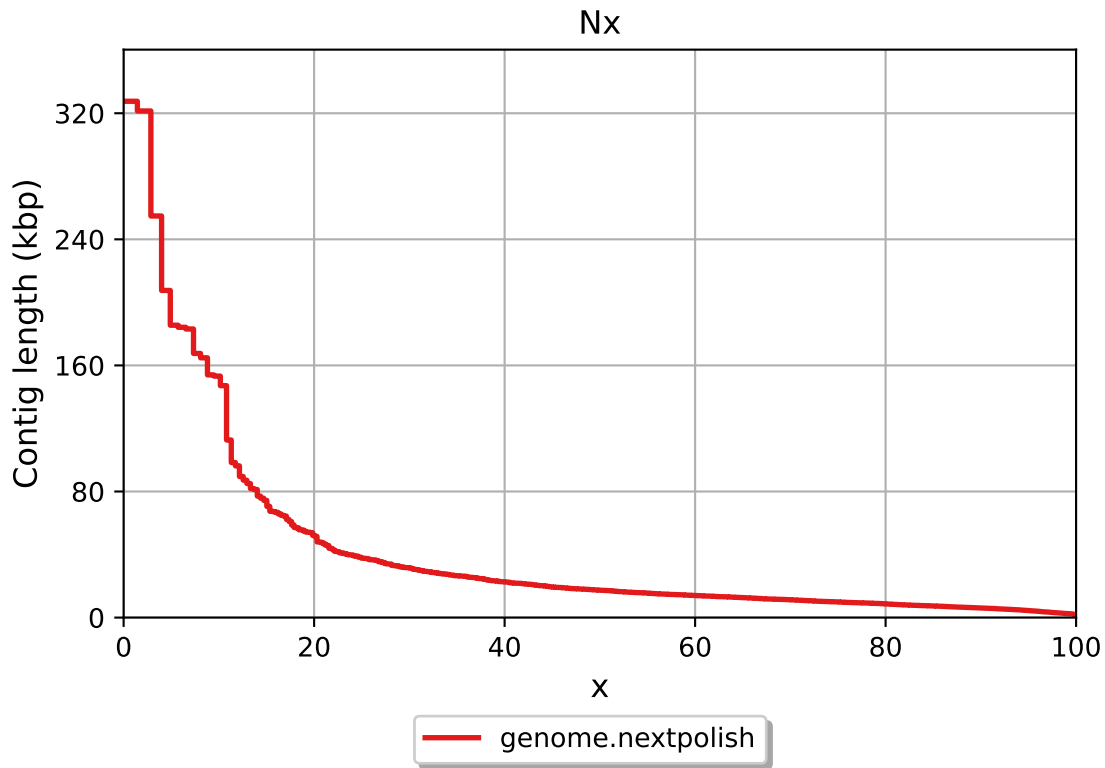


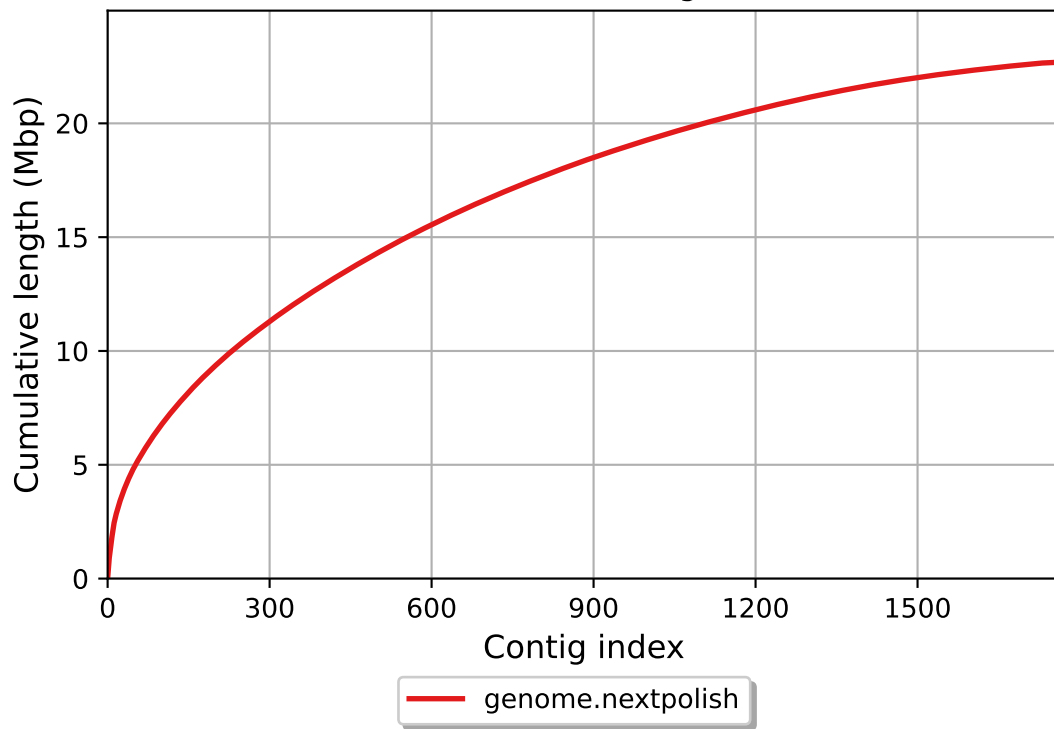
# Report

	genome.nextpolish
# contigs (>= 0 bp)	1764
# contigs (>= 1000 bp)	1745
# contigs (>= 5000 bp)	1320
# contigs (>= 10000 bp)	733
# contigs (>= 25000 bp)	160
# contigs (>= 50000 bp)	43
Total length (>= 0 bp)	22683549
Total length (>= 1000 bp)	22670529
Total length (>= 5000 bp)	21248753
