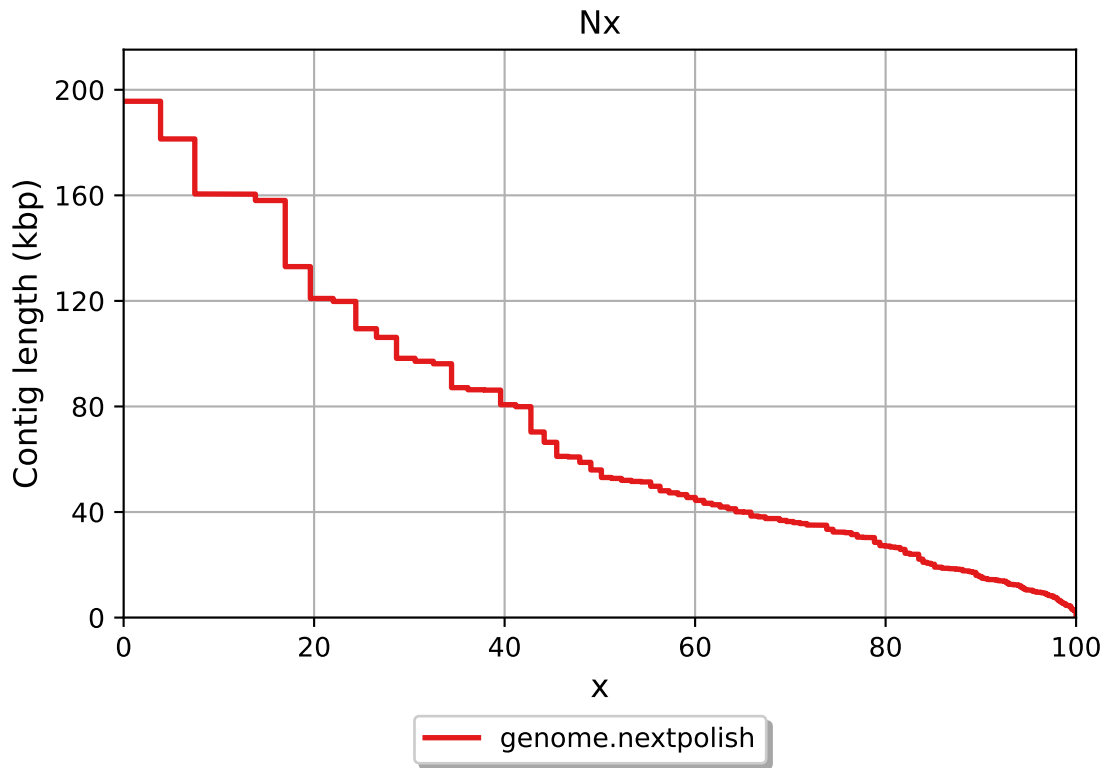


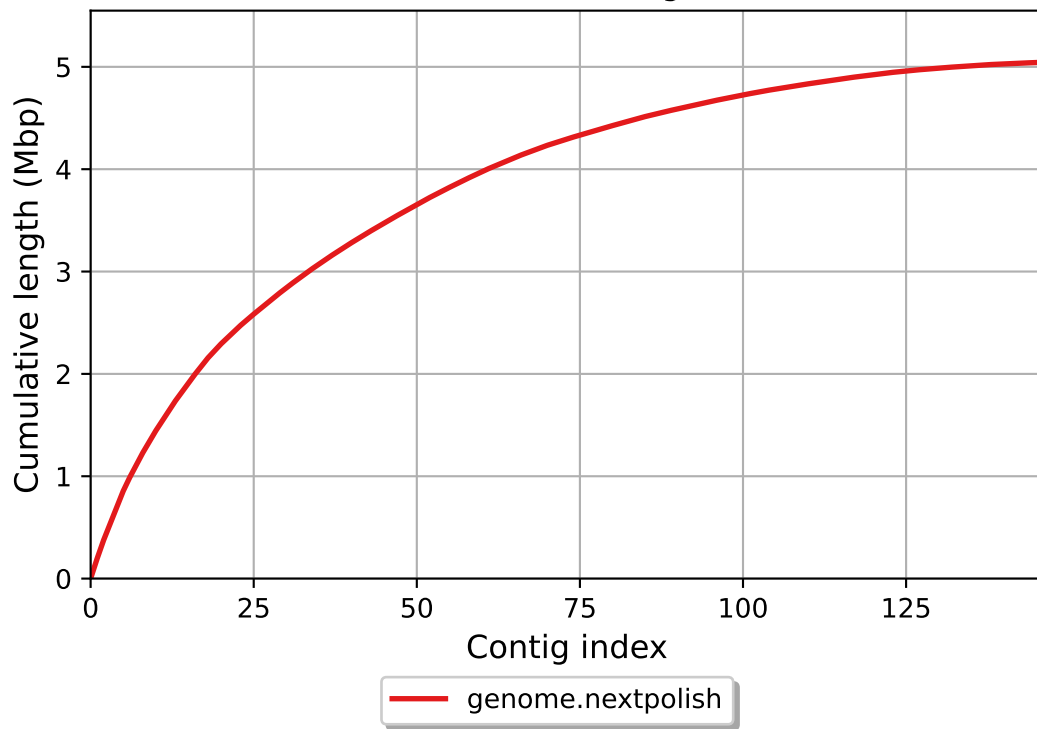
# Report

	genome.nextpolish
# contigs (>= 0 bp)	146
# contigs (>= 1000 bp)	146
# contigs (>= 5000 bp)	130
# contigs (>= 10000 bp)	108
# contigs (>= 25000 bp)	66
# contigs (>= 50000 bp)	29
Total length (>= 0 bp)	5044709
Total length (>= 1000 bp)	5044709
Total length (>= 5000 bp)	4987820
Total length (>= 10000 bp)	4814232
