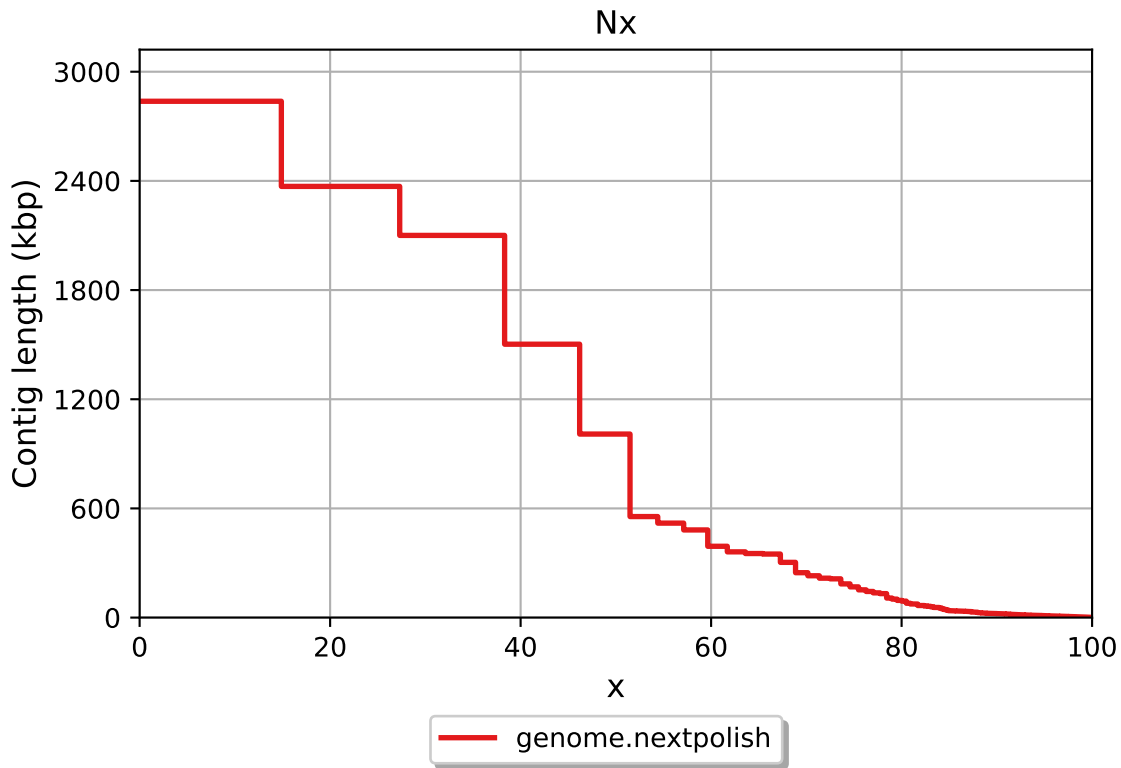


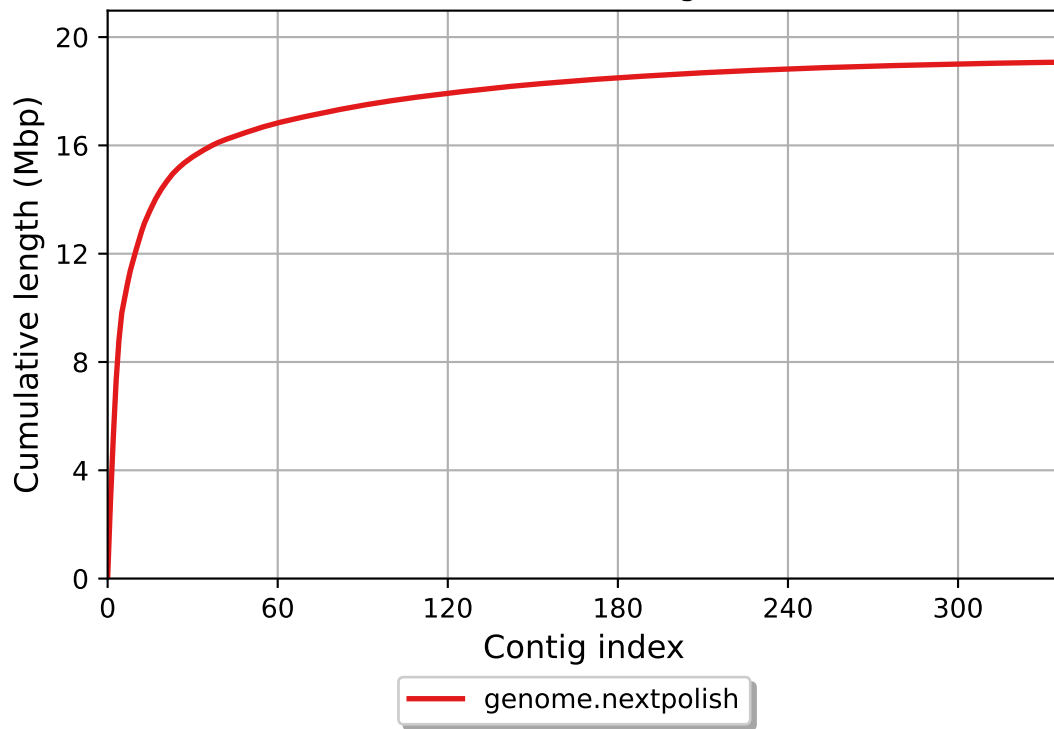
# Report

	genome.nextpolish
# contigs (>= 0 bp)	336
# contigs (>= 1000 bp)	330
# contigs (>= 5000 bp)	218
# contigs (>= 10000 bp)	144
# contigs (>= 25000 bp)	62
# contigs (>= 50000 bp)	38
Total length (>= 0 bp)	19070394
Total length (>= 1000 bp)	19066530
Total length (>= 5000 bp)	18724498
Total length (>= 10000 bp)	18198070
