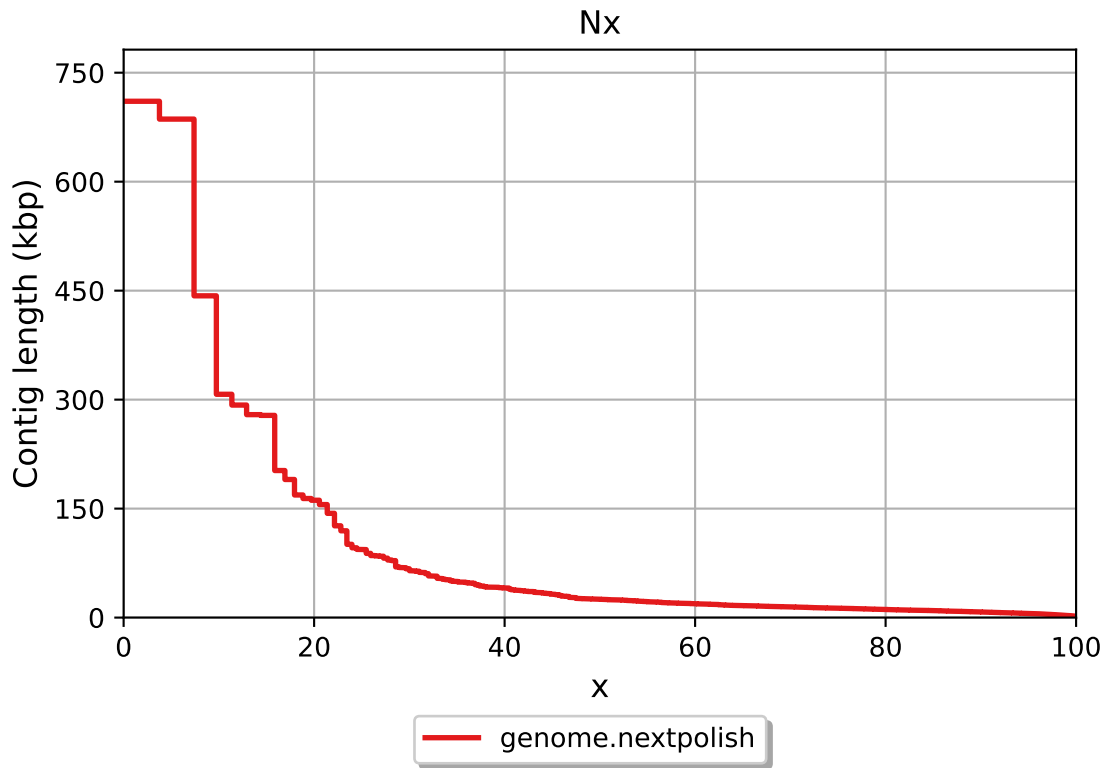


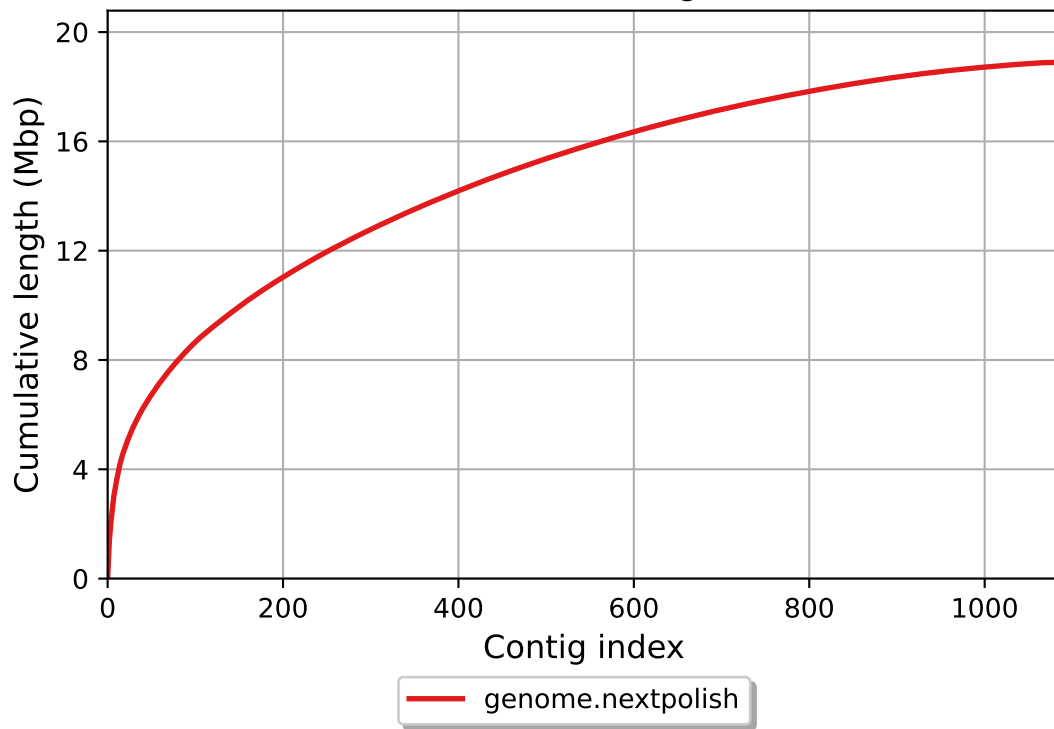
Report

	genome.nextpolish
# contigs (>= 0 bp)	1086
# contigs (>= 1000 bp)	1070
# contigs (>= 5000 bp)	870
# contigs (>= 10000 bp)	541
# contigs (>= 25000 bp)	134
# contigs (>= 50000 bp)	46
Total length (>= 0 bp)	18894921
Total length (>= 1000 bp)	18882992
Total length (>= 5000 bp)	18212527
