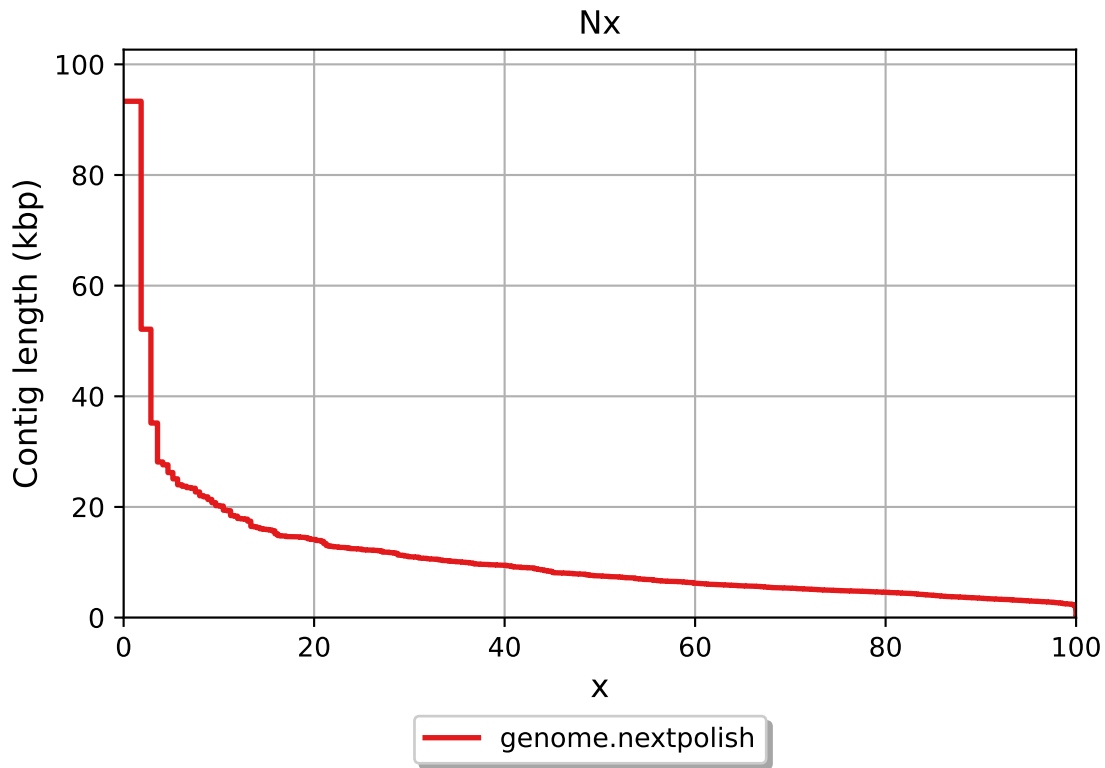


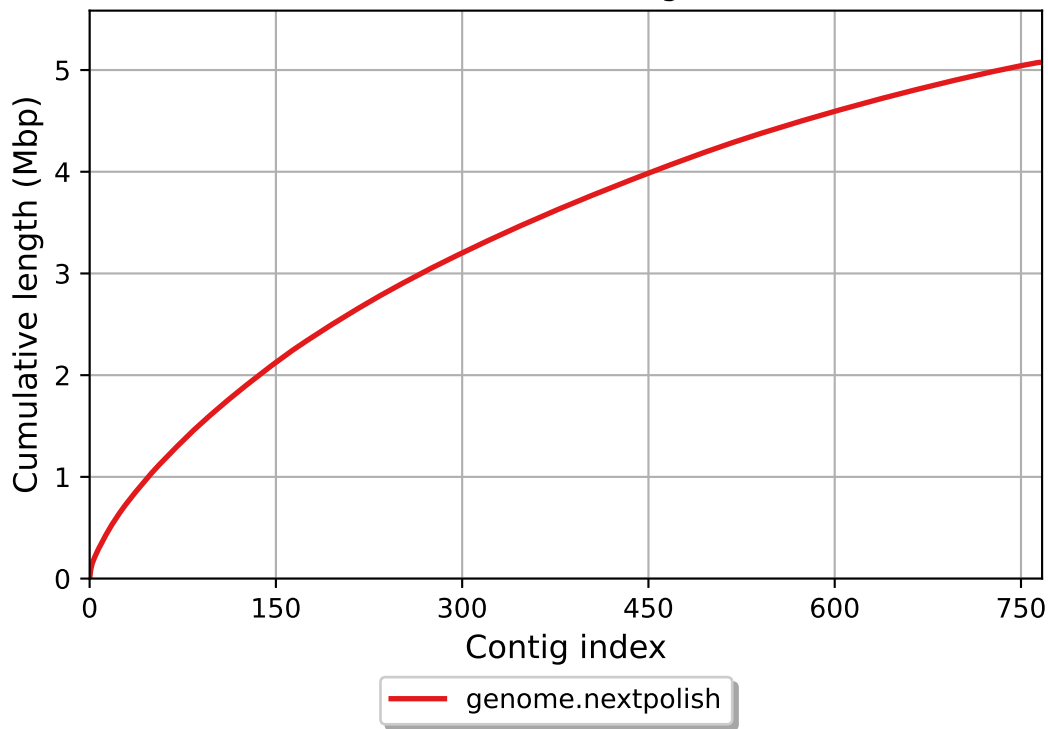
Report

	genome.nextpolish
# contigs (>= 0 bp)	767
# contigs (>= 1000 bp)	764
# contigs (>= 5000 bp)	395
# contigs (>= 10000 bp)	117
# contigs (>= 25000 bp)	7
# contigs (>= 50000 bp)	2
Total length (>= 0 bp)	5076403
Total length (>= 1000 bp)	5074480
Total length (>= 5000 bp)	3720270
