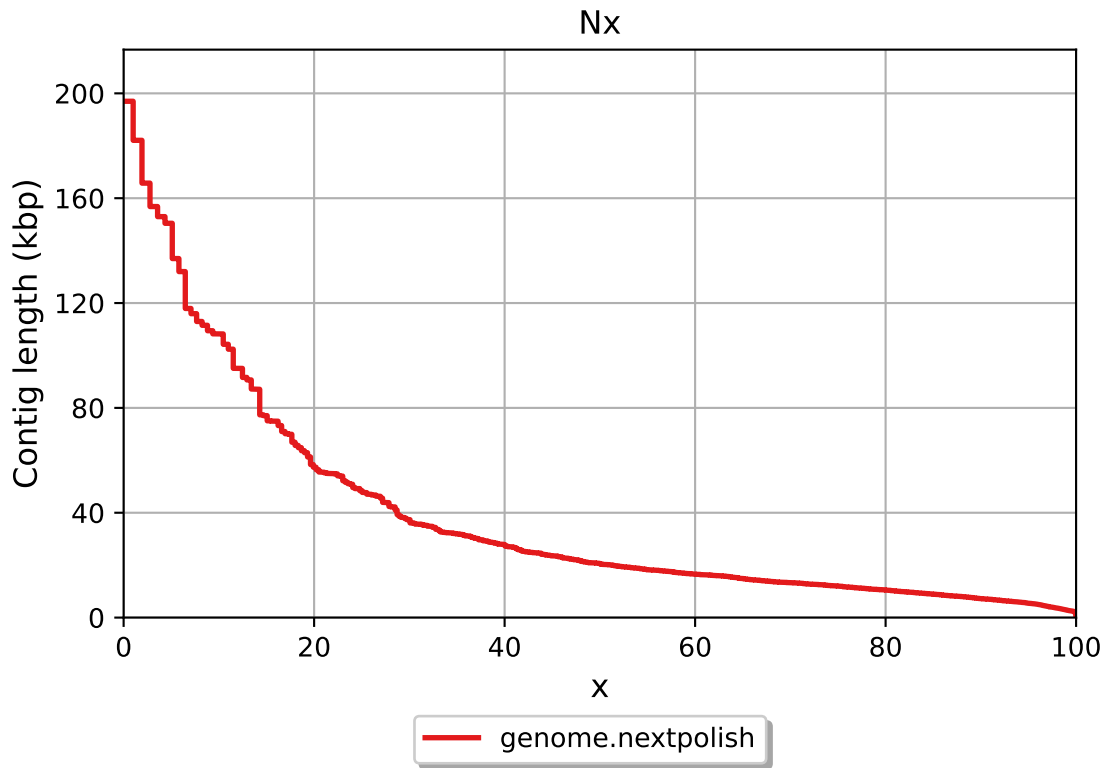


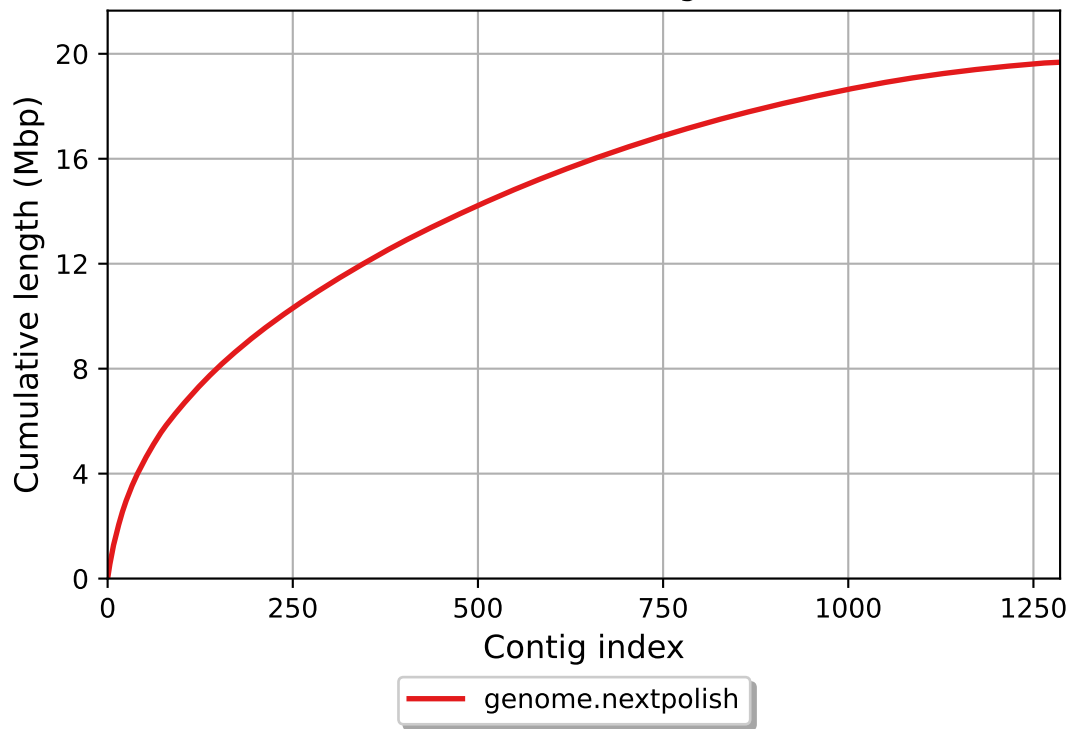
Report

	genome.nextpolish
# contigs (>= 0 bp)	1286
# contigs (>= 1000 bp)	1278
# contigs (>= 5000 bp)	1047
# contigs (>= 10000 bp)	660
# contigs (>= 25000 bp)	162
# contigs (>= 50000 bp)	54
Total length (>= 0 bp)	19677180
Total length (>= 1000 bp)	19671859
Total length (>= 5000 bp)	18895773
