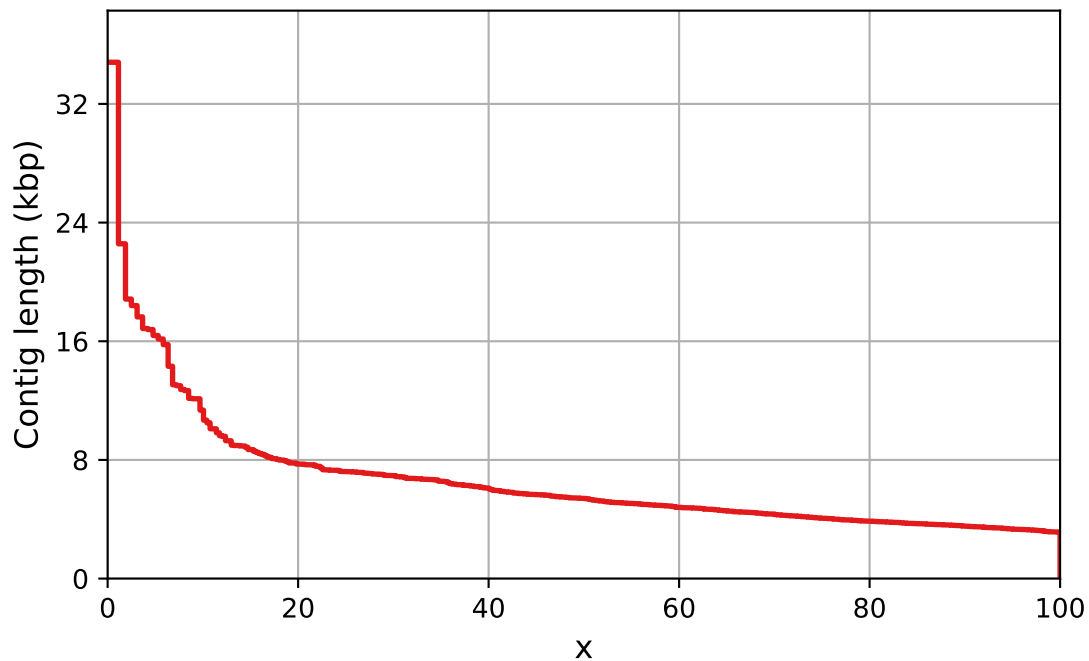


Report

	genome.nextpolish
# contigs (>= 0 bp)	573
# contigs (>= 1000 bp)	573
# contigs (>= 5000 bp)	235
# contigs (>= 10000 bp)	23
# contigs (>= 25000 bp)	1
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	3057549
Total length (>= 1000 bp)	3057549
Total length (>= 5000 bp)	1726403
Total length (>= 10000 bp)	349166
Total length (>= 25000 bp)	34807
Total length (>= 50000 bp)	0
# contigs	573
Largest contig	34807
Total length	3057549
GC (%)	59.30
N50	5411
N90	3536
auN	6703.0
L50	197
L90	482
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	4 + 1 part

