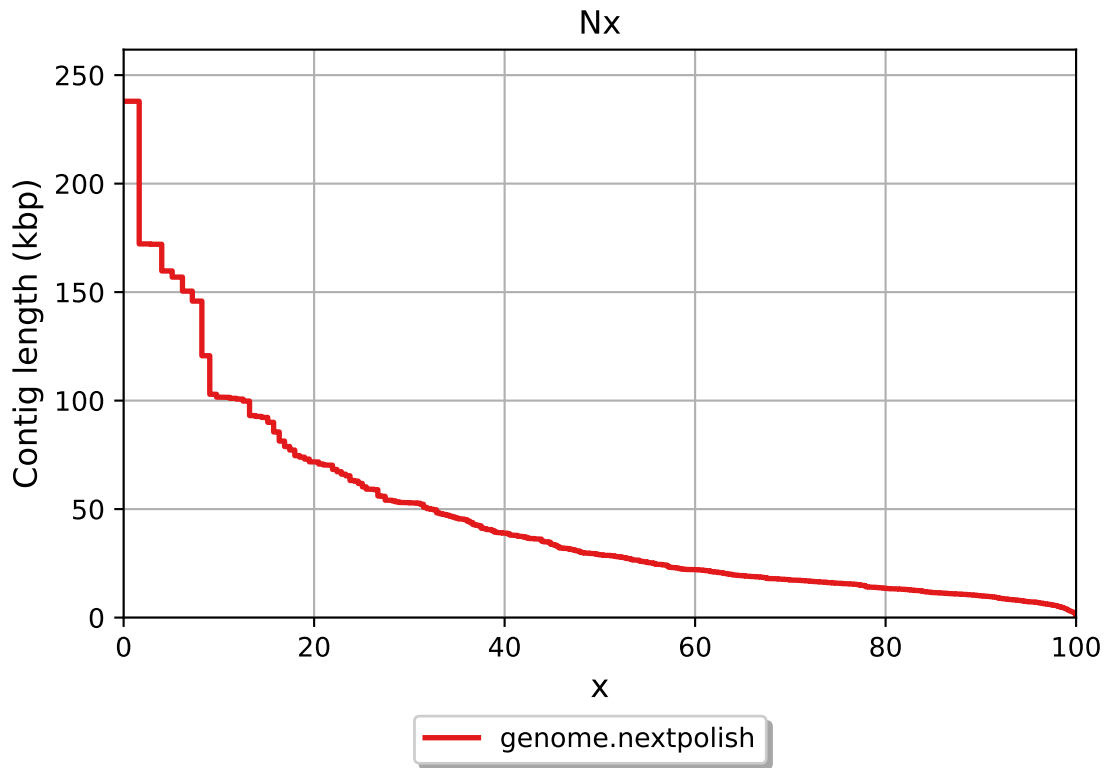


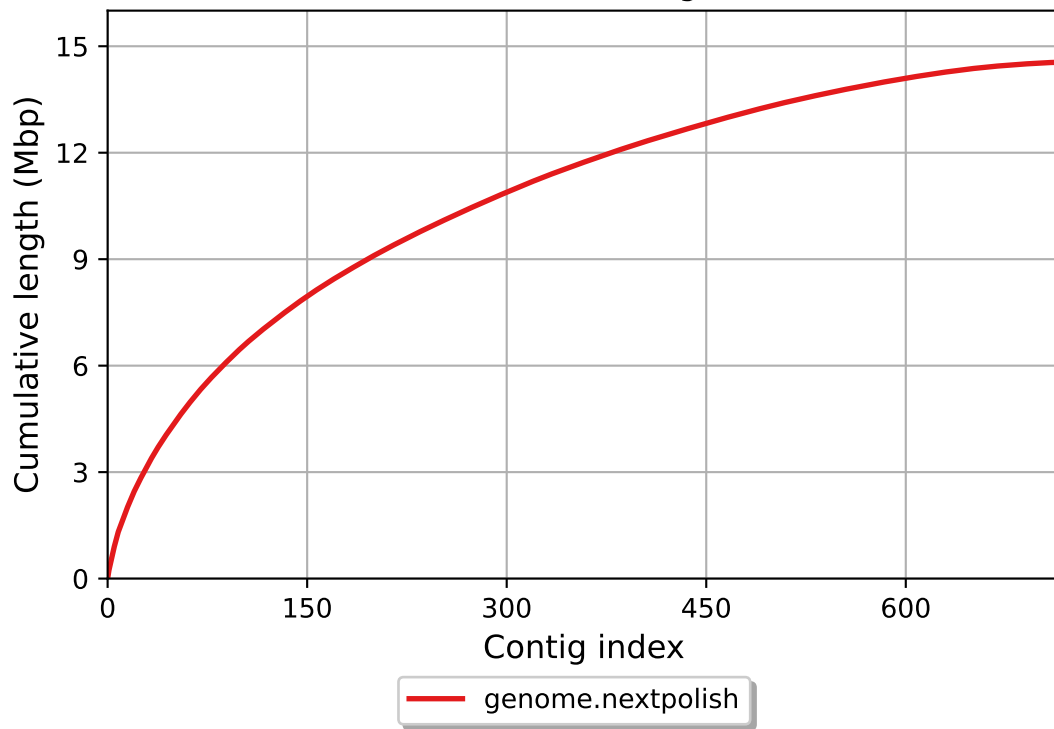
# Report

	genome.nextpolish
# contigs (>= 0 bp)	716
# contigs (>= 1000 bp)	714
# contigs (>= 5000 bp)	634
# contigs (>= 10000 bp)	478
# contigs (>= 25000 bp)	156
# contigs (>= 50000 bp)	56
Total length (>= 0 bp)	14548613
Total length (>= 1000 bp)	14547353
Total length (>= 5000 bp)	14293864
