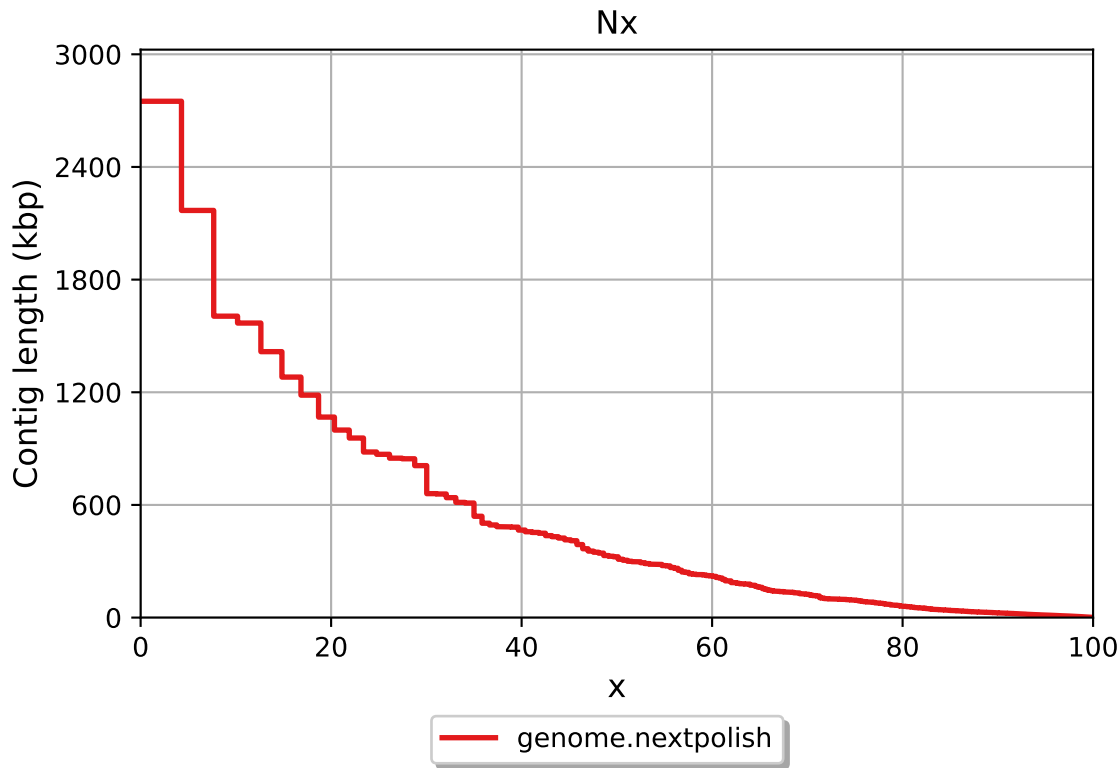


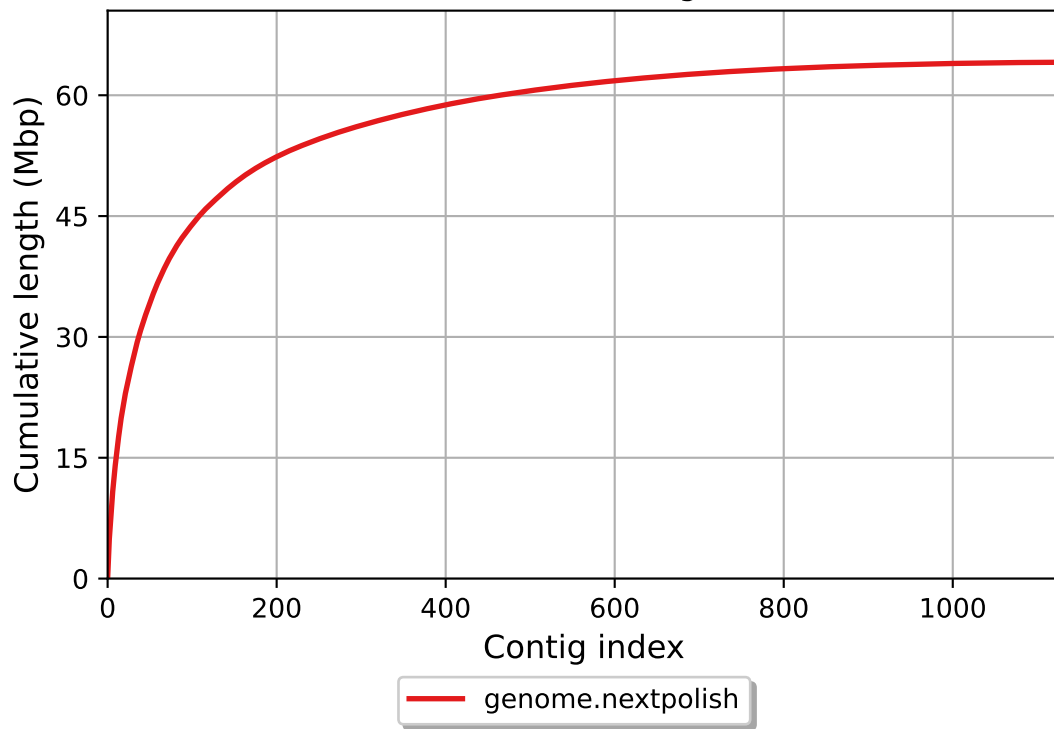
Report

	genome.nextpolish
# contigs (>= 0 bp)	1137
# contigs (>= 1000 bp)	1080
# contigs (>= 5000 bp)	793
# contigs (>= 10000 bp)	609
# contigs (>= 25000 bp)	352
# contigs (>= 50000 bp)	209
Total length (>= 0 bp)	64097286
Total length (>= 1000 bp)	64062029
Total length (>= 5000 bp)	63264089
