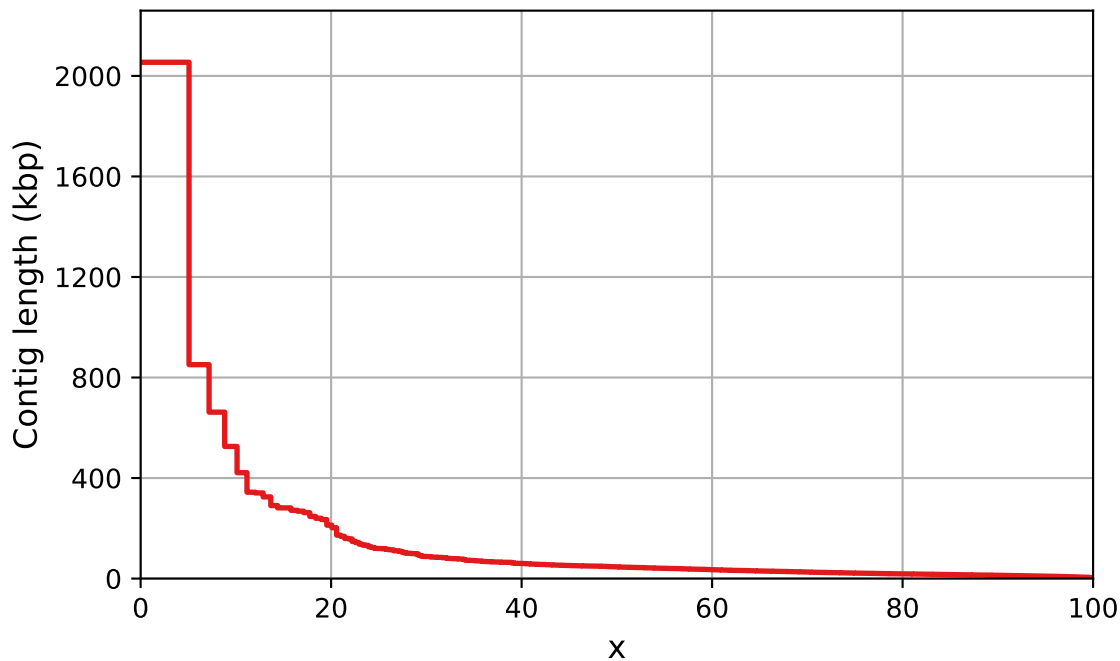


# Report

	genome.nextpolish
# contigs (>= 0 bp)	1342
# contigs (>= 1000 bp)	1338
# contigs (>= 5000 bp)	1286
# contigs (>= 10000 bp)	1018
# contigs (>= 25000 bp)	430
# contigs (>= 50000 bp)	161
Total length (>= 0 bp)	40426821
Total length (>= 1000 bp)	40424225
Total length (>= 5000 bp)	40243455
Total length (>= 10000 bp)	38227526
Total length (>= 25000 bp)	28637157
Total length (>= 50000 bp)	19094374
# contigs	1342
Largest contig	2054280
Total length	40426821
GC (%)	48.79
N50	46646
N90	13114
auN	206676.5
L50	185
L90	858
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	119 + 9 part

