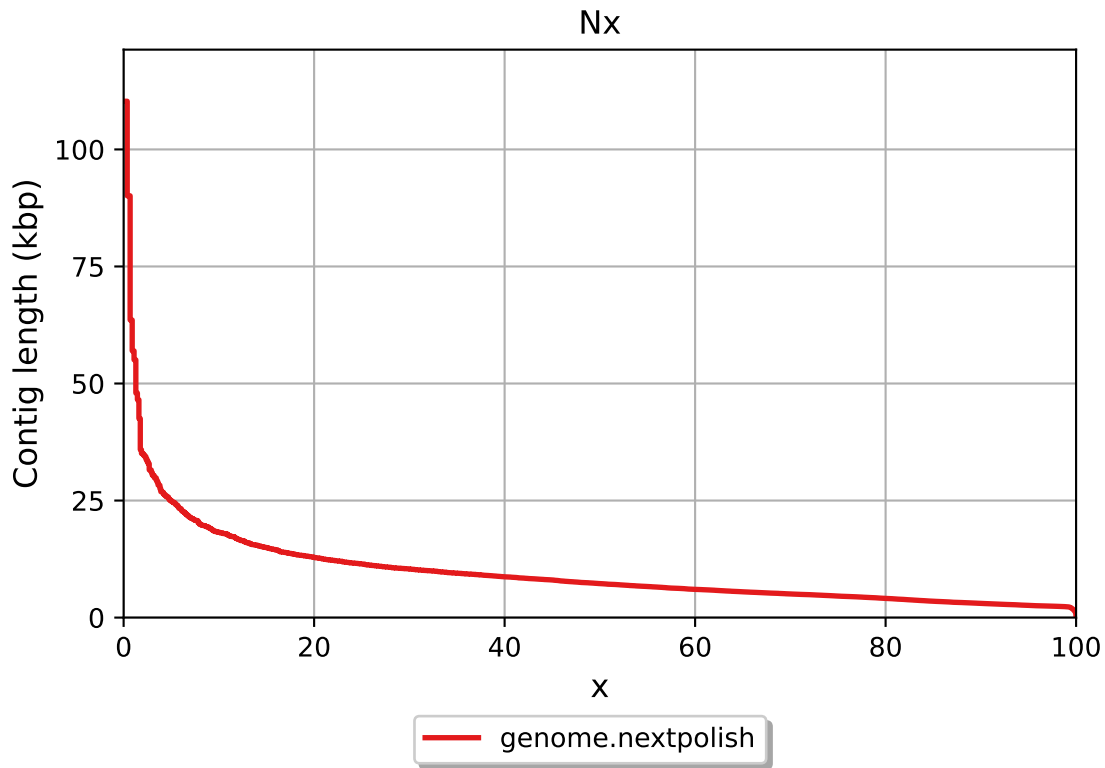


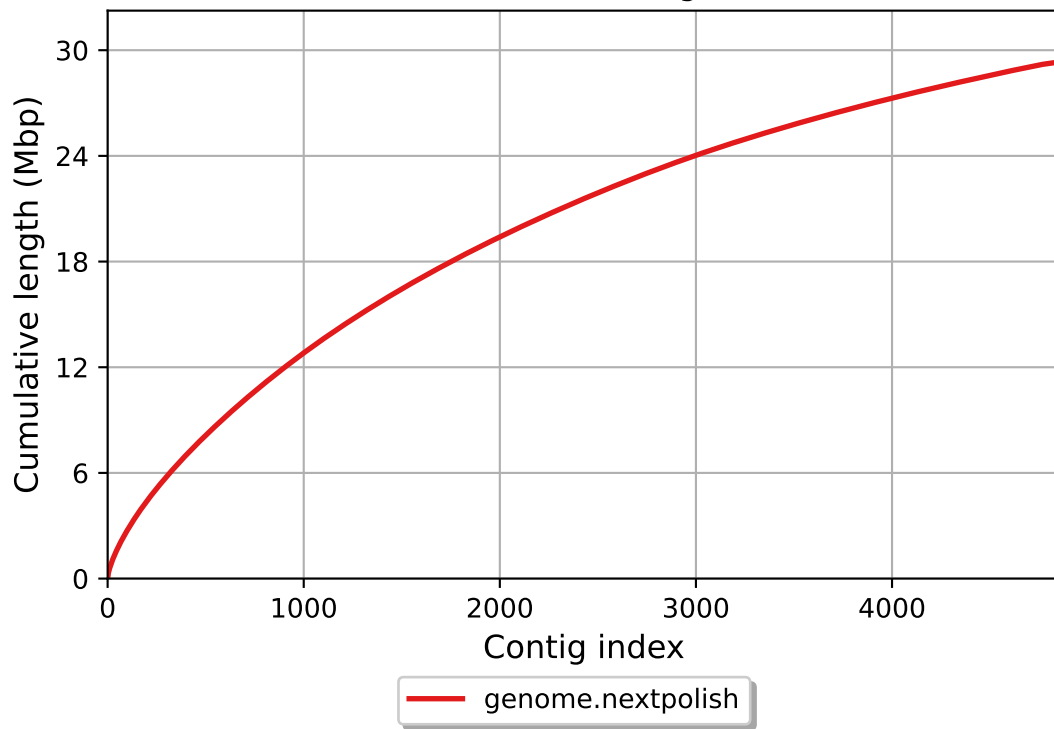
Report

	genome.nextpolish
# contigs (>= 0 bp)	4857
# contigs (>= 1000 bp)	4838
# contigs (>= 5000 bp)	2244
# contigs (>= 10000 bp)	627
# contigs (>= 25000 bp)	39
# contigs (>= 50000 bp)	5
Total length (>= 0 bp)	29315315
Total length (>= 1000 bp)	29301878
Total length (>= 5000 bp)	20668408
