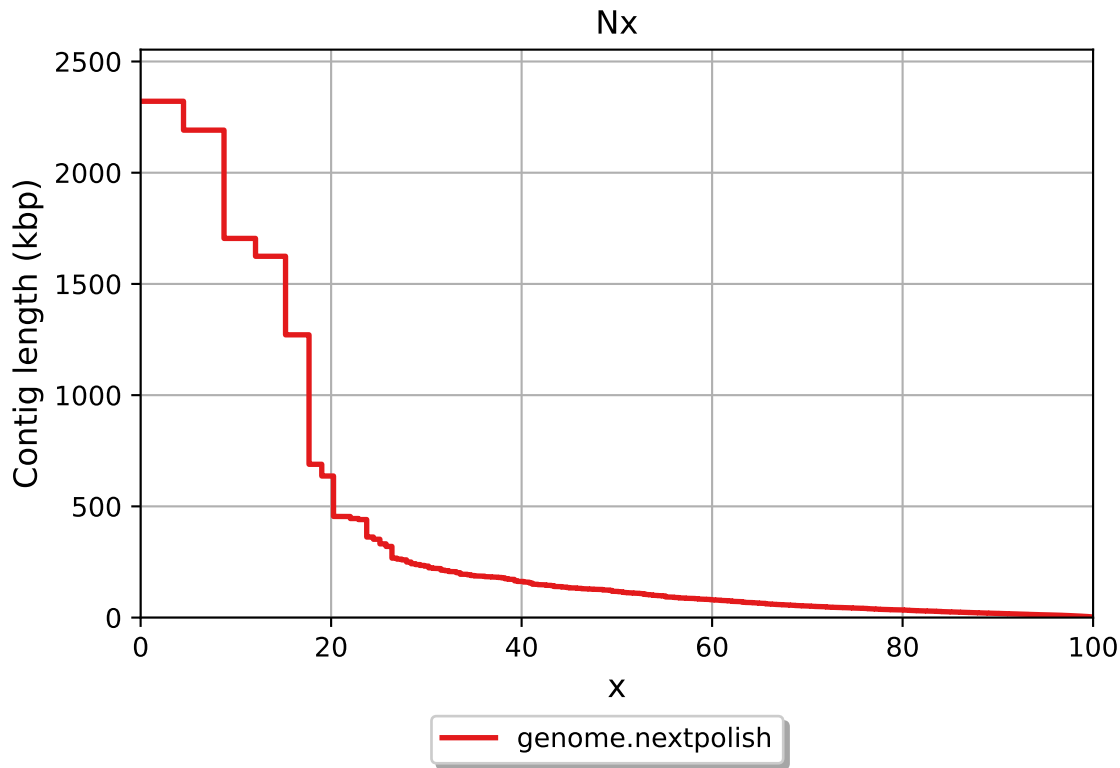


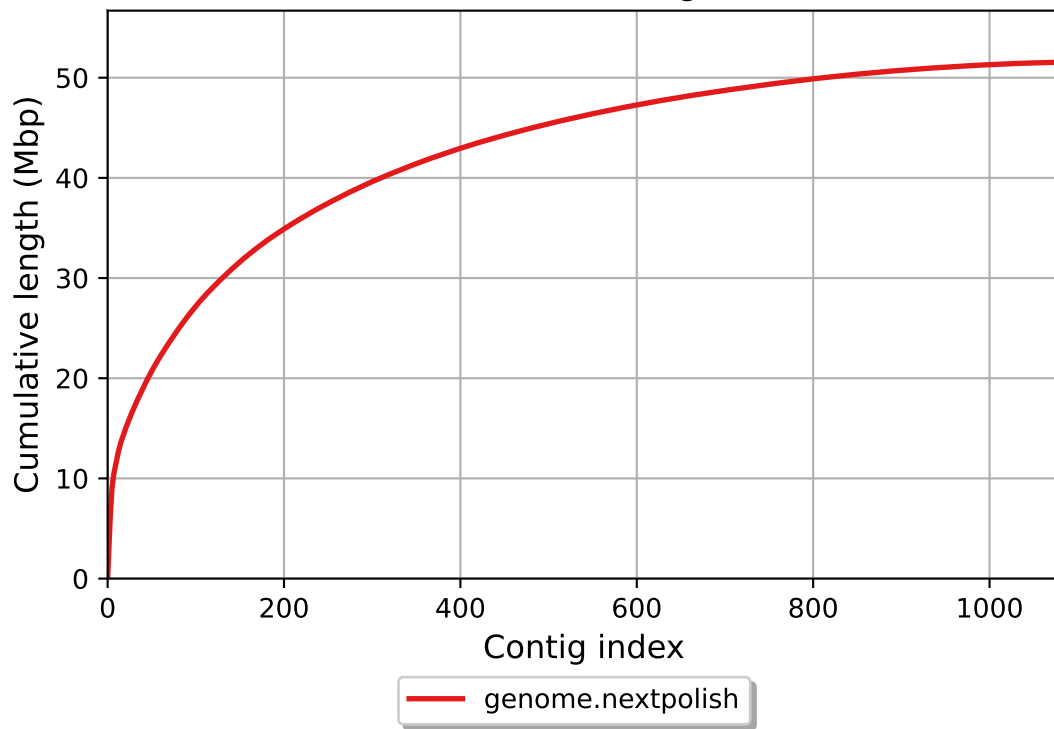
Report

	genome.nextpolish
# contigs (>= 0 bp)	1080
# contigs (>= 1000 bp)	1077
# contigs (>= 5000 bp)	974
# contigs (>= 10000 bp)	804
# contigs (>= 25000 bp)	442
# contigs (>= 50000 bp)	231
Total length (>= 0 bp)	51540563
Total length (>= 1000 bp)	51538781
Total length (>= 5000 bp)	51183307
