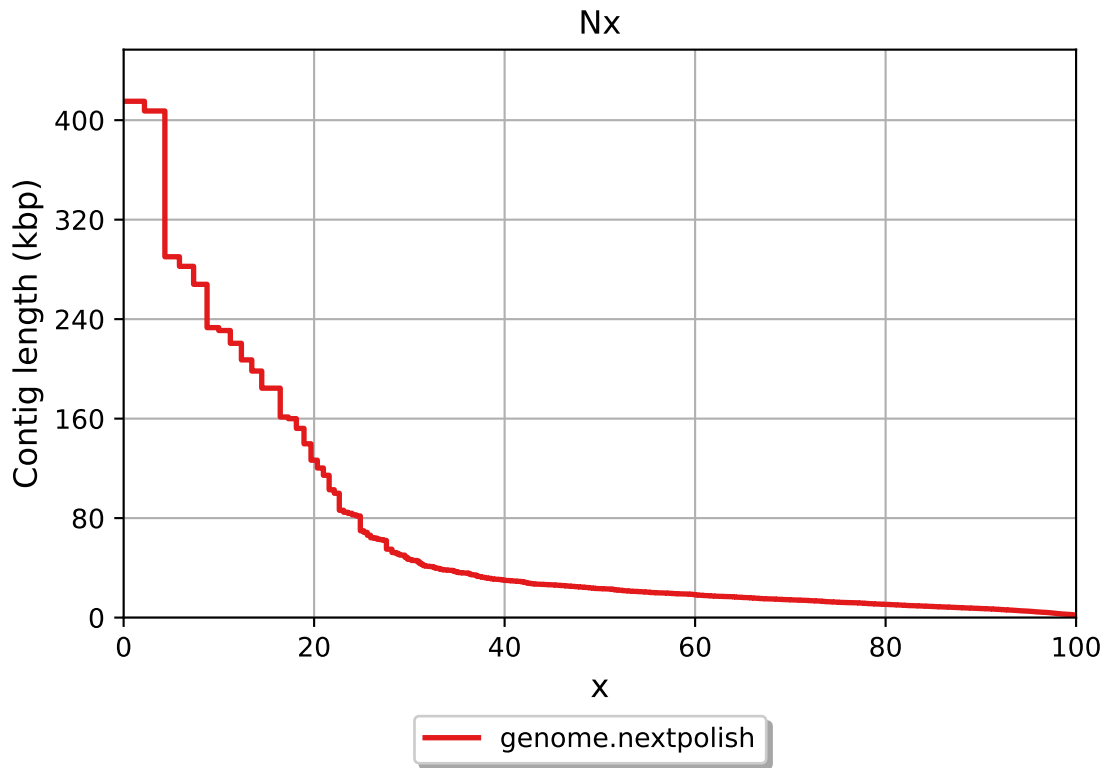


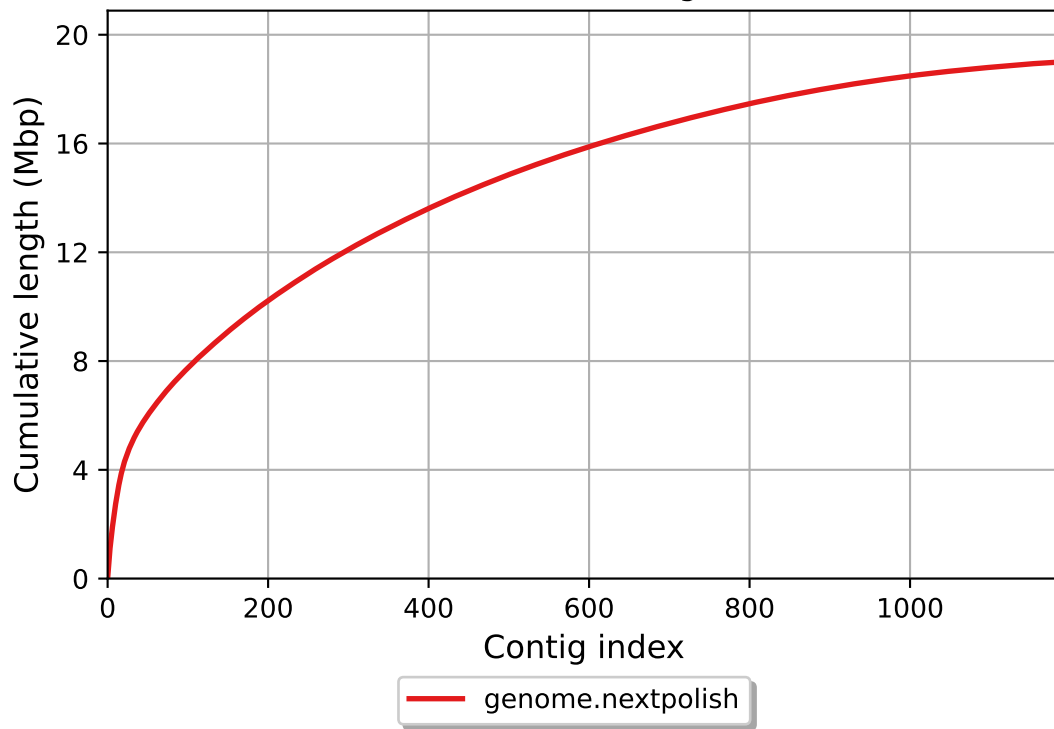
Report

	genome.nextpolish
# contigs (>= 0 bp)	1187
# contigs (>= 1000 bp)	1179
# contigs (>= 5000 bp)	911
# contigs (>= 10000 bp)	563
# contigs (>= 25000 bp)	146
# contigs (>= 50000 bp)	41
Total length (>= 0 bp)	18987623
Total length (>= 1000 bp)	18981893
Total length (>= 5000 bp)	18100167
