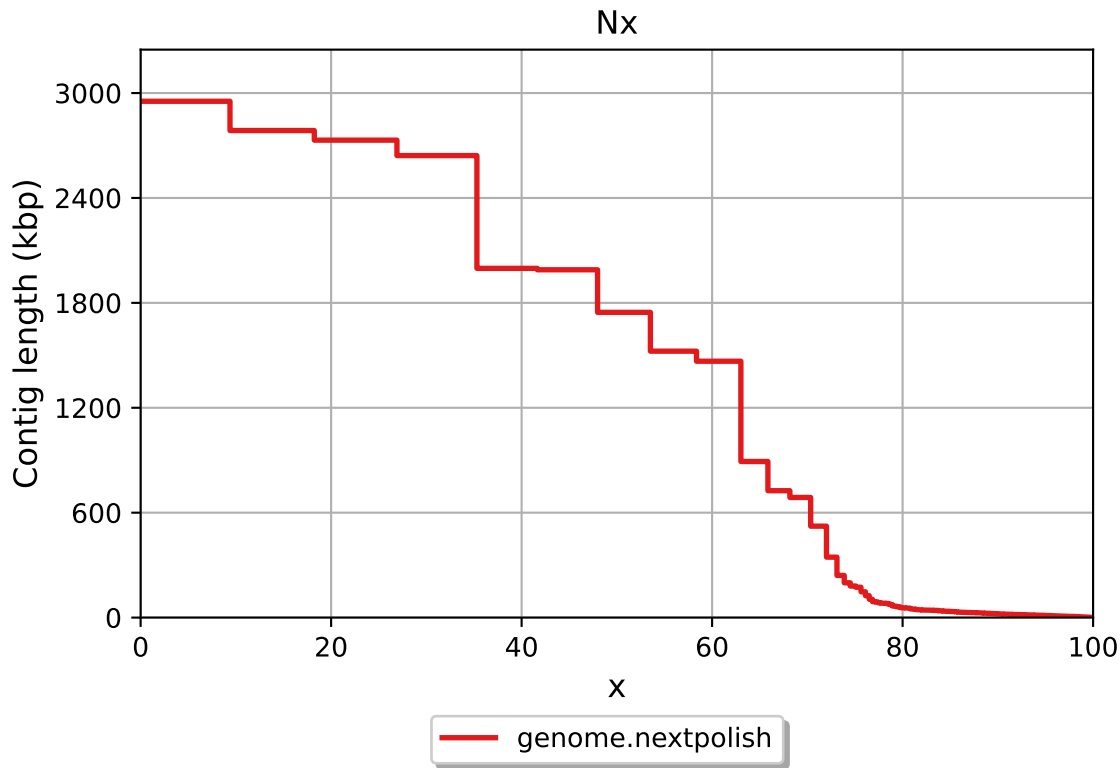


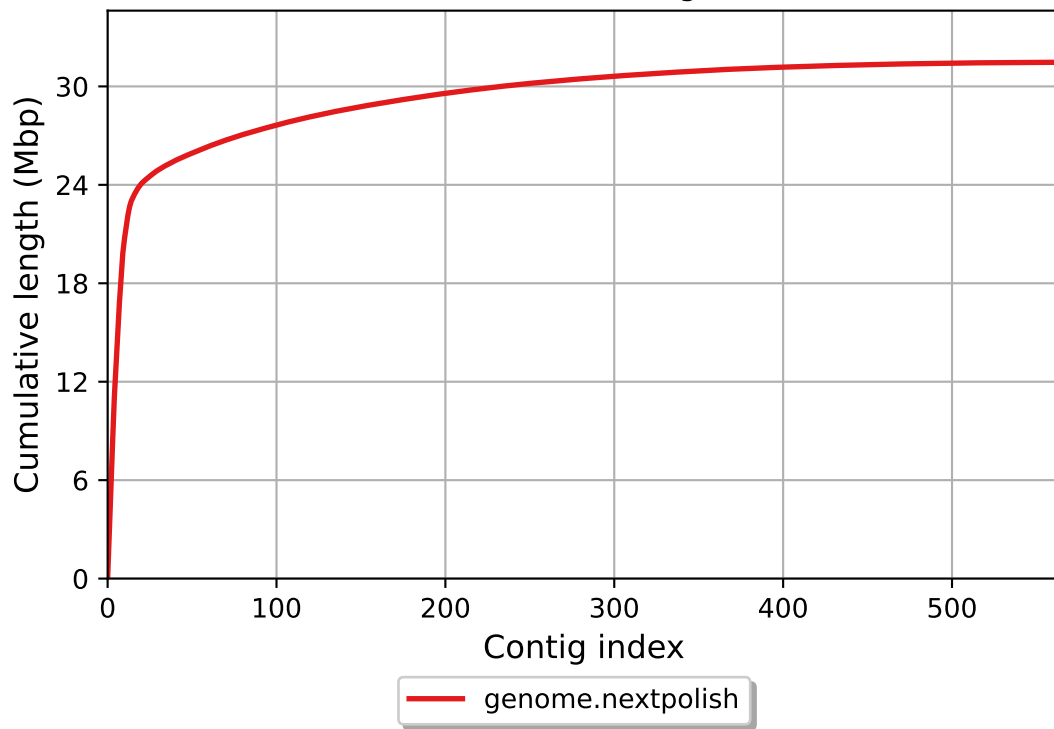
Report

	genome.nextpolish
# contigs (>= 0 bp)	566
# contigs (>= 1000 bp)	518
# contigs (>= 5000 bp)	364
# contigs (>= 10000 bp)	248
# contigs (>= 25000 bp)	110
# contigs (>= 50000 bp)	39
Total length (>= 0 bp)	31471733
Total length (>= 1000 bp)	31440363
Total length (>= 5000 bp)	31023273
