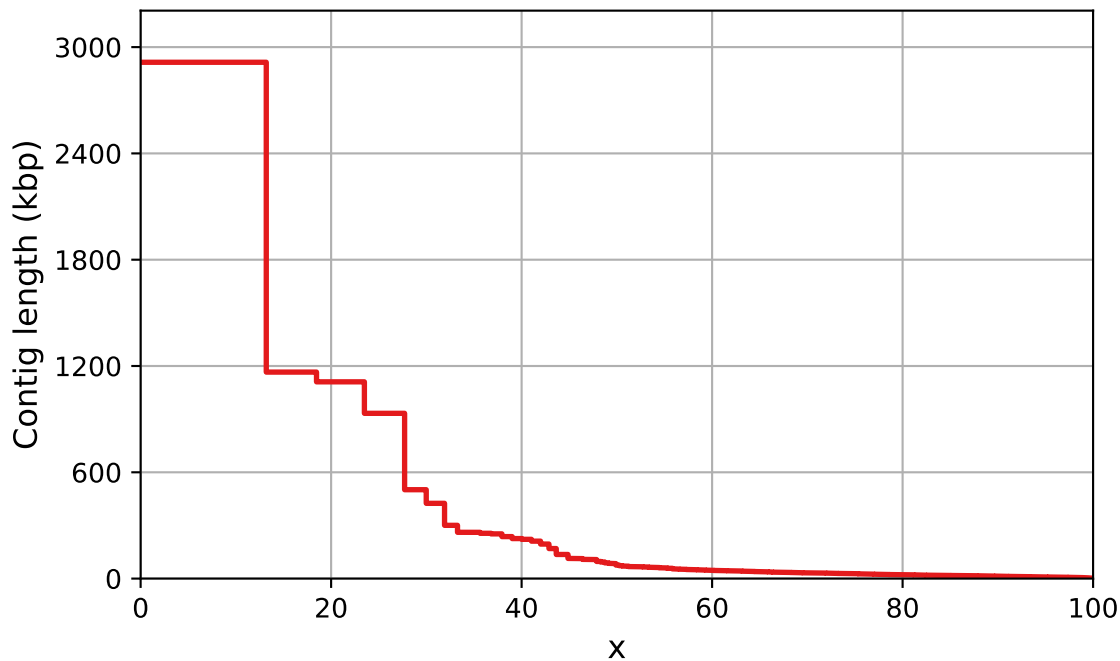


Report

	genome.nextpolish
# contigs (>= 0 bp)	613
# contigs (>= 1000 bp)	611
# contigs (>= 5000 bp)	559
# contigs (>= 10000 bp)	406
# contigs (>= 25000 bp)	178
# contigs (>= 50000 bp)	60
Total length (>= 0 bp)	22093807
Total length (>= 1000 bp)	22092461
Total length (>= 5000 bp)	21903292
Total length (>= 10000 bp)	20727007
Total length (>= 25000 bp)	16986271
Total length (>= 50000 bp)	12857866
# contigs	613
Largest contig	2914443
Total length	22093807
GC (%)	53.89
N50	76085
N90	13648
auN	611095.6
L50	31
L90	335
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	50 + 1 part

