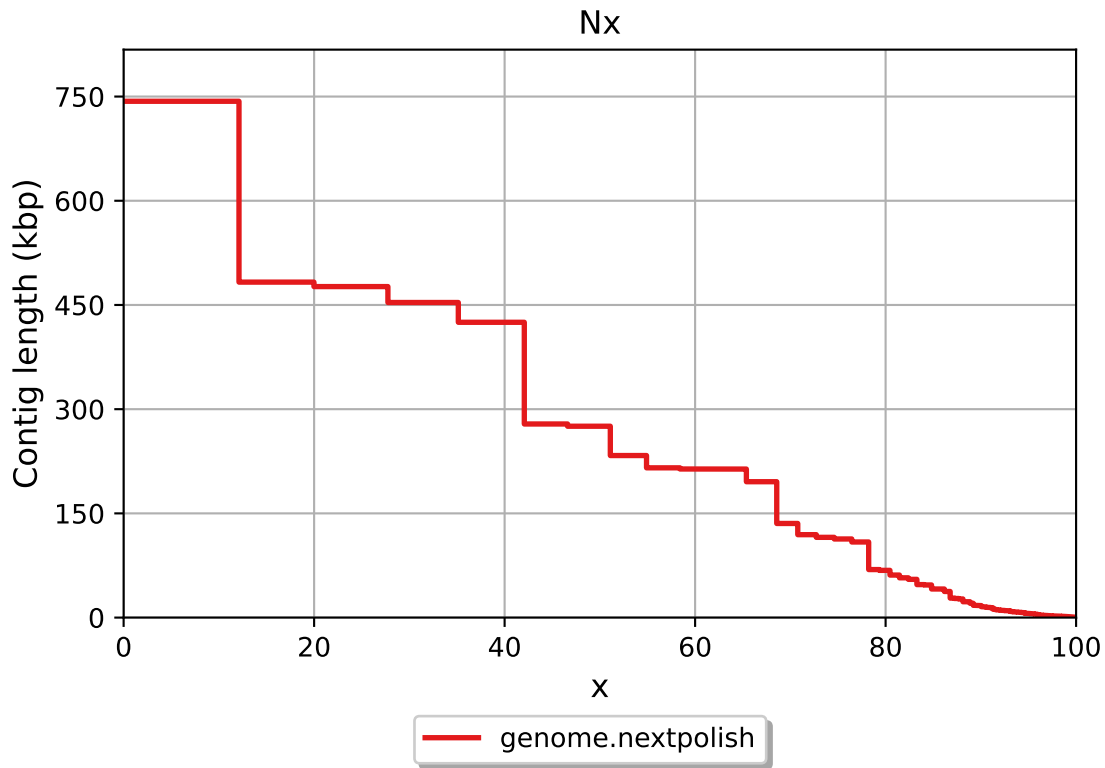


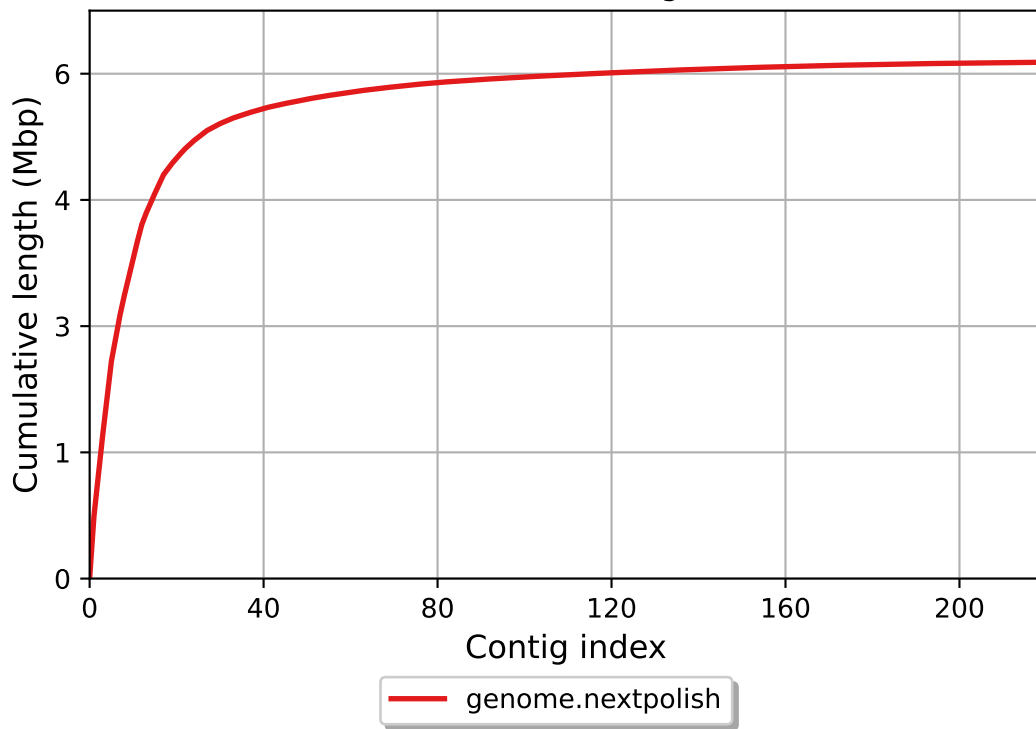
Report

	genome.nextpolish
# contigs (>= 0 bp)	223
# contigs (>= 1000 bp)	183
# contigs (>= 5000 bp)	75
# contigs (>= 10000 bp)	49
# contigs (>= 25000 bp)	30
# contigs (>= 50000 bp)	22
Total length (>= 0 bp)	6137050
Total length (>= 1000 bp)	6111547
Total length (>= 5000 bp)	5871341
