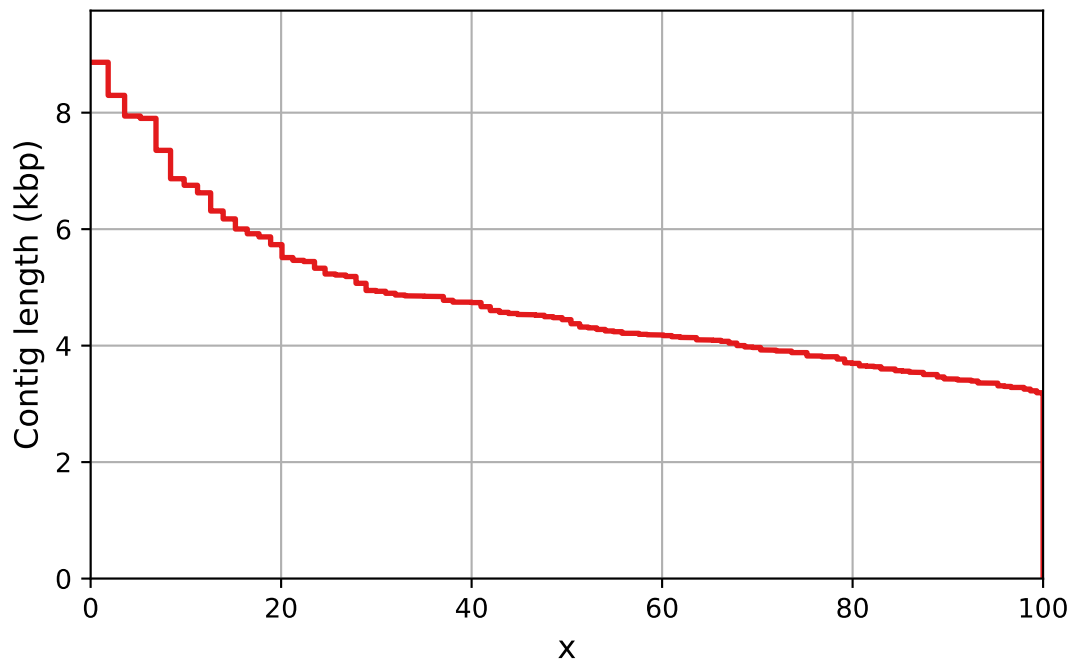


Report

	genome.nextpolish
# contigs (>= 0 bp)	107
# contigs (>= 1000 bp)	107
# contigs (>= 5000 bp)	22
# contigs (>= 10000 bp)	0
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	480872
Total length (>= 1000 bp)	480872
Total length (>= 5000 bp)	139063
Total length (>= 10000 bp)	0
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	107
Largest contig	8865
Total length	480872
GC (%)	59.10
N50	4446
N90	3429
auN	4786.4
L50	44
L90	93
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	1 + 1 part