Total length (>= 10000 bp)	15788712
Total length (>= 25000 bp)	9564124
Total length (>= 50000 bp)	6533774
# contigs	1086
Largest contig	710669
Total length	18894921
GC (%)	42.28
N50	25236
N90	7762
auN	112601.2
L50	130
L90	679
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	46 + 4 part

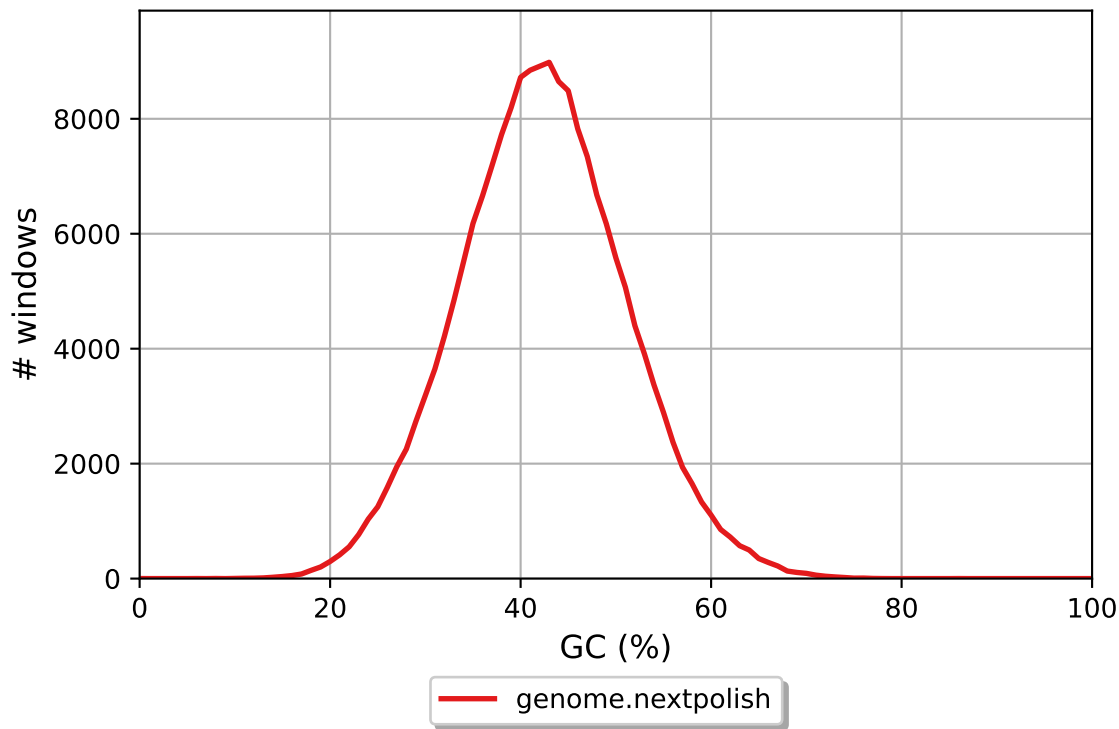
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

