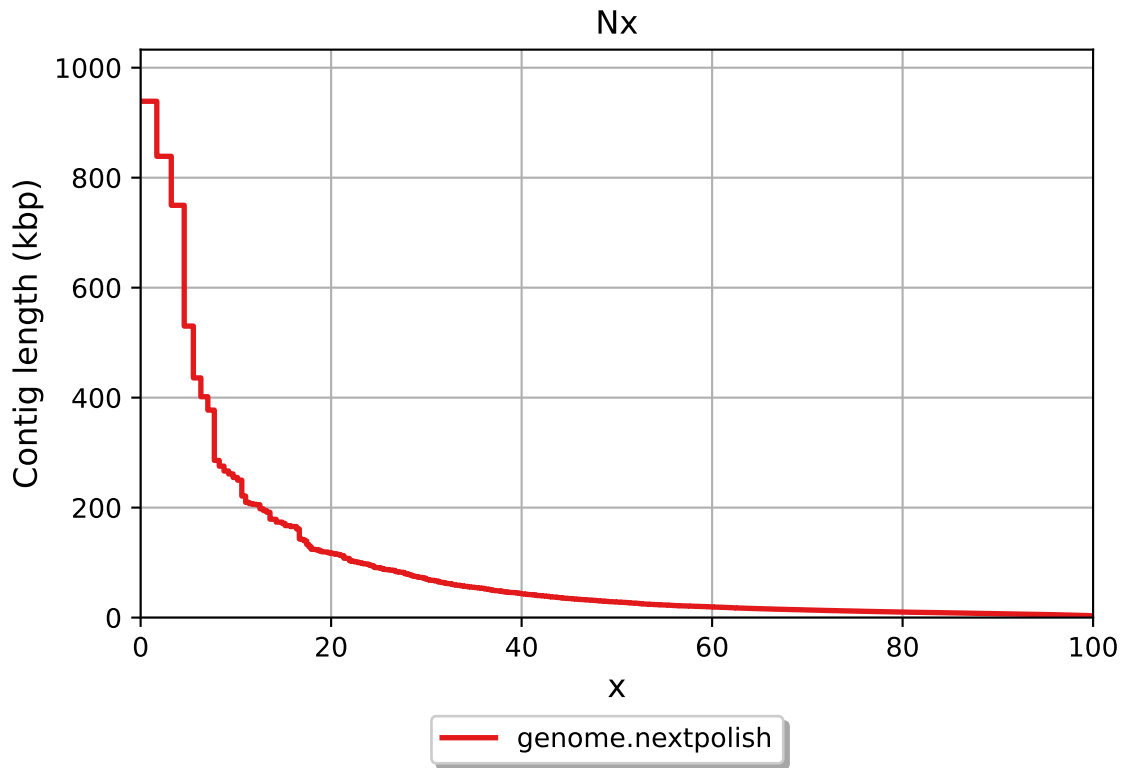


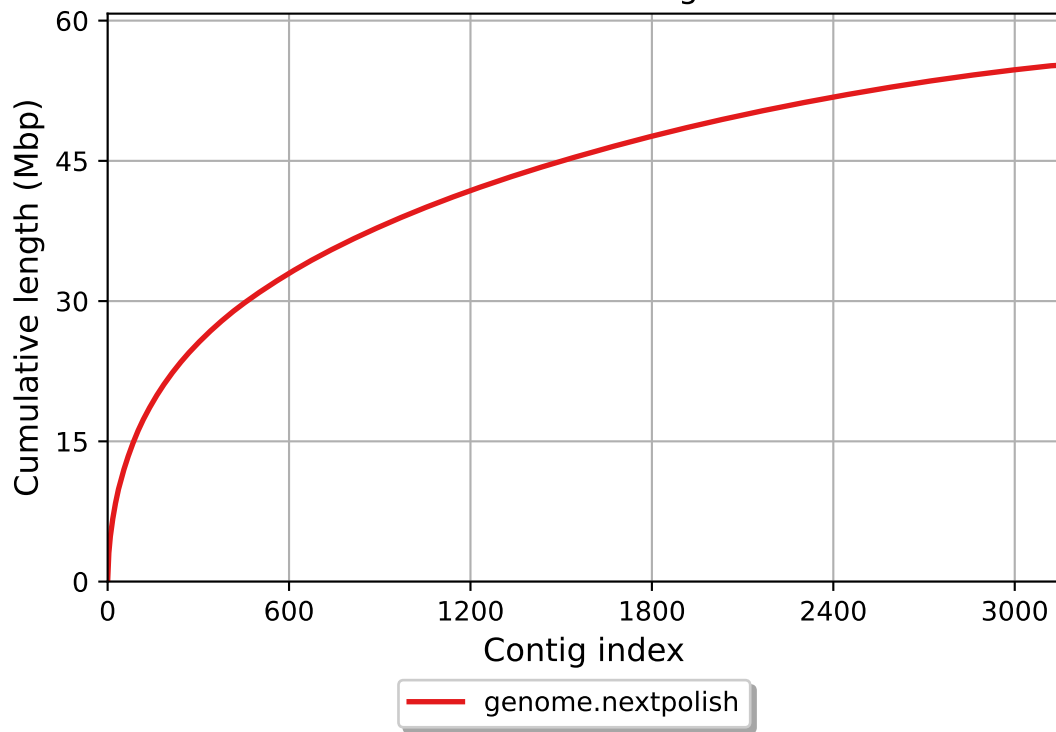
Report

	genome.nextpolish
# contigs (>= 0 bp)	3150
# contigs (>= 1000 bp)	3144
# contigs (>= 5000 bp)	2669
# contigs (>= 10000 bp)	1427
# contigs (>= 25000 bp)	422
# contigs (>= 50000 bp)	172
Total length (>= 0 bp)	55224556
Total length (>= 1000 bp)	55220039
Total length (>= 5000 bp)	53290384