Total length (>= 10000 bp)	16035937
Total length (>= 25000 bp)	8378887
Total length (>= 50000 bp)	4732475
# contigs	1286
Largest contig	196979
Total length	19677180
GC (%)	42.72
N50	20568
N90	7224
auN	38877.2
L50	227
L90	855
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	35 + 10 part

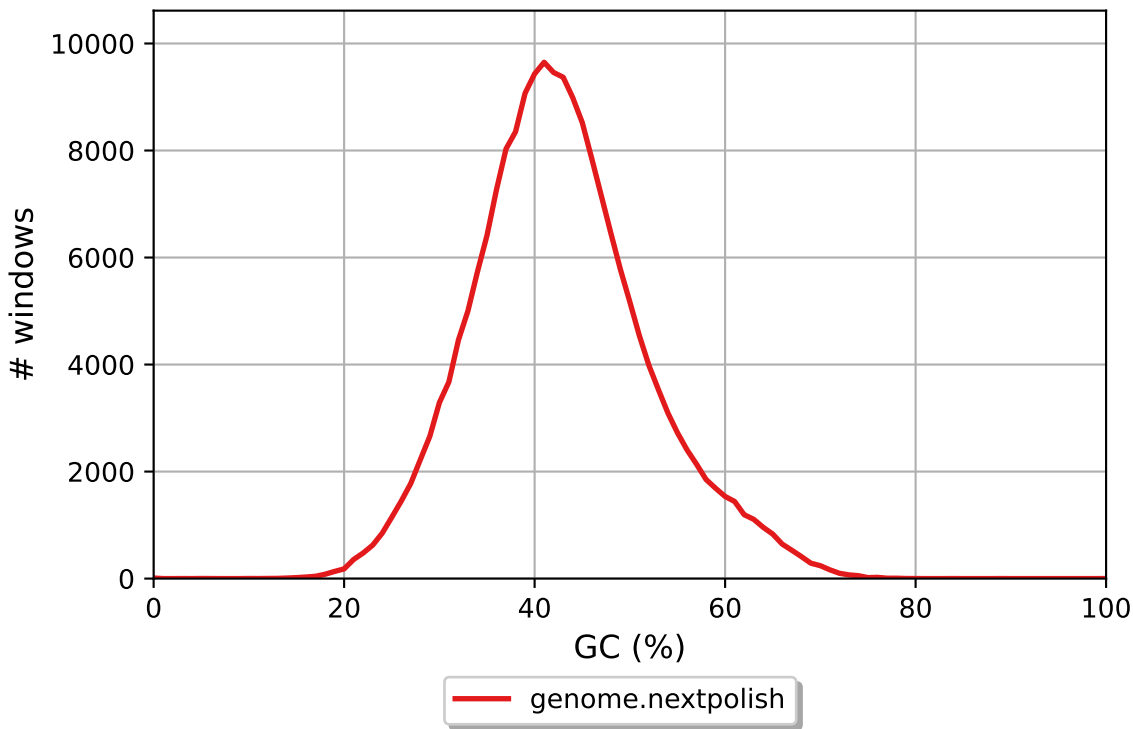
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

