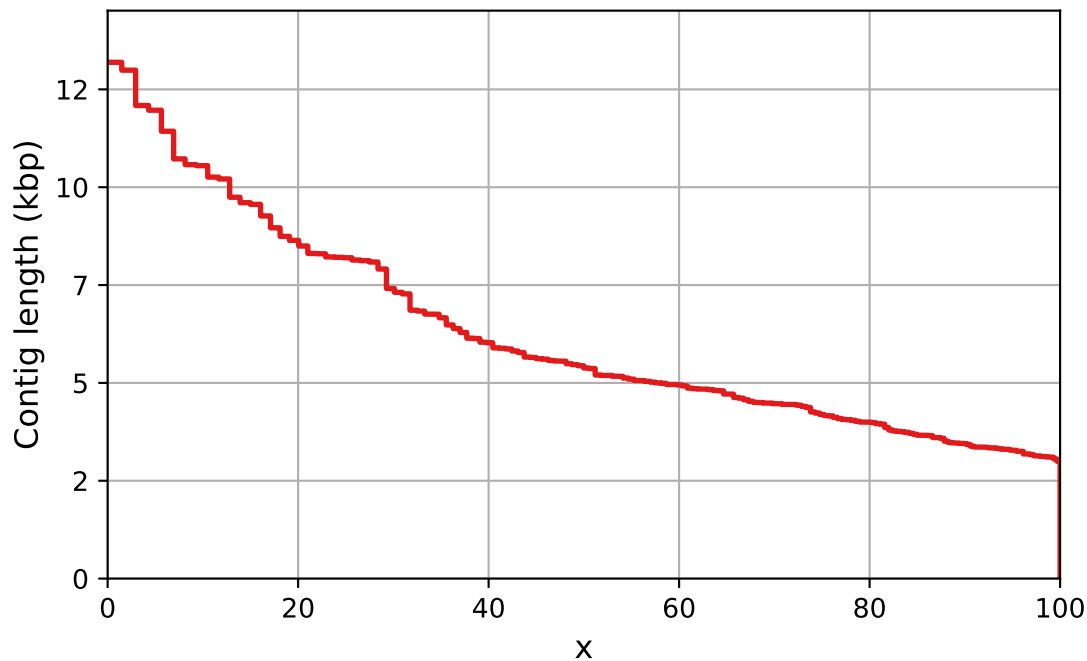


# Report

	genome.nextpolish
# contigs (>= 0 bp)	166
# contigs (>= 1000 bp)	166
# contigs (>= 5000 bp)	71
# contigs (>= 10000 bp)	10
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	889849
Total length (>= 1000 bp)	889849
Total length (>= 5000 bp)	516816
Total length (>= 10000 bp)	113990
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	166
Largest contig	13192
Total length	889849
GC (%)	50.79
N50	5380
N90	3450
auN	6281.6
L50	58
L90	139
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	2 + 0 part

