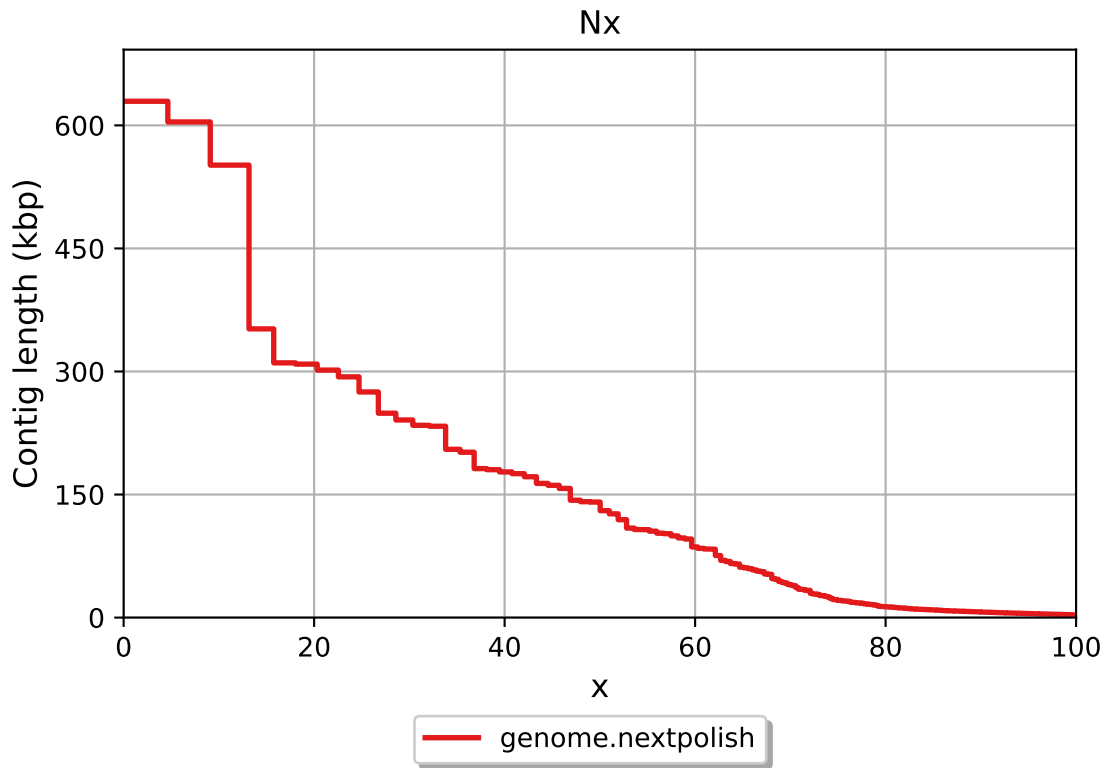


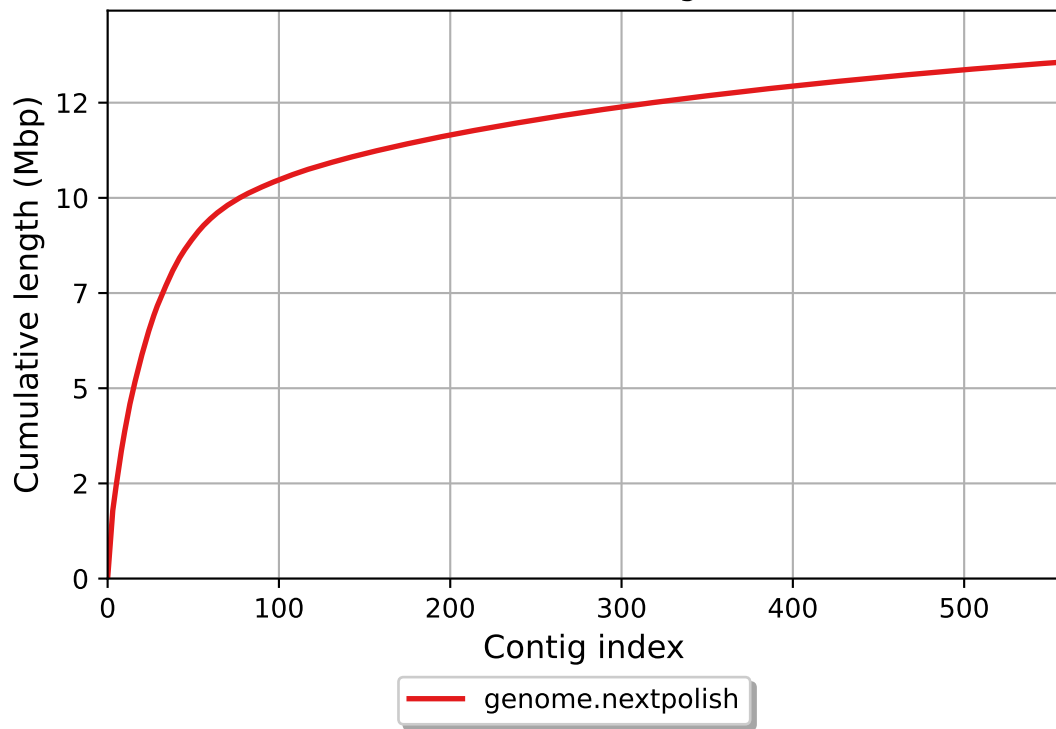
# Report

	genome.nextpolish
# contigs (>= 0 bp)	556
# contigs (>= 1000 bp)	556
# contigs (>= 5000 bp)	383
# contigs (>= 10000 bp)	169
# contigs (>= 25000 bp)	79
# contigs (>= 50000 bp)	55
Total length (>= 0 bp)	13561610
Total length (>= 1000 bp)	13561610
Total length (>= 5000 bp)	12854447
Total length (>= 10000 bp)	11360437
