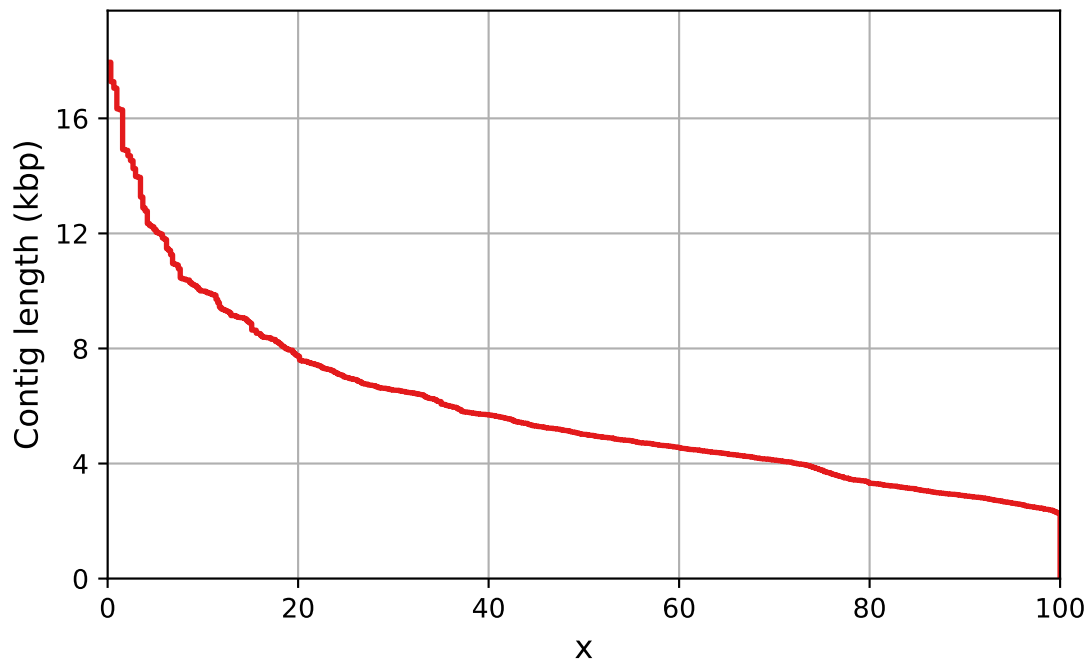


Report

	genome.nextpolish
# contigs (>= 0 bp)	1135
# contigs (>= 1000 bp)	1135
# contigs (>= 5000 bp)	375
# contigs (>= 10000 bp)	43
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	5395770
Total length (>= 1000 bp)	5395770
Total length (>= 5000 bp)	2716620
Total length (>= 10000 bp)	534520
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	1135
Largest contig	17950
Total length	5395770
GC (%)	42.05
N50	5013
N90	2884
auN	5871.0
L50	372
L90	928
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	18 + 3 part

