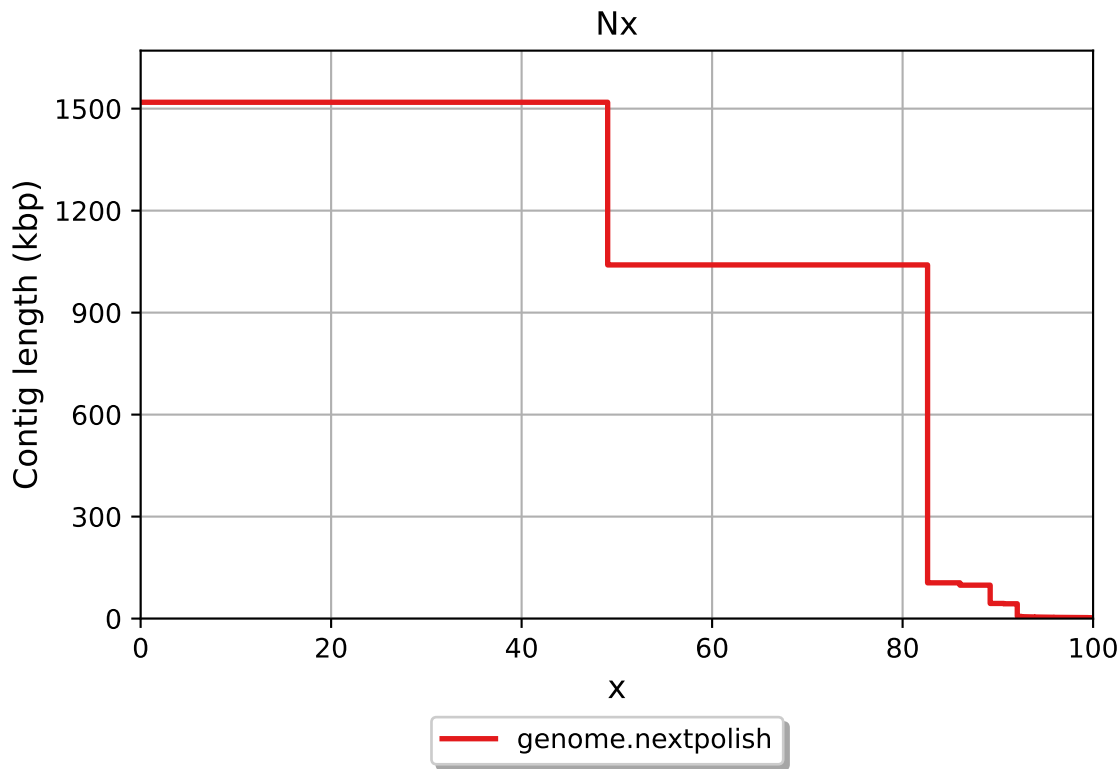


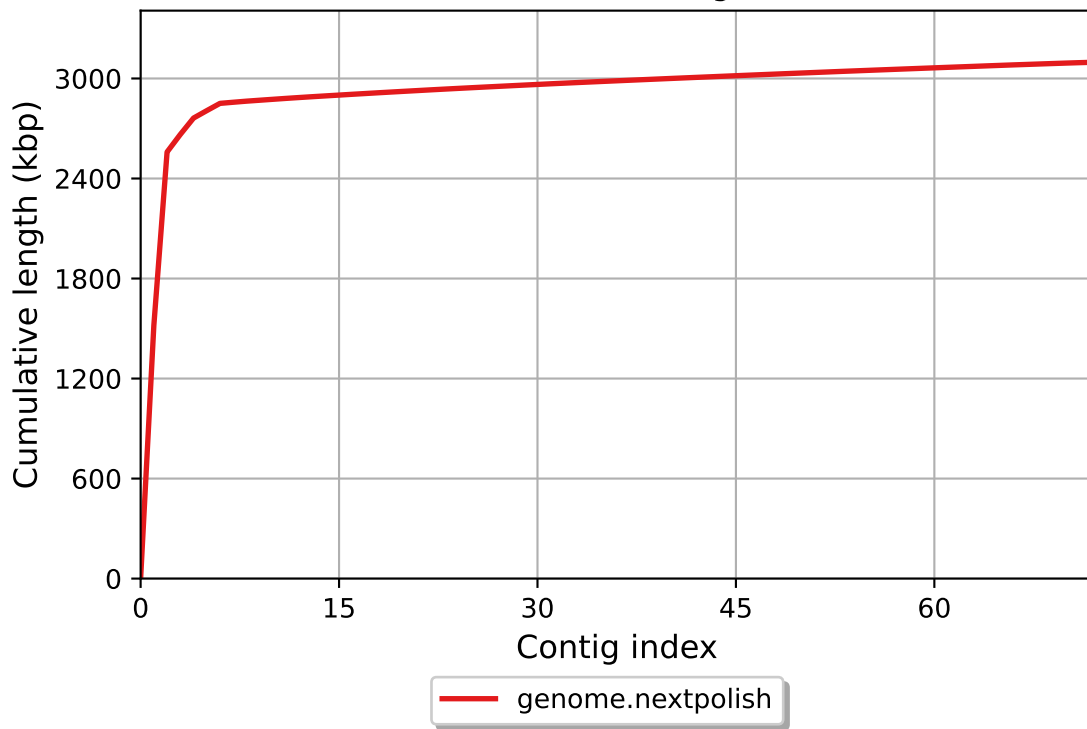
# Report

	genome.nextpolish
# contigs (>= 0 bp)	72
# contigs (>= 1000 bp)	72
# contigs (>= 5000 bp)	13
# contigs (>= 10000 bp)	6
# contigs (>= 25000 bp)	6
# contigs (>= 50000 bp)	4
Total length (>= 0 bp)	3097275
Total length (>= 1000 bp)	3097275
Total length (>= 5000 bp)	2890873
