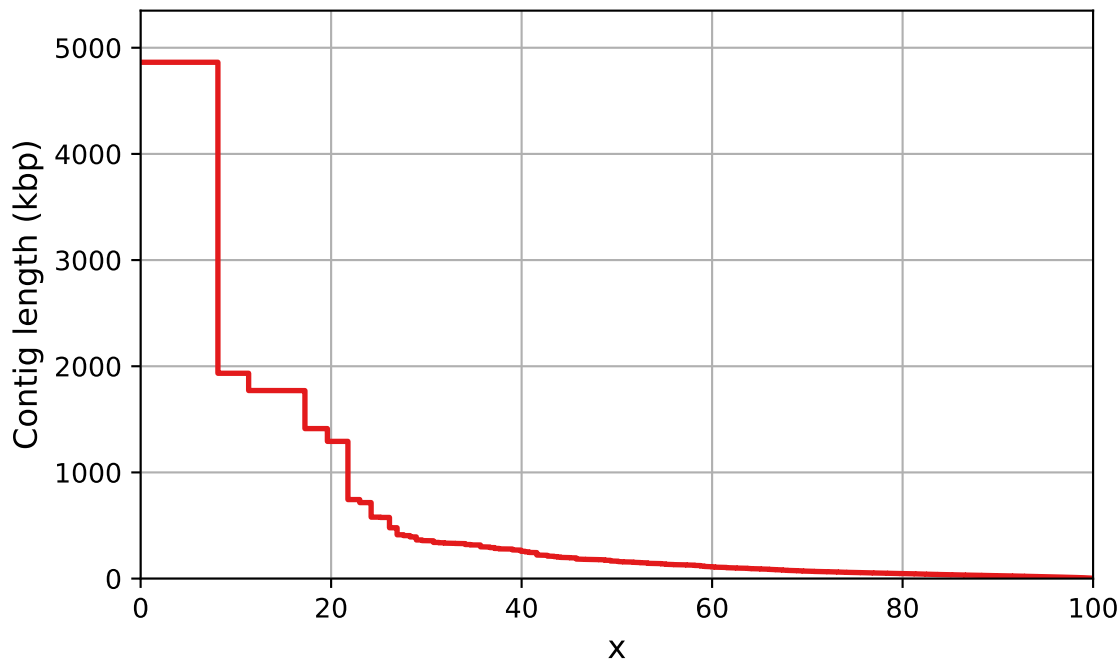


Report

	genome.nextpolish
# contigs (>= 0 bp)	902
# contigs (>= 1000 bp)	900
# contigs (>= 5000 bp)	850
# contigs (>= 10000 bp)	732
# contigs (>= 25000 bp)	481
# contigs (>= 50000 bp)	263
Total length (>= 0 bp)	59933535
Total length (>= 1000 bp)	59932291
Total length (>= 5000 bp)	59765111
Total length (>= 10000 bp)	58892542
Total length (>= 25000 bp)	54736003
Total length (>= 50000 bp)	47013830
# contigs	902
Largest contig	4863176
Total length	59933535
GC (%)	53.42
N50	163767
N90	27227
auN	752672.1
L50	66
L90	451
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	128 + 1 part