Total length (>= 10000 bp)	1811418
Total length (>= 25000 bp)	287655
Total length (>= 50000 bp)	145453
# contigs	767
Largest contig	93327
Total length	5076403
GC (%)	39.64
N50	7533
N90	3504
auN	11214.1
L50	201
L90	593
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	16 + 2 part

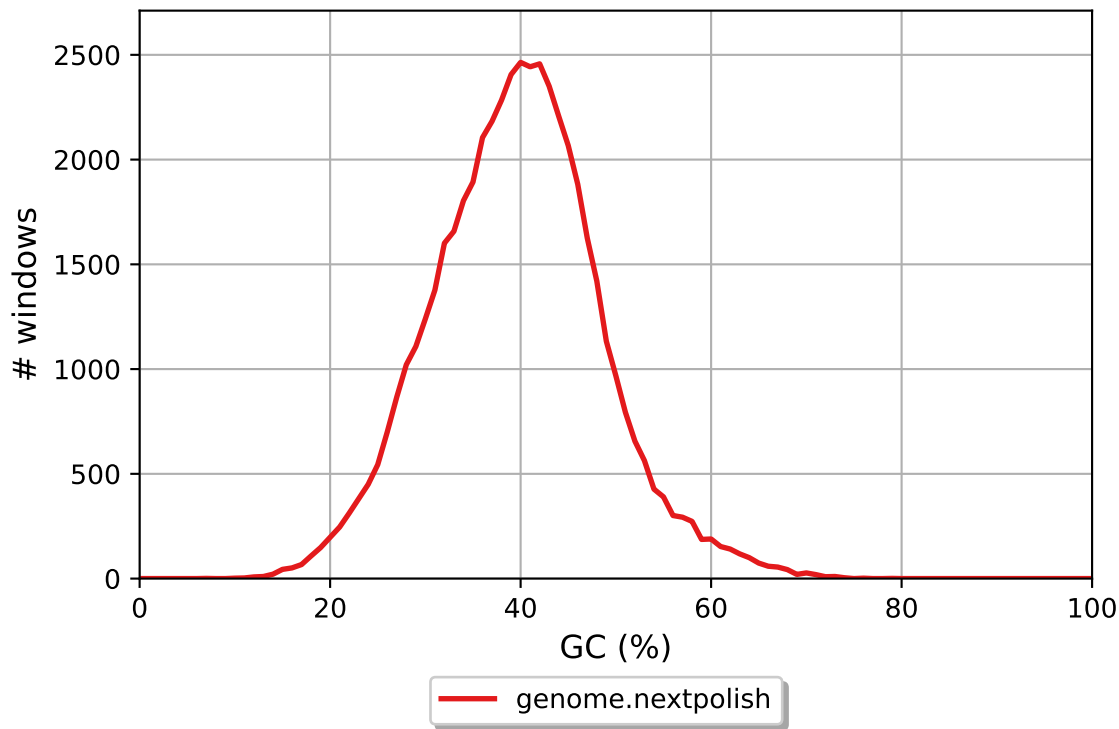
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

