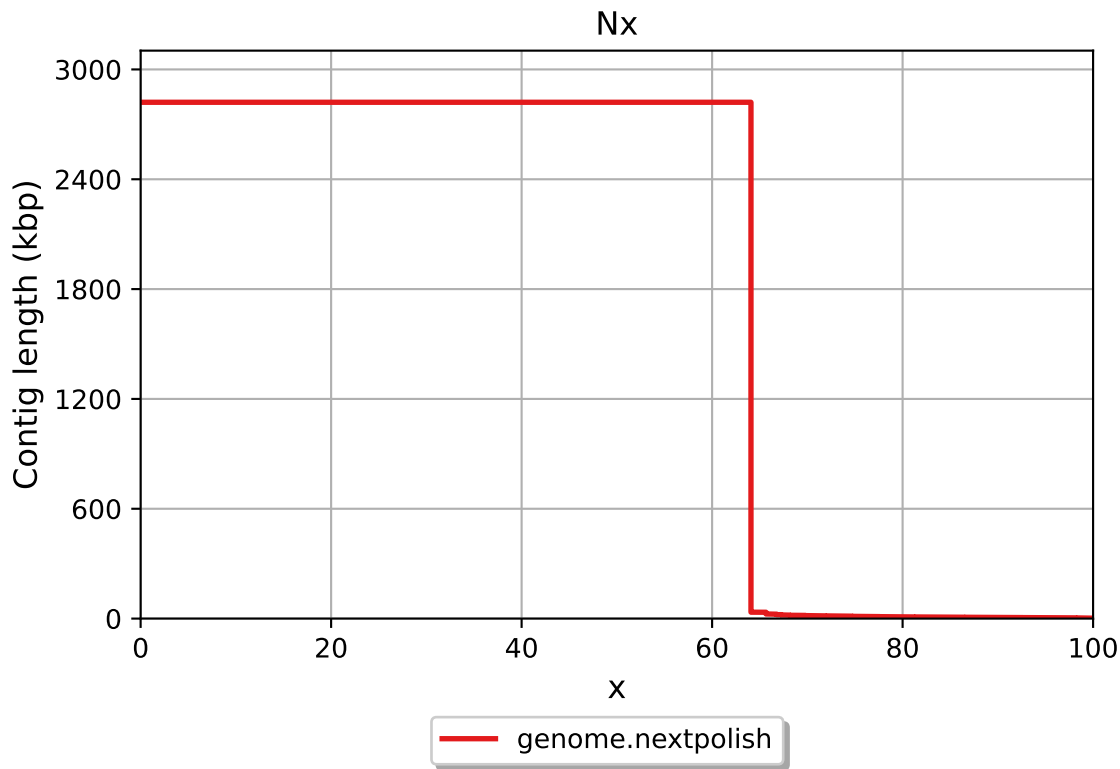


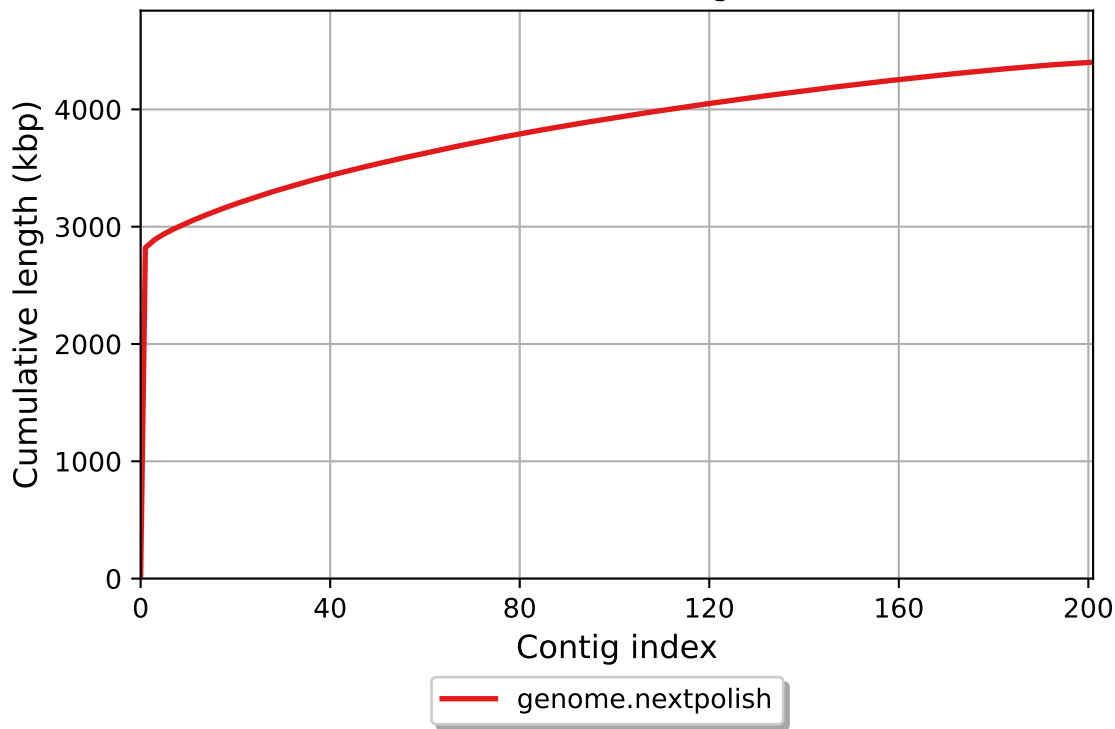
Report

	genome.nextpolish
# contigs (>= 0 bp)	201
# contigs (>= 1000 bp)	201
# contigs (>= 5000 bp)	143
# contigs (>= 10000 bp)	45
# contigs (>= 25000 bp)	4
# contigs (>= 50000 bp)	1
Total length (>= 0 bp)	4401662
Total length (>= 1000 bp)	4401662
Total length (>= 5000 bp)	4173078
