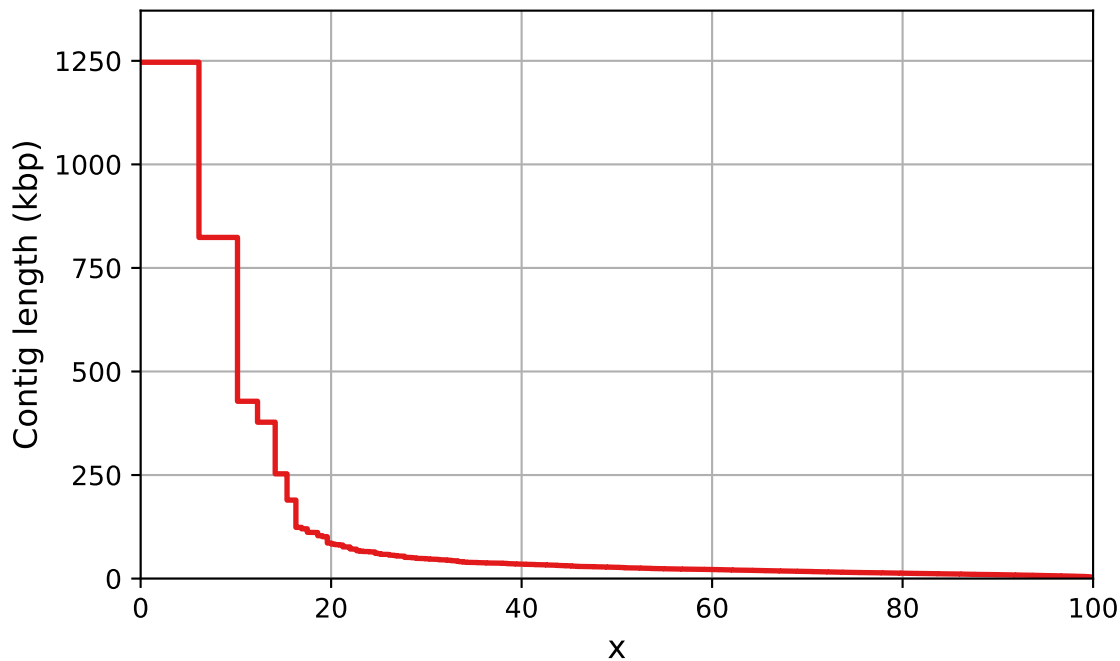


# Report

	genome.nextpolish
# contigs (>= 0 bp)	977
# contigs (>= 1000 bp)	977
# contigs (>= 5000 bp)	935
# contigs (>= 10000 bp)	633
# contigs (>= 25000 bp)	186
# contigs (>= 50000 bp)	41
Total length (>= 0 bp)	20354014
Total length (>= 1000 bp)	20354014
Total length (>= 5000 bp)	20181603
Total length (>= 10000 bp)	17912583
Total length (>= 25000 bp)	10751981
Total length (>= 50000 bp)	5841796
# contigs	977
Largest contig	1246541
Total length	20354014
GC (%)	51.60
N50	27107
N90	9409
auN	155673.7
L50	164
L90	675
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	68 + 5 part

