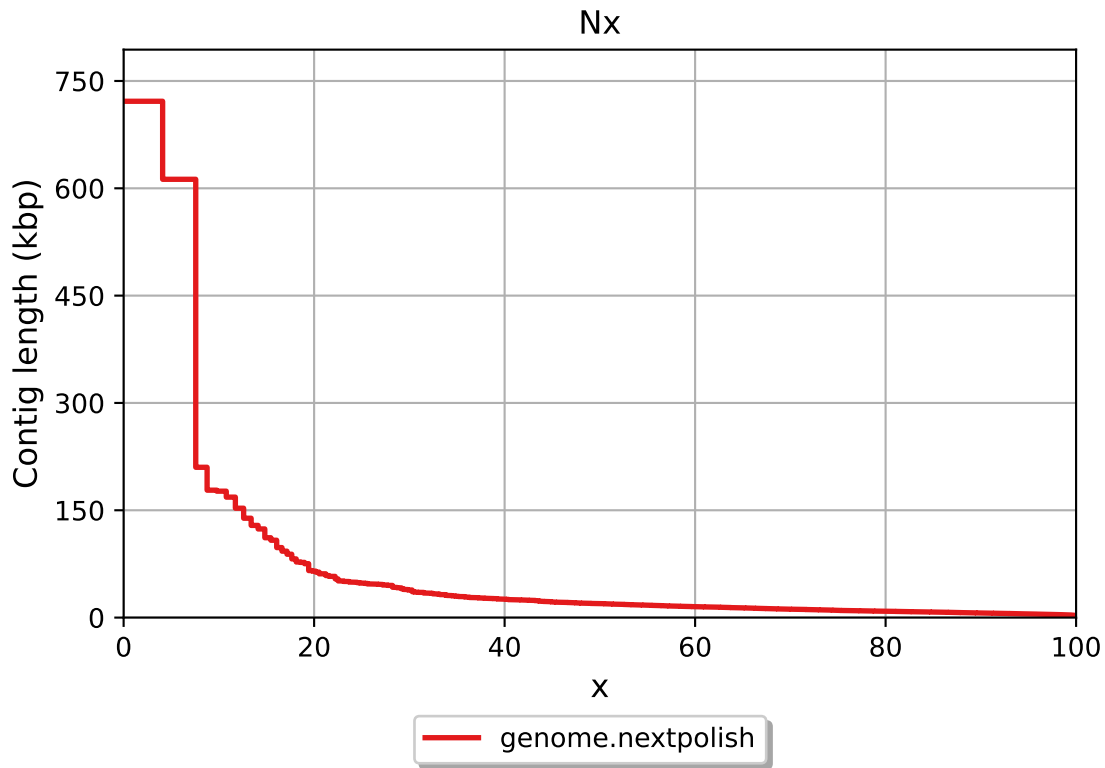


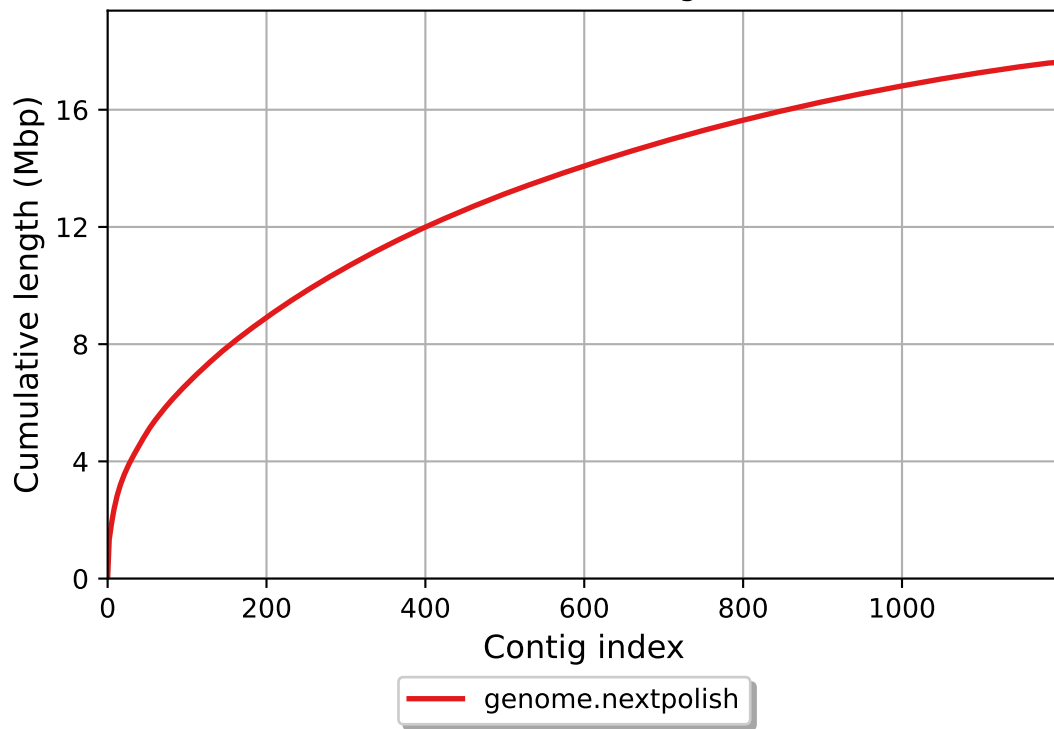
Report

	genome.nextpolish
# contigs (>= 0 bp)	1199
# contigs (>= 1000 bp)	1197
# contigs (>= 5000 bp)	1001
# contigs (>= 10000 bp)	512
# contigs (>= 25000 bp)	122
# contigs (>= 50000 bp)	32
Total length (>= 0 bp)	17619869
Total length (>= 1000 bp)	17618670
Total length (>= 5000 bp)	16817406
