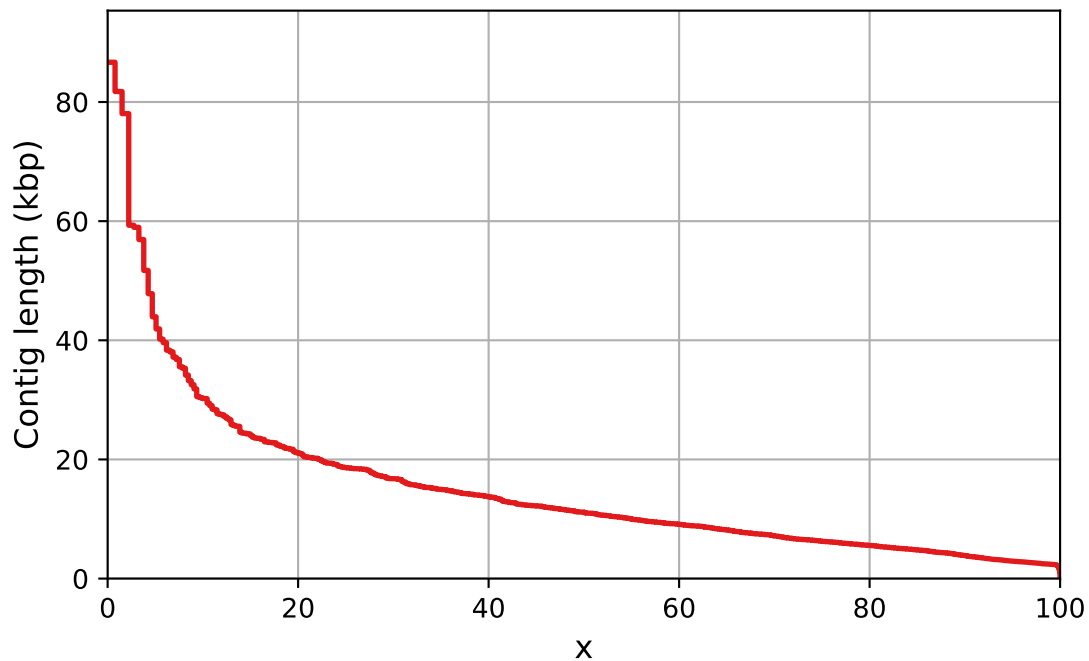


Report

	genome.nextpolish
# contigs (>= 0 bp)	1343
# contigs (>= 1000 bp)	1342
# contigs (>= 5000 bp)	799
# contigs (>= 10000 bp)	347
# contigs (>= 25000 bp)	40
# contigs (>= 50000 bp)	7
Total length (>= 0 bp)	11116569
Total length (>= 1000 bp)	11116113
Total length (>= 5000 bp)	9309502
Total length (>= 10000 bp)	6113835
Total length (>= 25000 bp)	1542670
Total length (>= 50000 bp)	473404
# contigs	1342
Largest contig	86669
Total length	11116113
GC (%)	44.93
N50	11131
N90	3872
auN	15439.5
L50	295
L90	956
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	15 + 3 part

