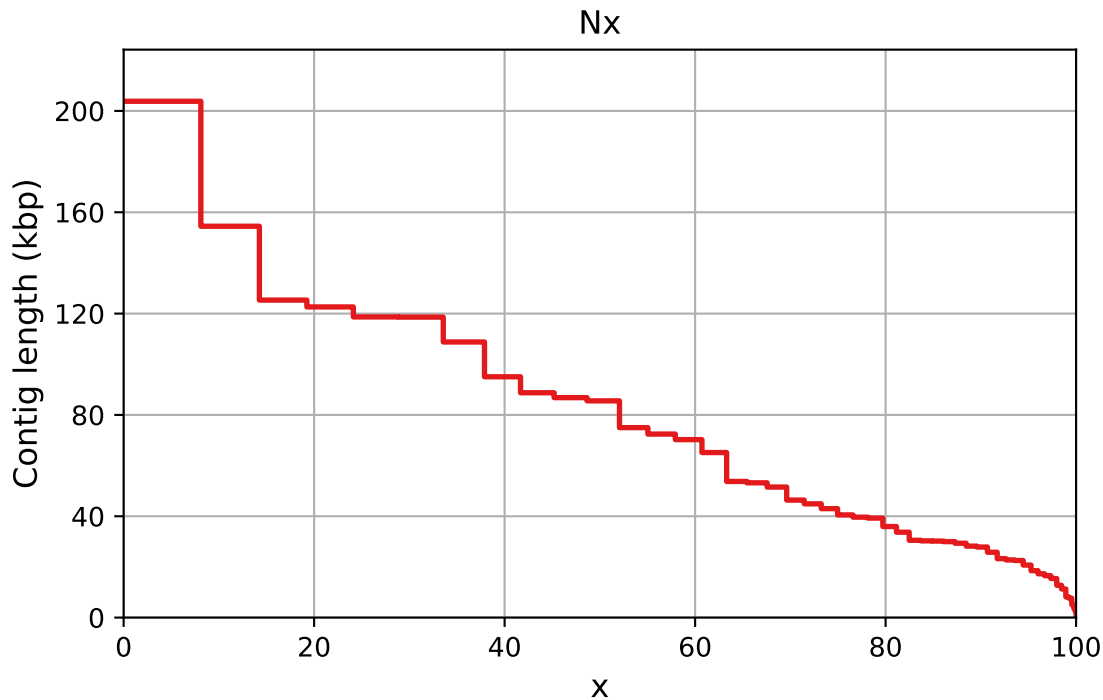


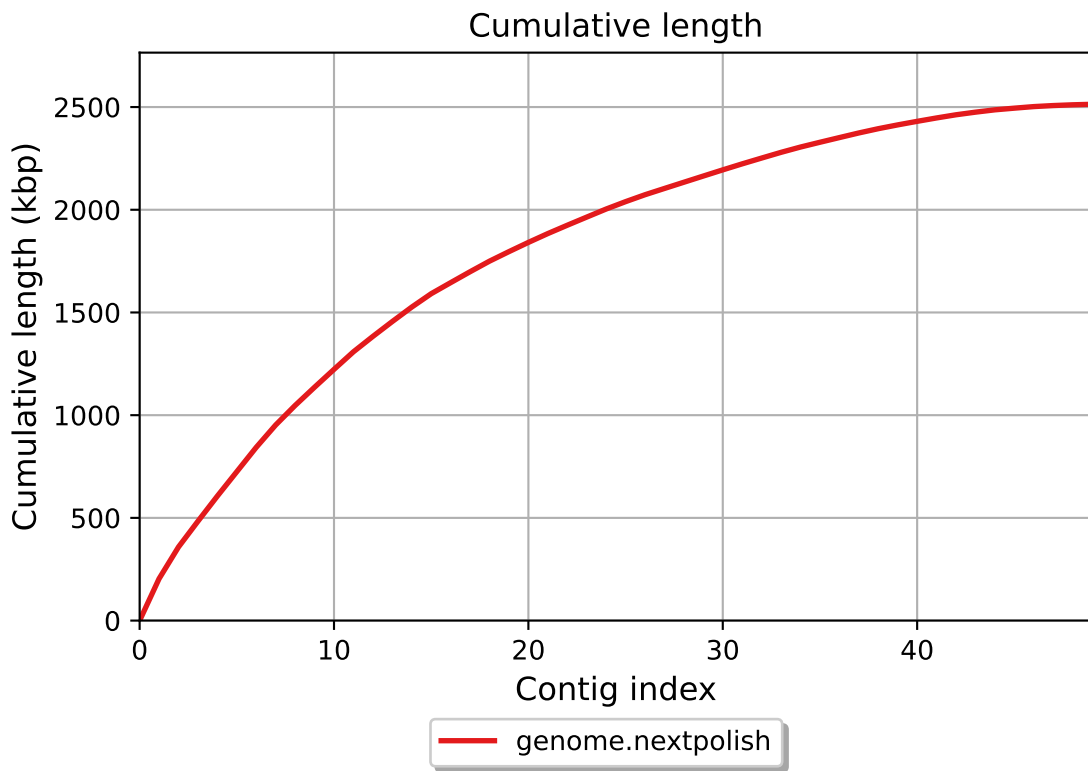
Report

	genome.nextpolish
# contigs (>= 0 bp)	49
# contigs (>= 1000 bp)	49
# contigs (>= 5000 bp)	47
# contigs (>= 10000 bp)	44
# contigs (>= 25000 bp)	34
# contigs (>= 50000 bp)	18
Total length (>= 0 bp)	2513542
Total length (>= 1000 bp)	2513542
Total length (>= 5000 bp)	2507308
Total length (>= 10000 bp)	2486436