Total length (>= 10000 bp)	49927924
Total length (>= 25000 bp)	44067723
Total length (>= 50000 bp)	36558163
# contigs	1080
Largest contig	2321208
Total length	51540563
GC (%)	54.79
N50	117417
N90	18763
auN	444750.9
L50	88
L90	550
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	92 + 2 part

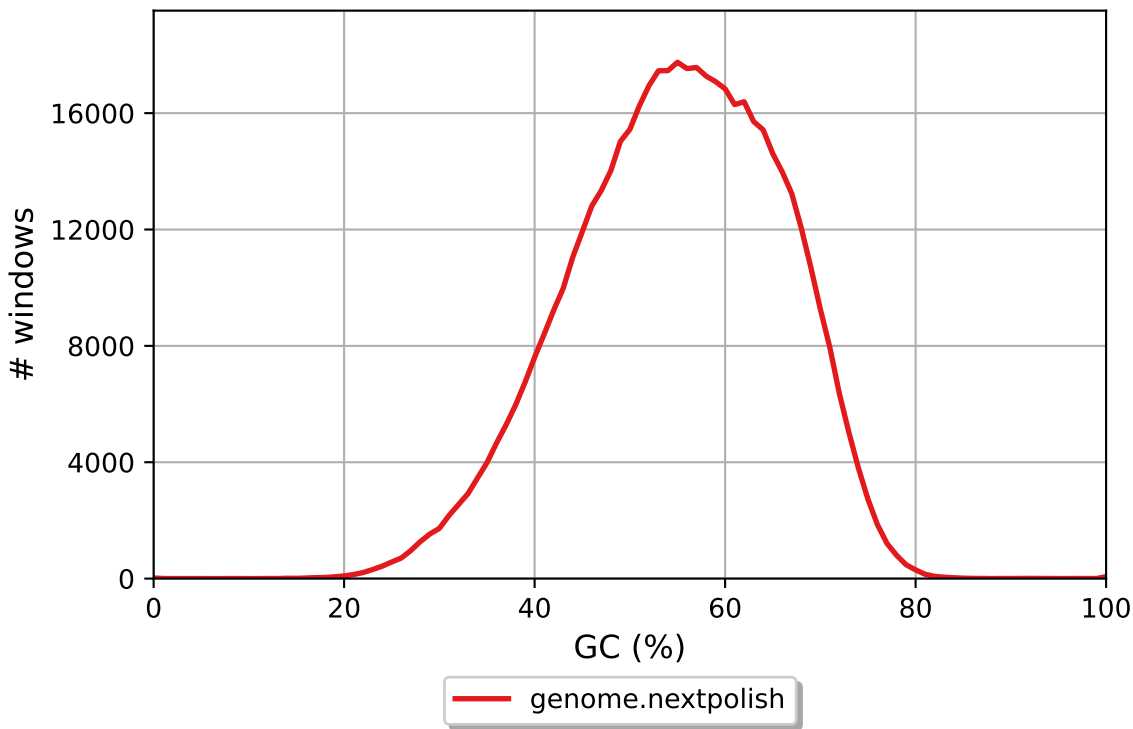
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

