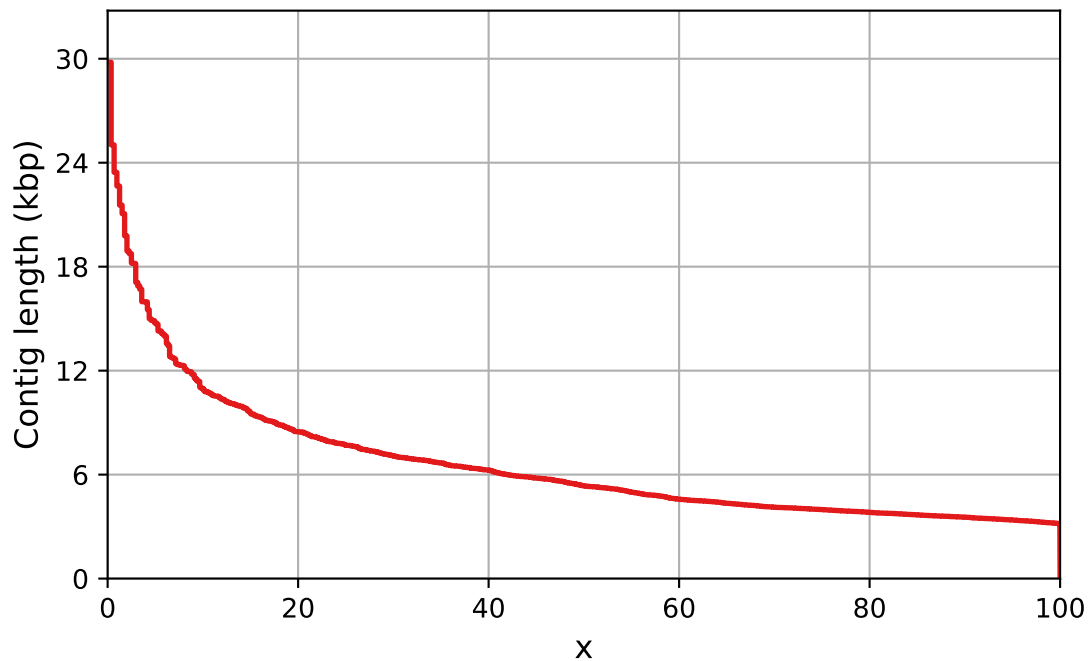


Report

	genome.nextpolish
# contigs (>= 0 bp)	1509
# contigs (>= 1000 bp)	1509
# contigs (>= 5000 bp)	578
# contigs (>= 10000 bp)	81
# contigs (>= 25000 bp)	2
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	8046034
Total length (>= 1000 bp)	8046034
Total length (>= 5000 bp)	4418939
Total length (>= 10000 bp)	1091885
Total length (>= 25000 bp)	54828
Total length (>= 50000 bp)	0
# contigs	1509
Largest contig	29796
Total length	8046034
GC (%)	37.98
N50	5357
N90	3540
auN	6628.6
L50	502
L90	1270
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	15 + 1 part

