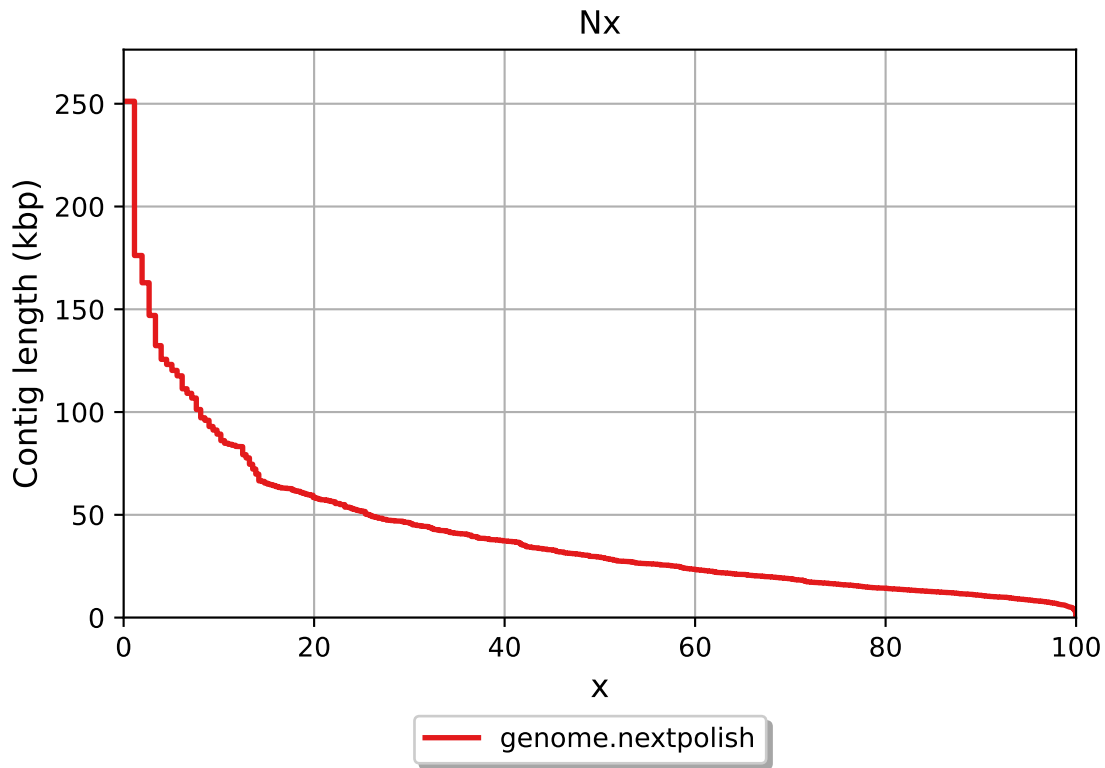


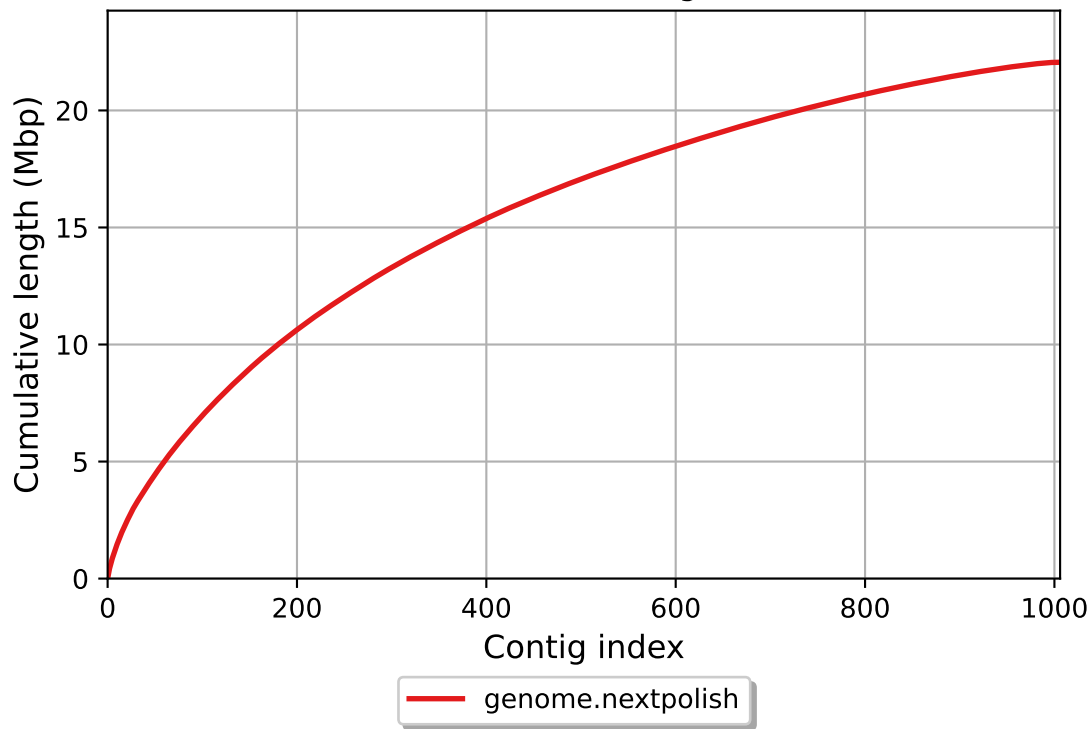
# Report

	genome.nextpolish
# contigs (>= 0 bp)	1006
# contigs (>= 1000 bp)	999
# contigs (>= 5000 bp)	968
# contigs (>= 10000 bp)	756
# contigs (>= 25000 bp)	279
# contigs (>= 50000 bp)	73
Total length (>= 0 bp)	22057989
Total length (>= 1000 bp)	22053553
Total length (>= 5000 bp)	21932546
Total length (>= 10000 bp)	20270421
