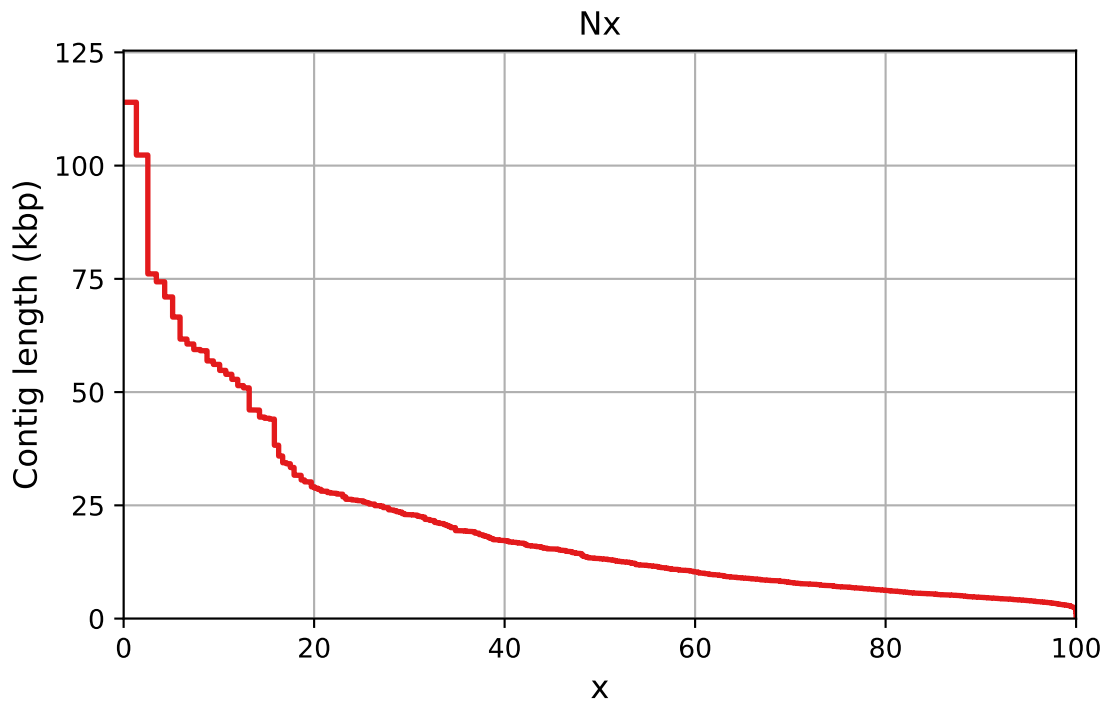


Report

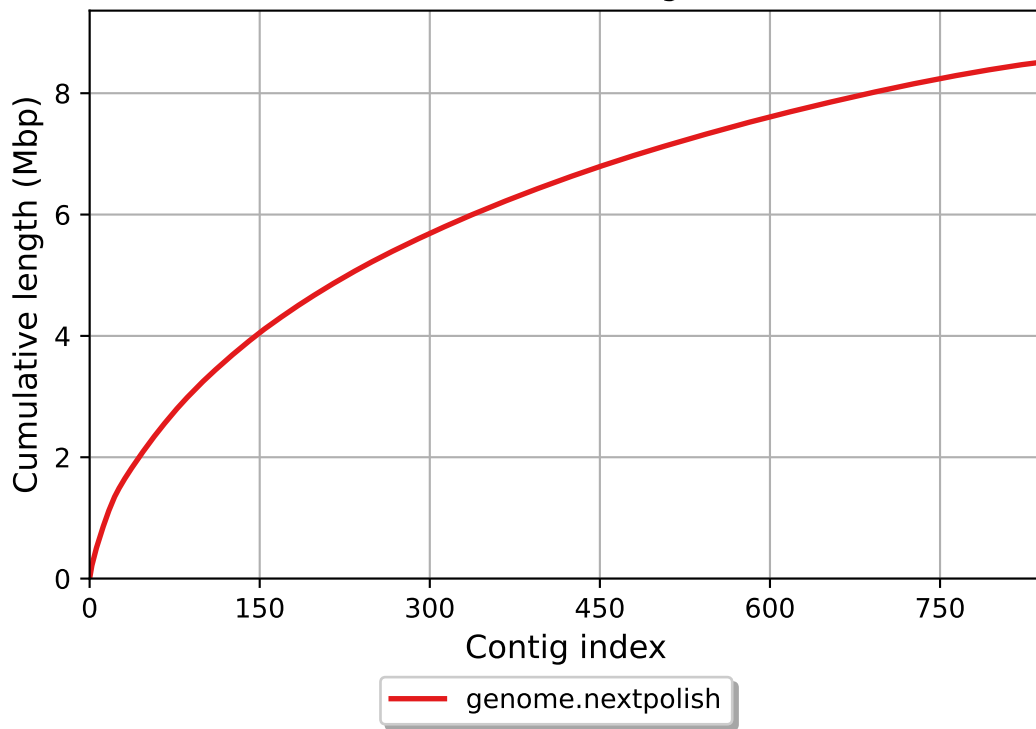
	genome.nextpolish
# contigs (>= 0 bp)	840
# contigs (>= 1000 bp)	839
# contigs (>= 5000 bp)	578
# contigs (>= 10000 bp)	246
# contigs (>= 25000 bp)	53
# contigs (>= 50000 bp)	17
Total length (>= 0 bp)	8510078
Total length (>= 1000 bp)	8509570
Total length (>= 5000 bp)	7503024
Total length (>= 10000 bp)	5191798
Total length (>= 25000 bp)	2243753
Total length (>= 50000 bp)	1122051
# contigs	840
Largest contig	113982
Total length	8510078
GC (%)	45.53
N50	13250
N90	4710
auN	21687.4
L50	165
L90	611
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	17 + 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).