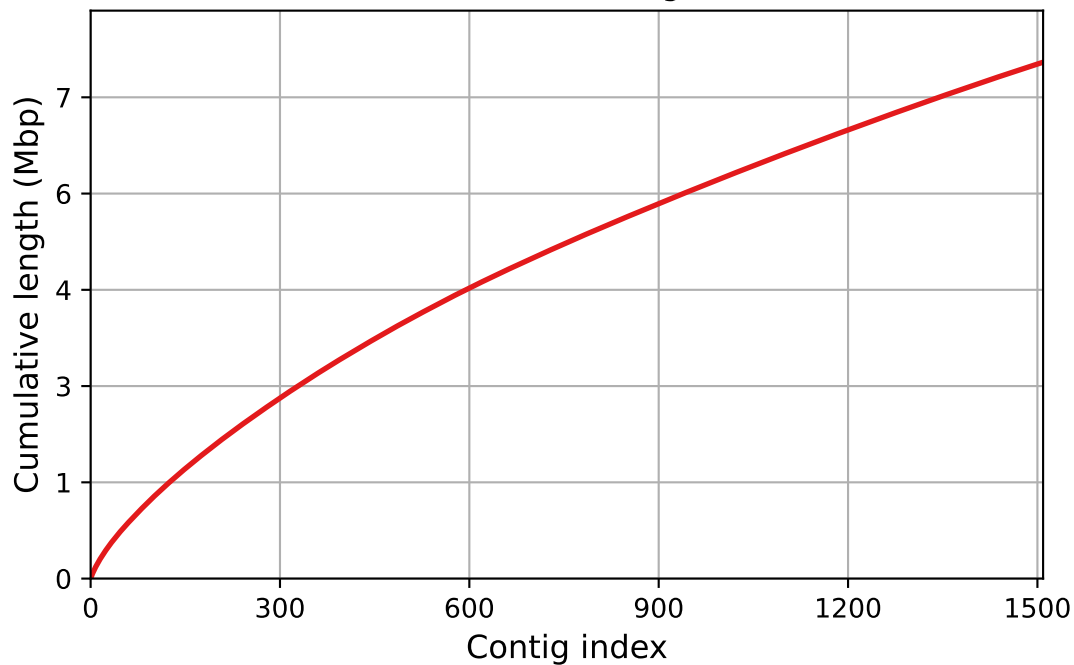
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx



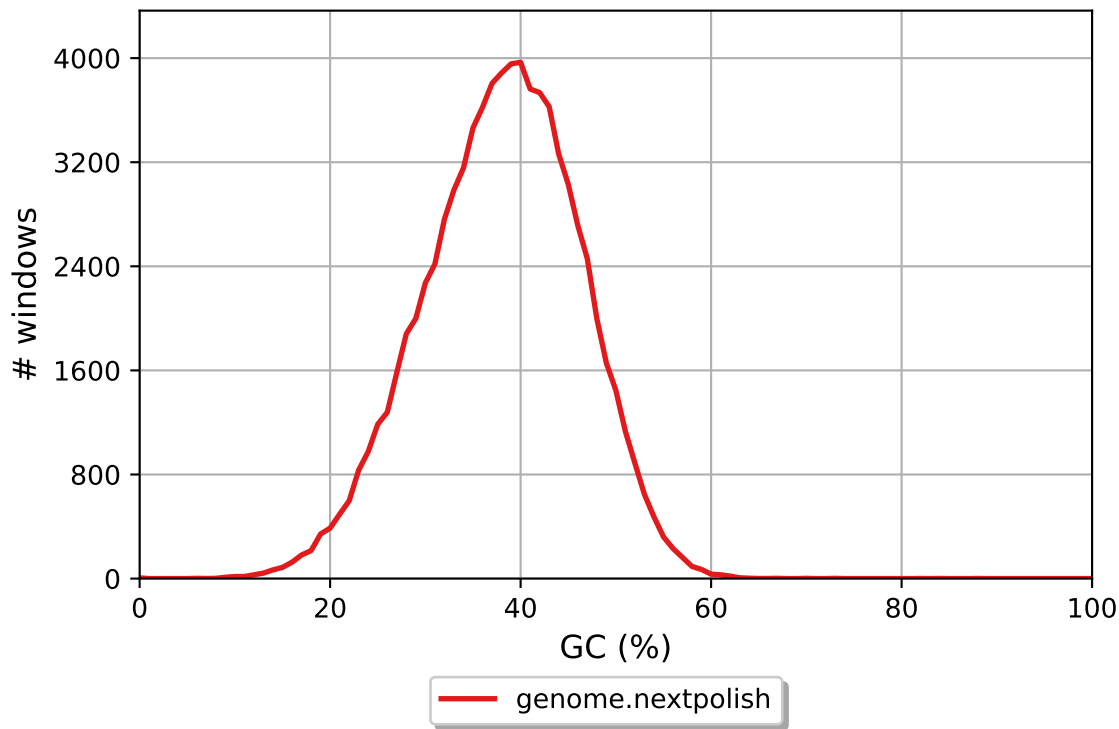
— genome.nextpolish

Cumulative length



— genome.nextpolish

GC content



genome.nextpolish GC content

