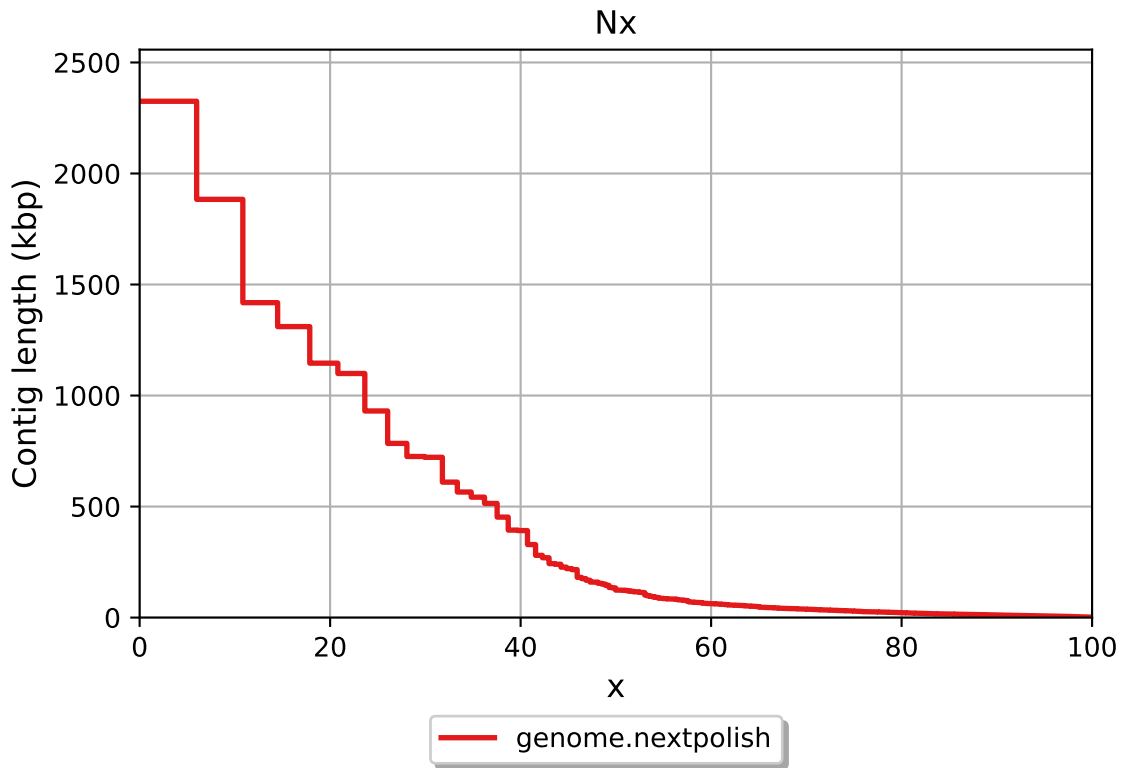


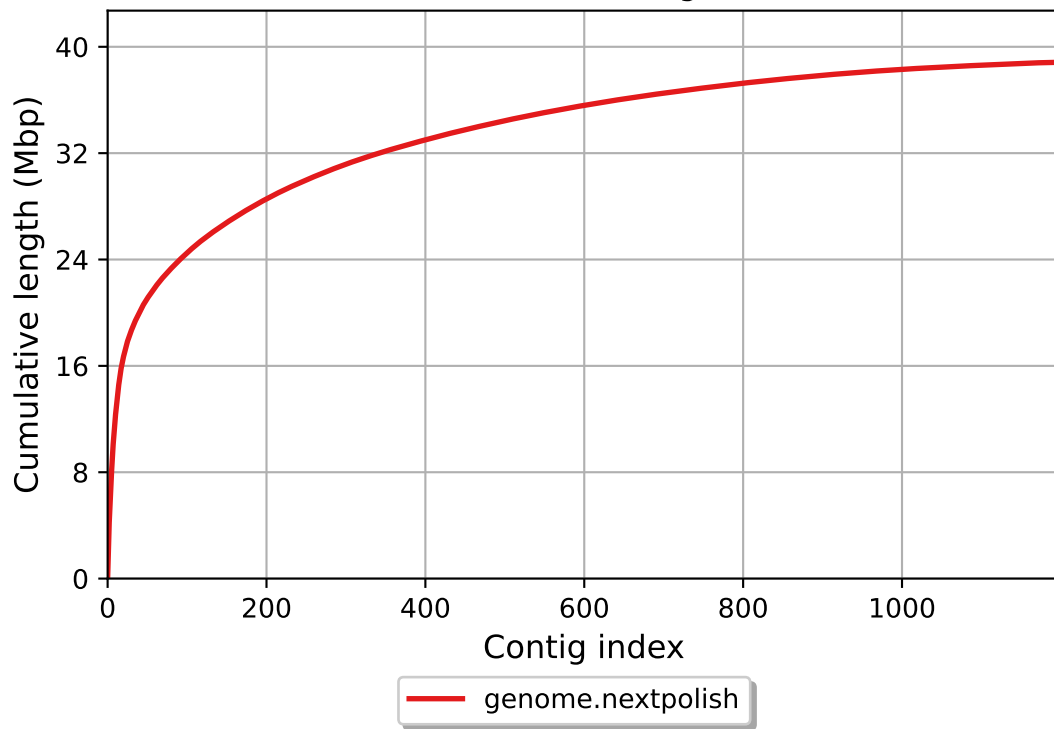
# Report

	genome.nextpolish
# contigs (>= 0 bp)	1200
# contigs (>= 1000 bp)	1194
# contigs (>= 5000 bp)	916
# contigs (>= 10000 bp)	612
# contigs (>= 25000 bp)	259
# contigs (>= 50000 bp)	113
Total length (>= 0 bp)	38836840
Total length (>= 1000 bp)	38832999
Total length (>= 5000 bp)	37938955
