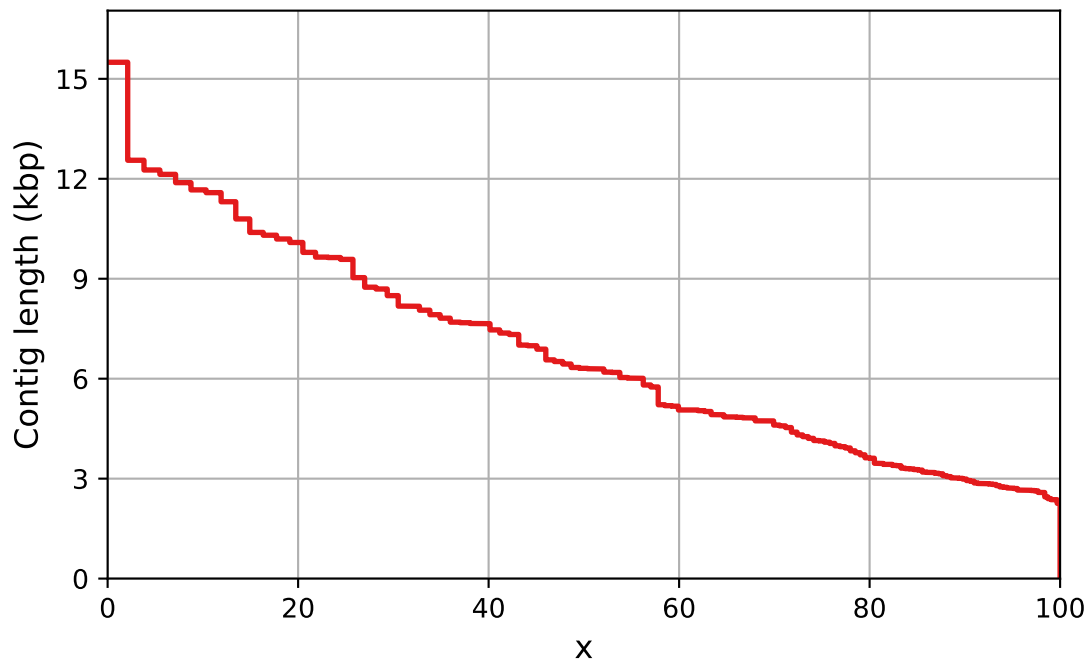


# Report

	genome.nextpolish
# contigs (>= 0 bp)	136
# contigs (>= 1000 bp)	136
# contigs (>= 5000 bp)	58
# contigs (>= 10000 bp)	13
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	735065
Total length (>= 1000 bp)	735065
Total length (>= 5000 bp)	465748
Total length (>= 10000 bp)	150668
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	136
Largest contig	15497
Total length	735065
GC (%)	42.89
N50	6311
N90	2987
auN	6858.3
L50	41
L90	109
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	8 + 1 part

