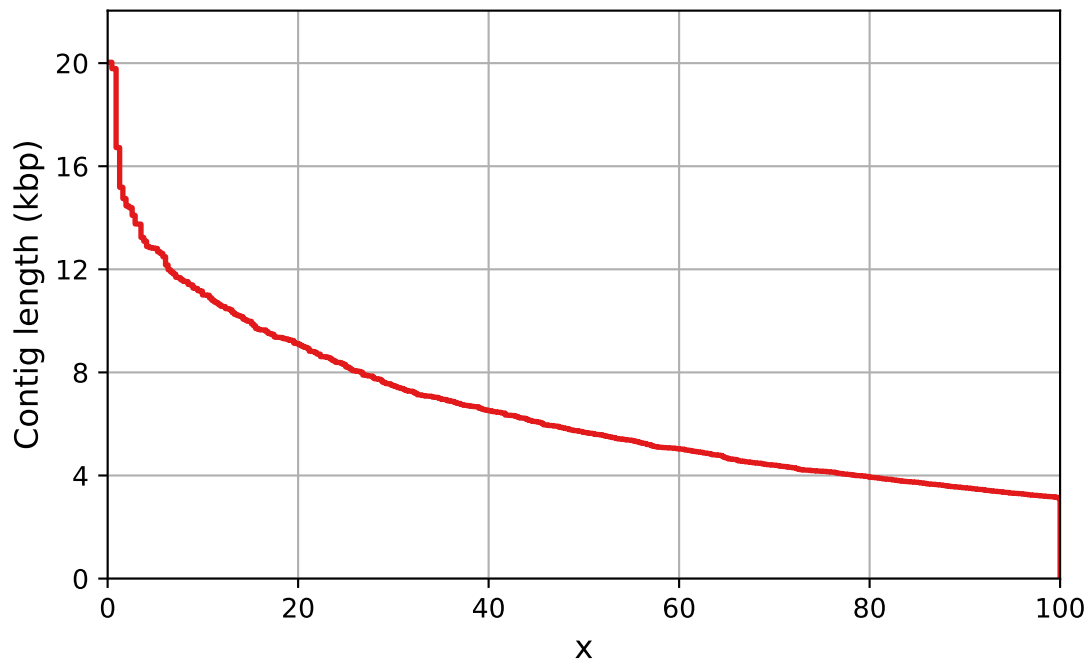


Report

	genome.nextpolish
# contigs (>= 0 bp)	814
# contigs (>= 1000 bp)	814
# contigs (>= 5000 bp)	361
# contigs (>= 10000 bp)	54
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	4479967
Total length (>= 1000 bp)	4479967
Total length (>= 5000 bp)	2715059
Total length (>= 10000 bp)	656421
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	814
Largest contig	20033
Total length	4479967
GC (%)	54.50
N50	5676
N90	3522
auN	6602.1
L50	272
L90	680
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	8 + 0 part

