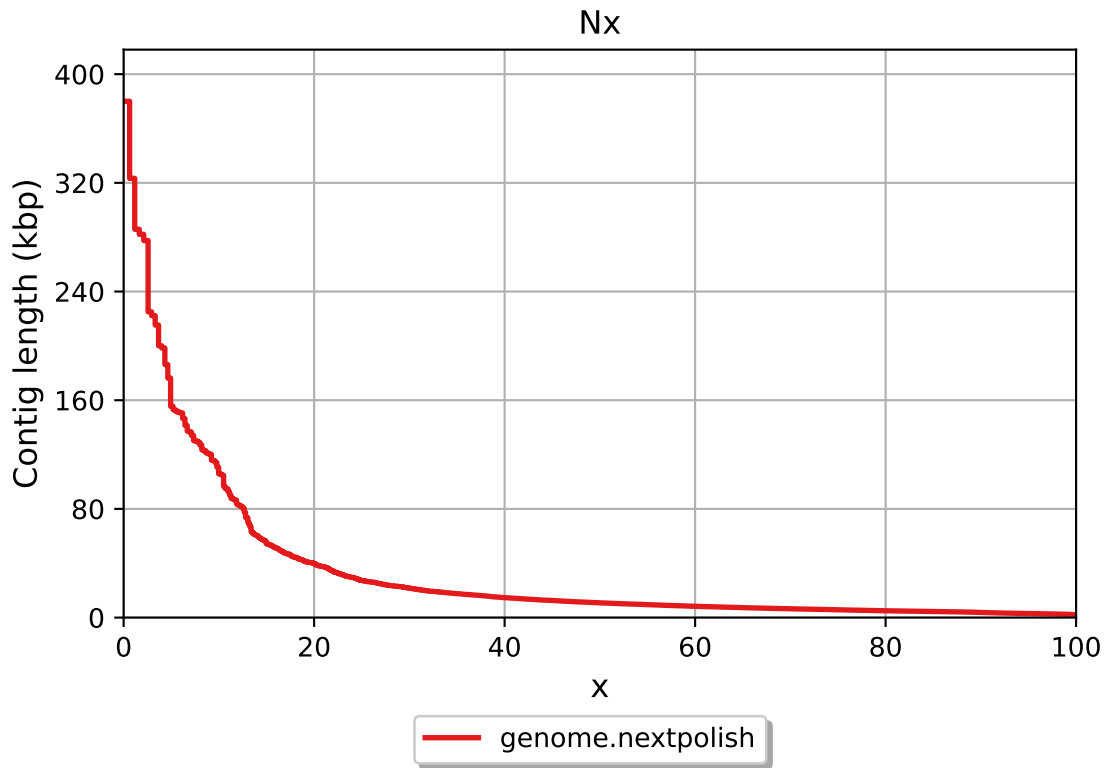


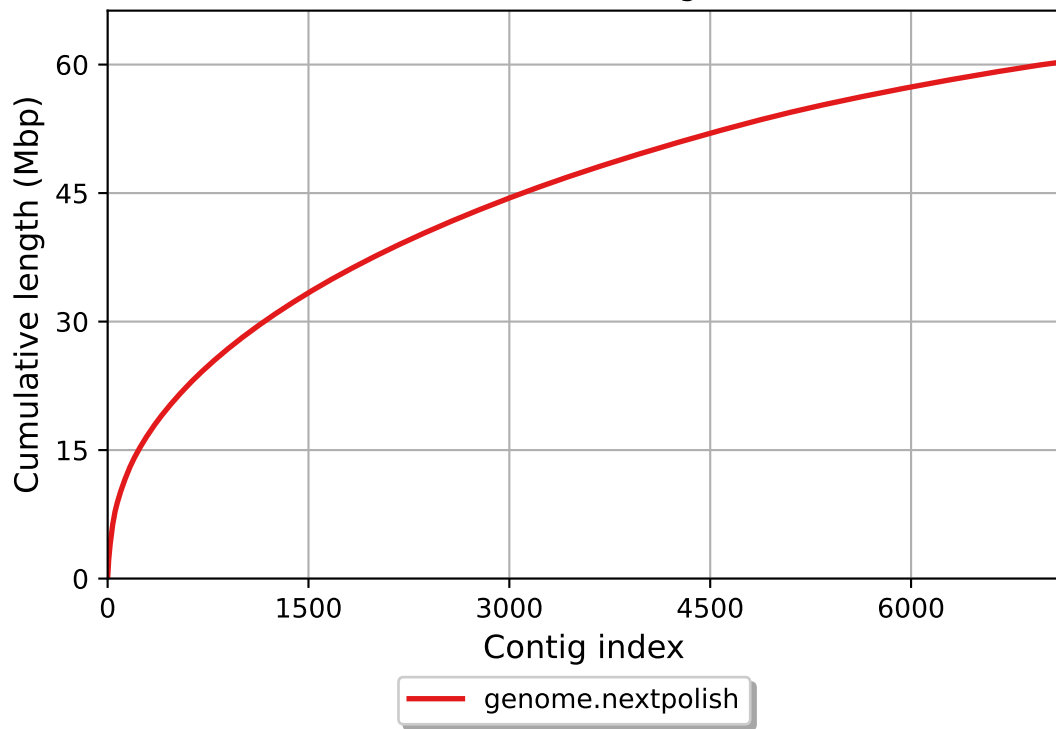
Report

	genome.nextpolish
# contigs (>= 0 bp)	7113
# contigs (>= 1000 bp)	7105
# contigs (>= 5000 bp)	3709
# contigs (>= 10000 bp)	1364
# contigs (>= 25000 bp)	278
# contigs (>= 50000 bp)	91
Total length (>= 0 bp)	60270416
Total length (>= 1000 bp)	60264557
Total length (>= 5000 bp)	48288949