Total length (>= 10000 bp)	16979130
Total length (>= 25000 bp)	8438499
Total length (>= 50000 bp)	4605436
# contigs	1764
Largest contig	327600
Total length	22683549
GC (%)	49.17
N50	17399
N90	6100
auN	44280.2
L50	303
L90	1171
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	71 + 8 part

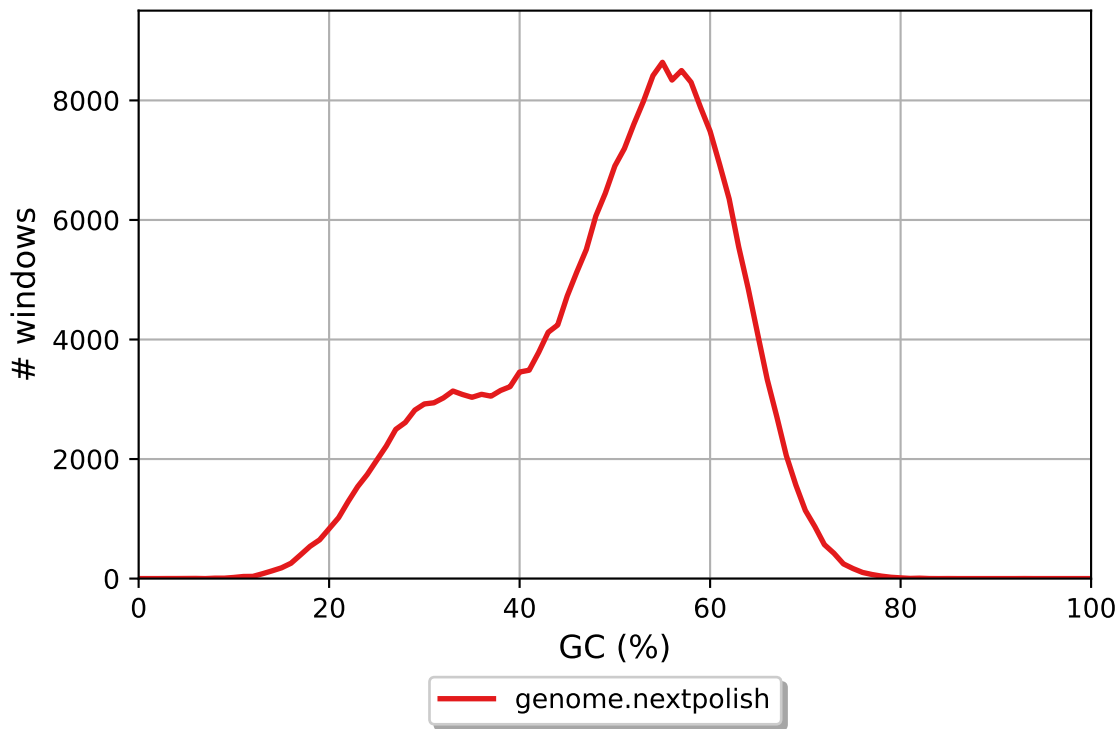
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

