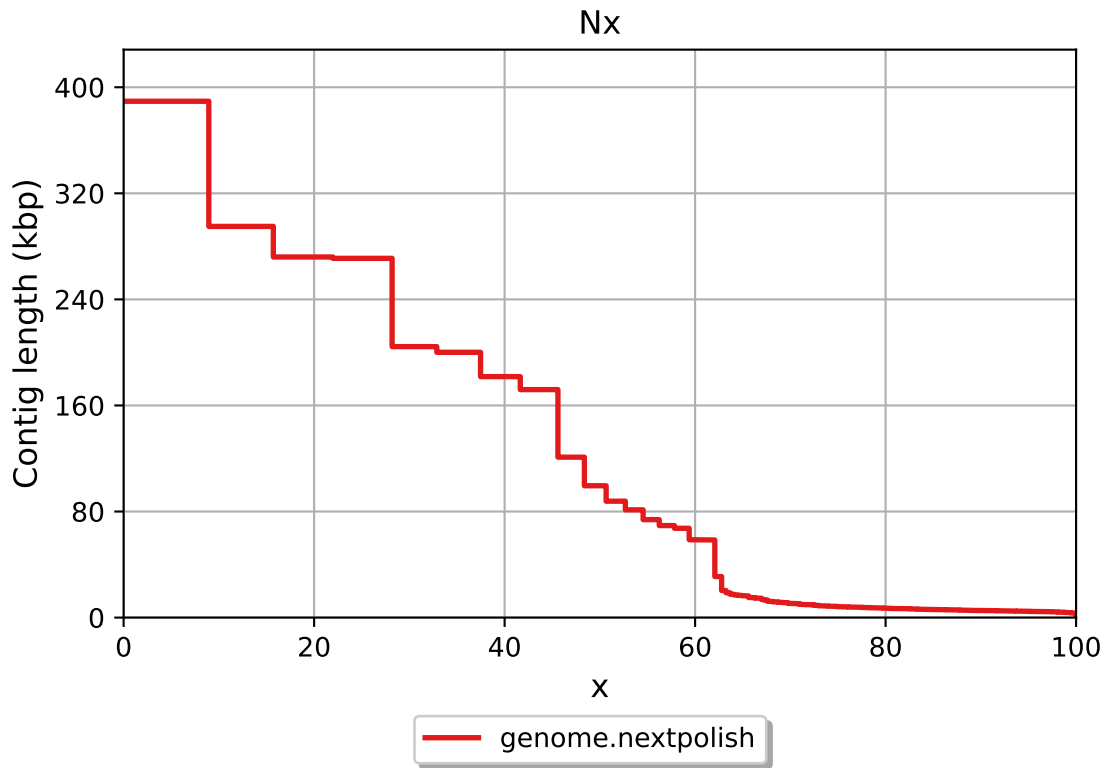


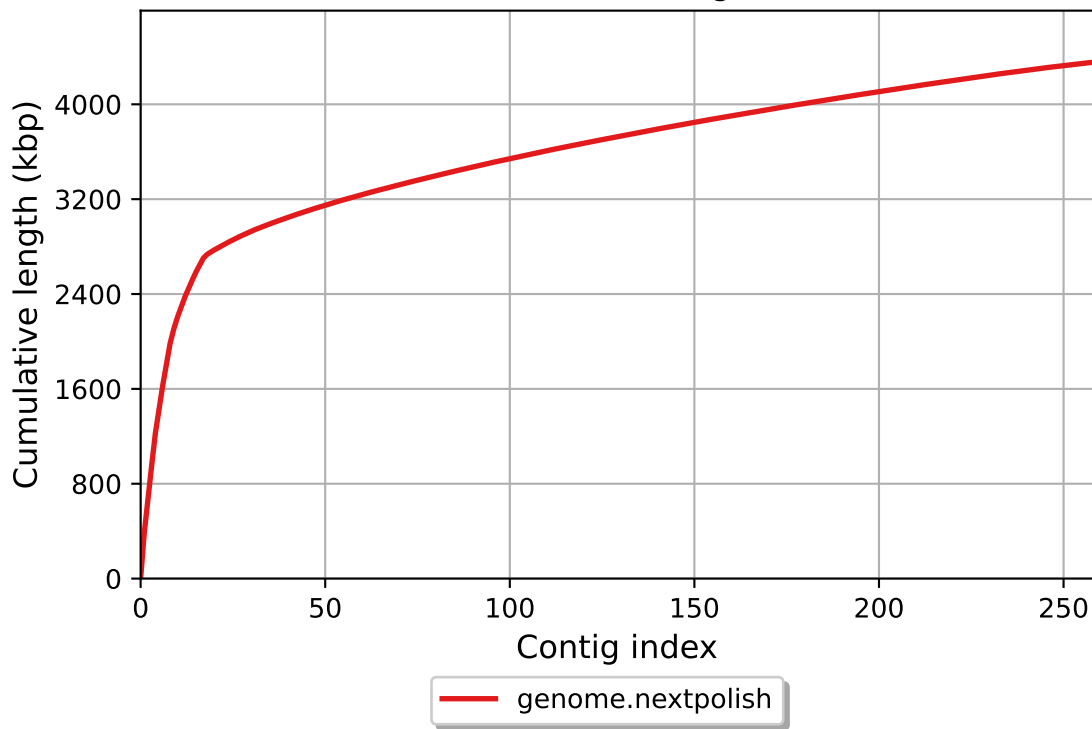
Report

	genome.nextpolish
# contigs (>= 0 bp)	258
# contigs (>= 1000 bp)	258
# contigs (>= 5000 bp)	187
# contigs (>= 10000 bp)	45
# contigs (>= 25000 bp)	18
# contigs (>= 50000 bp)	17
Total length (>= 0 bp)	4354578
Total length (>= 1000 bp)	4354578
Total length (>= 5000 bp)	4043169