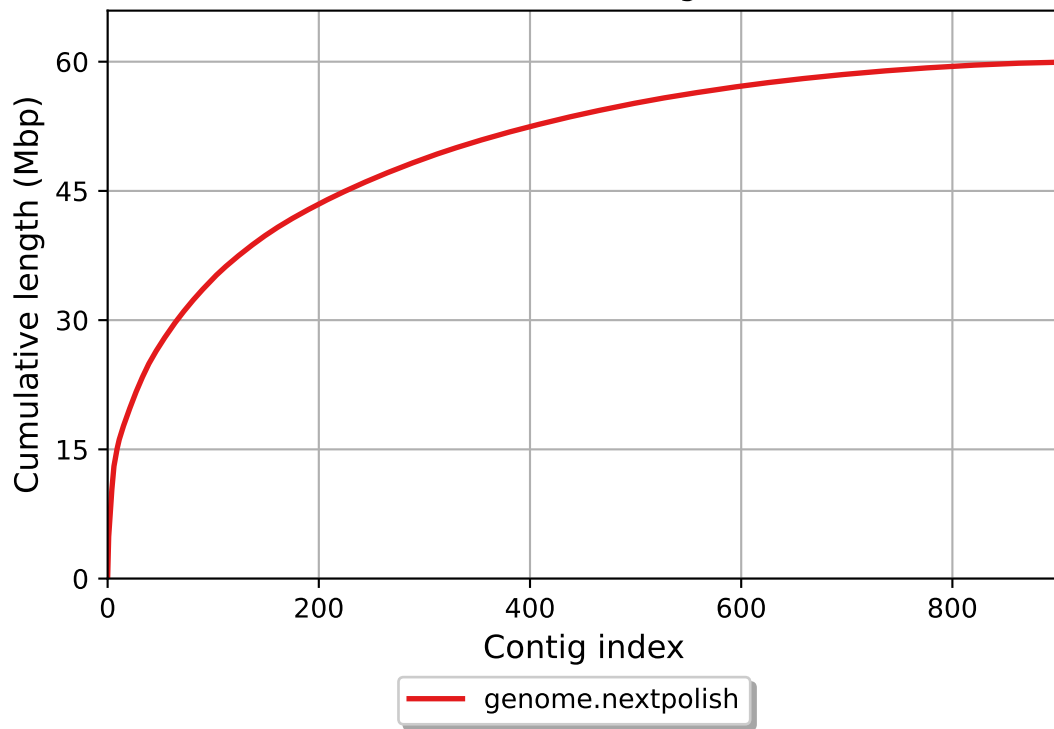
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

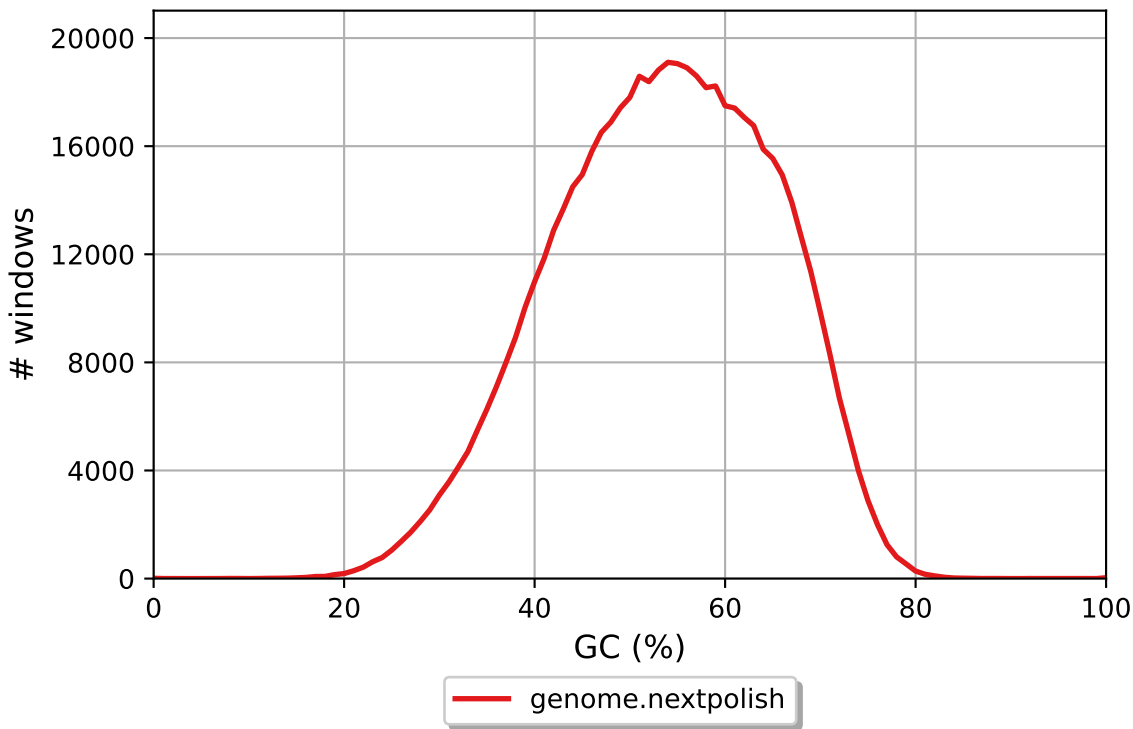


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