Total length (>= 10000 bp)	13118870
Total length (>= 25000 bp)	8106866
Total length (>= 50000 bp)	4680937
# contigs	716
Largest contig	237951
Total length	14548613
GC (%)	48.12
N50	28946
N90	10062
auN	47354.5
L50	126
L90	476
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	42 + 5 part

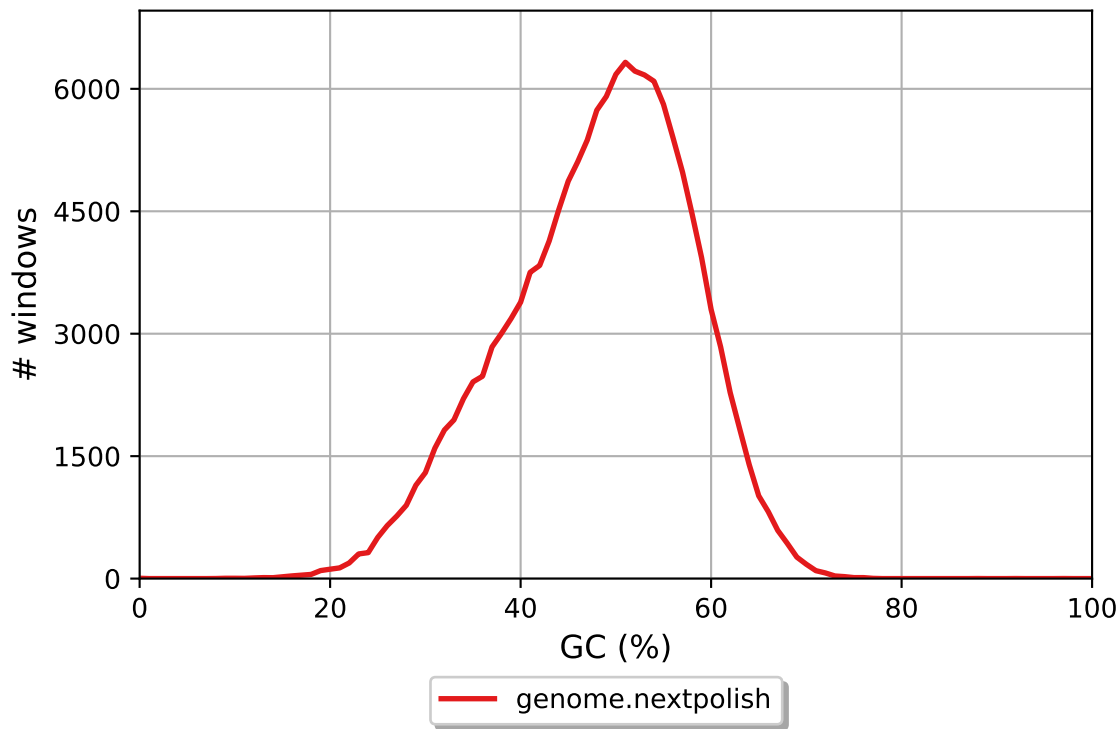
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