Total length (>= 10000 bp)	3488221
Total length (>= 25000 bp)	2915319
Total length (>= 50000 bp)	2820606
# contigs	201
Largest contig	2820606
Total length	4401662
GC (%)	34.70
N50	2820606
N90	6232
auN	1811416.6
L50	1
L90	106
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	13 + 6 part

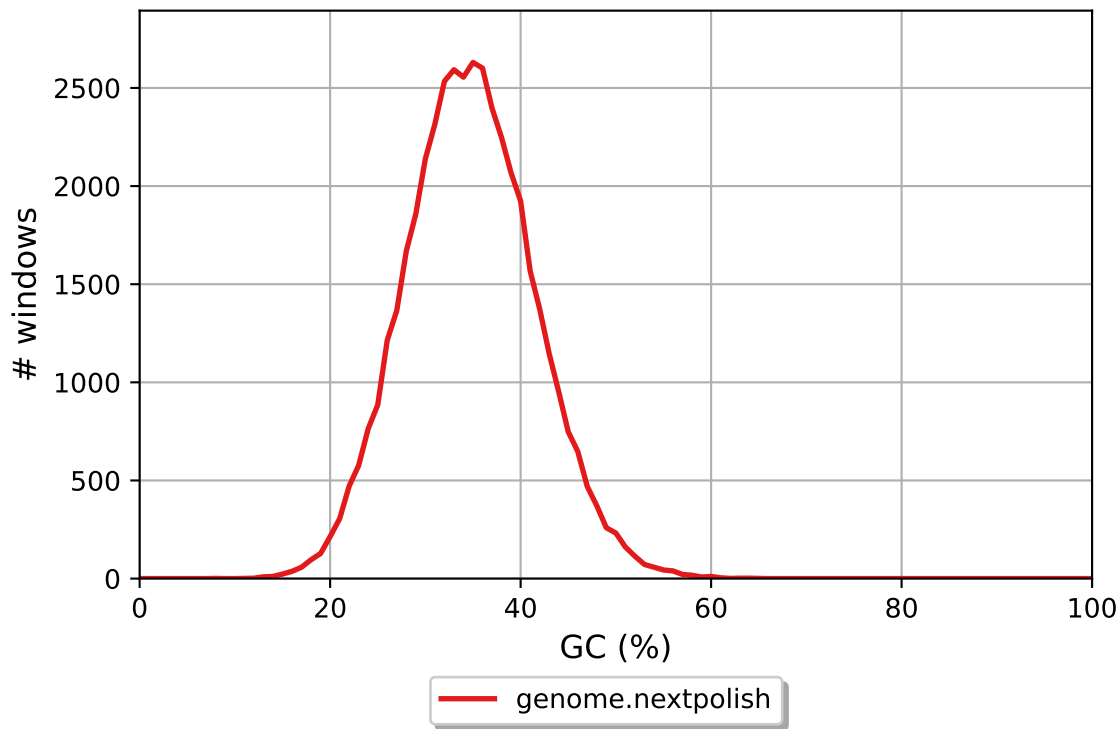
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

