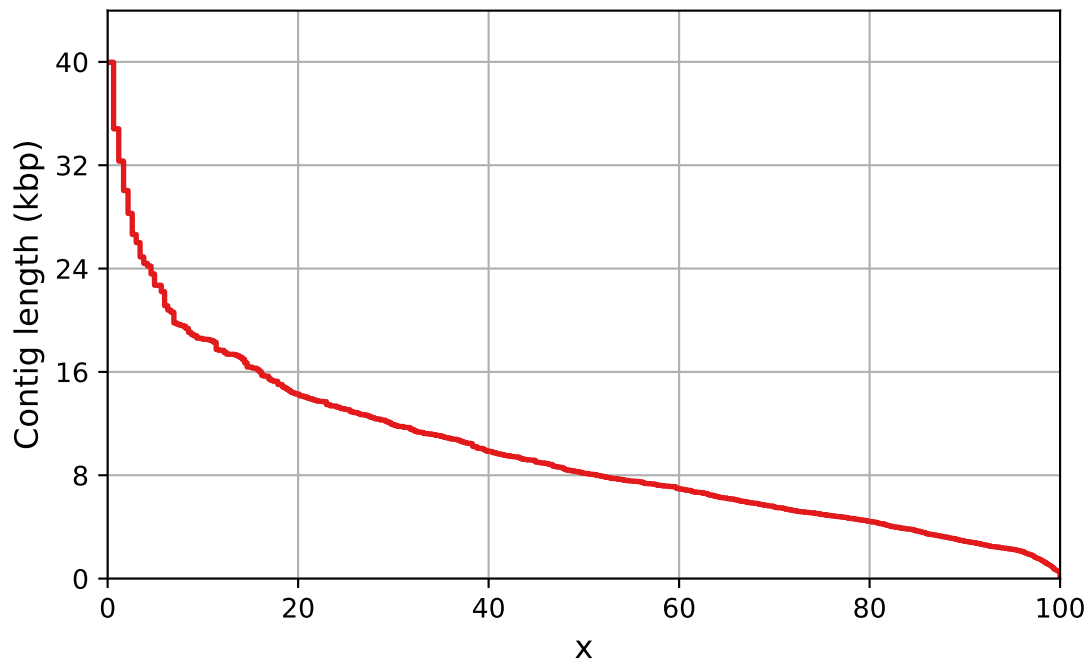


Report

	genome.nextpolish
# contigs (>= 0 bp)	1115
# contigs (>= 1000 bp)	1020
# contigs (>= 5000 bp)	490
# contigs (>= 10000 bp)	171
# contigs (>= 25000 bp)	7
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	6401314
Total length (>= 1000 bp)	6337386
Total length (>= 5000 bp)	4781468
Total length (>= 10000 bp)	2532491
Total length (>= 25000 bp)	218060
Total length (>= 50000 bp)	0
# contigs	1111
Largest contig	39969
Total length	6399809
GC (%)	54.10
N50	8153
N90	2890
auN	9884.3
L50	245
L90	739
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	12 + 3 part