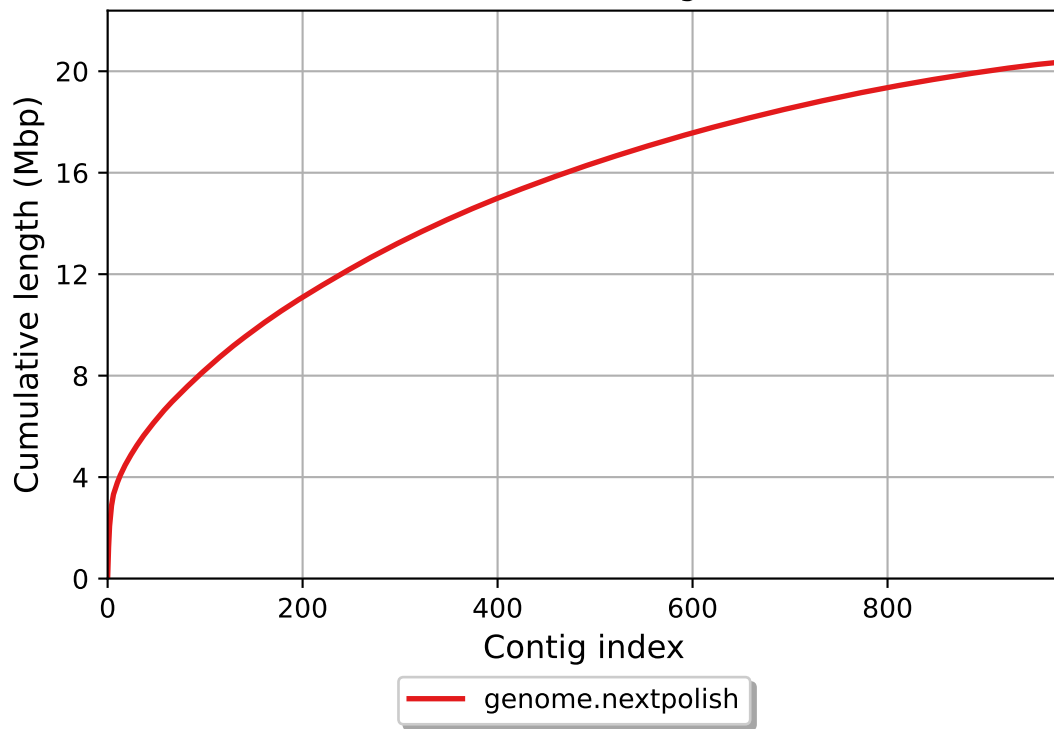
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

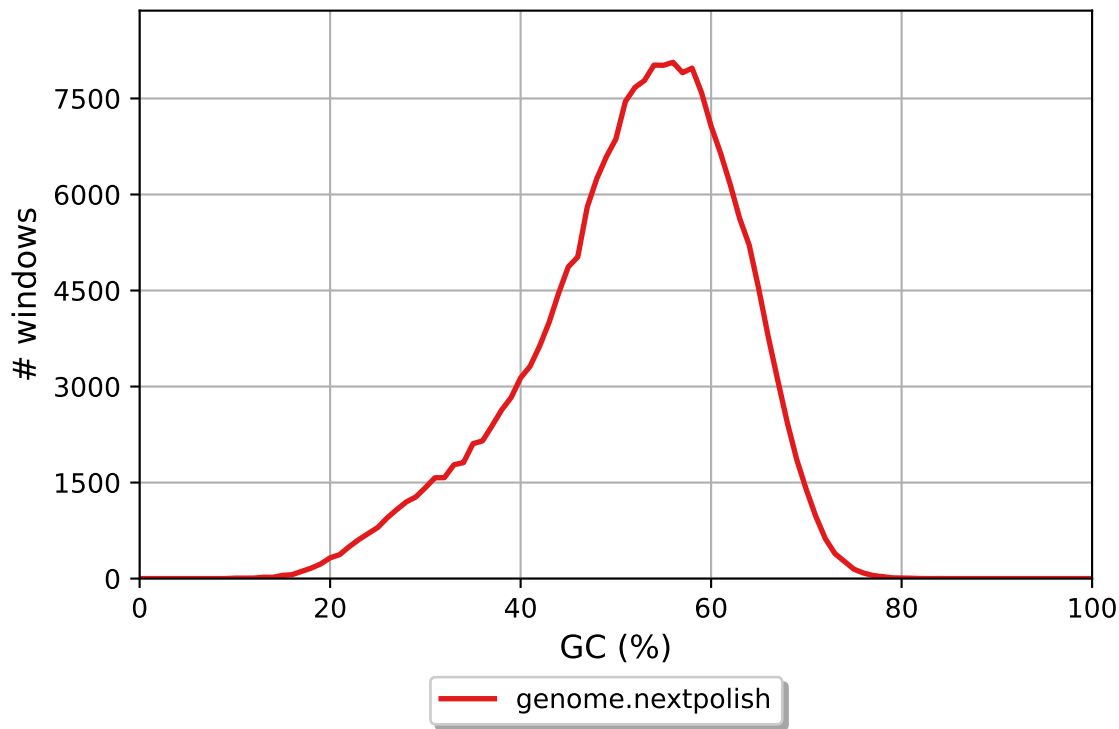


genome.nextpolish

Cumulative length



## GC content



genome.nextpolish GC content

