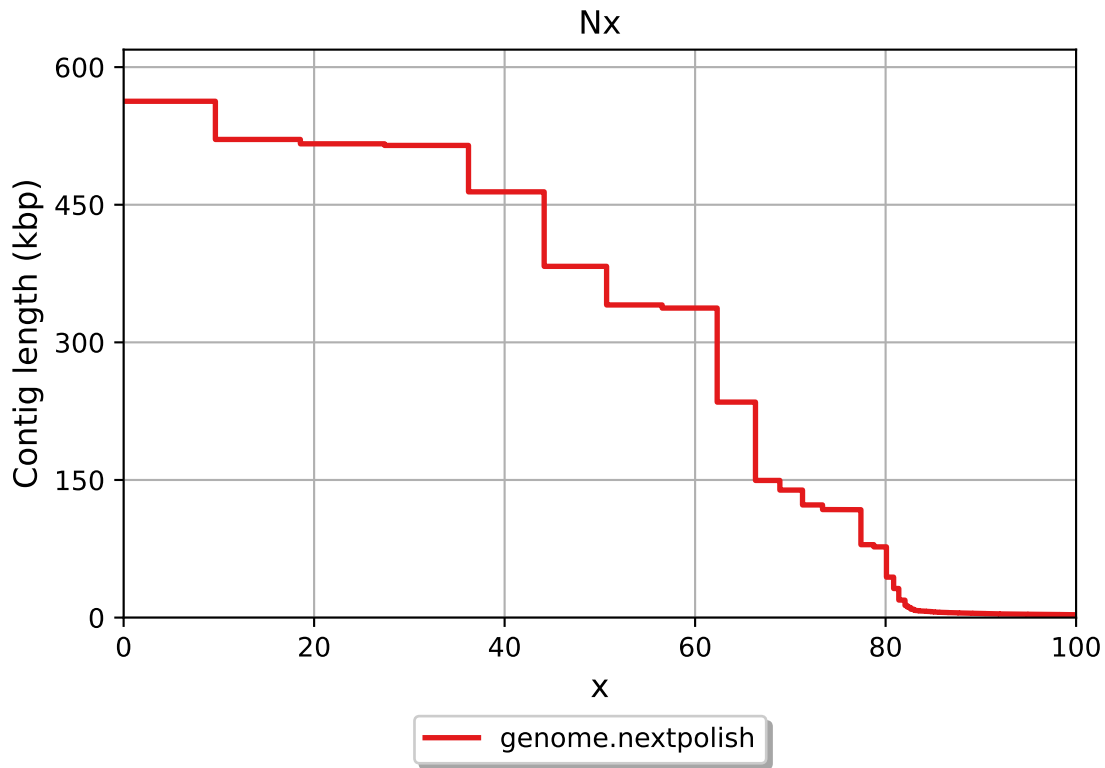


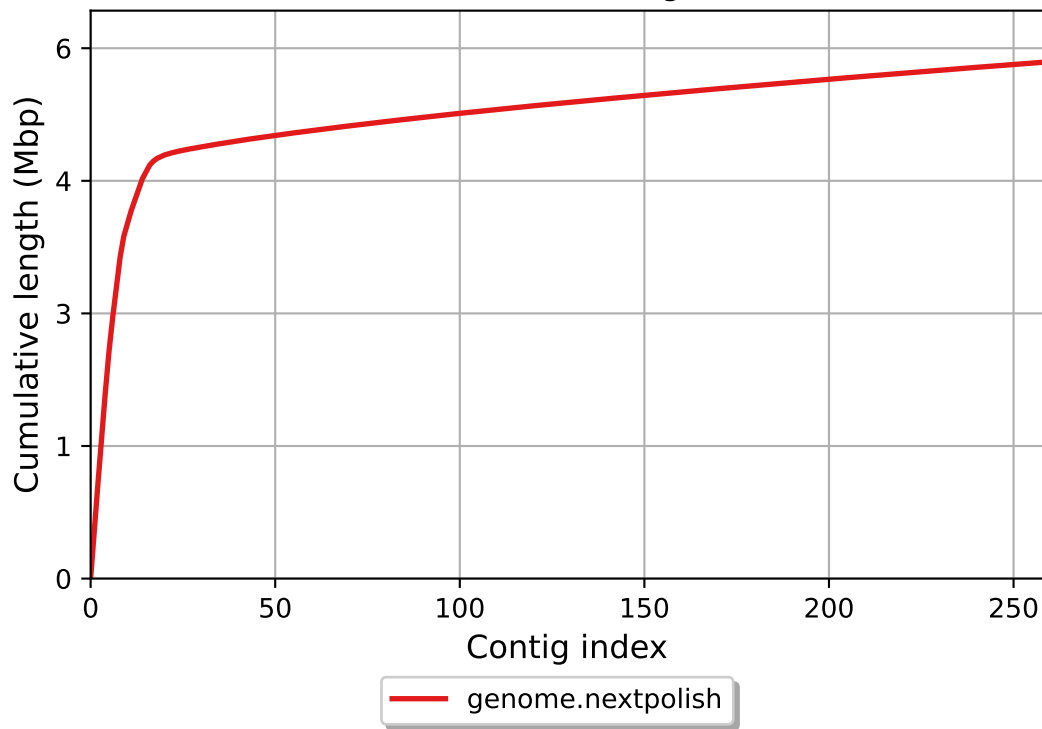
# Report

	genome.nextpolish
# contigs (>= 0 bp)	258
# contigs (>= 1000 bp)	258
# contigs (>= 5000 bp)	72
# contigs (>= 10000 bp)	23
# contigs (>= 25000 bp)	18
# contigs (>= 50000 bp)	16
Total length (>= 0 bp)	5841179
Total length (>= 1000 bp)	5841179
Total length (>= 5000 bp)	5131302
Total length (>= 10000 bp)	4827636