Total length (>= 25000 bp)	2305359
Total length (>= 50000 bp)	1749804
# contigs	49
Largest contig	203812
Total length	2513542
GC (%)	41.03
N50	85546
N90	27856
auN	87485.8
L50	11
L90	33
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	4 + 0 part

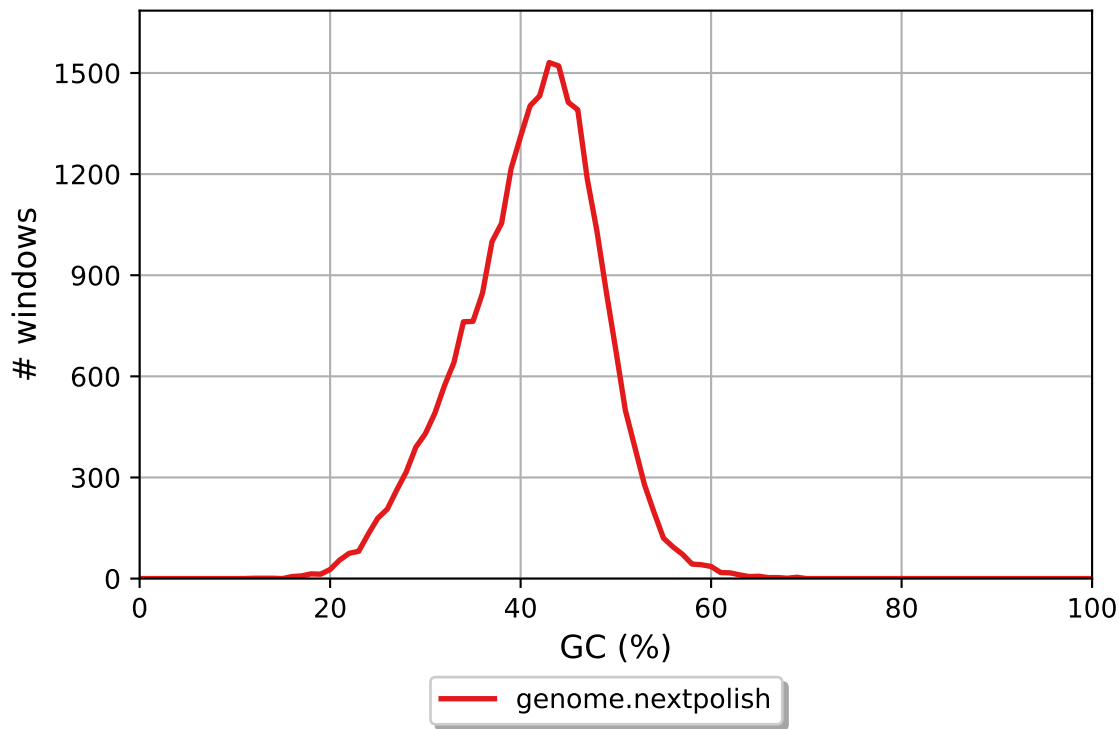
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



— genome.nextpolish



GC content



genome.nextpolish GC content

