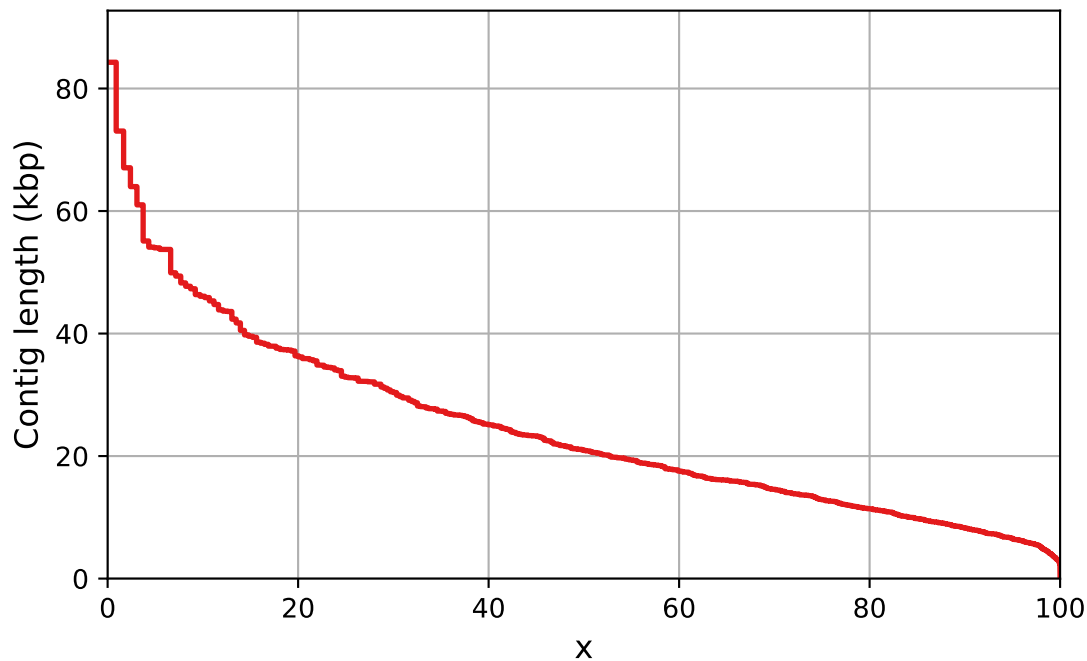


# Report

	genome.nextpolish
# contigs (>= 0 bp)	579
# contigs (>= 1000 bp)	579
# contigs (>= 5000 bp)	532
# contigs (>= 10000 bp)	360
# contigs (>= 25000 bp)	104
# contigs (>= 50000 bp)	10
Total length (>= 0 bp)	9370212
Total length (>= 1000 bp)	9370212
Total length (>= 5000 bp)	9192740
Total length (>= 10000 bp)	7893544
Total length (>= 25000 bp)	3794405
Total length (>= 50000 bp)	620056
# contigs	579
Largest contig	84283
Total length	9370212
GC (%)	47.78
N50	20995
N90	8254
auN	24716.1
L50	143
L90	420
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	23 + 5 part