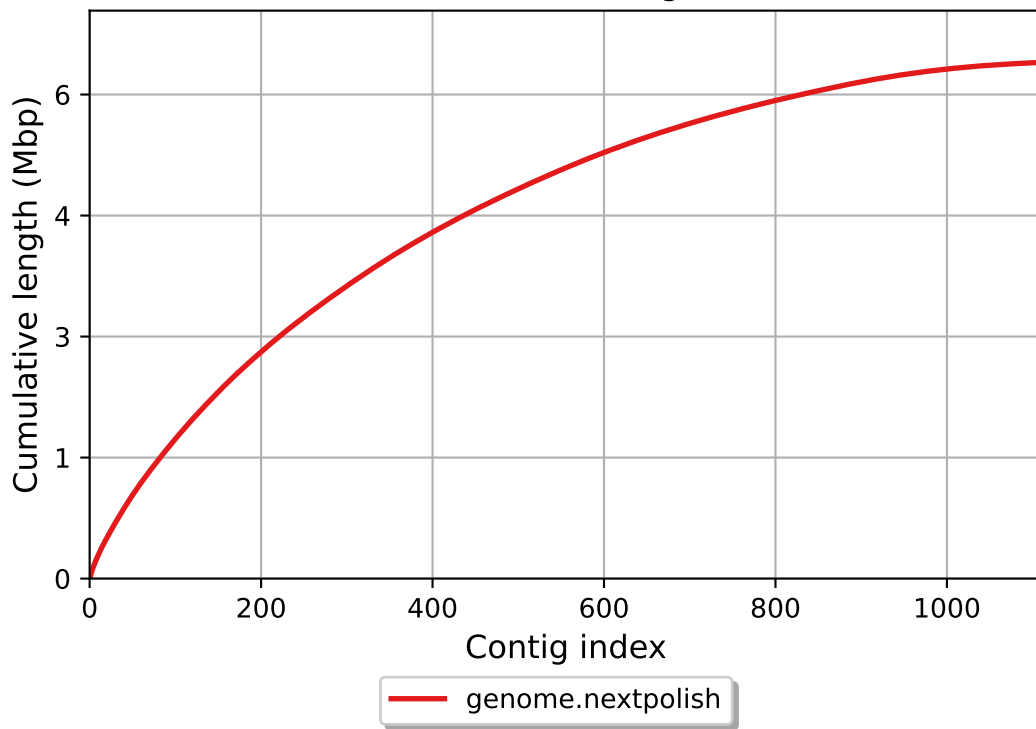
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

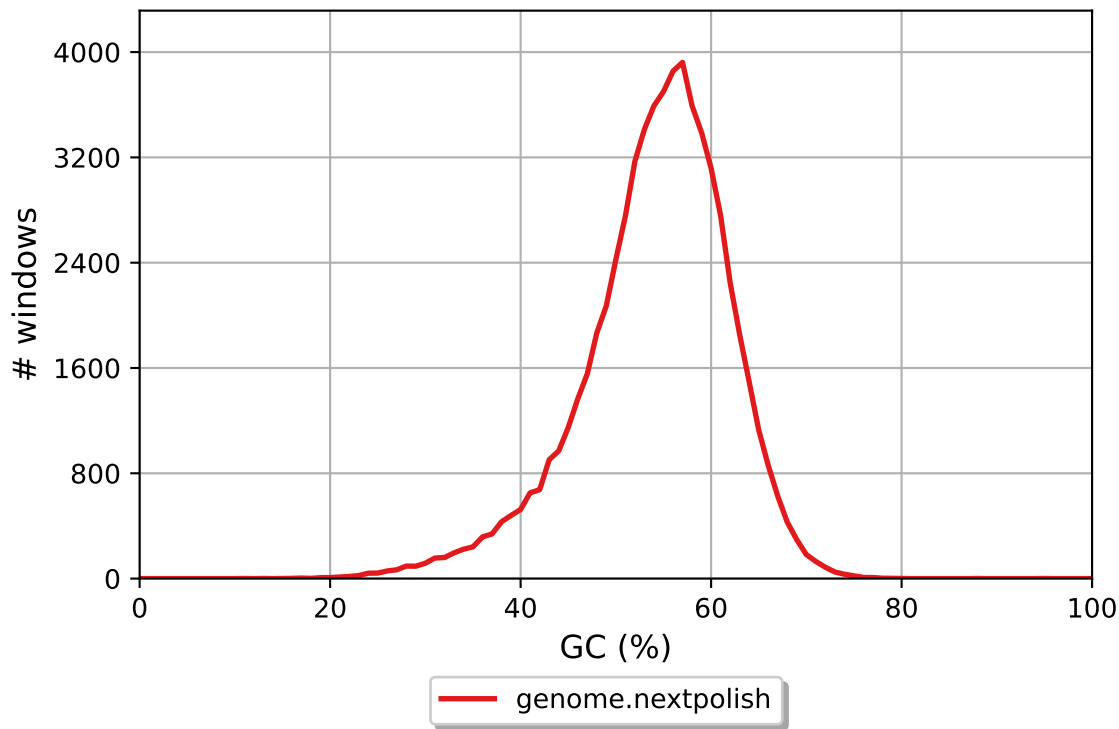


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