Total length (>= 25000 bp)	10050076
Total length (>= 50000 bp)	9228311
# contigs	556
Largest contig	629390
Total length	13561610
GC (%)	51.98
N50	140808
N90	6803
auN	186257.4
L50	26
L90	272
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	28 + 3 part

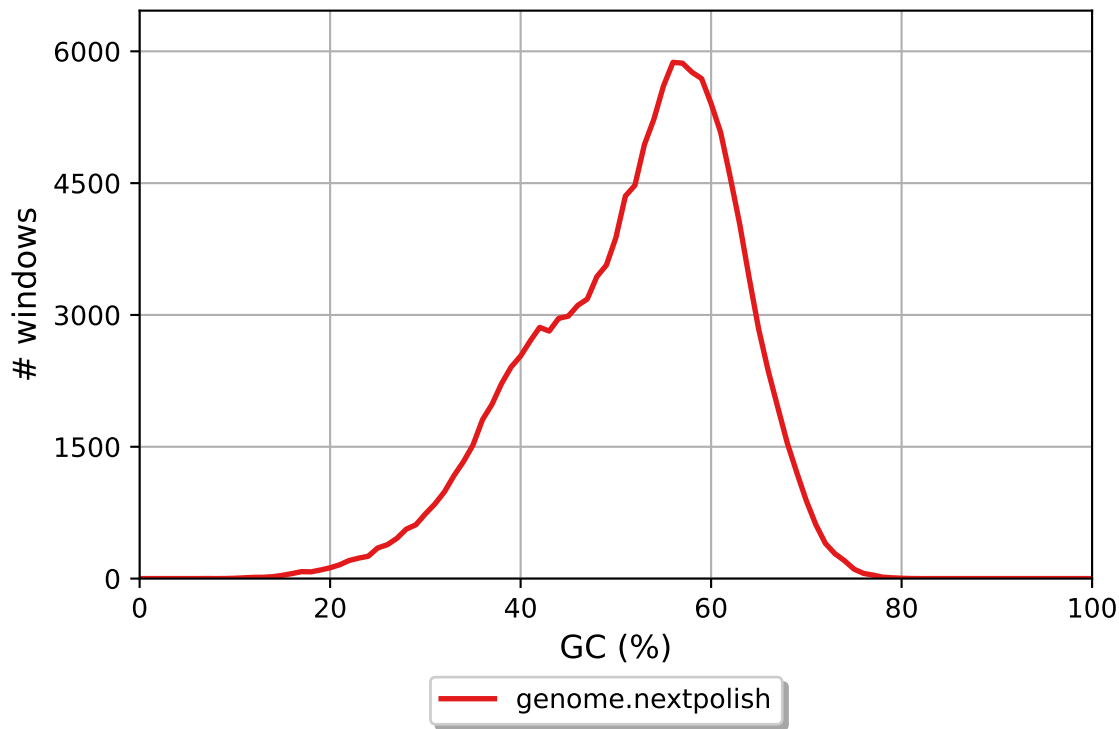
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



## GC content



genome.nextpolish GC content

