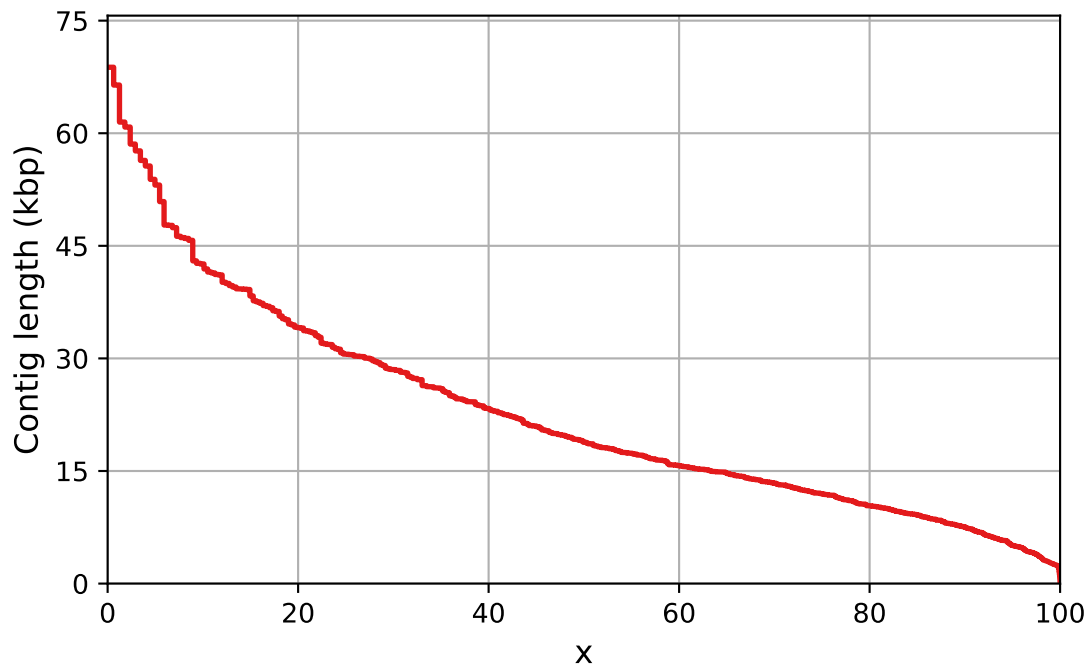


# Report

	genome.nextpolish
# contigs (>= 0 bp)	776
# contigs (>= 1000 bp)	769
# contigs (>= 5000 bp)	619
# contigs (>= 10000 bp)	427
# contigs (>= 25000 bp)	109
# contigs (>= 50000 bp)	11
Total length (>= 0 bp)	10861120
Total length (>= 1000 bp)	10856198
Total length (>= 5000 bp)	10340652
Total length (>= 10000 bp)	8887506
Total length (>= 25000 bp)	3923251
Total length (>= 50000 bp)	643573
# contigs	776
Largest contig	68788
Total length	10861120
GC (%)	48.98
N50	18827
N90	7557
auN	22766.5
L50	179
L90	529
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	22 + 5 part

