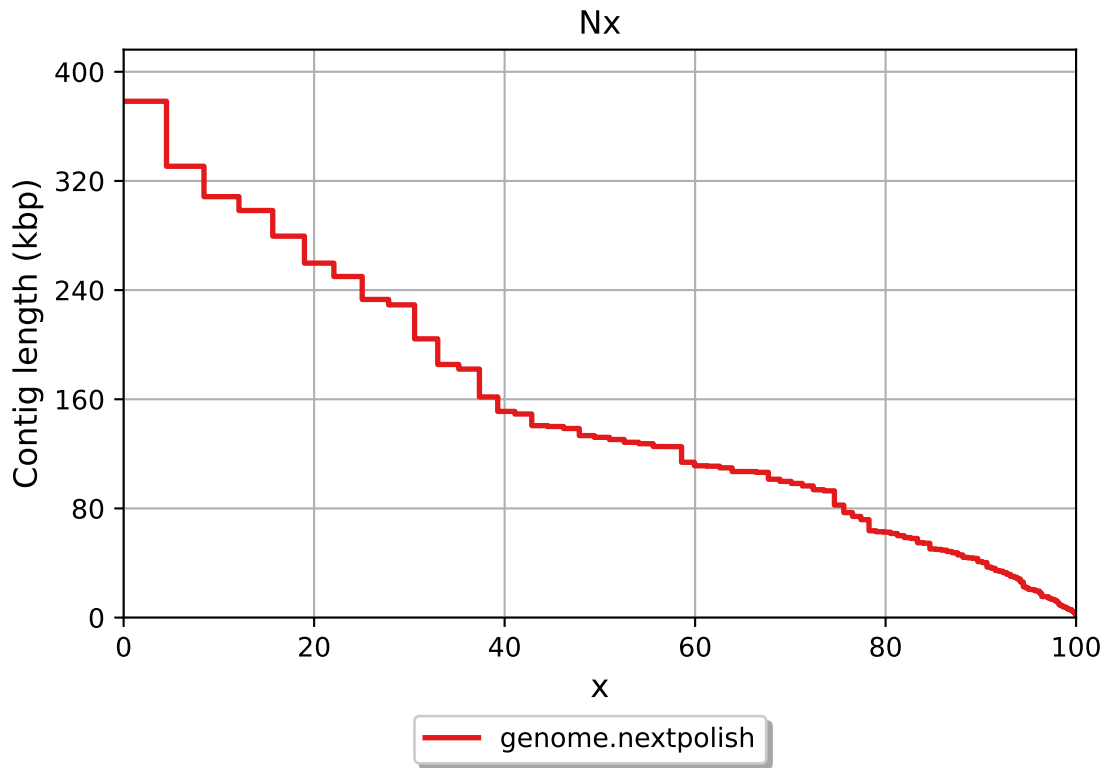


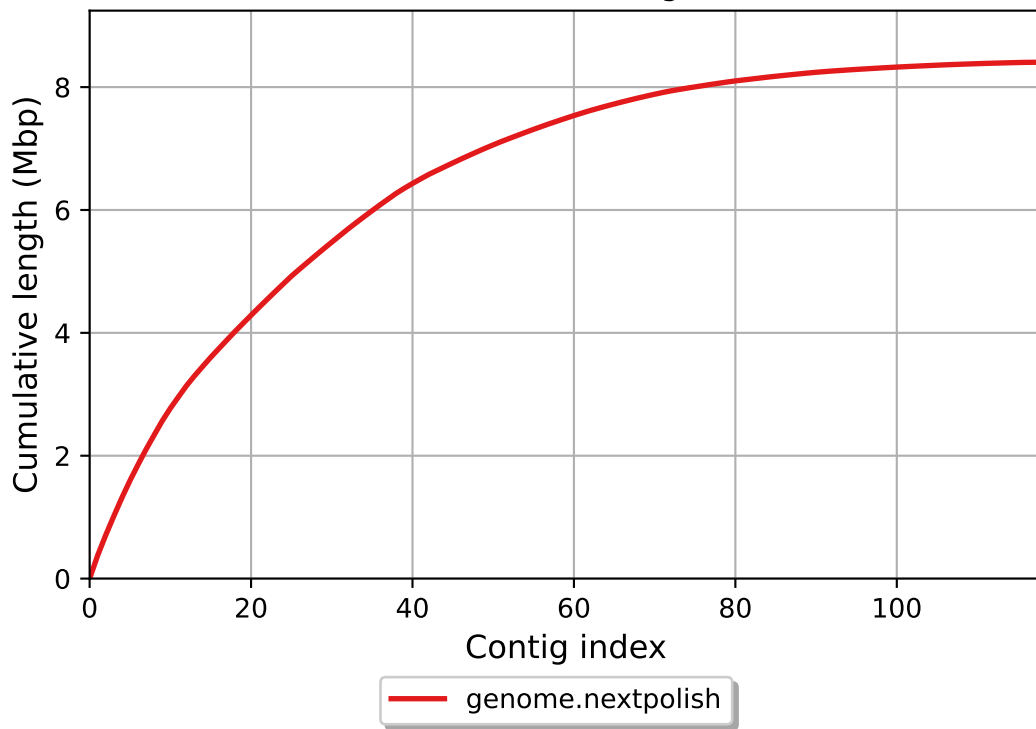
Report

	genome.nextpolish
# contigs (>= 0 bp)	119
# contigs (>= 1000 bp)	117
# contigs (>= 5000 bp)	107
# contigs (>= 10000 bp)	91
# contigs (>= 25000 bp)	72
# contigs (>= 50000 bp)	52
Total length (>= 0 bp)	8405061
Total length (>= 1000 bp)	8404065
Total length (>= 5000 bp)	8367995