Total length (>= 10000 bp)	9448236
Total length (>= 25000 bp)	1432020
Total length (>= 50000 bp)	375992
# contigs	4857
Largest contig	110287
Total length	29315315
GC (%)	41.78
N50	7269
N90	3032
auN	9980.5
L50	1240
L90	3692
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	26 + 4 part

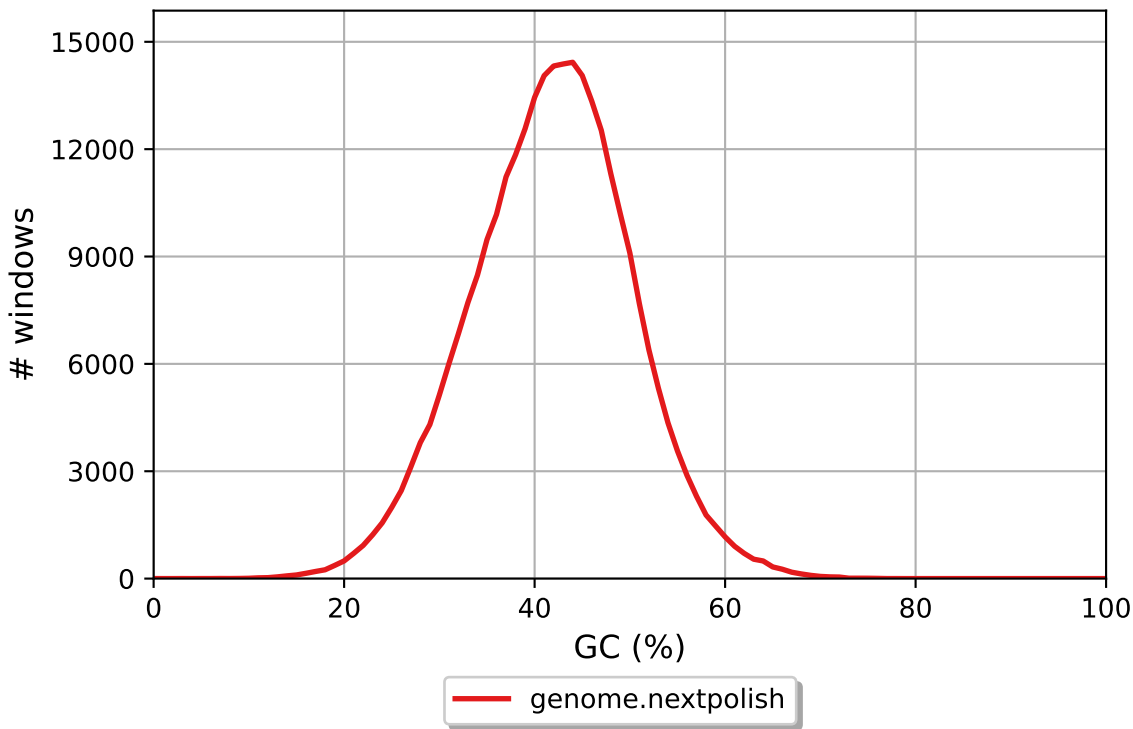
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

