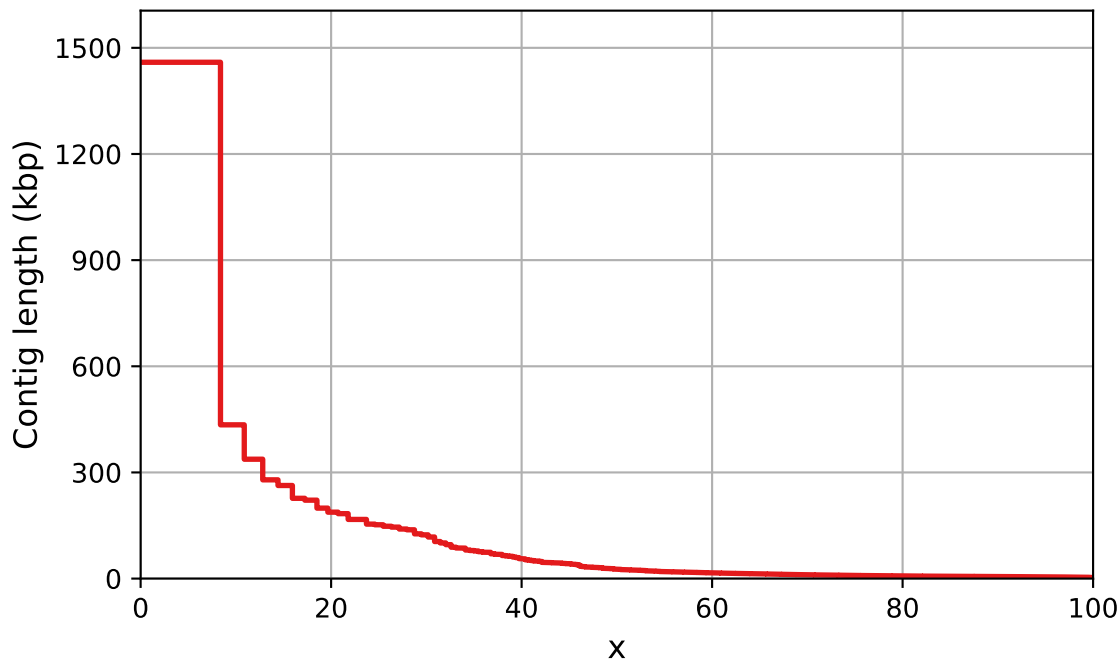


Report

	genome.nextpolish
# contigs (>= 0 bp)	1140
# contigs (>= 1000 bp)	1140
# contigs (>= 5000 bp)	911
# contigs (>= 10000 bp)	338
# contigs (>= 25000 bp)	96
# contigs (>= 50000 bp)	46
Total length (>= 0 bp)	17404016
Total length (>= 1000 bp)	17404016
Total length (>= 5000 bp)	16423456
Total length (>= 10000 bp)	12511457
Total length (>= 25000 bp)	8924676
Total length (>= 50000 bp)	7184848
# contigs	1140
Largest contig	1459196
Total length	17404016
GC (%)	47.77
N50	26137
N90	5739
auN	190007.2
L50	88
L90	770
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	52 + 4 part