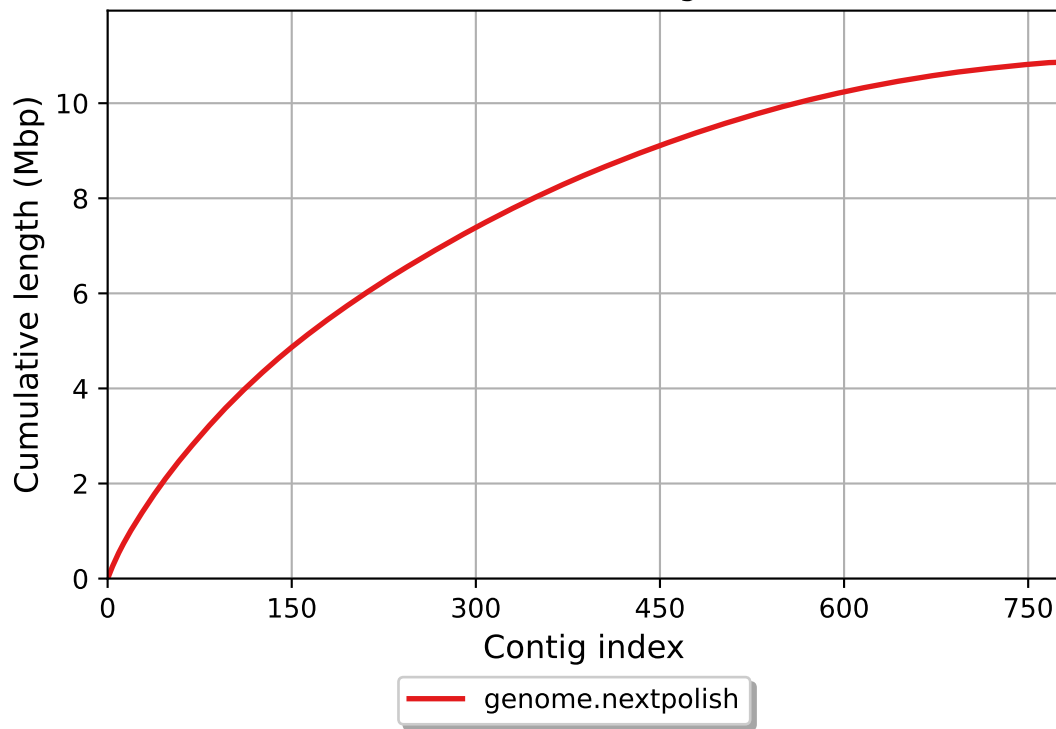
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

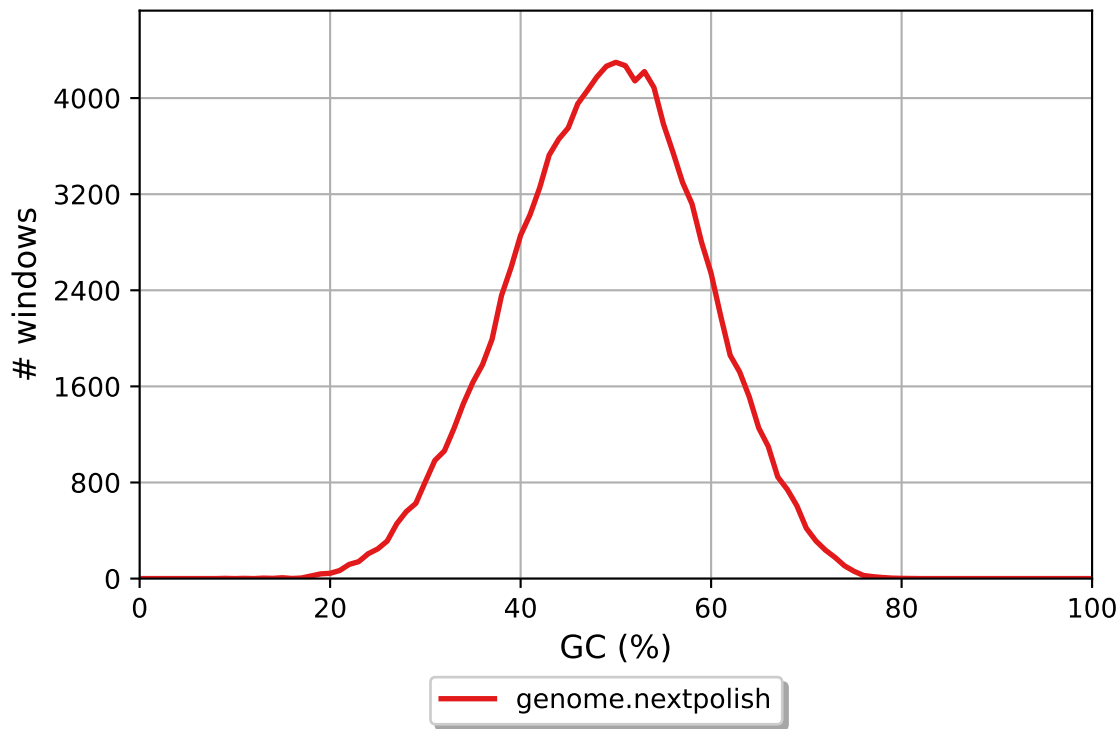


— genome.nextpolish

Cumulative length



## GC content



genome.nextpolish GC content