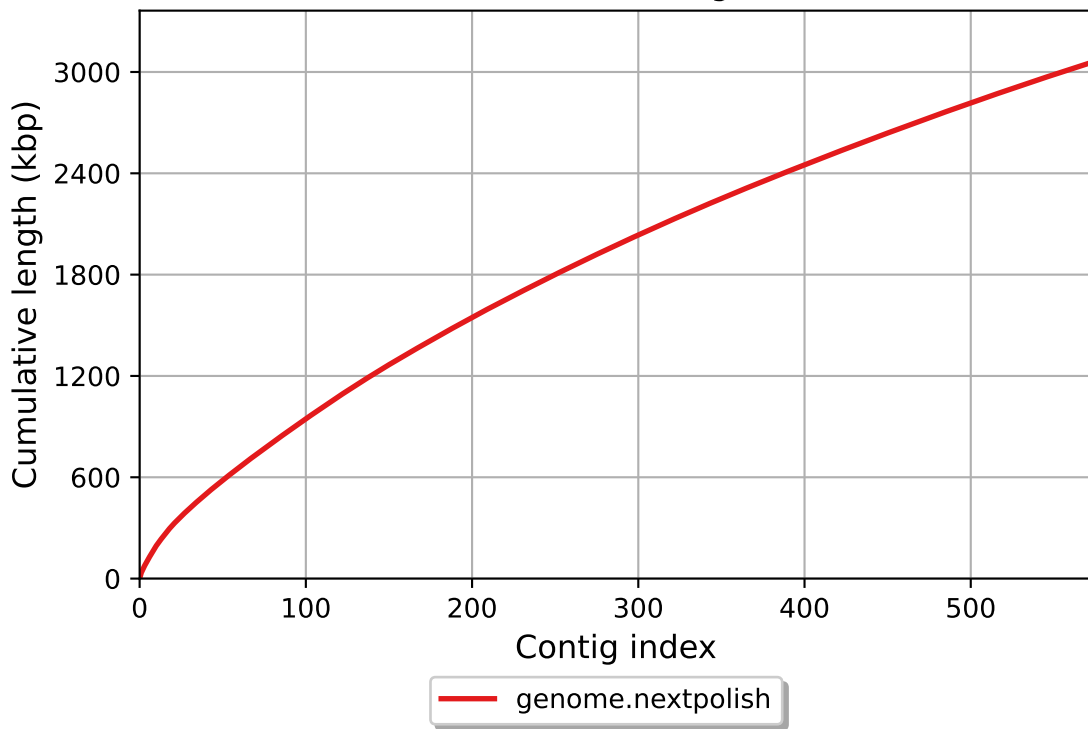
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

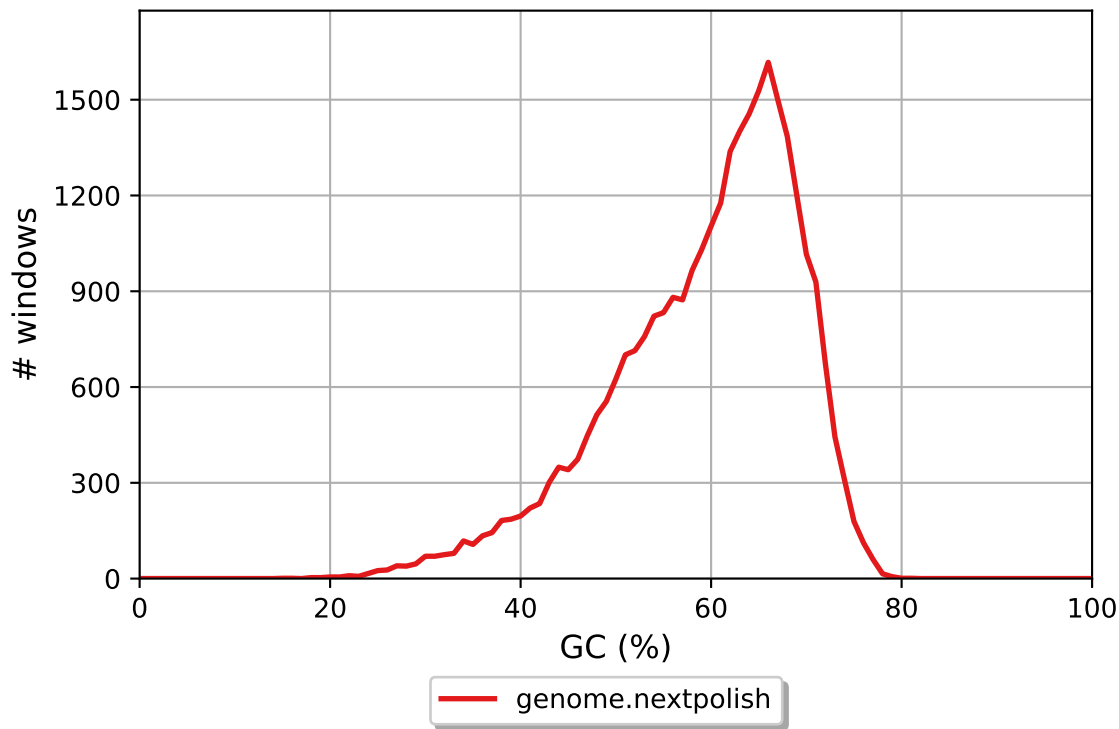


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

