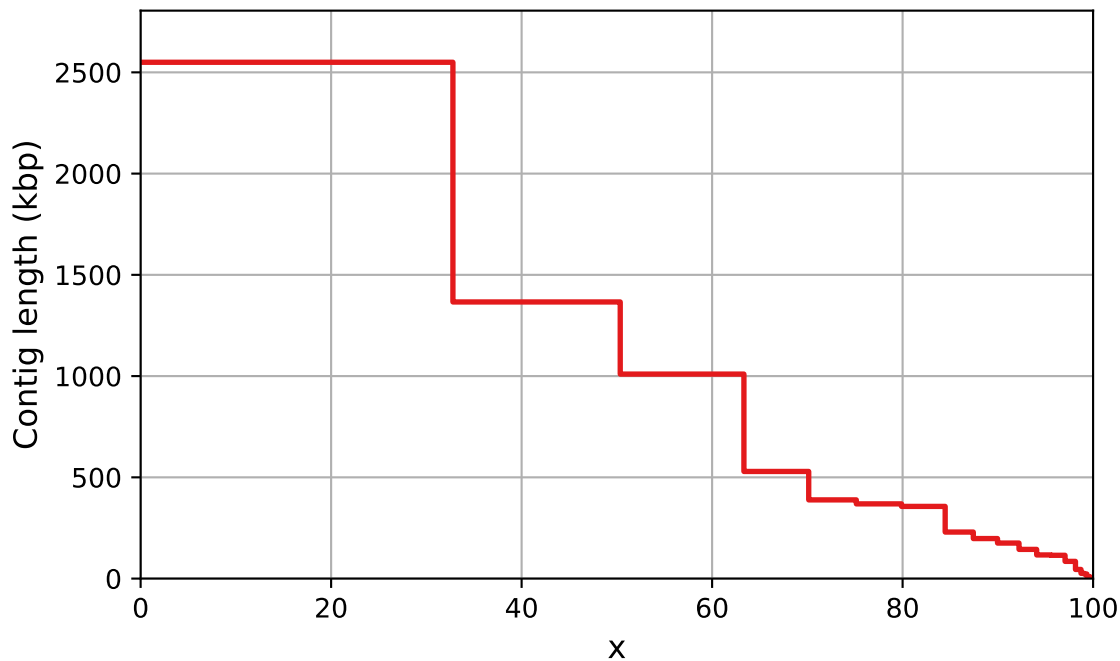


# Report

	genome.nextpolish
# contigs (>= 0 bp)	28
# contigs (>= 1000 bp)	26
# contigs (>= 5000 bp)	22
# contigs (>= 10000 bp)	17
# contigs (>= 25000 bp)	16
# contigs (>= 50000 bp)	14
Total length (>= 0 bp)	7776764
Total length (>= 1000 bp)	7775451
Total length (>= 5000 bp)	7762793
Total length (>= 10000 bp)	7726342
Total length (>= 25000 bp)	7703874
Total length (>= 50000 bp)	7632850
# contigs	28
Largest contig	2549795
Total length	7776764
GC (%)	53.68
N50	1366118
N90	175423
auN	1319681.5
L50	2
L90	10
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	31 + 1 part

