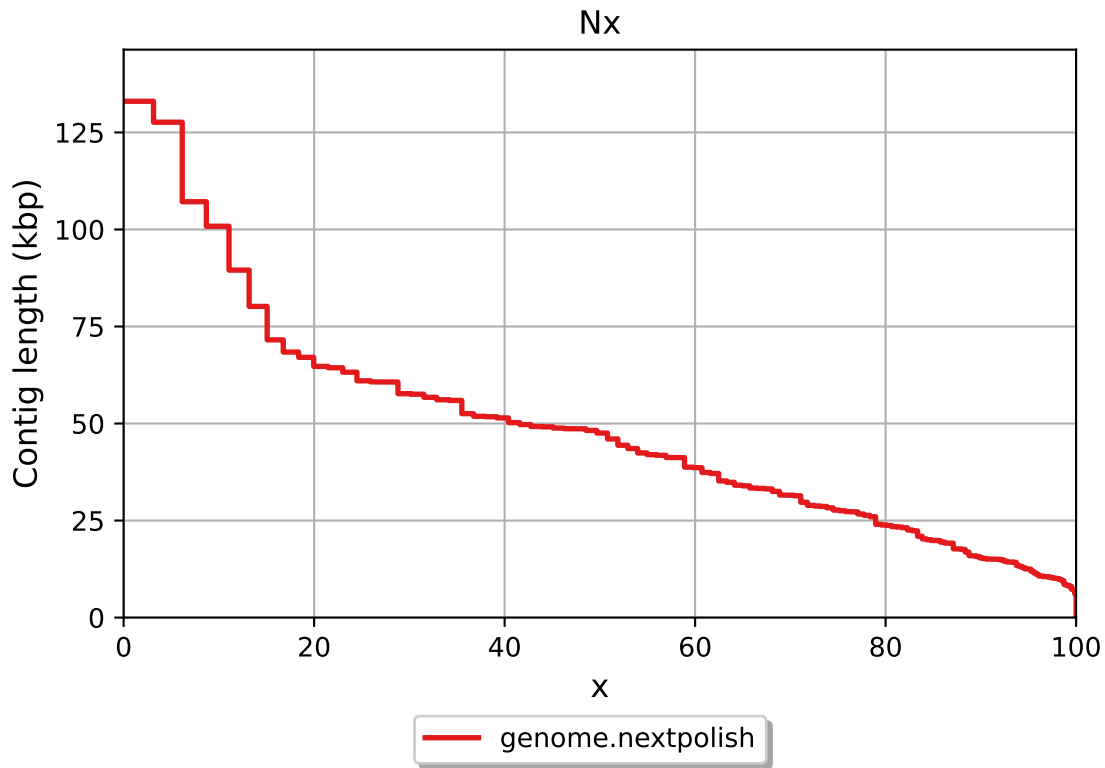


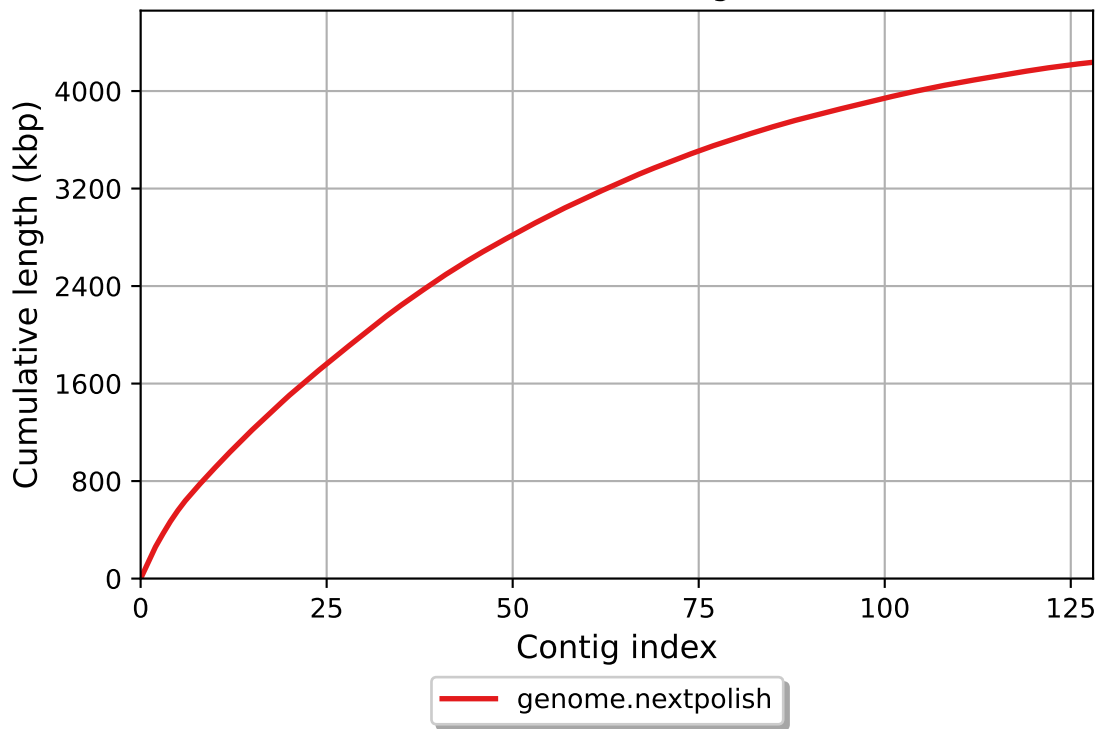
Report

	genome.nextpolish
# contigs (>= 0 bp)	128
# contigs (>= 1000 bp)	128
# contigs (>= 5000 bp)	128
# contigs (>= 10000 bp)	118
# contigs (>= 25000 bp)	68
# contigs (>= 50000 bp)	25
Total length (>= 0 bp)	4236108
Total length (>= 1000 bp)	4236108
Total length (>= 5000 bp)	4236108
