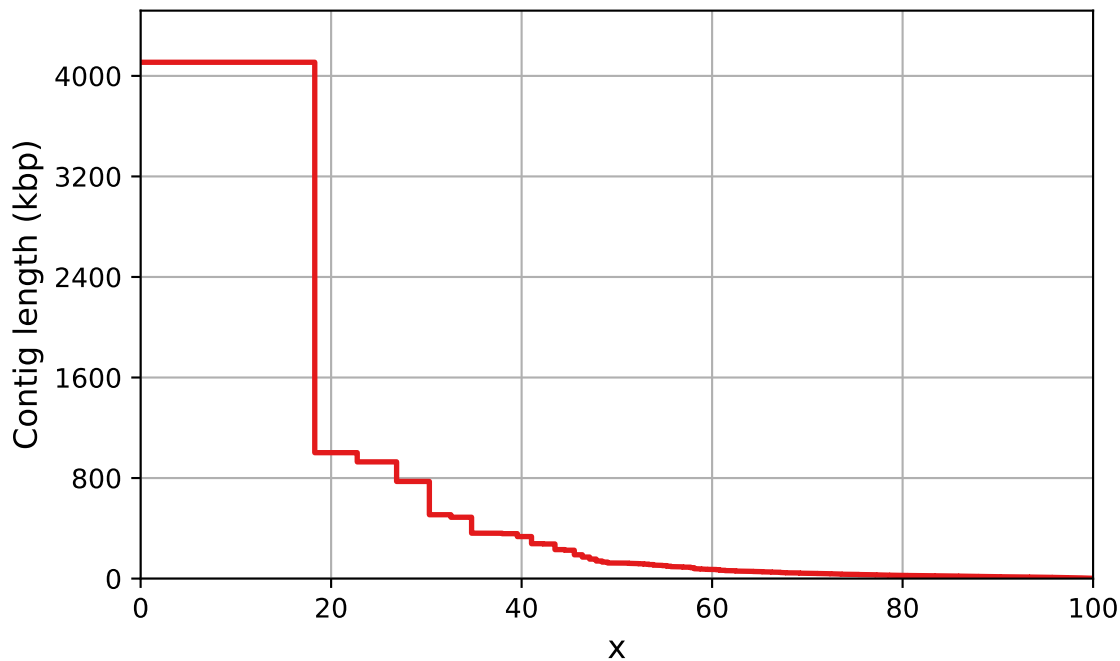


# Report

	genome.nextpolish
# contigs (>= 0 bp)	549
# contigs (>= 1000 bp)	545
# contigs (>= 5000 bp)	474
# contigs (>= 10000 bp)	358
# contigs (>= 25000 bp)	154
# contigs (>= 50000 bp)	71
Total length (>= 0 bp)	22471217
Total length (>= 1000 bp)	22467829
Total length (>= 5000 bp)	22213647
Total length (>= 10000 bp)	21355525
Total length (>= 25000 bp)	17966607
Total length (>= 50000 bp)	15089295
# contigs	549
Largest contig	4108414
Total length	22471217
GC (%)	56.82
N50	123638
N90	14845
auN	945379.8
L50	21
L90	269
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	63 + 2 part