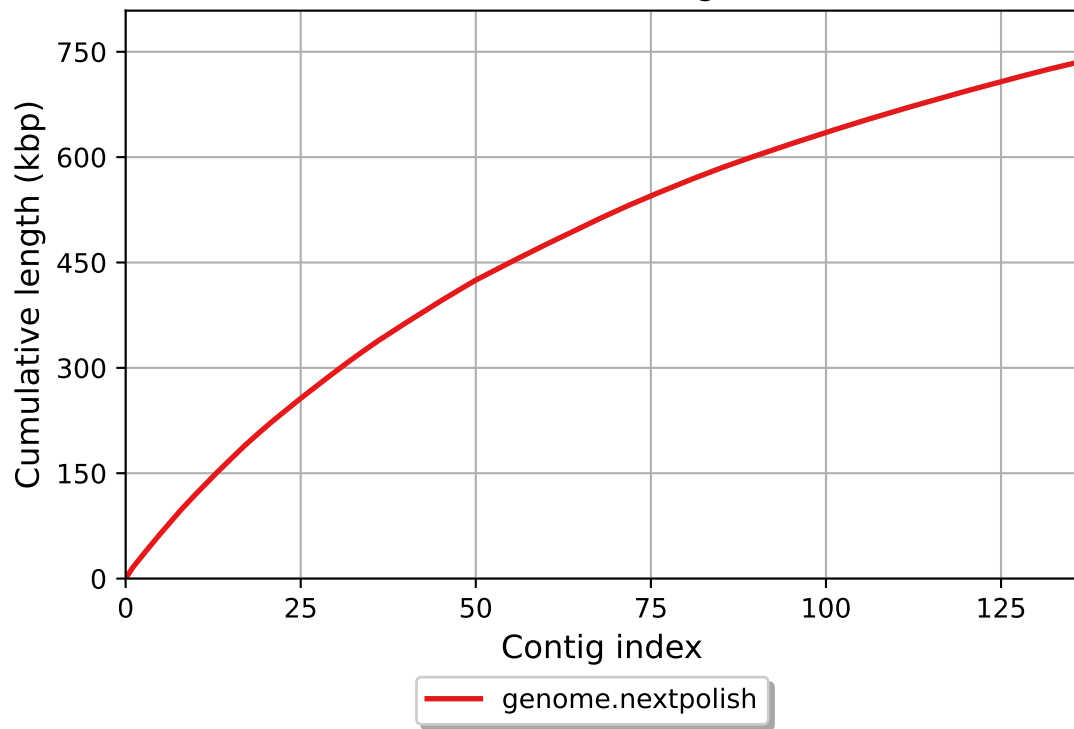
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

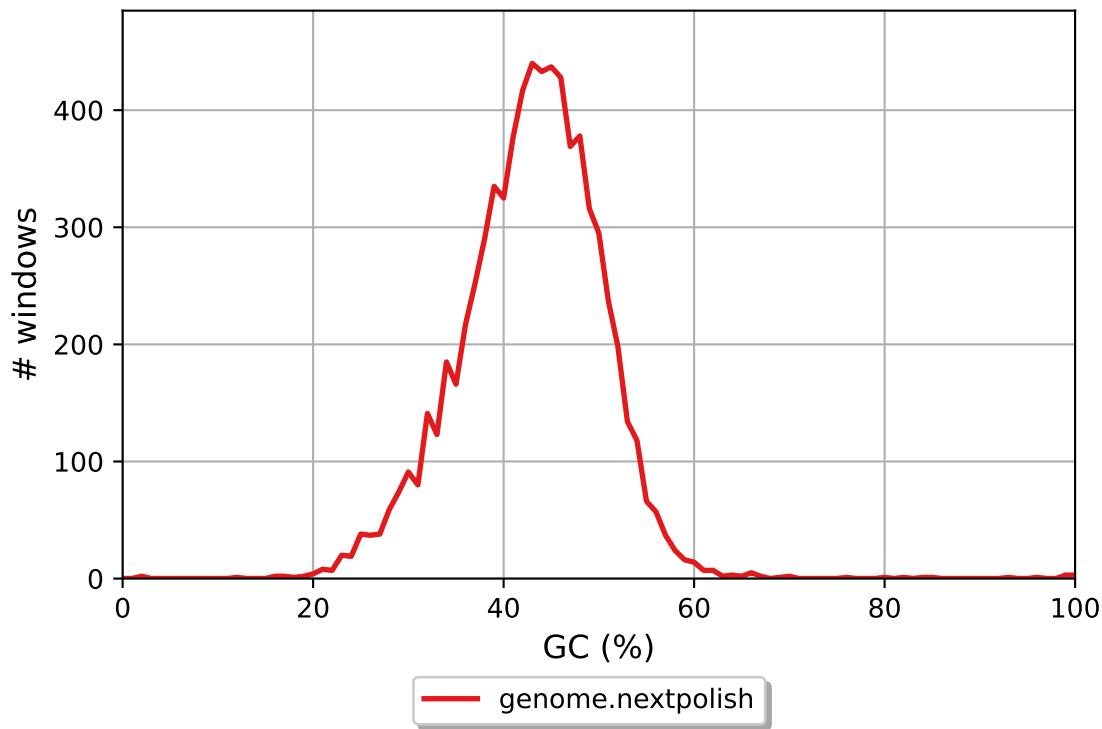


— genome.nextpolish

Cumulative length



## GC content



genome.nextpolish GC content