Total length (>= 10000 bp)	15525152
Total length (>= 25000 bp)	8984297
Total length (>= 50000 bp)	5605106
# contigs	1187
Largest contig	415202
Total length	18987623
GC (%)	42.46
N50	23286
N90	7273
auN	72283.0
L50	168
L90	746
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	39 + 2 part

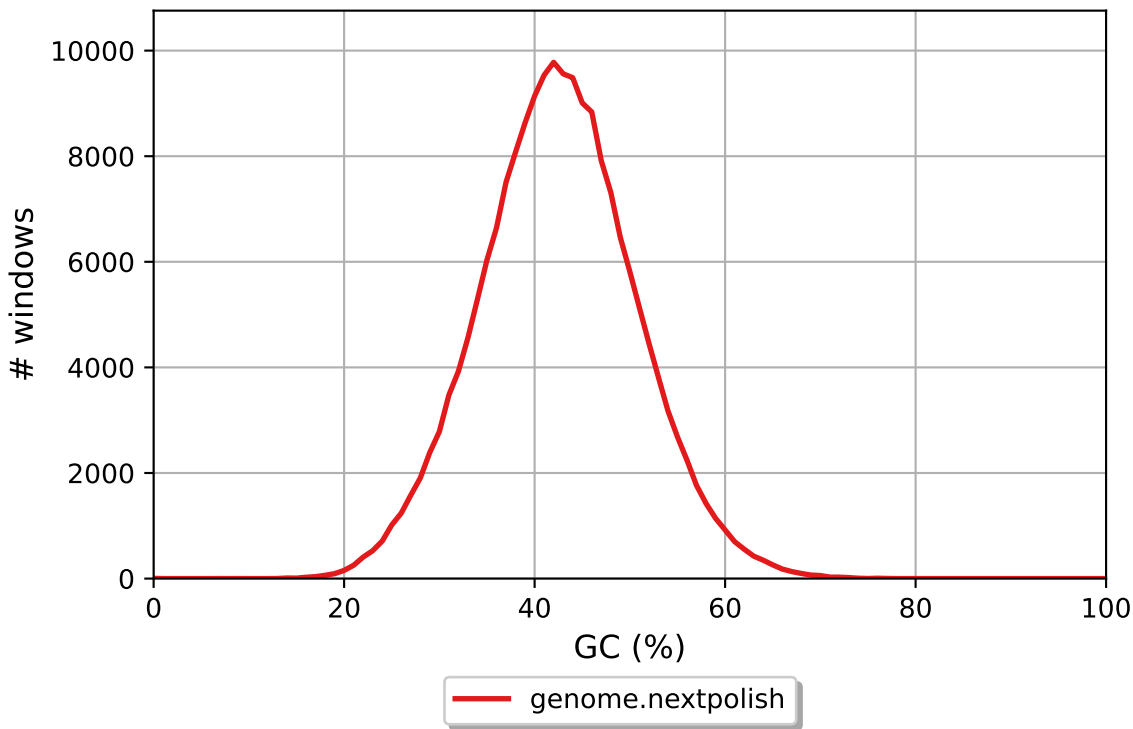
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