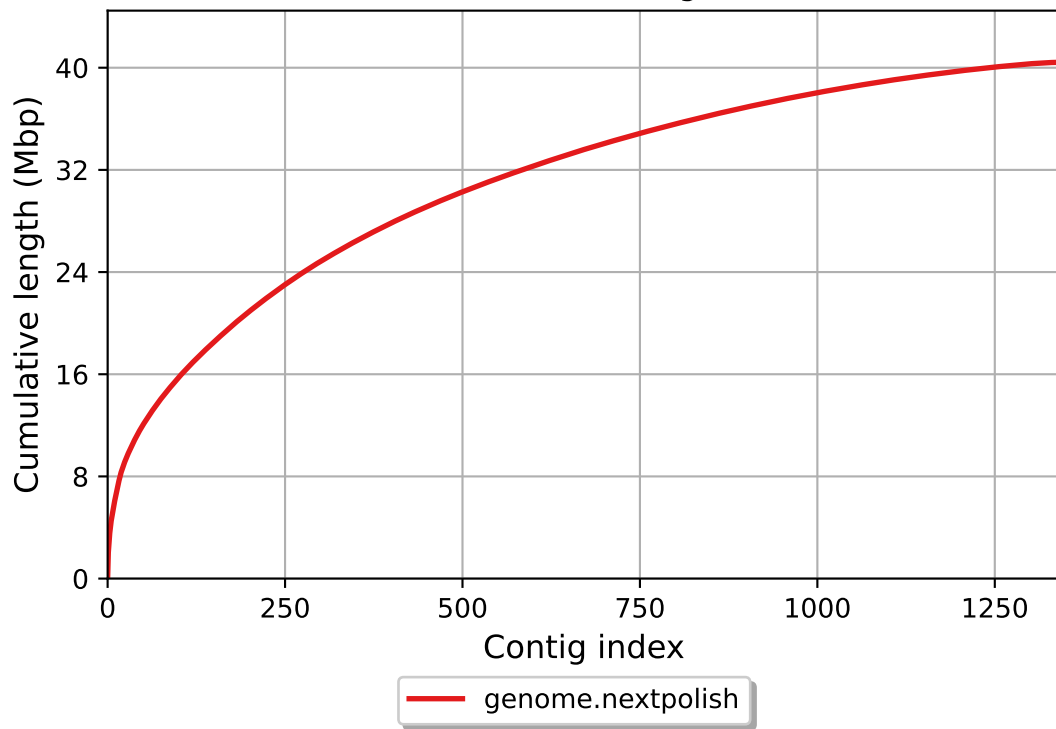
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

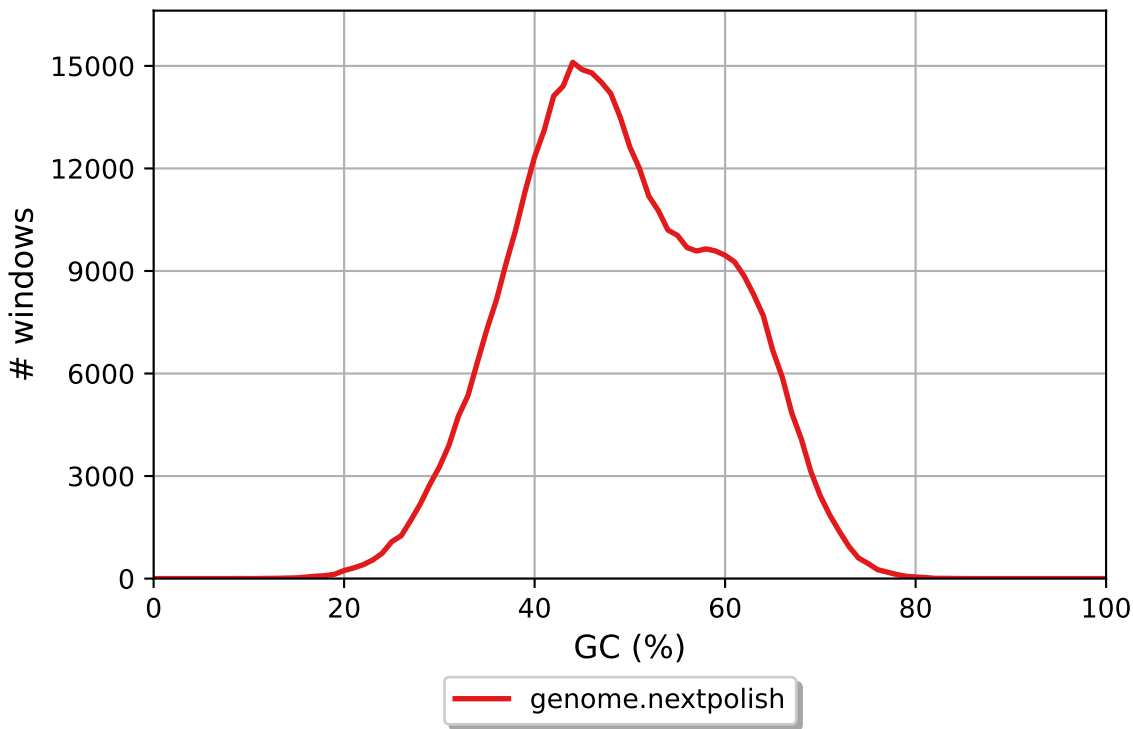


— genome.nextpolish

Cumulative length



## GC content



genome.nextpolish GC content