Total length (>= 10000 bp)	8252256
Total length (>= 25000 bp)	7938720
Total length (>= 50000 bp)	7164441
# contigs	118
Largest contig	378389
Total length	8404578
GC (%)	57.90
N50	132089
N90	41076
auN	159835.3
L50	20
L90	61
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	34 + 4 part

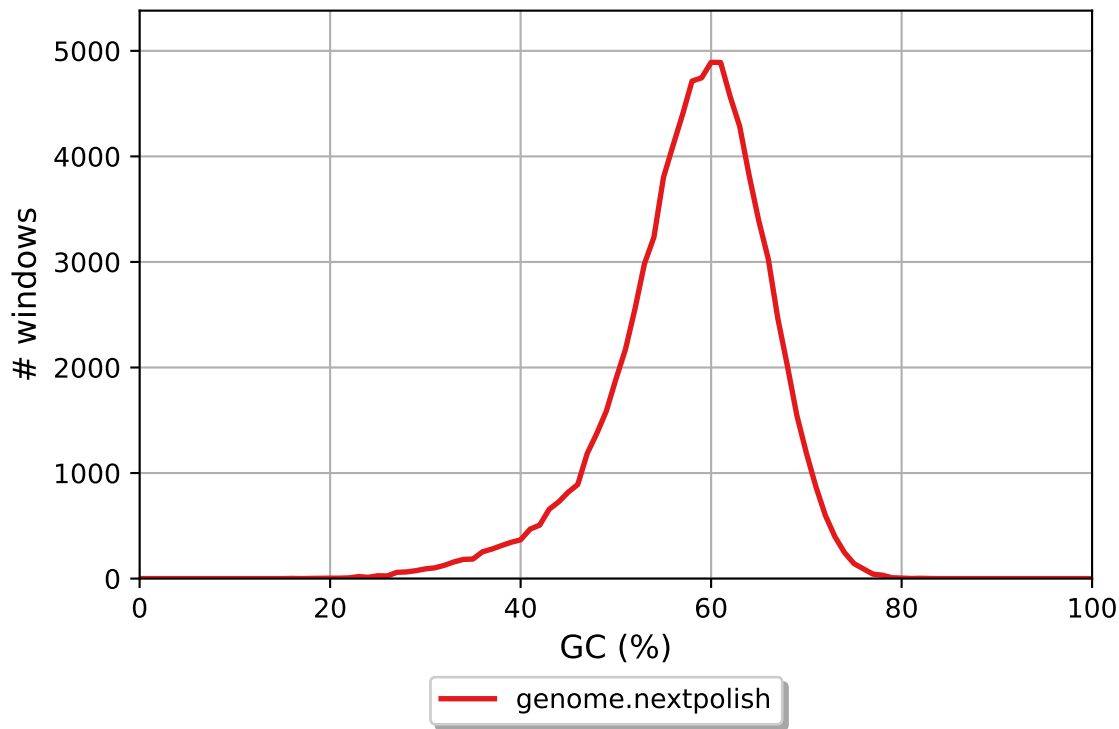
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