Total length (>= 25000 bp)	4753851
Total length (>= 50000 bp)	4678083
# contigs	258
Largest contig	562783
Total length	5841179
GC (%)	56.24
N50	382850
N90	4448
auN	320550.8
L50	6
L90	99
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	18 + 2 part

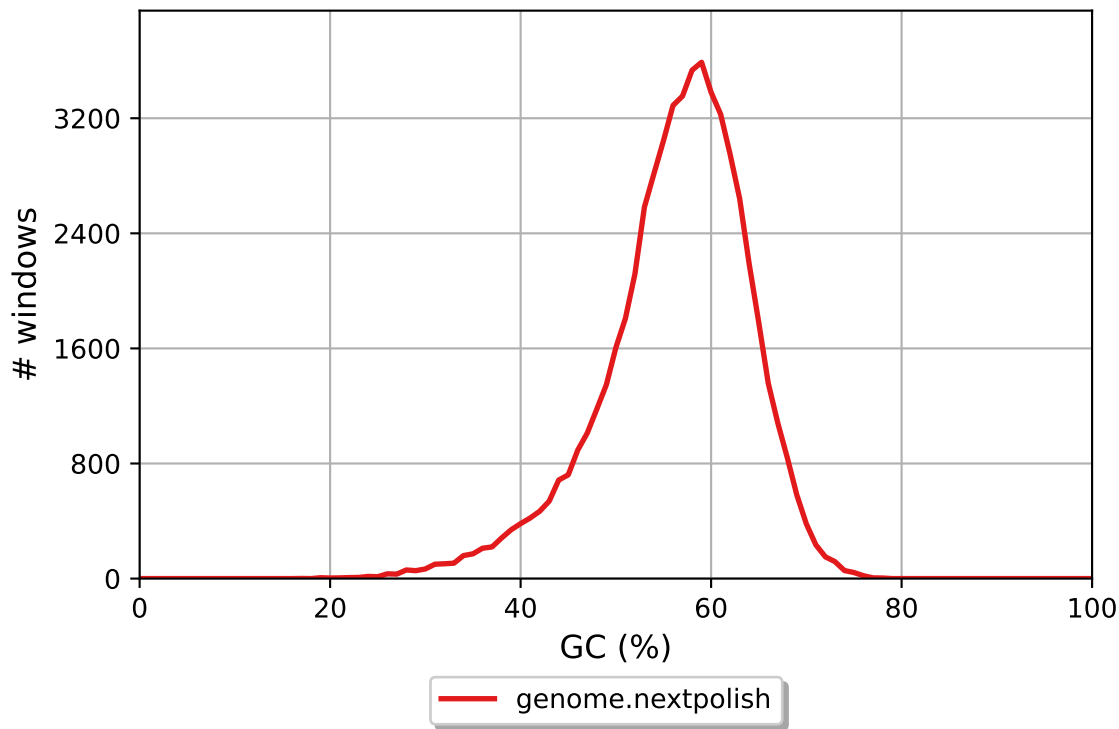
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



## GC content



genome.nextpolish GC content

