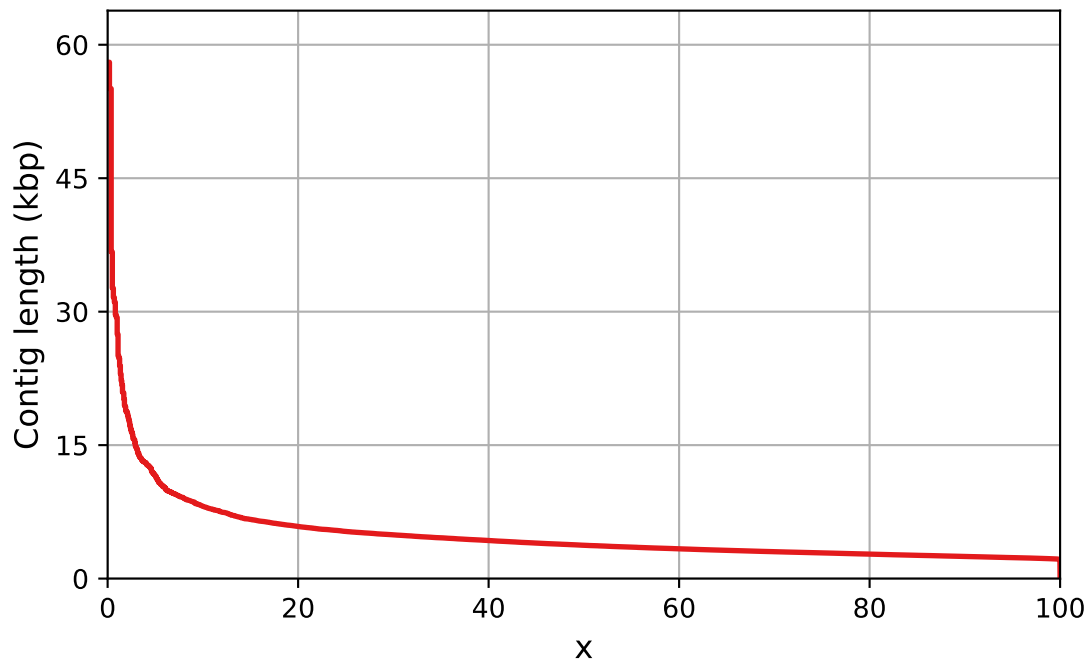


# Report

	genome.nextpolish
# contigs (>= 0 bp)	7999
# contigs (>= 1000 bp)	7999
# contigs (>= 5000 bp)	1194
# contigs (>= 10000 bp)	119
# contigs (>= 25000 bp)	10
# contigs (>= 50000 bp)	2
Total length (>= 0 bp)	30282332
Total length (>= 1000 bp)	30282332
Total length (>= 5000 bp)	8714335
Total length (>= 10000 bp)	1858265
Total length (>= 25000 bp)	356623
Total length (>= 50000 bp)	113098
# contigs	7999
Largest contig	58030
Total length	30282332
GC (%)	37.99
N50	3744
N90	2486
auN	5058.7
L50	2689
L90	6708
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	5 + 6 part