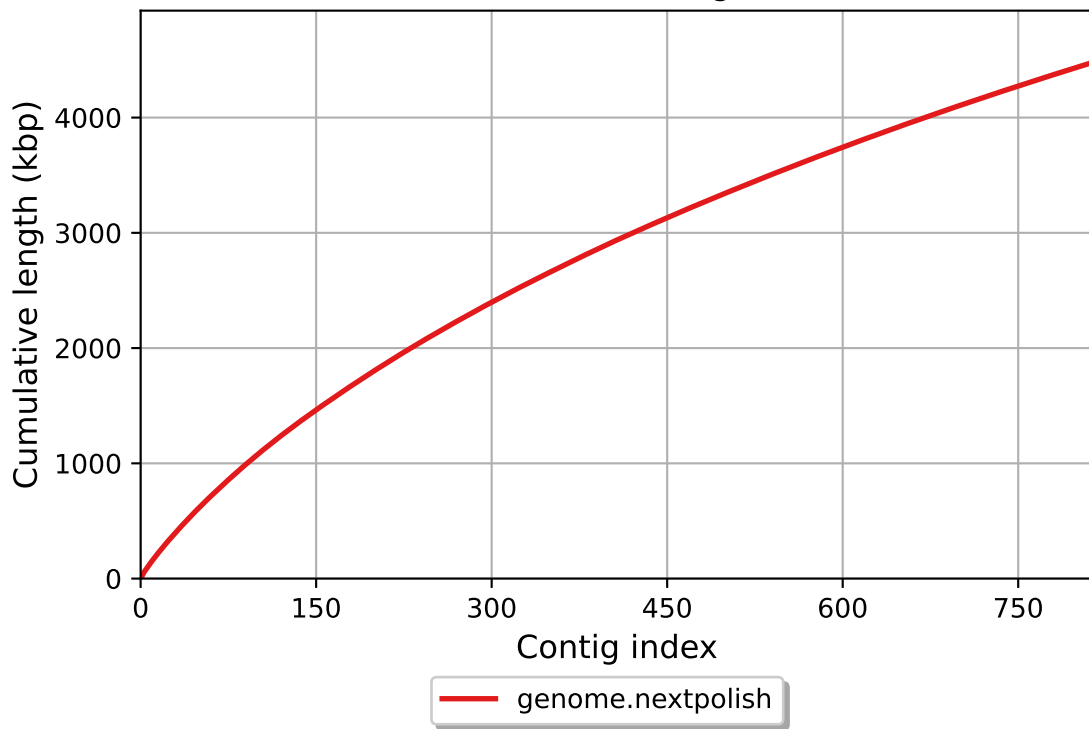
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

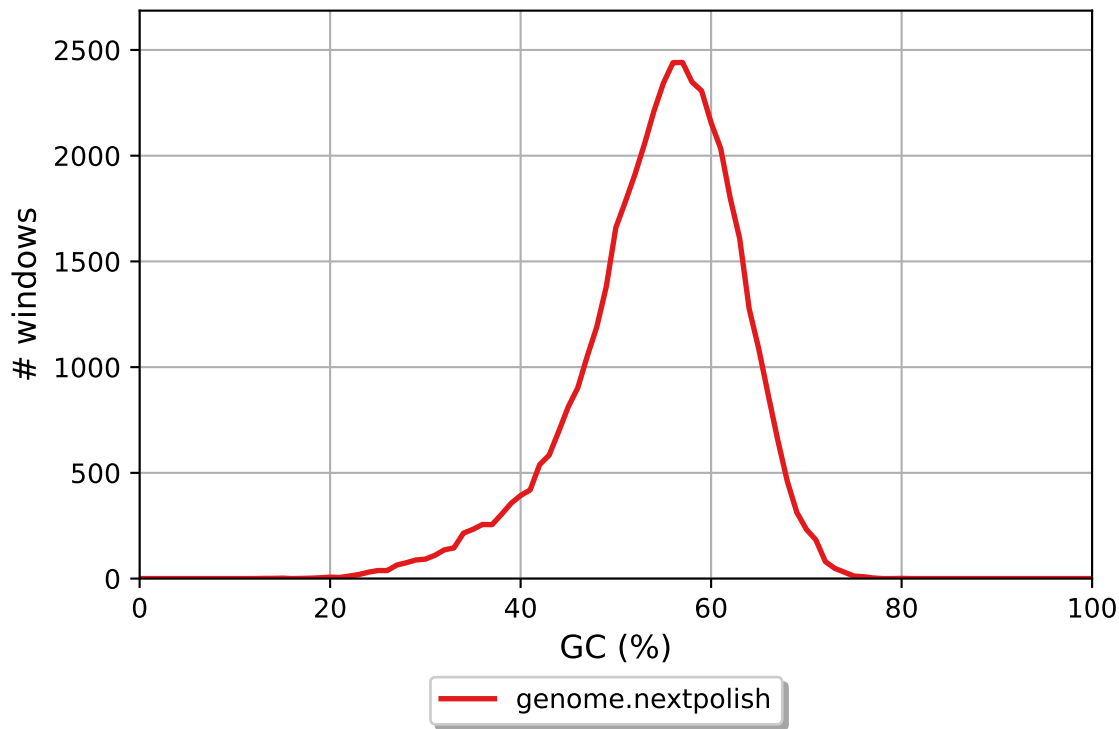


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