Total length (>= 10000 bp)	30163632
Total length (>= 25000 bp)	27908105
Total length (>= 50000 bp)	25429721
# contigs	564
Largest contig	2952998
Total length	31471364
GC (%)	39.55
N50	1745768
N90	21782
auN	1557236.5
L50	7
L90	128
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	170 + 10 part

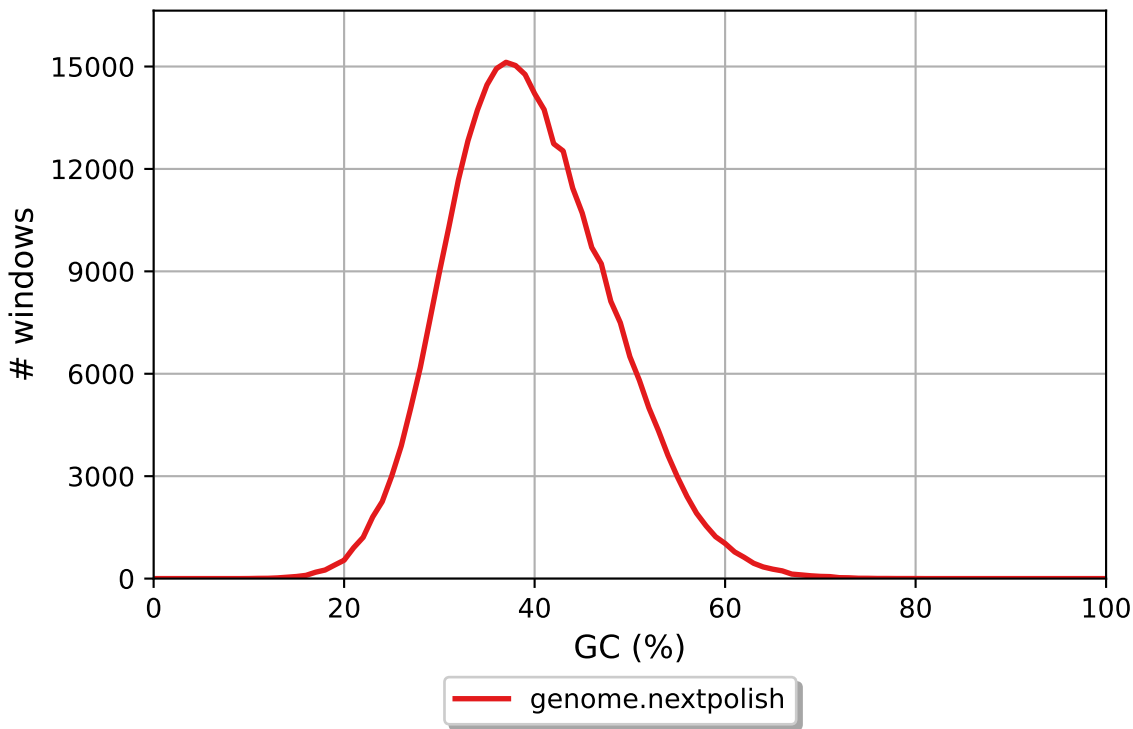
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