Total length (>= 10000 bp)	2850701
Total length (>= 25000 bp)	2850701
Total length (>= 50000 bp)	2762577
# contigs	72
Largest contig	1518760
Total length	3097275
GC (%)	34.33
N50	1040296
N90	44538
auN	1102402.2
L50	2
L90	5
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	8 + 7 part

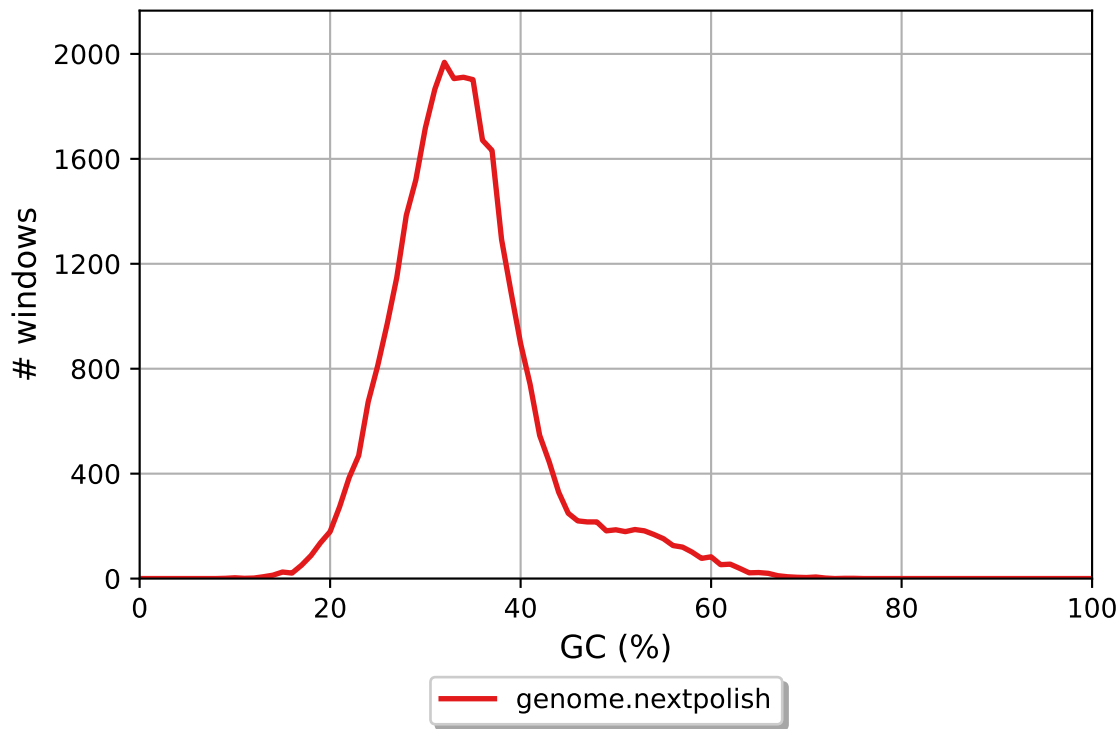
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