Total length (>= 10000 bp)	5686973
Total length (>= 25000 bp)	5406934
Total length (>= 50000 bp)	5110304
# contigs	219
Largest contig	743214
Total length	6135897
GC (%)	38.91
N50	275453
N90	17273
auN	308199.4
L50	7
L90	36
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	21 + 2 part

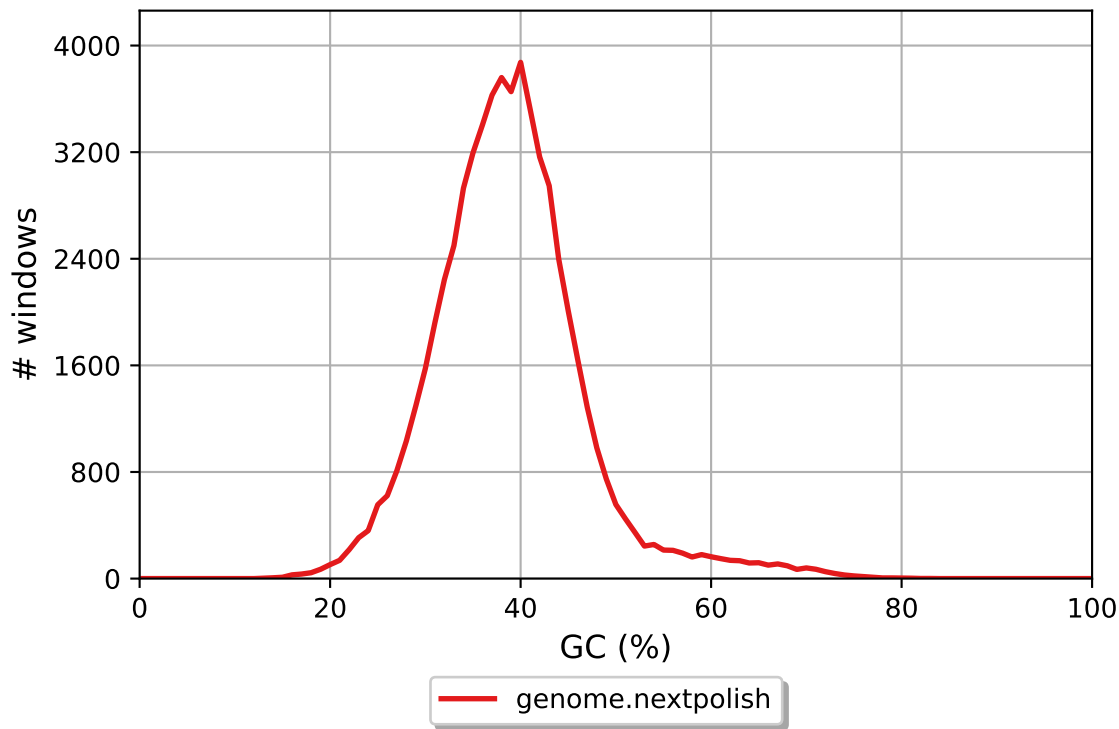
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

