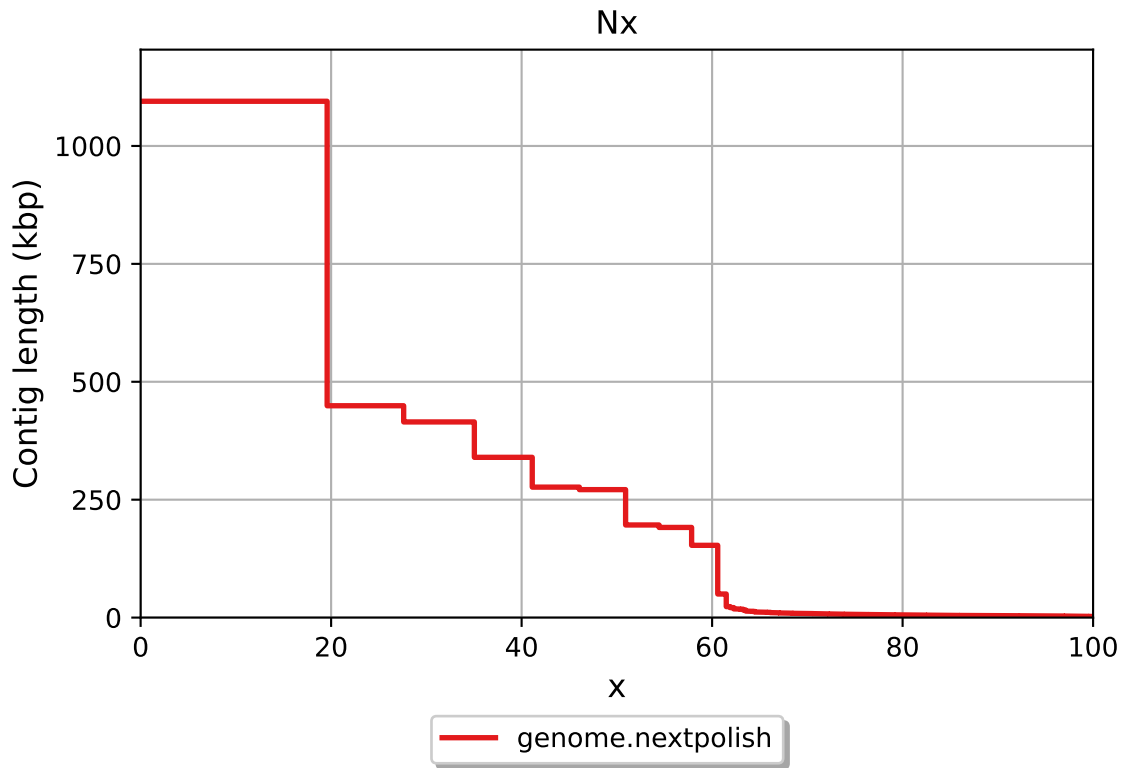


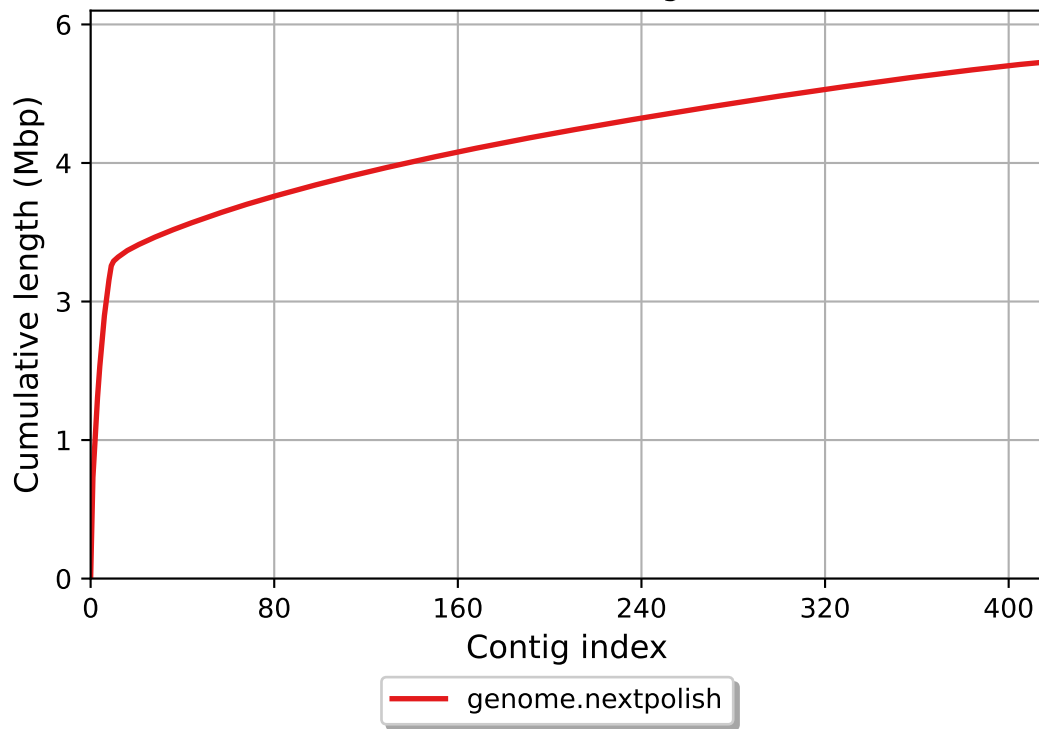
# Report

	genome.nextpolish
# contigs (>= 0 bp)	415
# contigs (>= 1000 bp)	414
# contigs (>= 5000 bp)	167
# contigs (>= 10000 bp)	33
# contigs (>= 25000 bp)	10
# contigs (>= 50000 bp)	9
Total length (>= 0 bp)	5590454
Total length (>= 1000 bp)	5589483
Total length (>= 5000 bp)	4654850
Total length (>= 10000 bp)	3748545