Total length (>= 10000 bp)	4153013
Total length (>= 25000 bp)	3344535
Total length (>= 50000 bp)	1762046
# contigs	128
Largest contig	133032
Total length	4236108
GC (%)	57.28
N50	47544
N90	15386
auN	50237.7
L50	33
L90	92
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	4 + 0 part

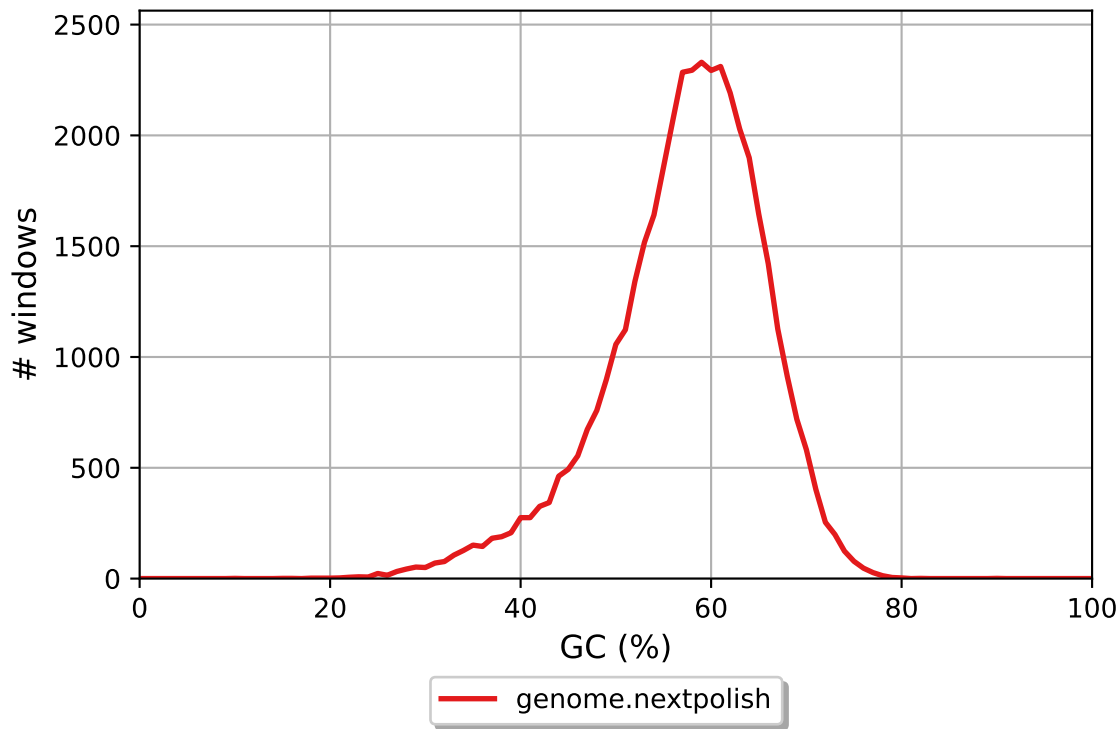
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