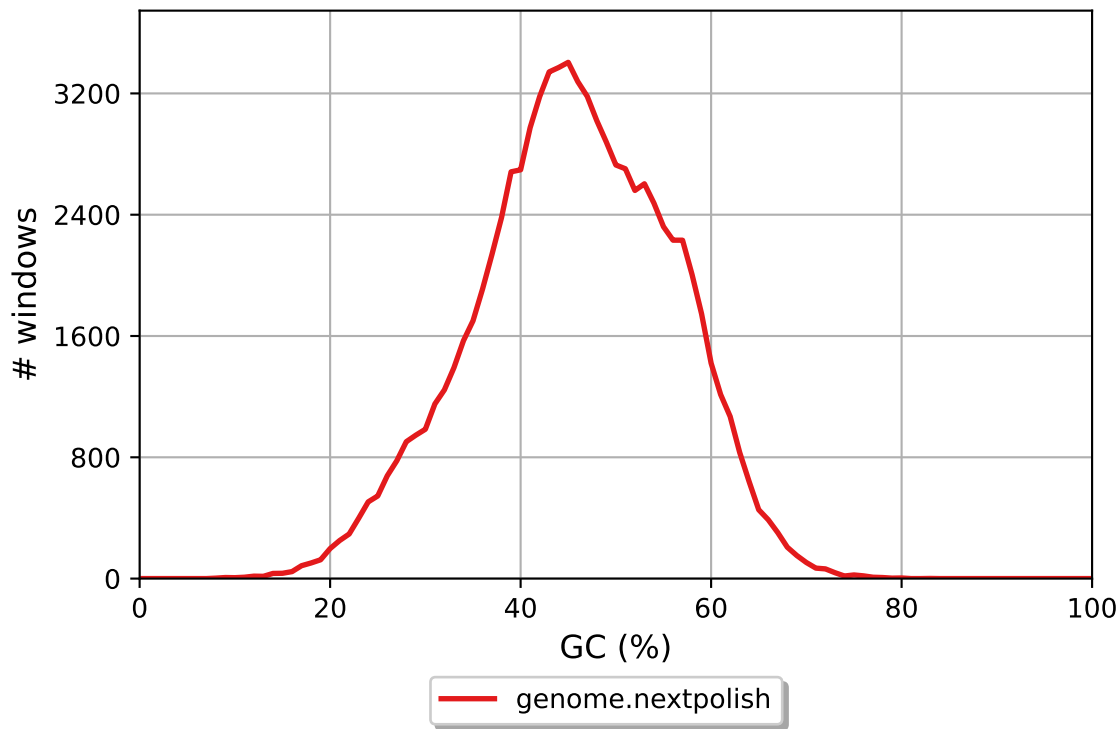


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