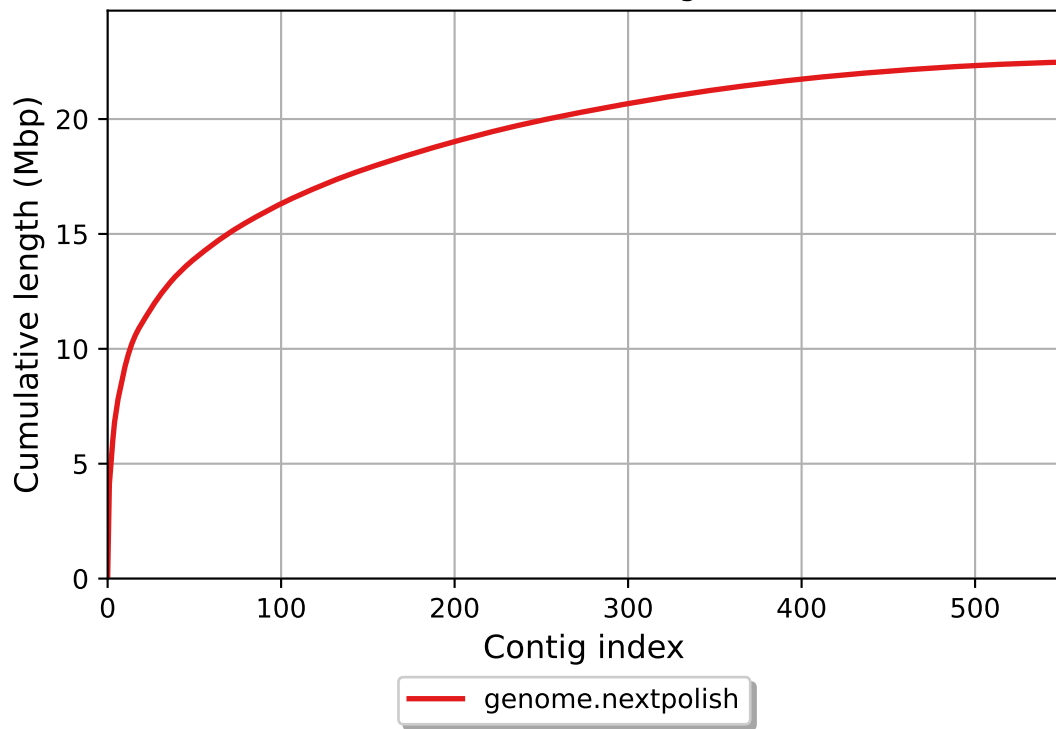
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

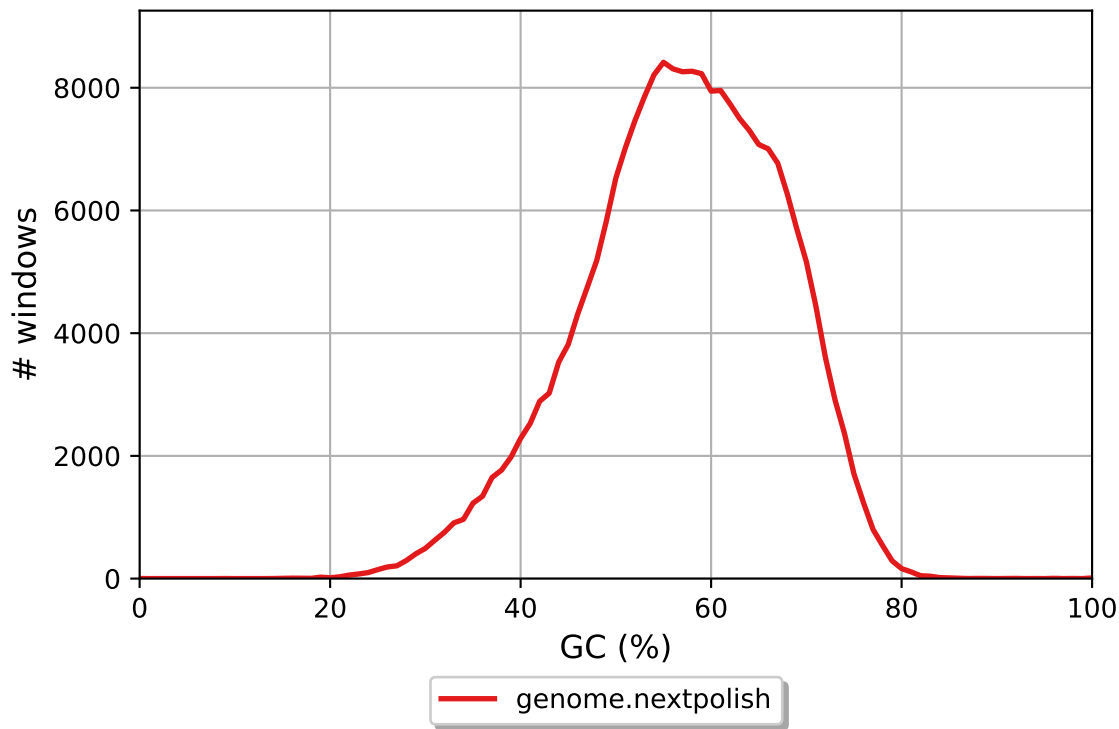


— genome.nextpolish

Cumulative length



## GC content



genome.nextpolish GC content