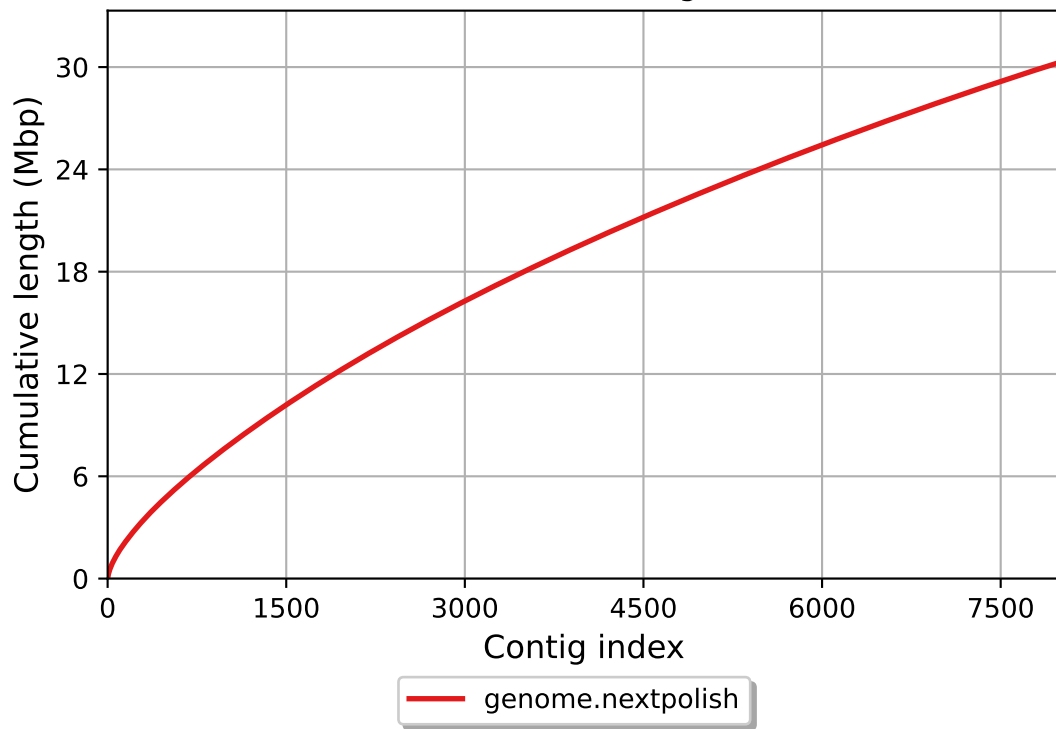
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

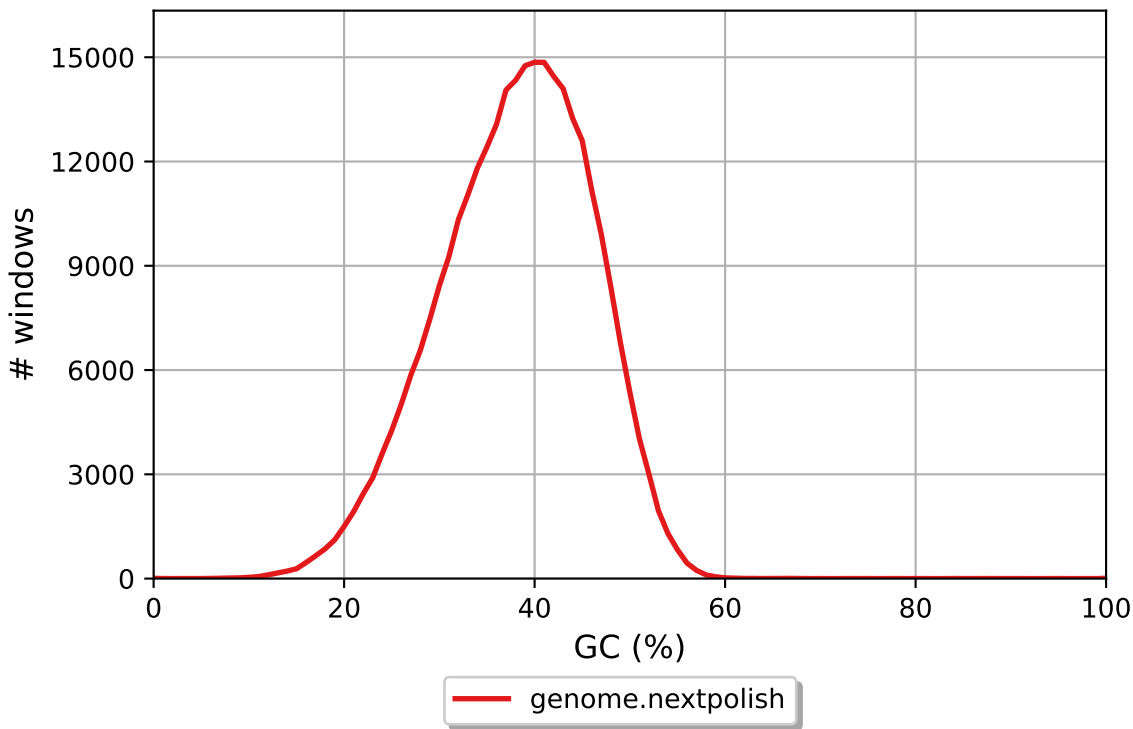


— genome.nextpolish

Cumulative length



# GC content



genome.nextpolish GC content