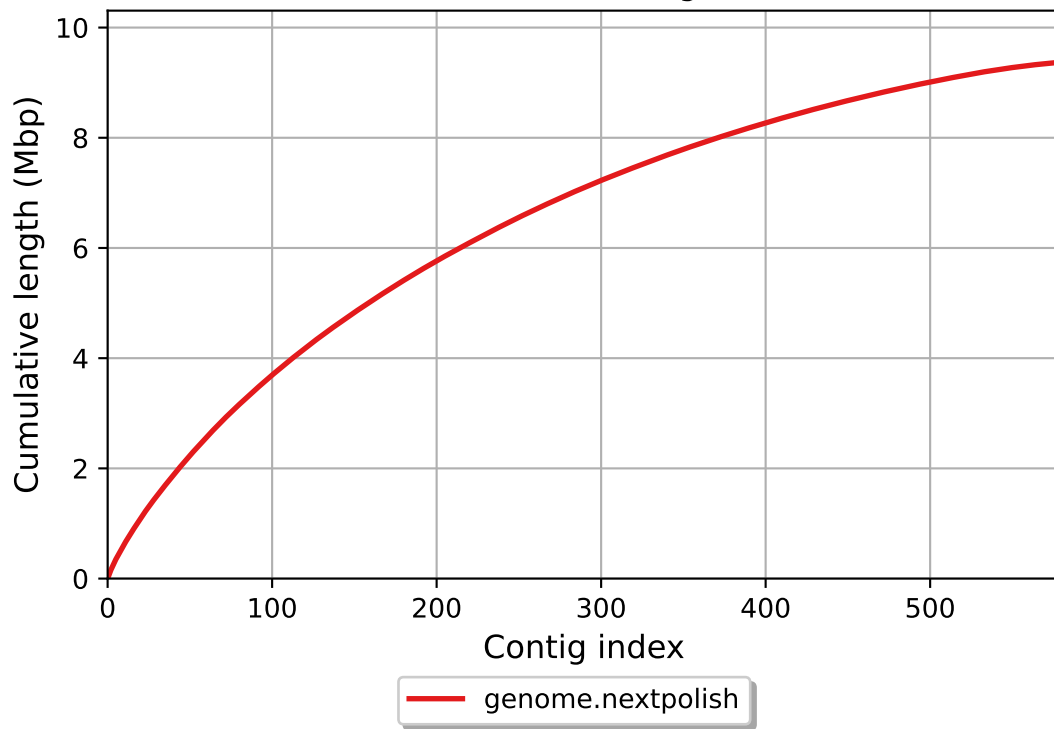
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

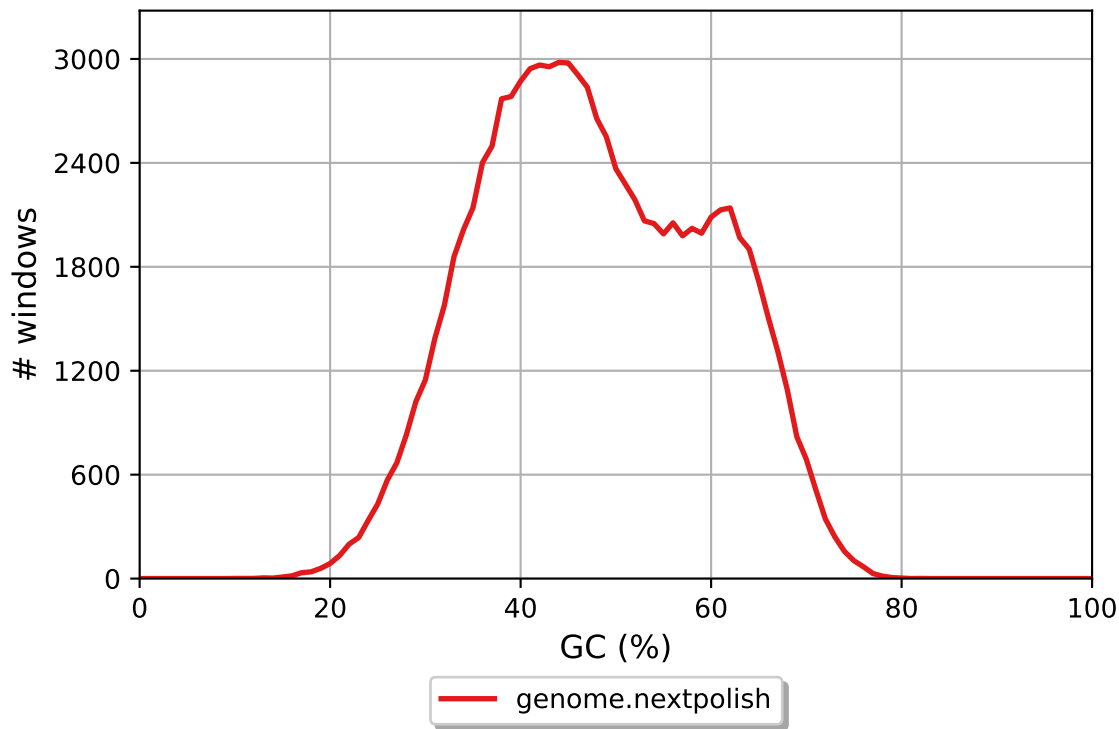


— genome.nextpolish

Cumulative length



## GC content



genome.nextpolish GC content