Total length (>= 10000 bp)	3100153
Total length (>= 25000 bp)	2733862
Total length (>= 50000 bp)	2702904
# contigs	258
Largest contig	389421
Total length	4354578
GC (%)	58.85
N50	99438
N90	5320
auN	138942.3
L50	10
L90	164
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	16 + 3 part

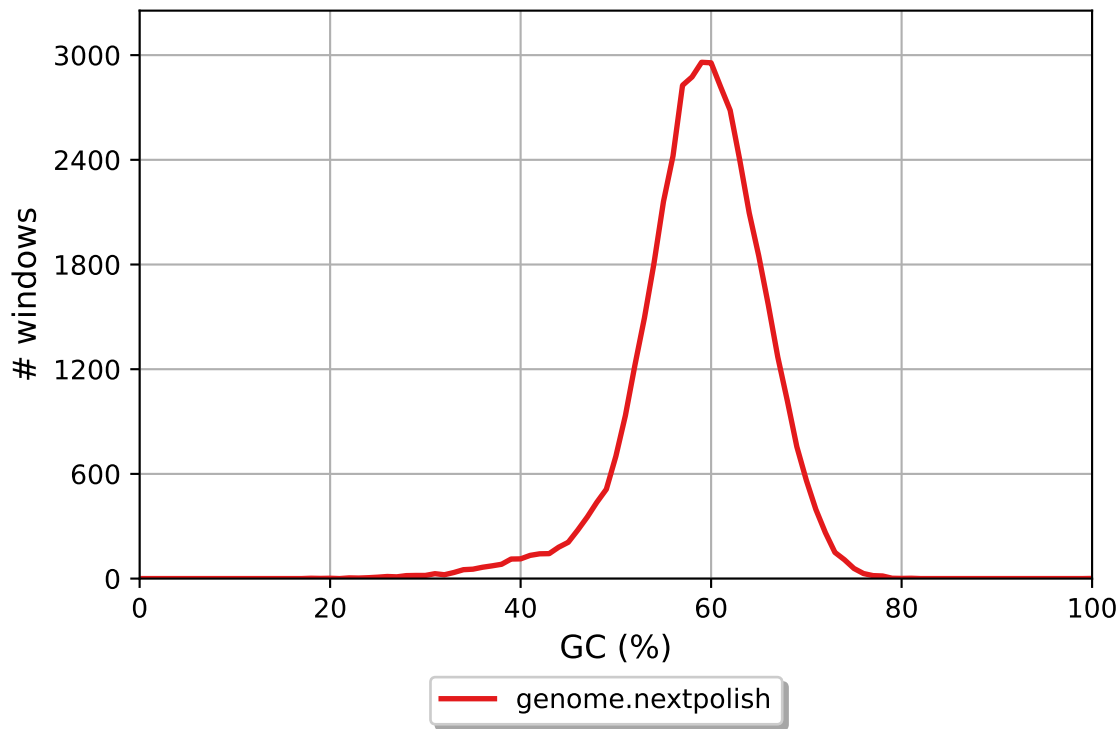
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