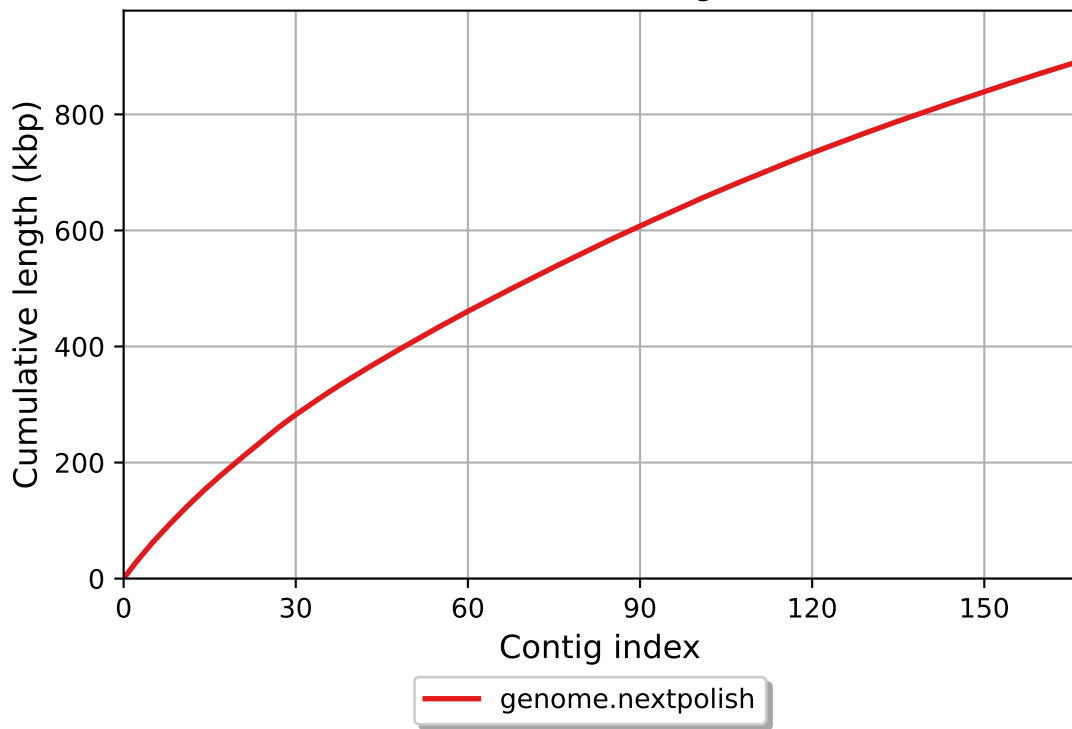
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

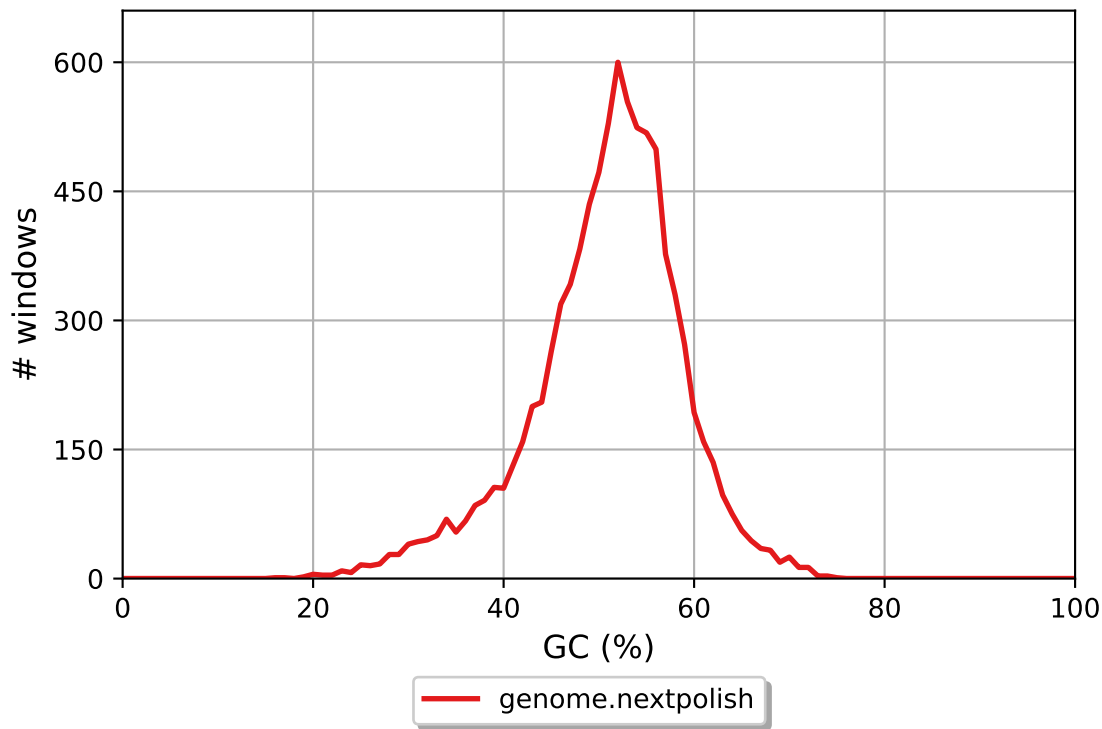


— genome.nextpolish

Cumulative length



## GC content



genome.nextpolish GC content

