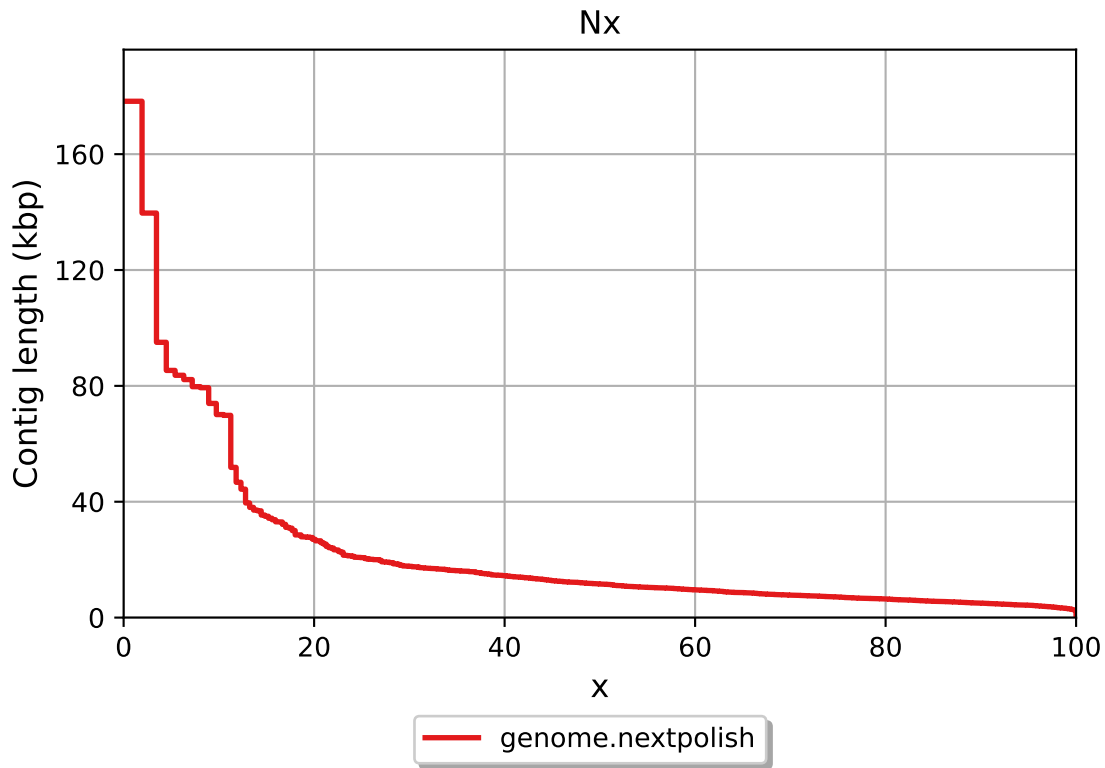


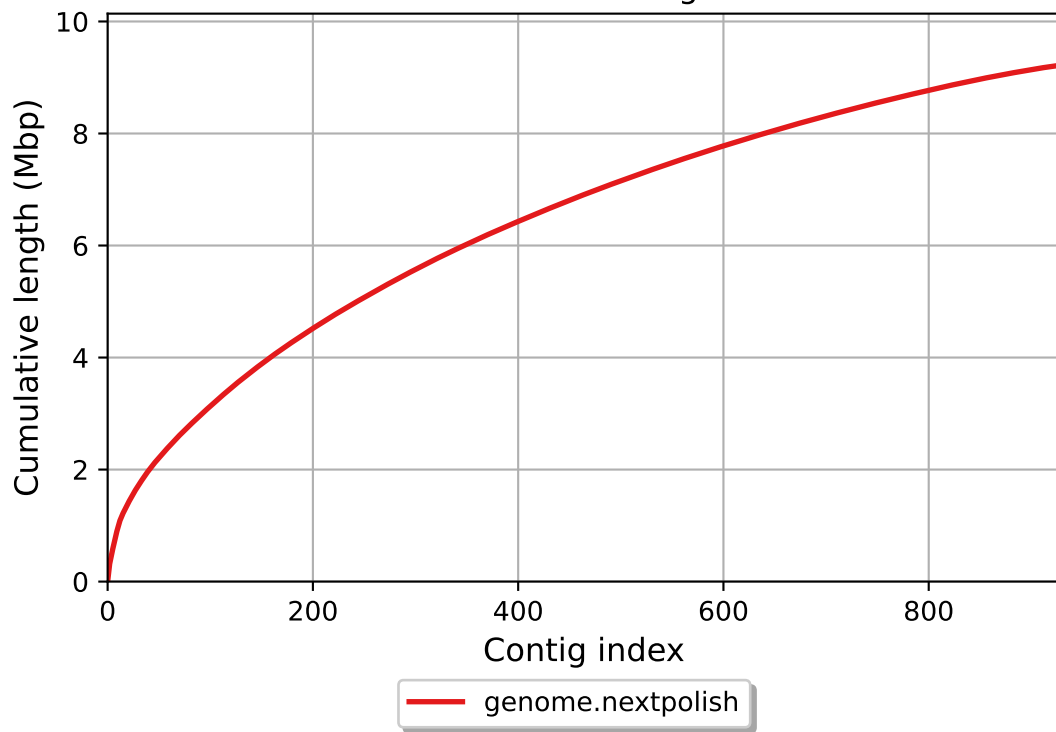
# Report

	genome.nextpolish
# contigs (>= 0 bp)	928
# contigs (>= 1000 bp)	928
# contigs (>= 5000 bp)	690
# contigs (>= 10000 bp)	277
# contigs (>= 25000 bp)	39
# contigs (>= 50000 bp)	12
Total length (>= 0 bp)	9218144
Total length (>= 1000 bp)	9218144
Total length (>= 5000 bp)	8263710
Total length (>= 10000 bp)	5350035