Total length (>= 25000 bp)	4138788
Total length (>= 50000 bp)	2791423
# contigs	146
Largest contig	195646
Total length	5044709
GC (%)	55.30
N50	55945
N90	15545
auN	75733.9
L50	24
L90	87
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	13 + 4 part

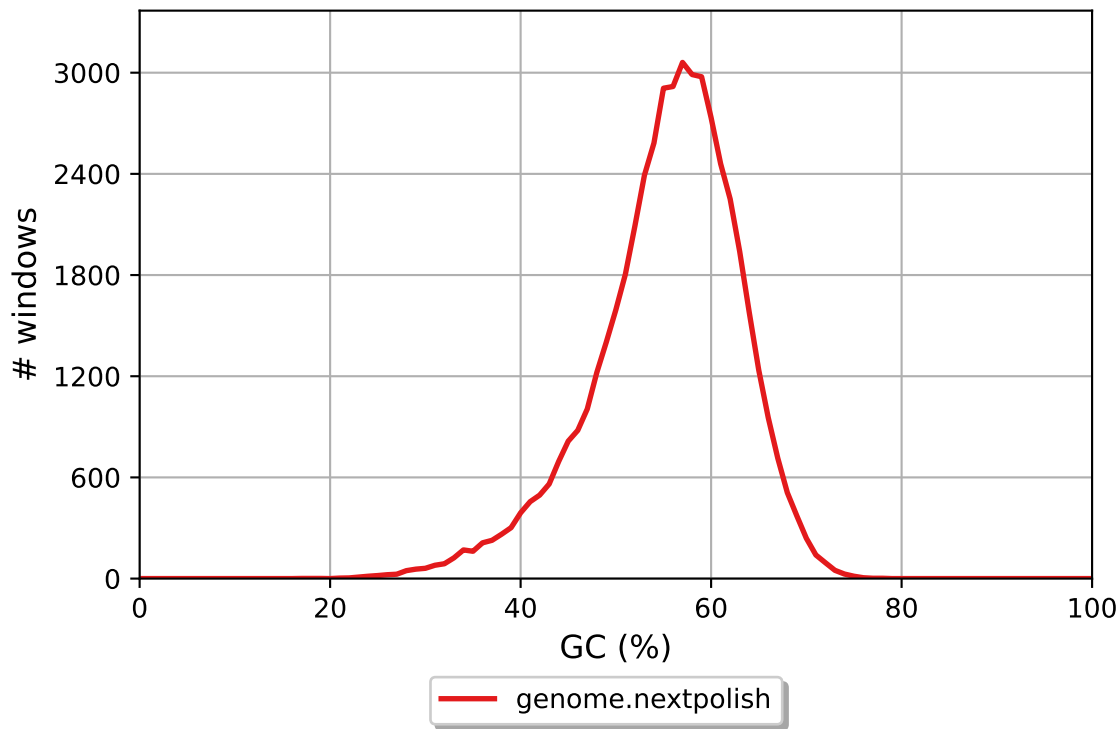
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



## GC content



genome.nextpolish GC content