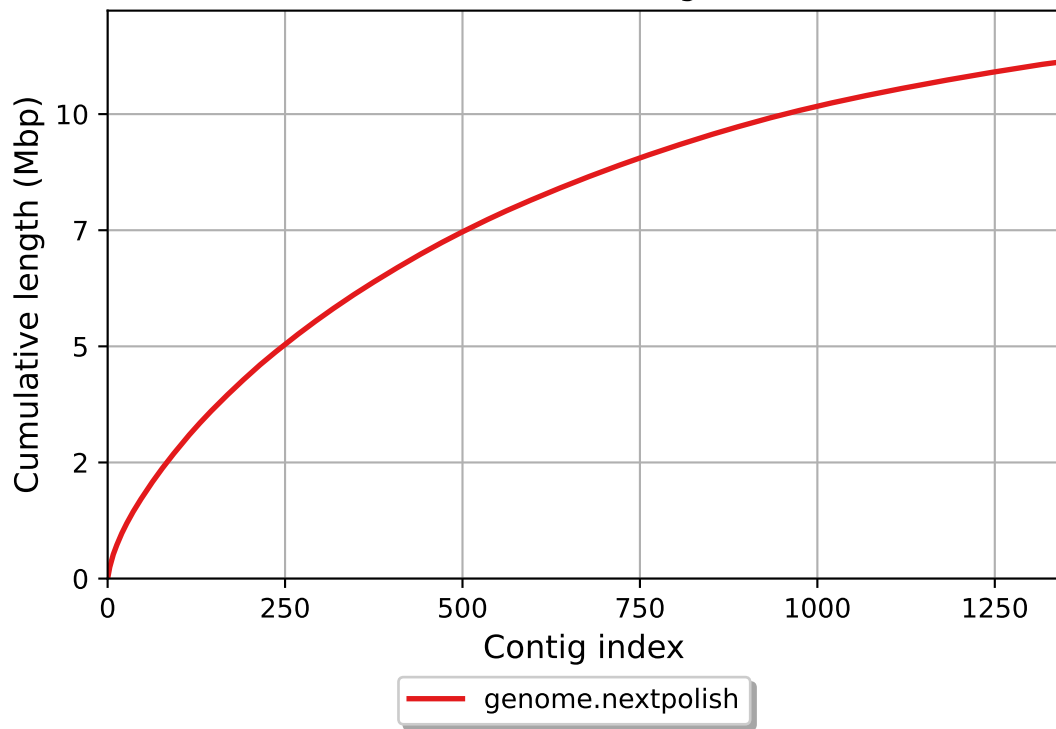
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

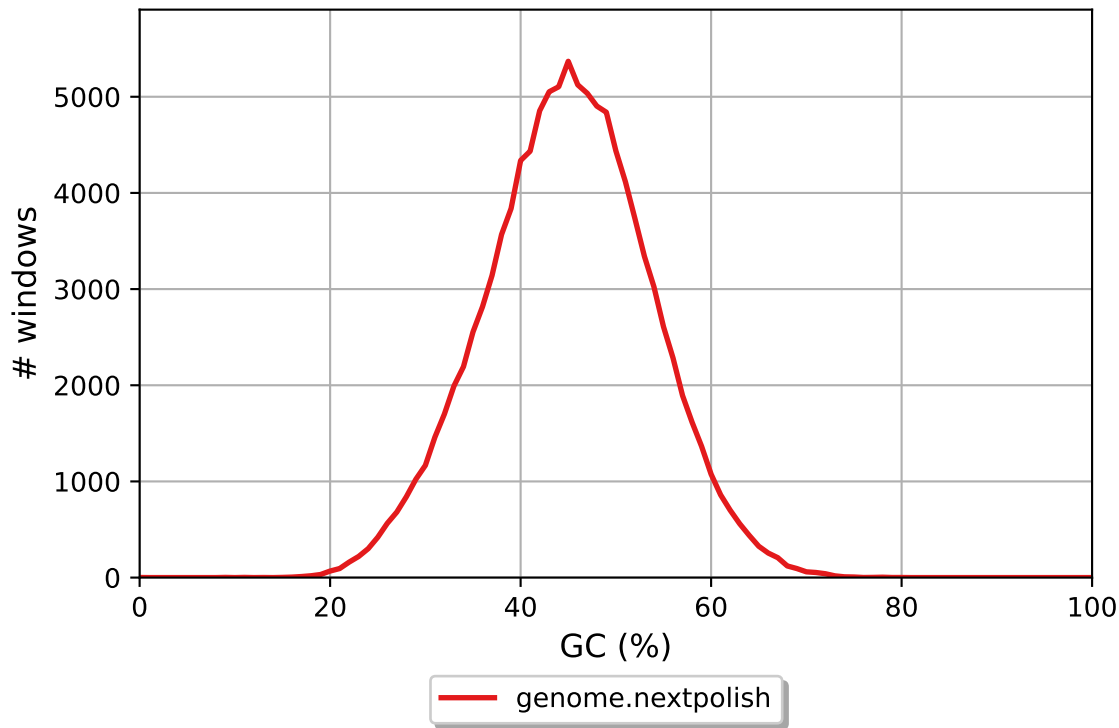


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