Total length (>= 25000 bp)	3437259
Total length (>= 50000 bp)	3387260
# contigs	415
Largest contig	1094760
Total length	5590454
GC (%)	35.72
N50	271315
N90	4092
auN	349445.7
L50	6
L90	251
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	19 + 3 part

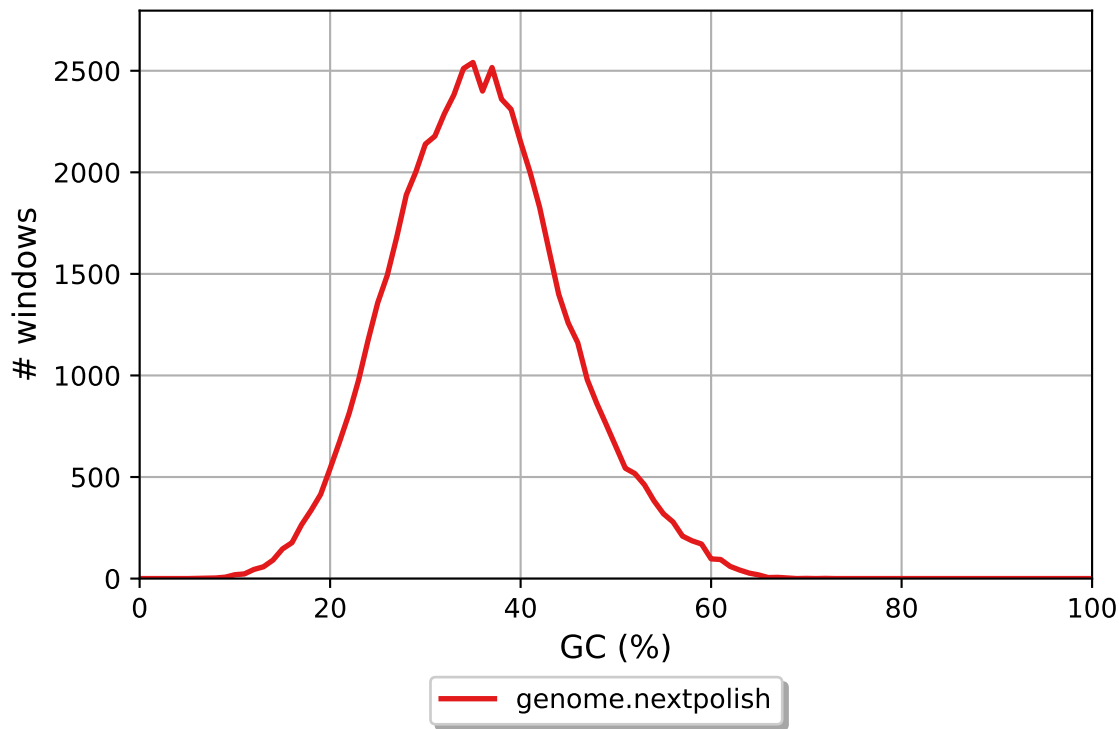
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



## GC content



genome.nextpolish GC content

