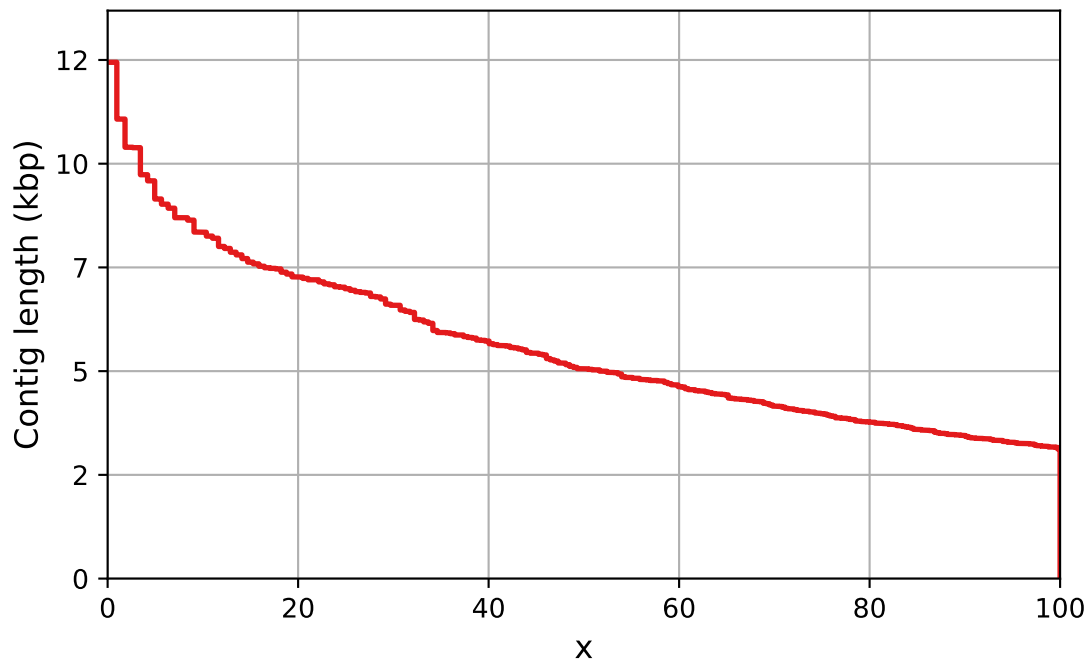


# Report

	genome.nextpolish
# contigs (>= 0 bp)	257
# contigs (>= 1000 bp)	257
# contigs (>= 5000 bp)	99
# contigs (>= 10000 bp)	4
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1287499
Total length (>= 1000 bp)	1287499
Total length (>= 5000 bp)	669959
Total length (>= 10000 bp)	44301
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	257
Largest contig	12444
Total length	1287499
GC (%)	52.32
N50	5059
N90	3448
auN	5599.2
L50	94
L90	218
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	4 + 0 part