Total length (>= 10000 bp)	44257291
Total length (>= 25000 bp)	29070546
Total length (>= 50000 bp)	20400296
# contigs	3150
Largest contig	938957
Total length	55224556
GC (%)	41.93
N50	28458
N90	7075
auN	100962.8
L50	368
L90	2071
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	136 + 22 part

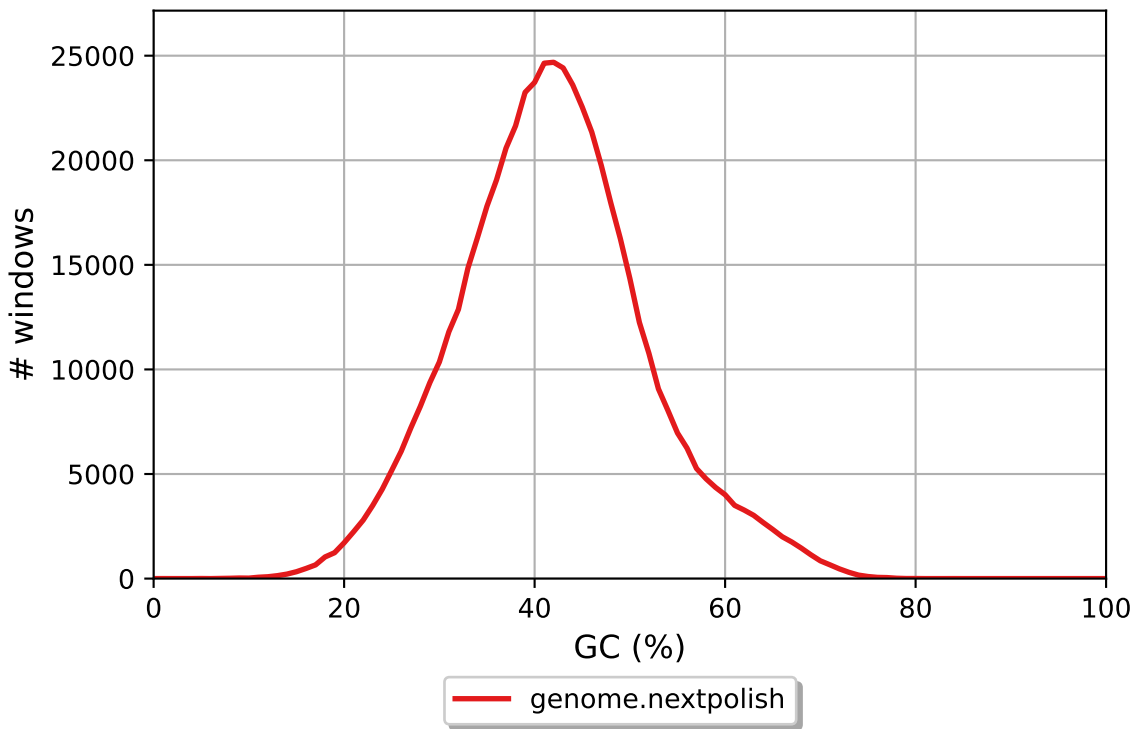
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