Total length (>= 25000 bp)	1959856
Total length (>= 50000 bp)	1088900
# contigs	928
Largest contig	178258
Total length	9218144
GC (%)	54.16
N50	11593
N90	4978
auN	23578.6
L50	208
L90	697
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	48 + 7 part

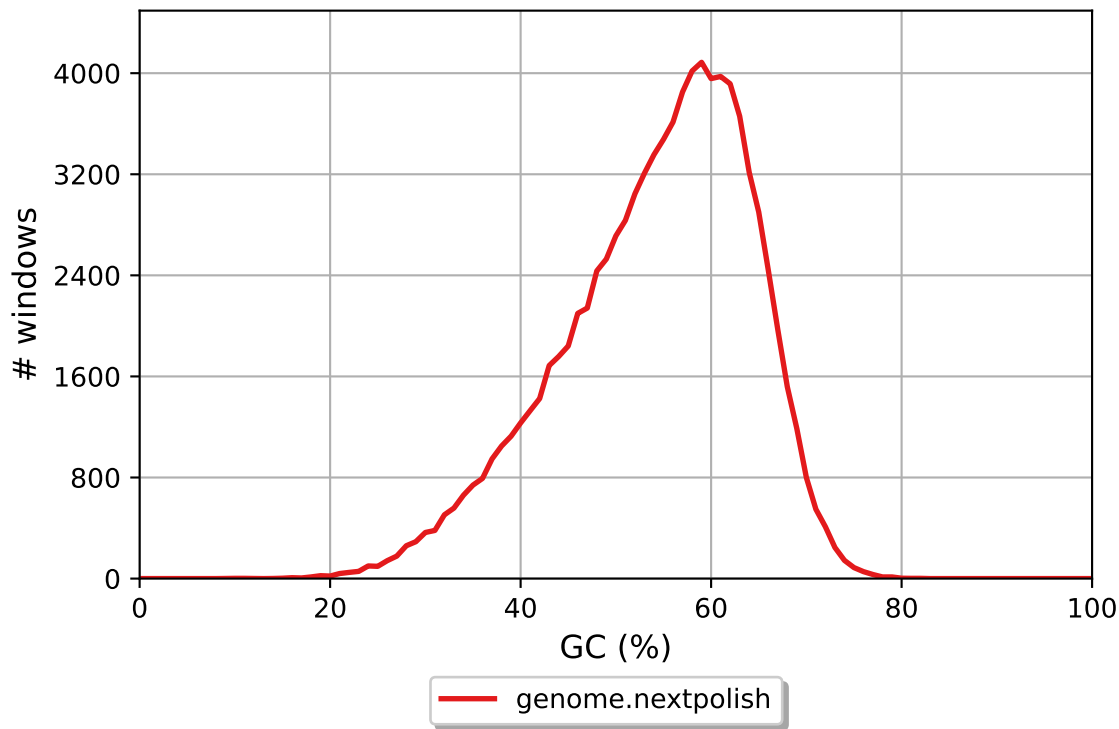
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



## GC content



genome.nextpolish GC content

