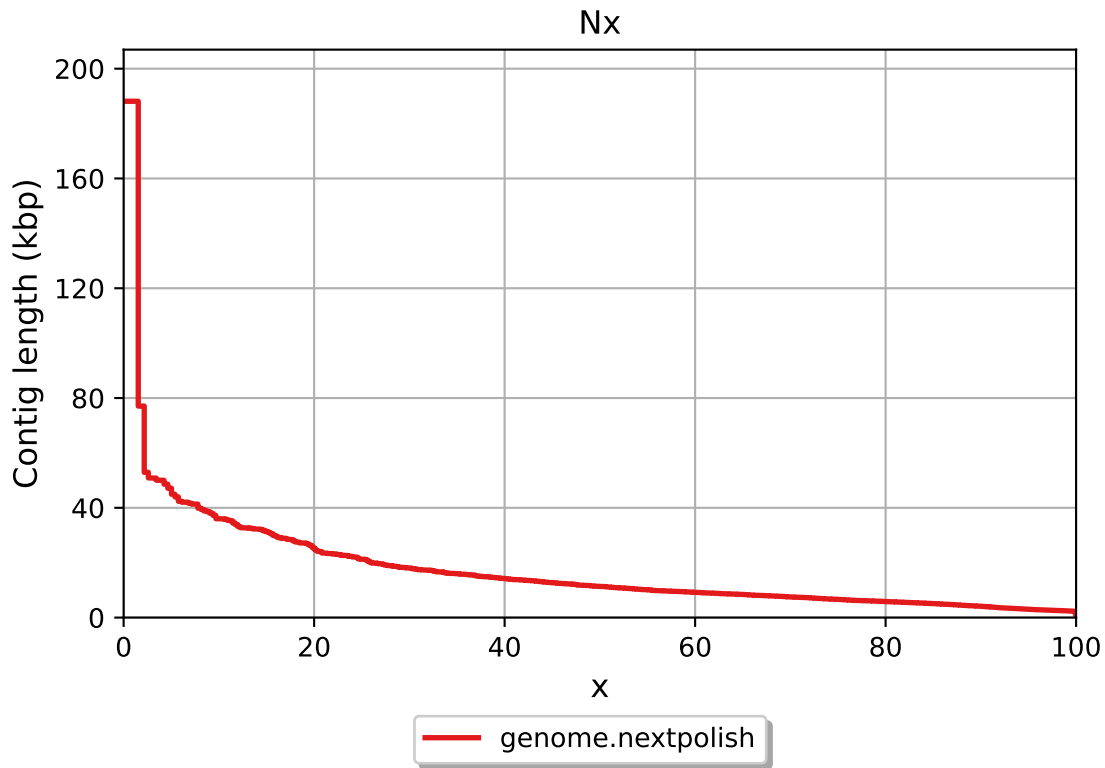


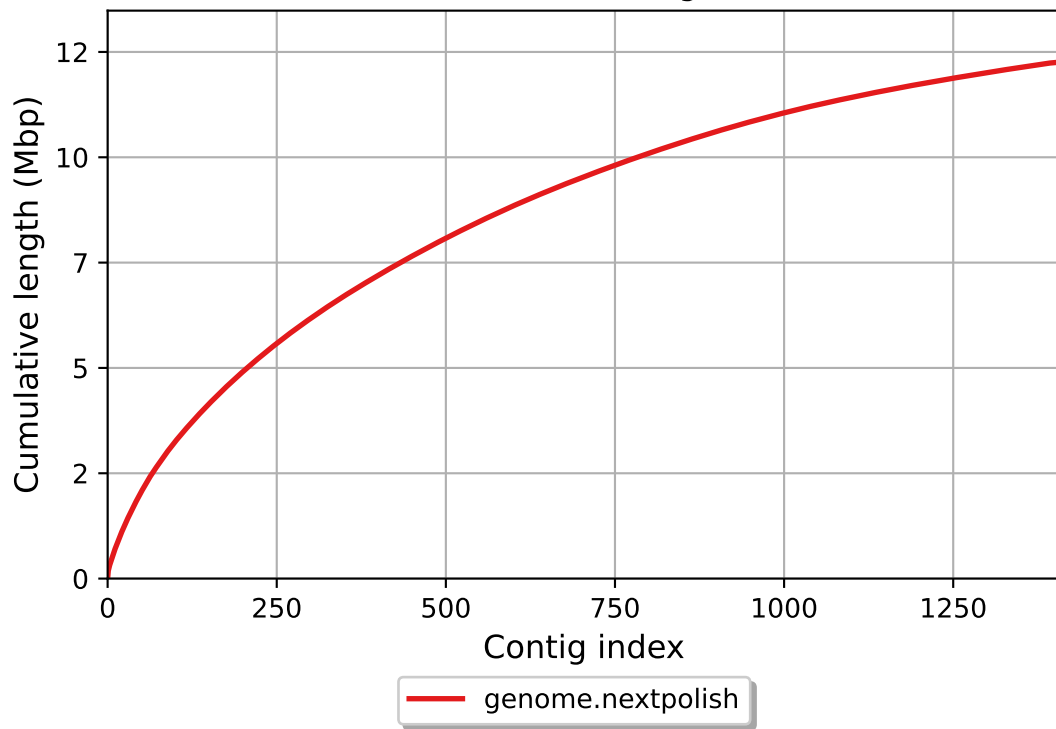
Report

	genome.nextpolish
# contigs (>= 0 bp)	1408
# contigs (>= 1000 bp)	1402
# contigs (>= 5000 bp)	877
# contigs (>= 10000 bp)	358
# contigs (>= 25000 bp)	64
# contigs (>= 50000 bp)	7
Total length (>= 0 bp)	12255079
Total length (>= 1000 bp)	12251419
Total length (>= 5000 bp)	10500146