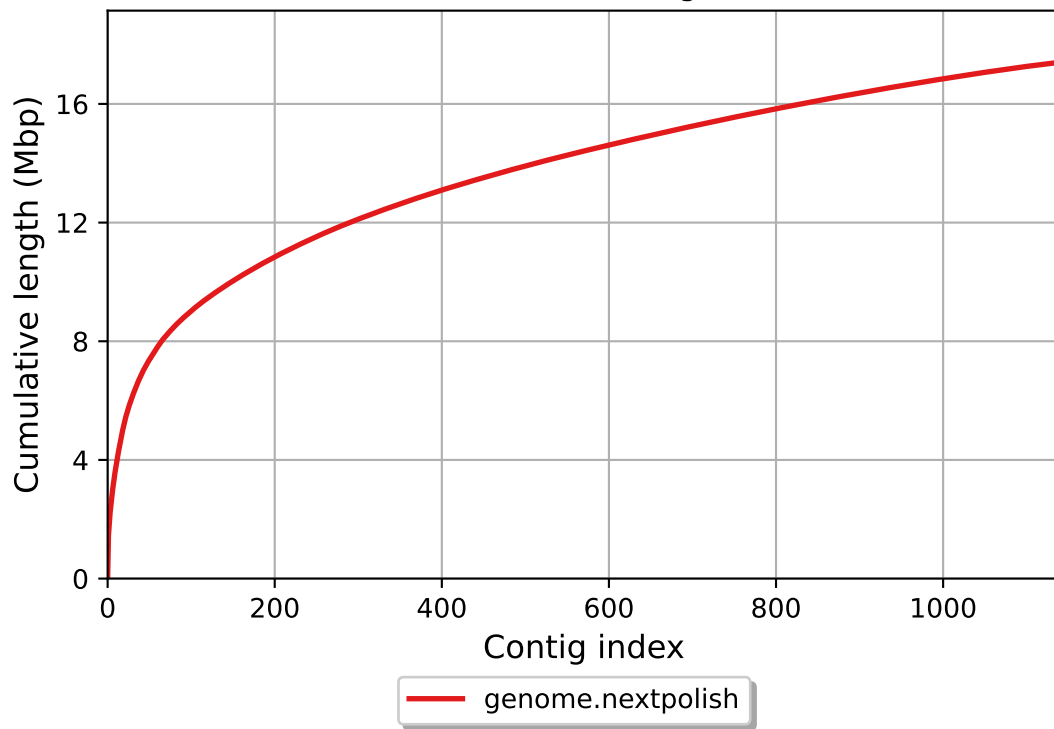
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

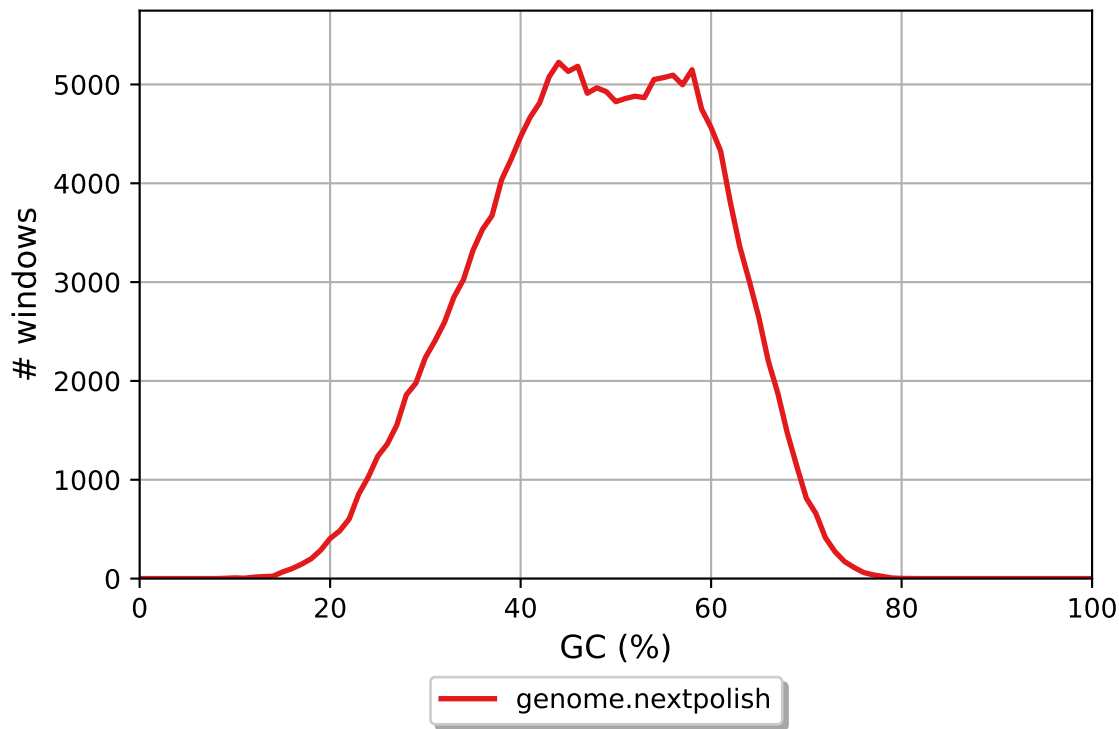


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

