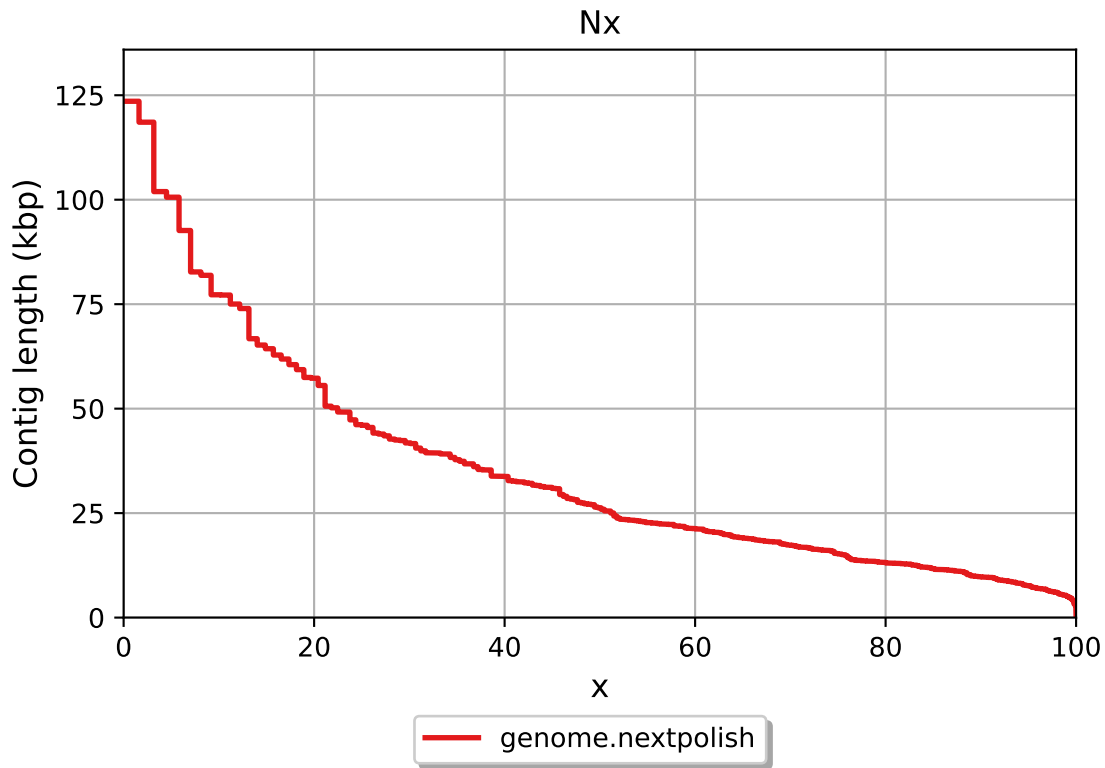


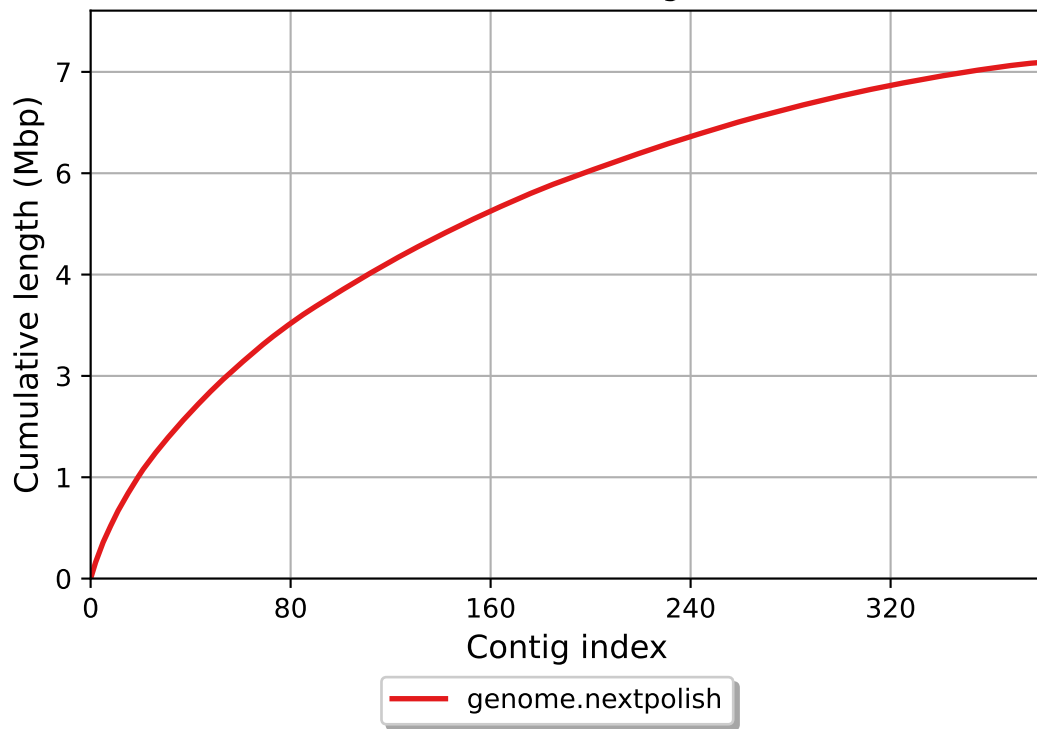
# Report

	genome.nextpolish
# contigs (>= 0 bp)	381
# contigs (>= 1000 bp)	381
# contigs (>= 5000 bp)	365
# contigs (>= 10000 bp)	263
# contigs (>= 25000 bp)	86
# contigs (>= 50000 bp)	23
Total length (>= 0 bp)	7642432
Total length (>= 1000 bp)	7642432
Total length (>= 5000 bp)	7579458
Total length (>= 10000 bp)	6797907