Total length (>= 10000 bp)	13252188
Total length (>= 25000 bp)	7220550
Total length (>= 50000 bp)	4170812
# contigs	1199
Largest contig	721637
Total length	17619869
GC (%)	42.74
N50	19513
N90	6450
auN	83003.4
L50	195
L90	833
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	46 + 5 part

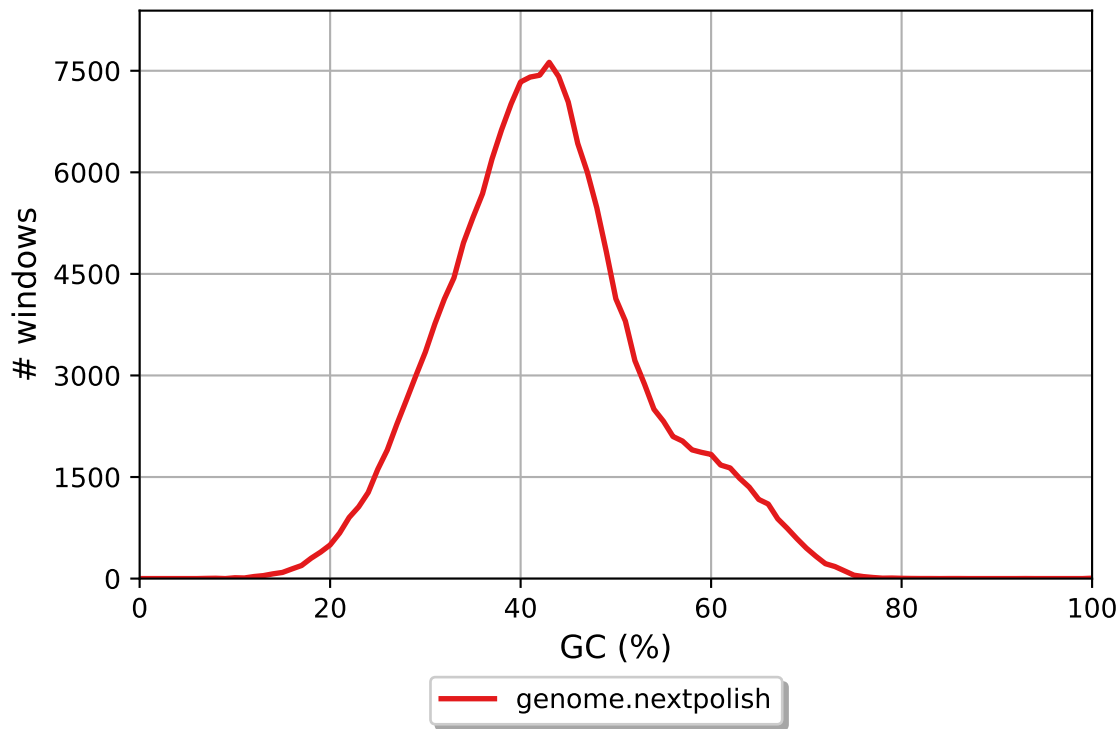
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