Total length (>= 10000 bp)	61898918
Total length (>= 25000 bp)	57699695
Total length (>= 50000 bp)	52822137
# contigs	1127
Largest contig	2749808
Total length	64093869
GC (%)	46.28
N50	323798
N90	25142
auN	607014.2
L50	43
L90	352
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	173 + 2 part

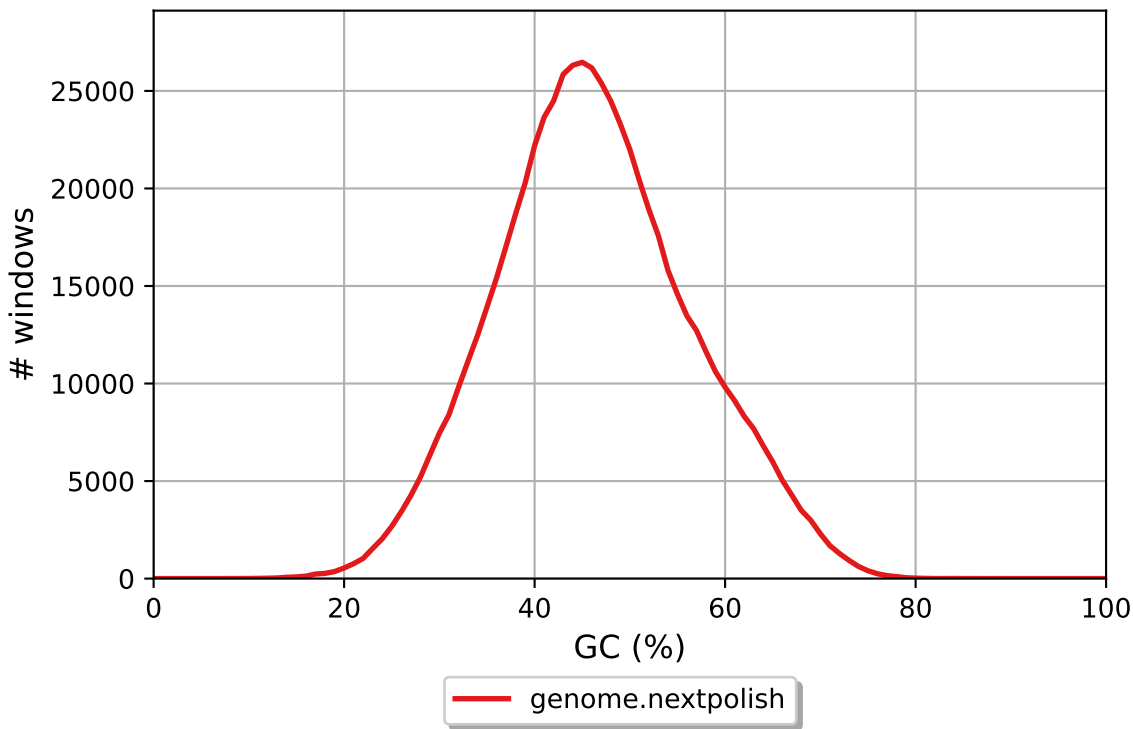
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