Total length (>= 25000 bp)	16882232
Total length (>= 50000 bp)	16060376
# contigs	336
Largest contig	2837624
Total length	19070394
GC (%)	43.91
N50	1008632
N90	21901
auN	1220561.3
L50	5
L90	75
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	90 + 1 part

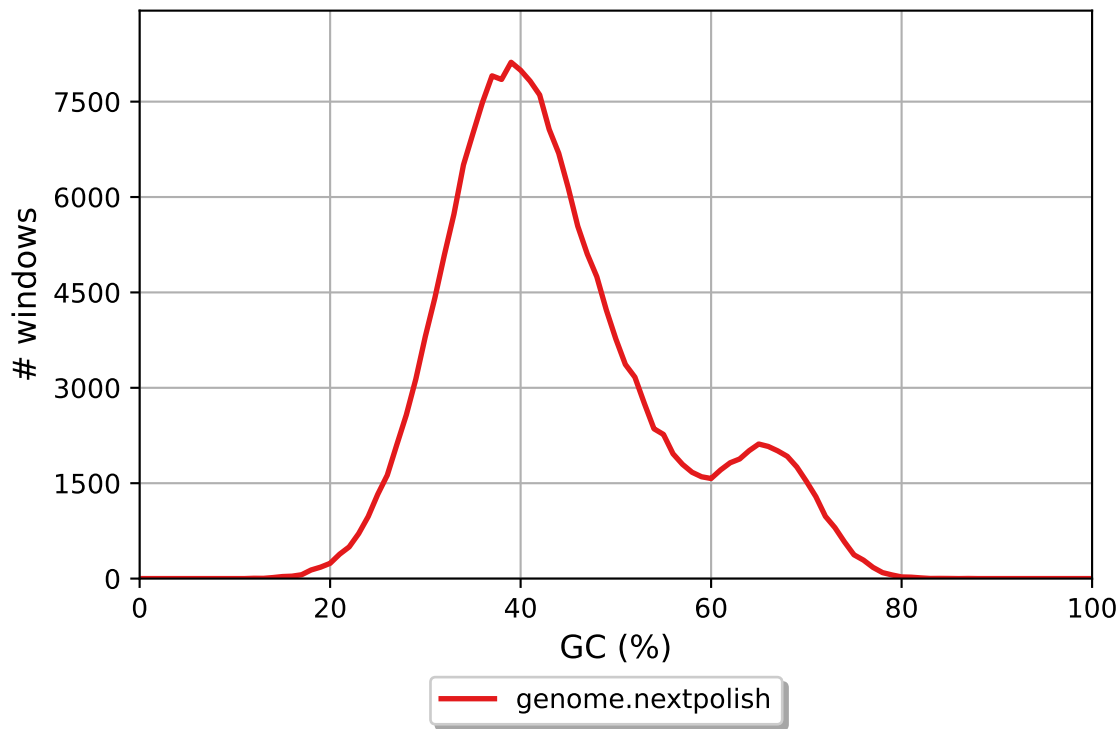
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



## GC content



genome.nextpolish GC content