Total length (>= 10000 bp)	35718034
Total length (>= 25000 bp)	30211262
Total length (>= 50000 bp)	25177210
# contigs	1199
Largest contig	2325272
Total length	38836363
GC (%)	47.91
N50	123894
N90	11686
auN	541301.3
L50	36
L90	542
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	67 + 5 part

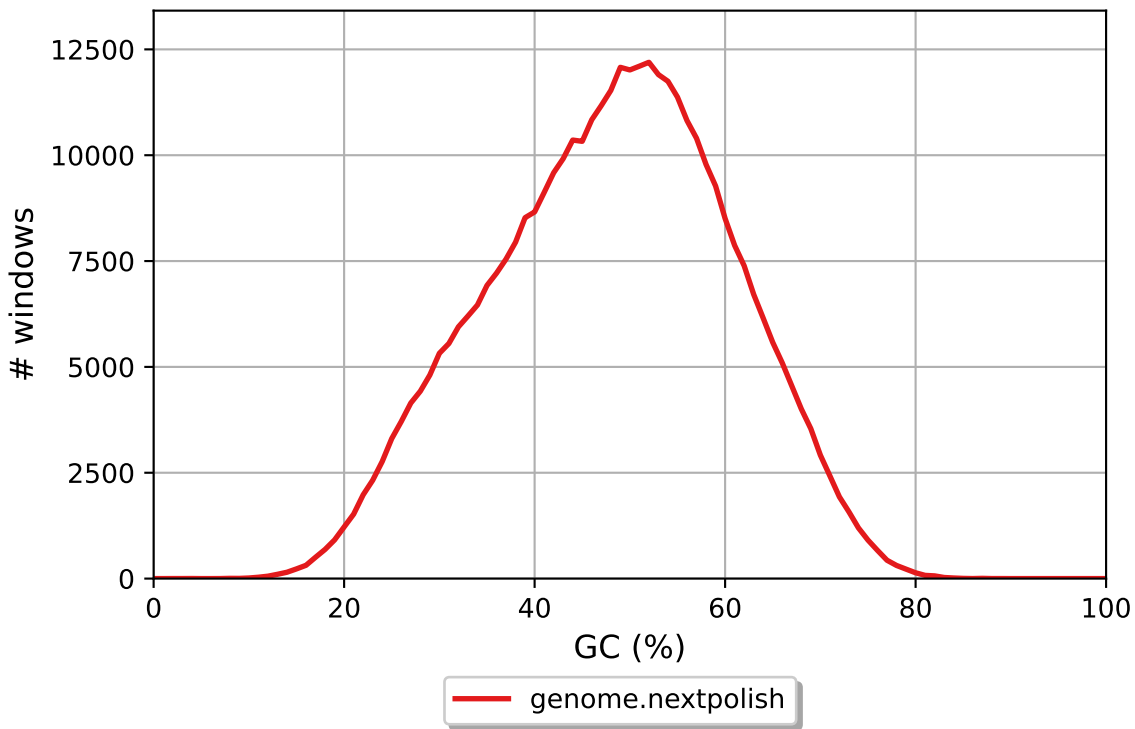
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

