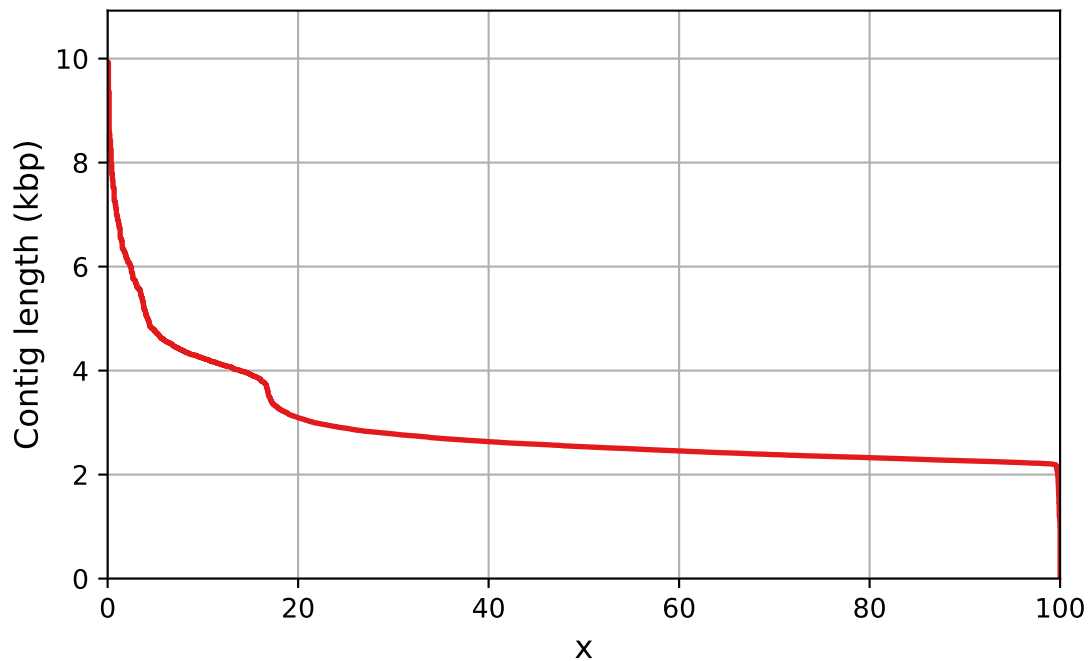


# Report

	genome.nextpolish
# contigs (>= 0 bp)	4494
# contigs (>= 1000 bp)	4494
# contigs (>= 5000 bp)	82
# contigs (>= 10000 bp)	0
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	12159865
Total length (>= 1000 bp)	12159865
Total length (>= 5000 bp)	511116
Total length (>= 10000 bp)	0
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	4494
Largest contig	9929
Total length	12159865
GC (%)	47.96
N50	2535
N90	2265
auN	2903.0
L50	1906
L90	3943
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	13 + 7 part