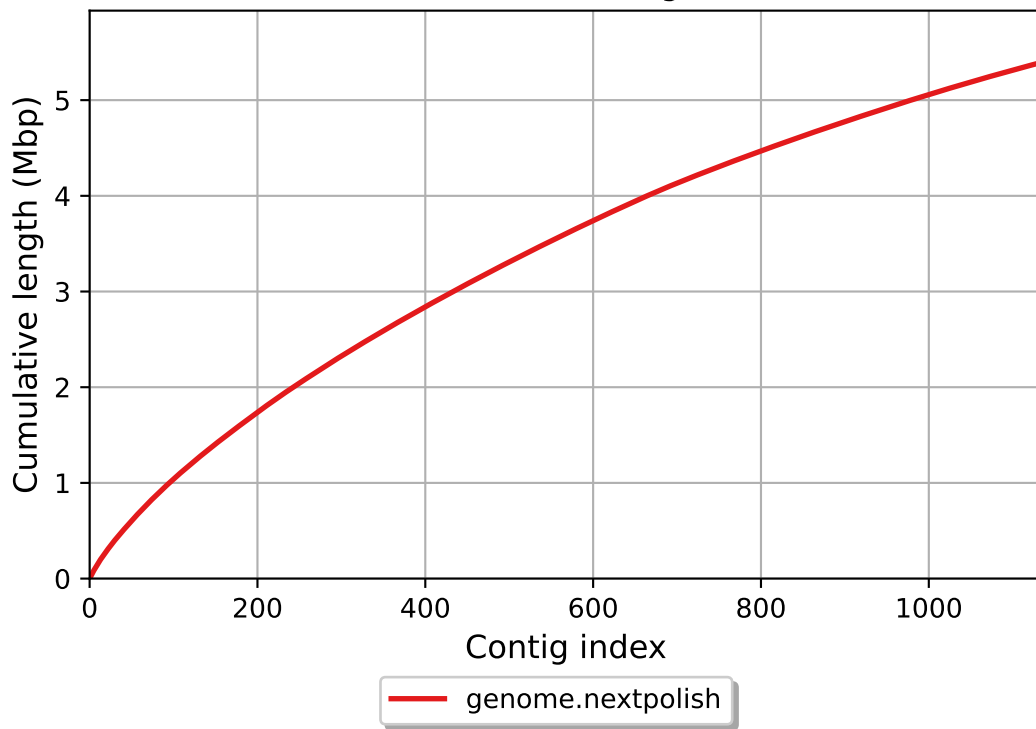
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

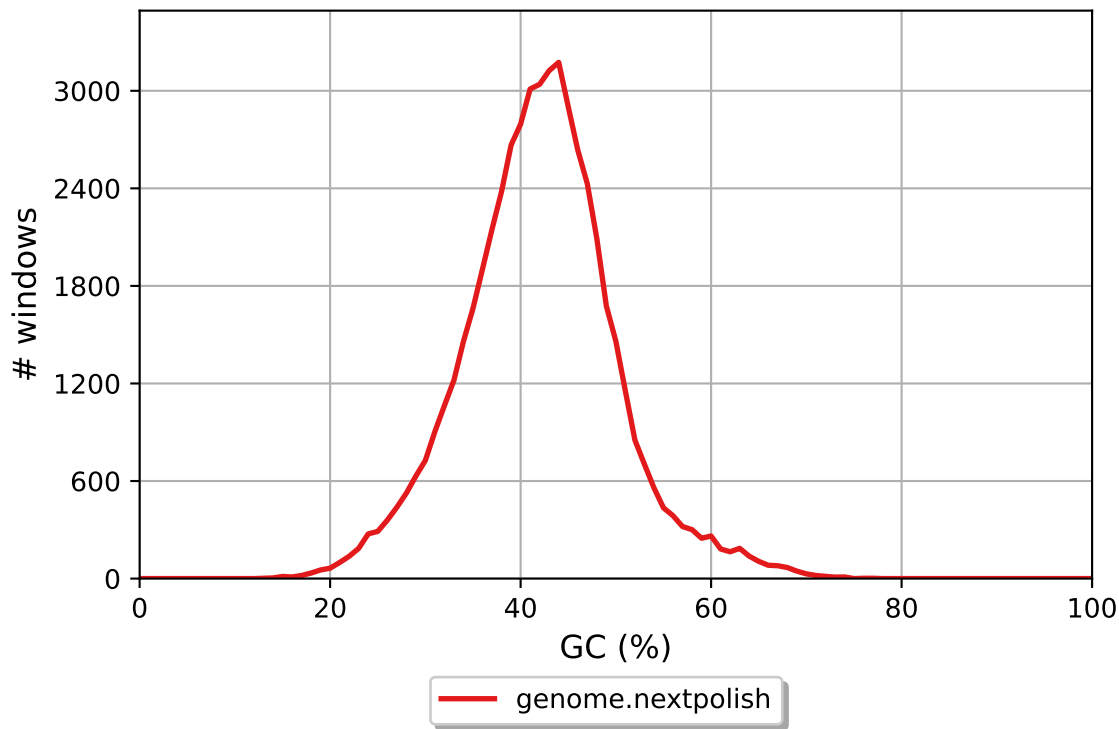


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