Total length (>= 25000 bp)	12786619
Total length (>= 50000 bp)	5697703
# contigs	1006
Largest contig	251204
Total length	22057989
GC (%)	48.75
N50	29427
N90	10828
auN	41865.6
L50	214
L90	716
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	42 + 6 part

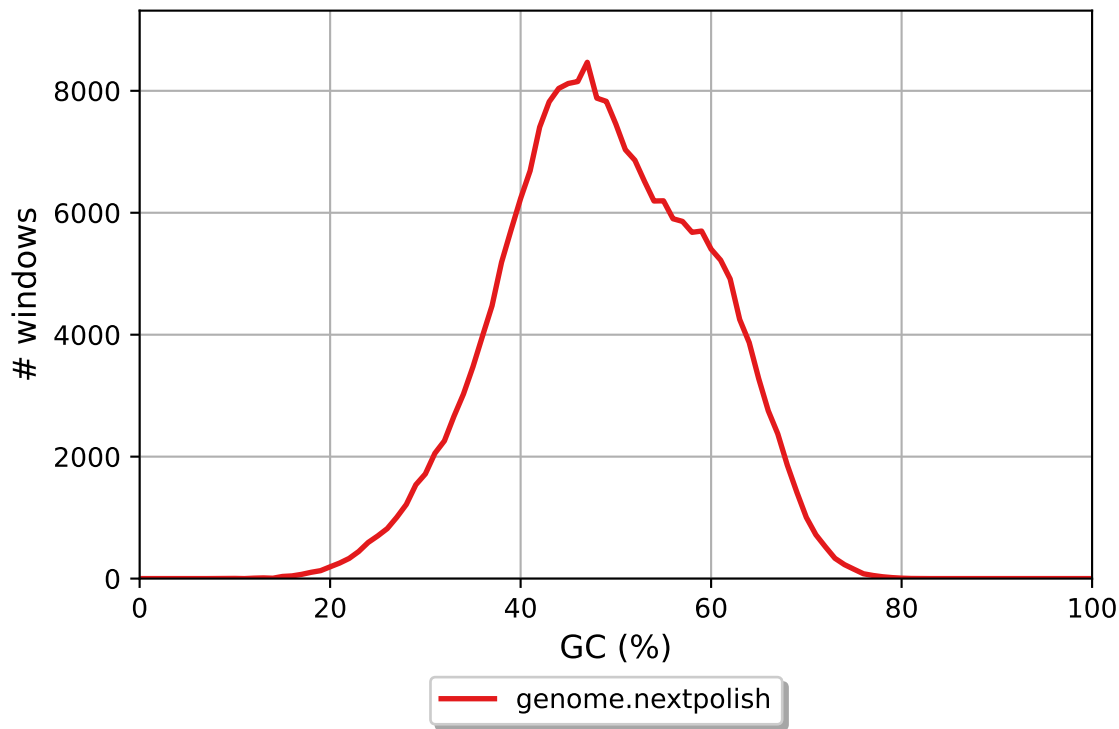
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



## GC content



genome.nextpolish GC content