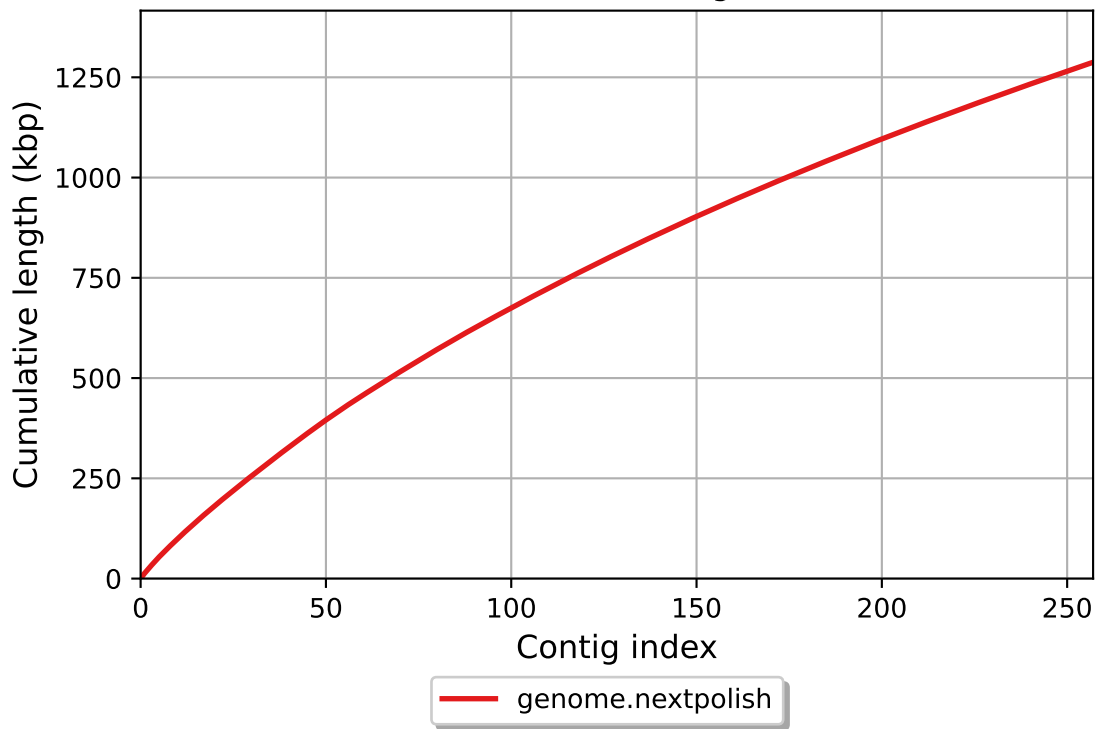
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

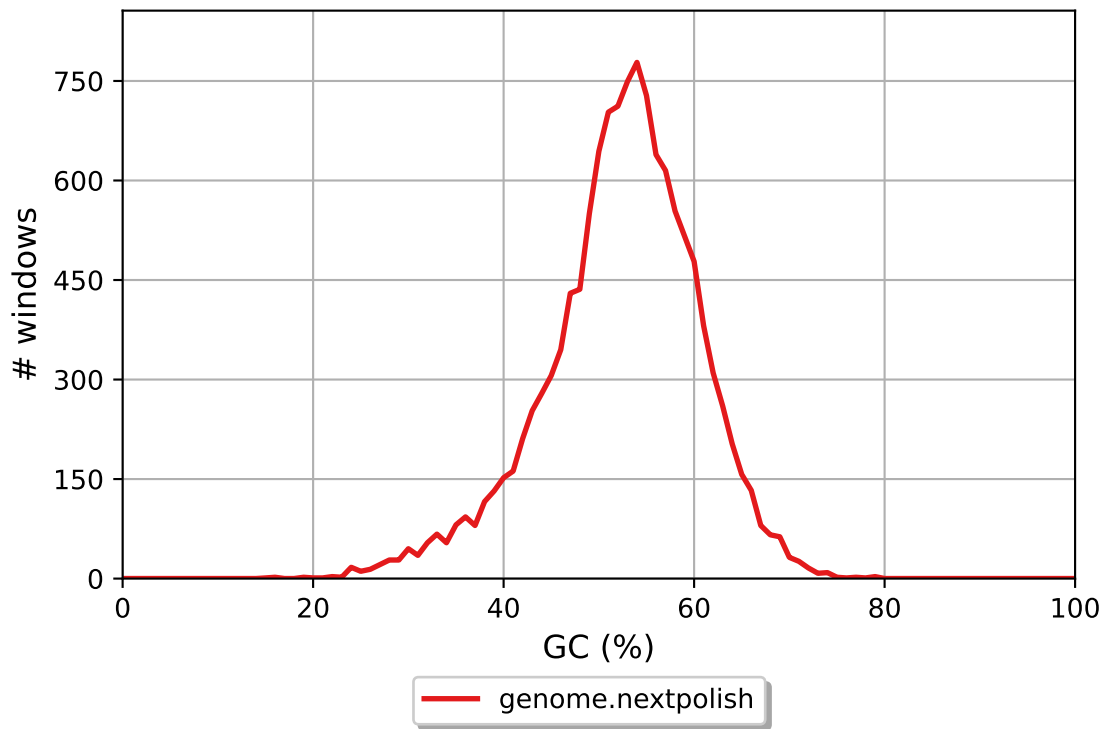


— genome.nextpolish

Cumulative length



## GC content



genome.nextpolish GC content