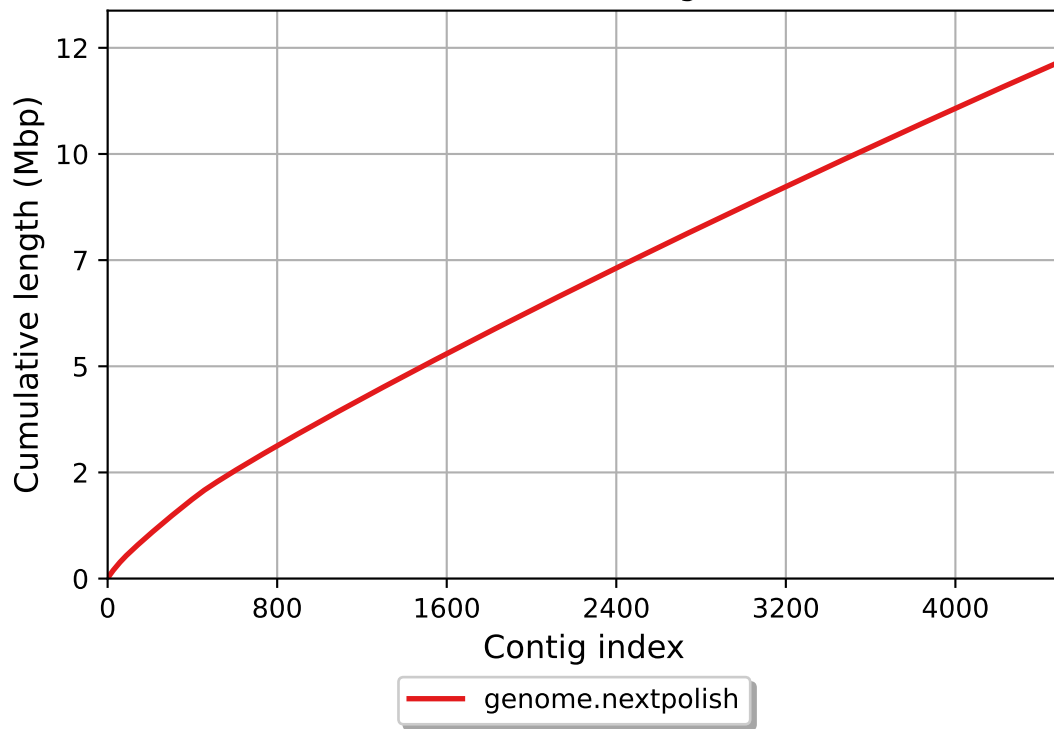
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

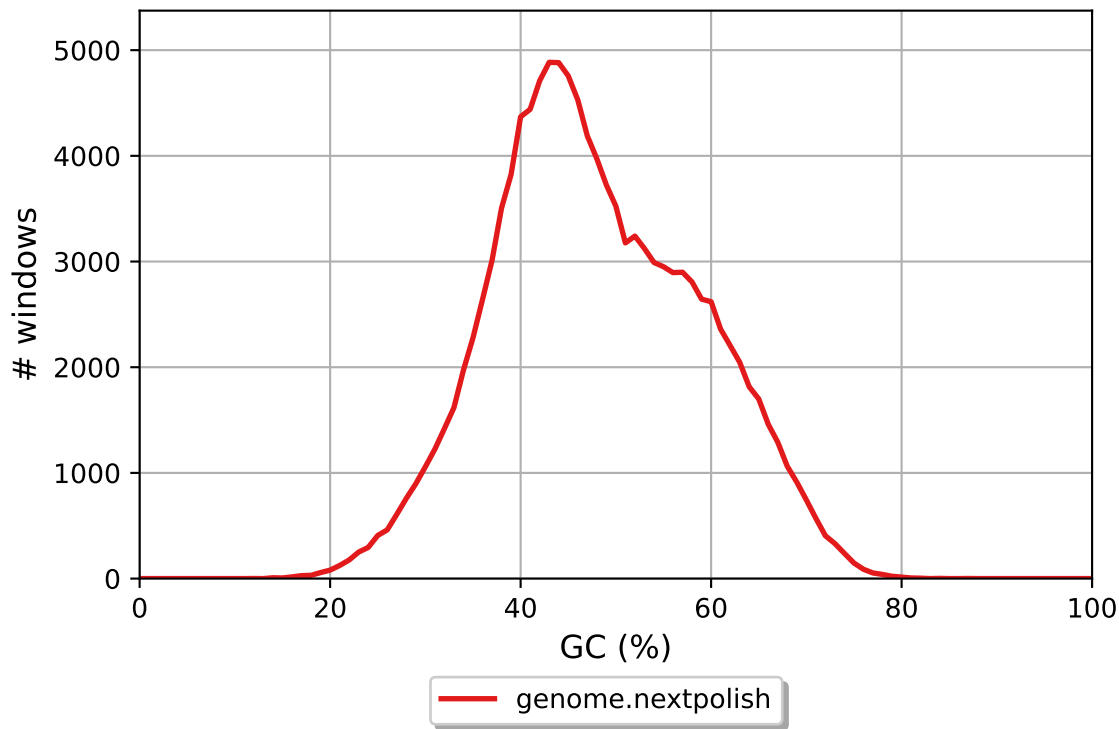


— genome.nextpolish

Cumulative length



## GC content



genome.nextpolish GC content