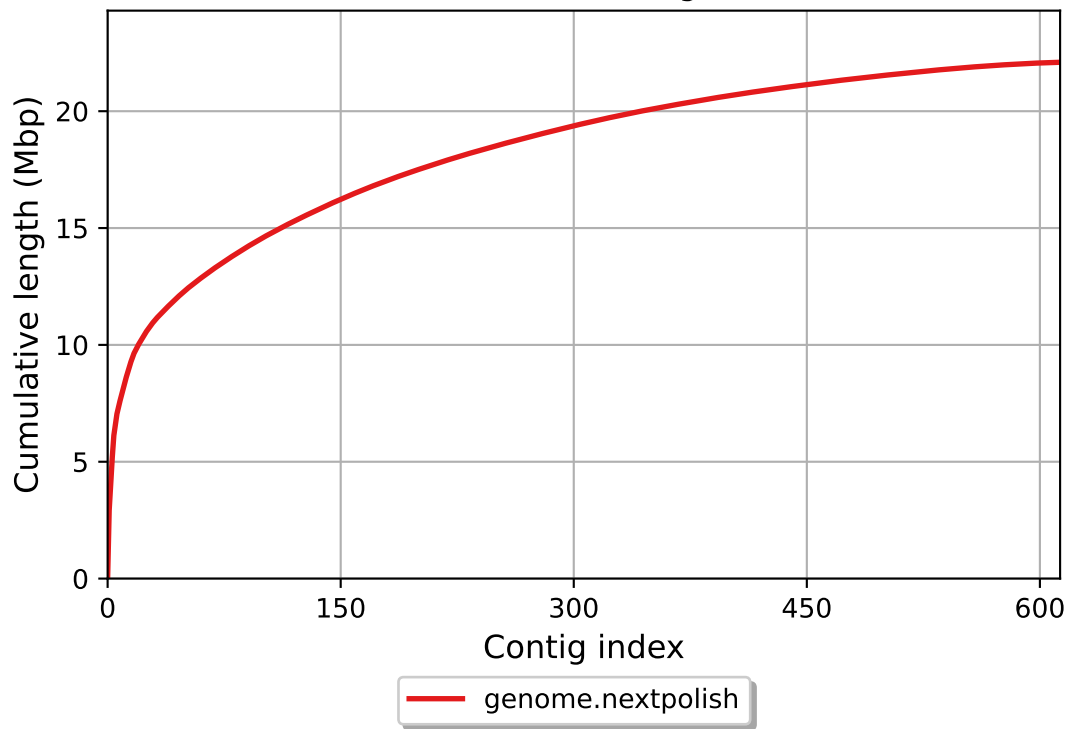
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

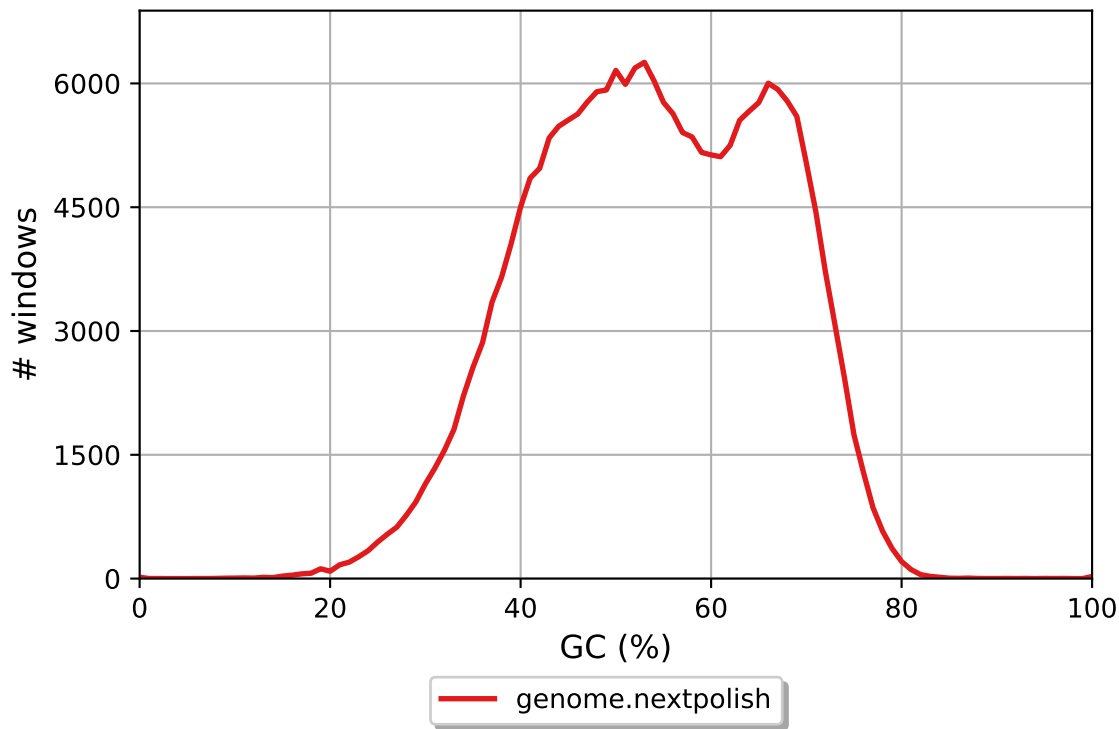


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