Total length (>= 10000 bp)	6791579
Total length (>= 25000 bp)	2450841
Total length (>= 50000 bp)	519901
# contigs	1408
Largest contig	188126
Total length	12255079
GC (%)	40.94
N50	11356
N90	4126
auN	18254.1
L50	296
L90	994
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	23 + 4 part

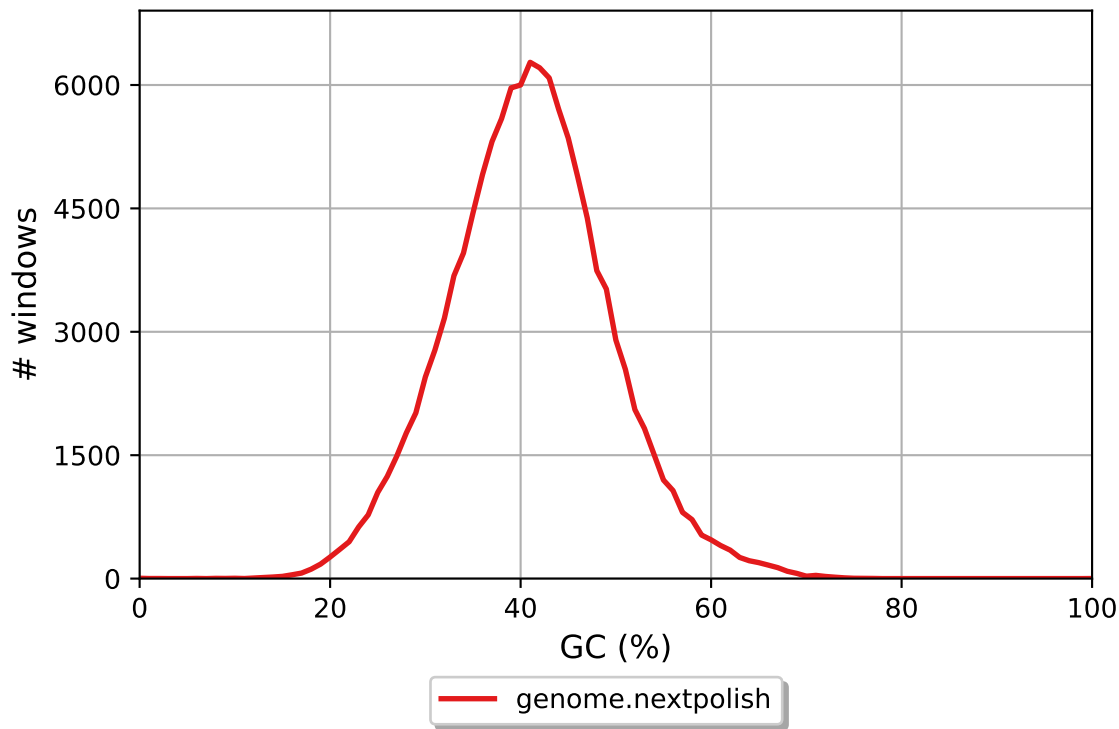
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