Total length (>= 10000 bp)	32043774
Total length (>= 25000 bp)	16247390
Total length (>= 50000 bp)	9856199
# contigs	7113
Largest contig	380049
Total length	60270416
GC (%)	42.53
N50	10881
N90	3792
auN	34833.8
L50	1181
L90	5050
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	89 + 8 part

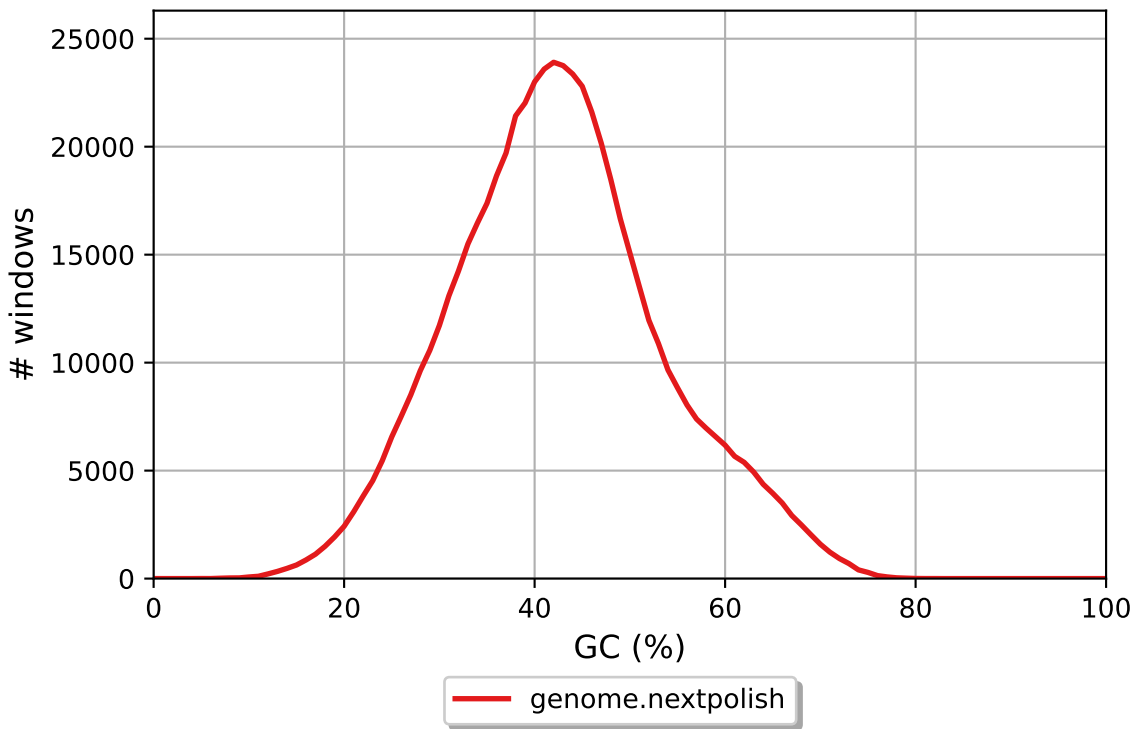
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