Total length (>= 25000 bp)	3932854
Total length (>= 50000 bp)	1717258
# contigs	381
Largest contig	123555
Total length	7642432
GC (%)	57.99
N50	26282
N90	9741
auN	35841.9
L50	82
L90	272
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	27 + 0 part

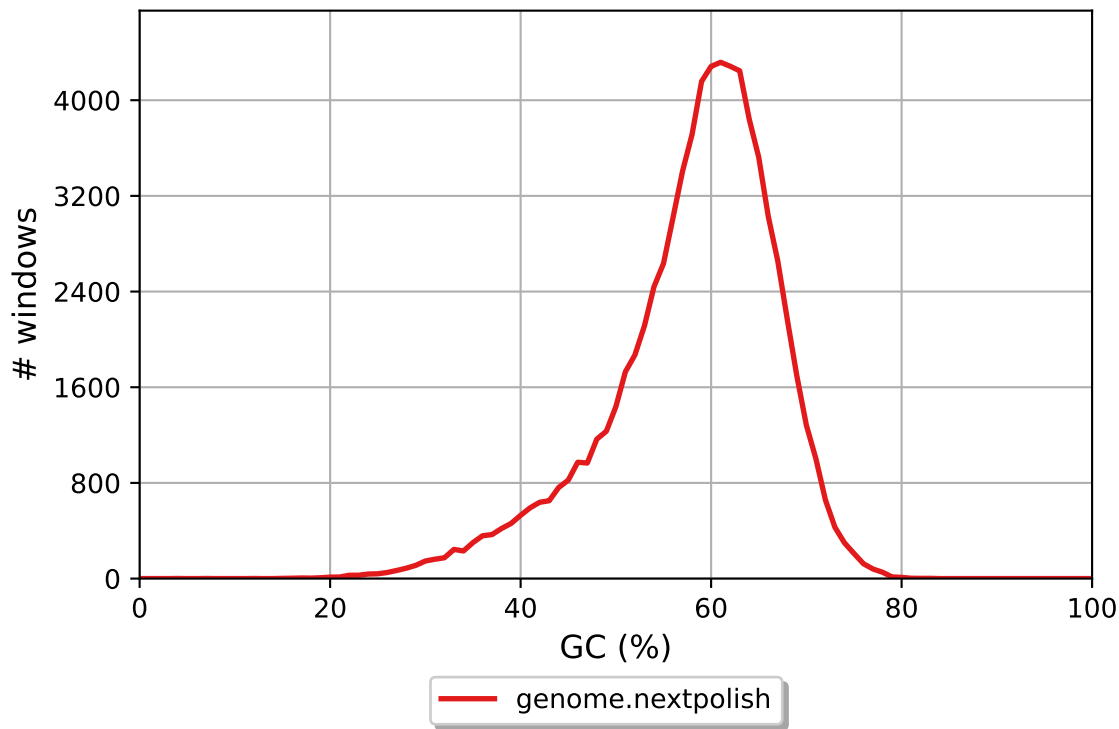
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



## GC content



genome.nextpolish GC content