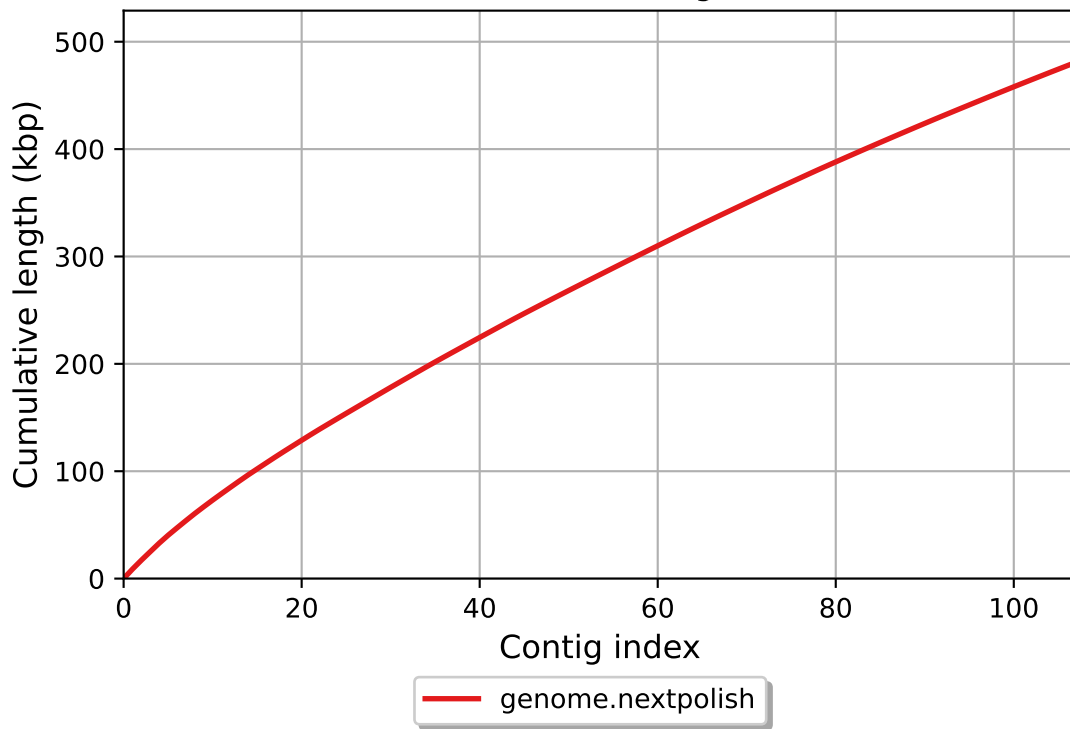
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

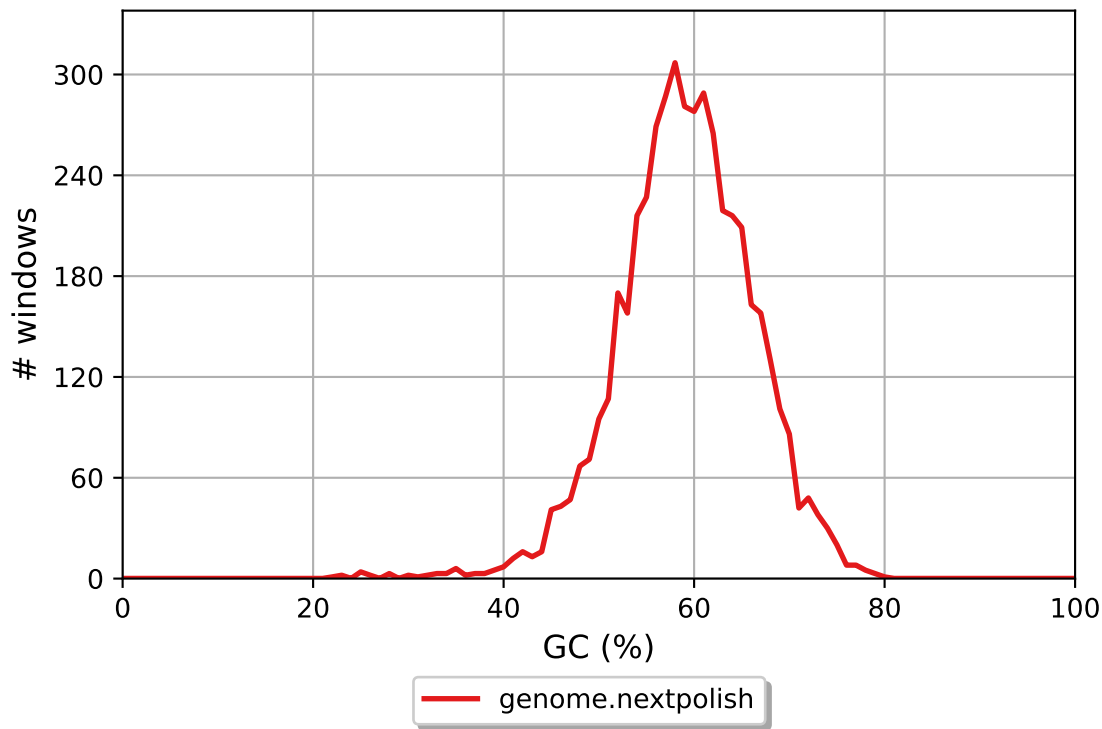


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