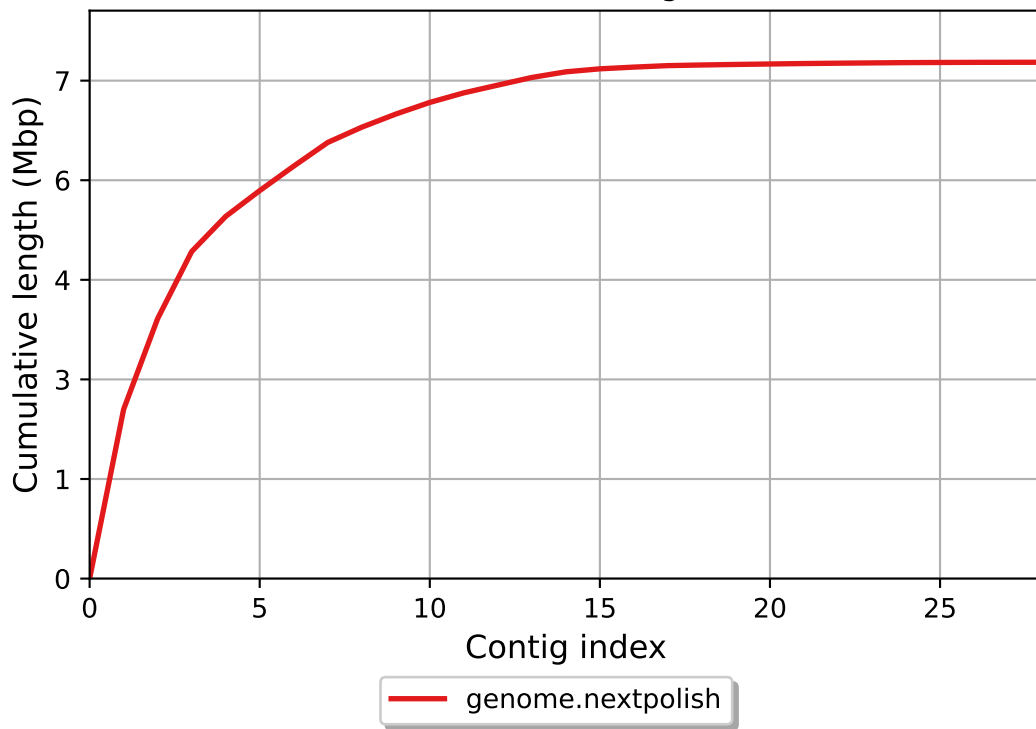
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

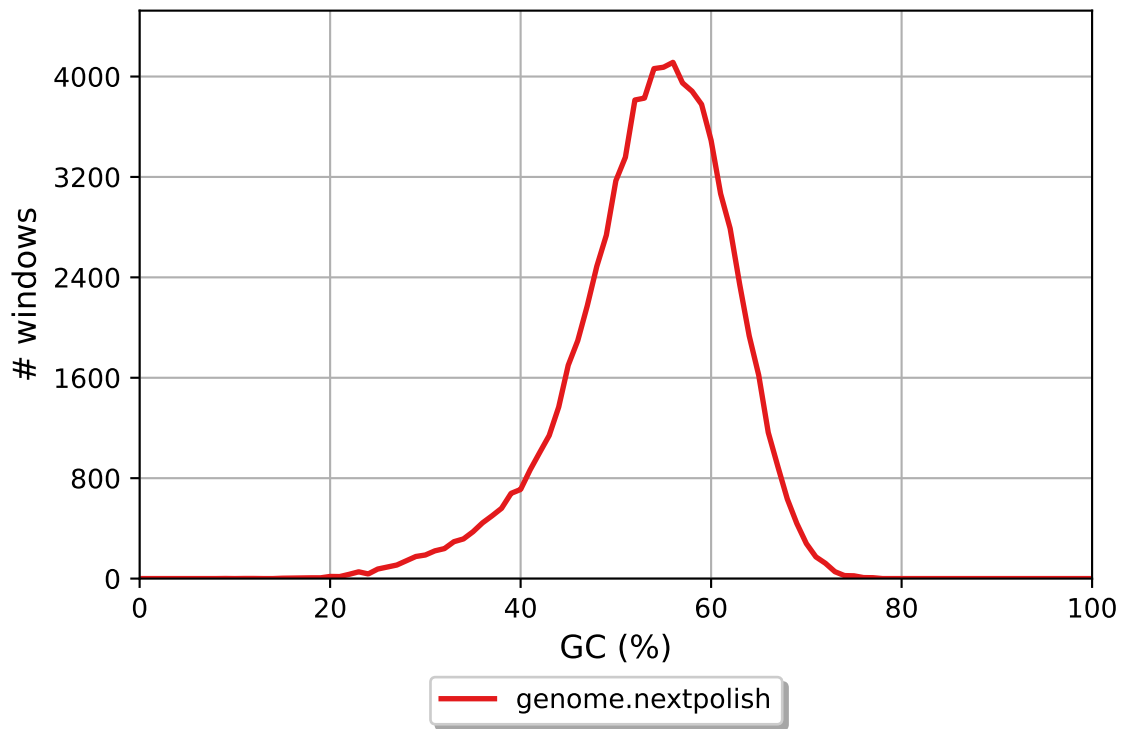


— genome.nextpolish

Cumulative length



## GC content



genome.nextpolish GC content

