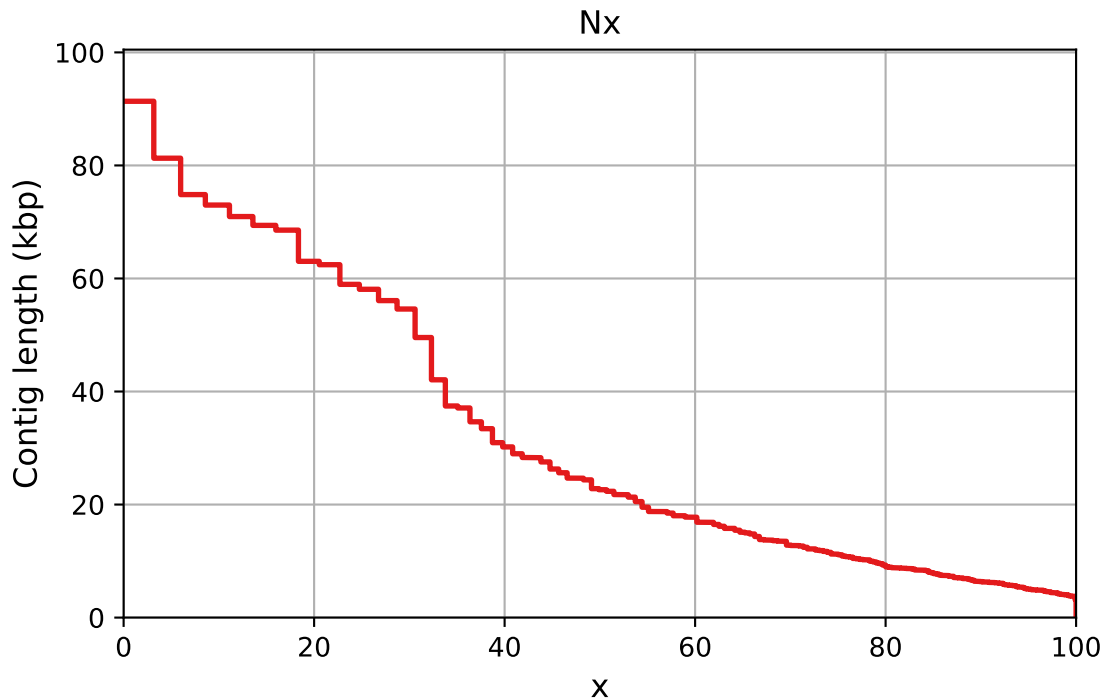


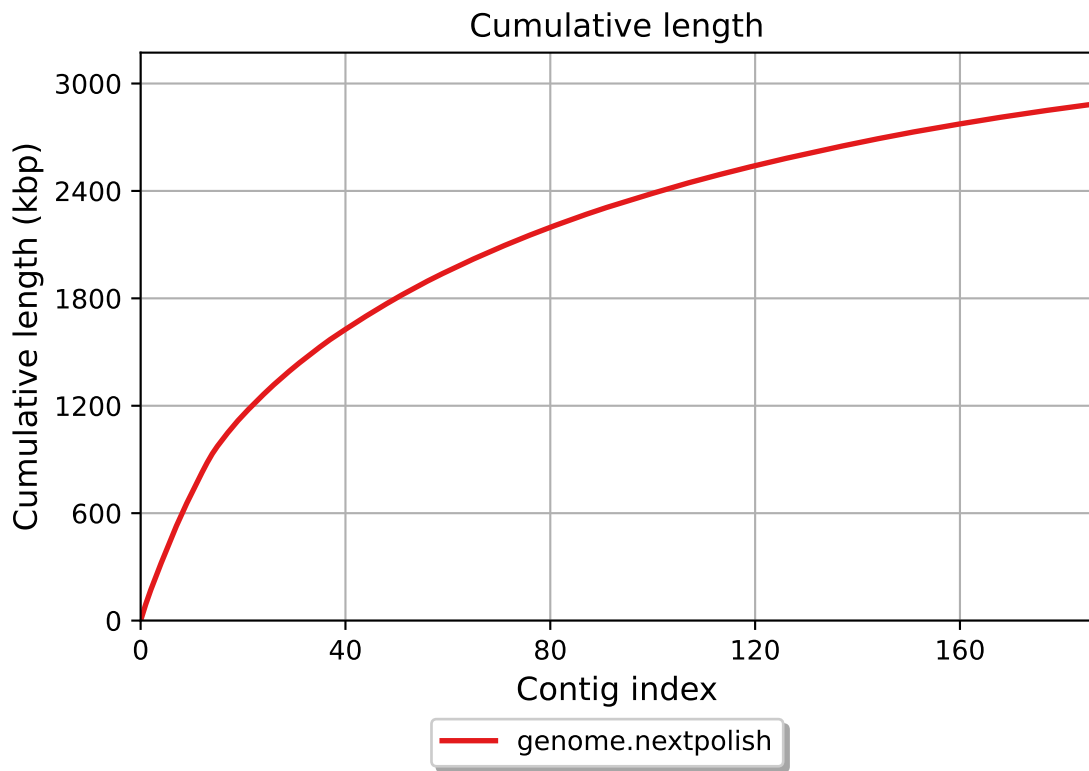
Report

	genome.nextpolish
# contigs (>= 0 bp)	186
# contigs (>= 1000 bp)	186
# contigs (>= 5000 bp)	154
# contigs (>= 10000 bp)	86
# contigs (>= 25000 bp)	27
# contigs (>= 50000 bp)	13
Total length (>= 0 bp)	2884089
Total length (>= 1000 bp)	2884089
Total length (>= 5000 bp)	2744990
Total length (>= 10000 bp)	2259302
Total length (>= 25000 bp)	1342984
Total length (>= 50000 bp)	882563
# contigs	186
Largest contig	91353
Total length	2884089
GC (%)	31.71
N50	22624
N90	6368
auN	33449.4
L50	32
L90	129
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	10 + 0 part

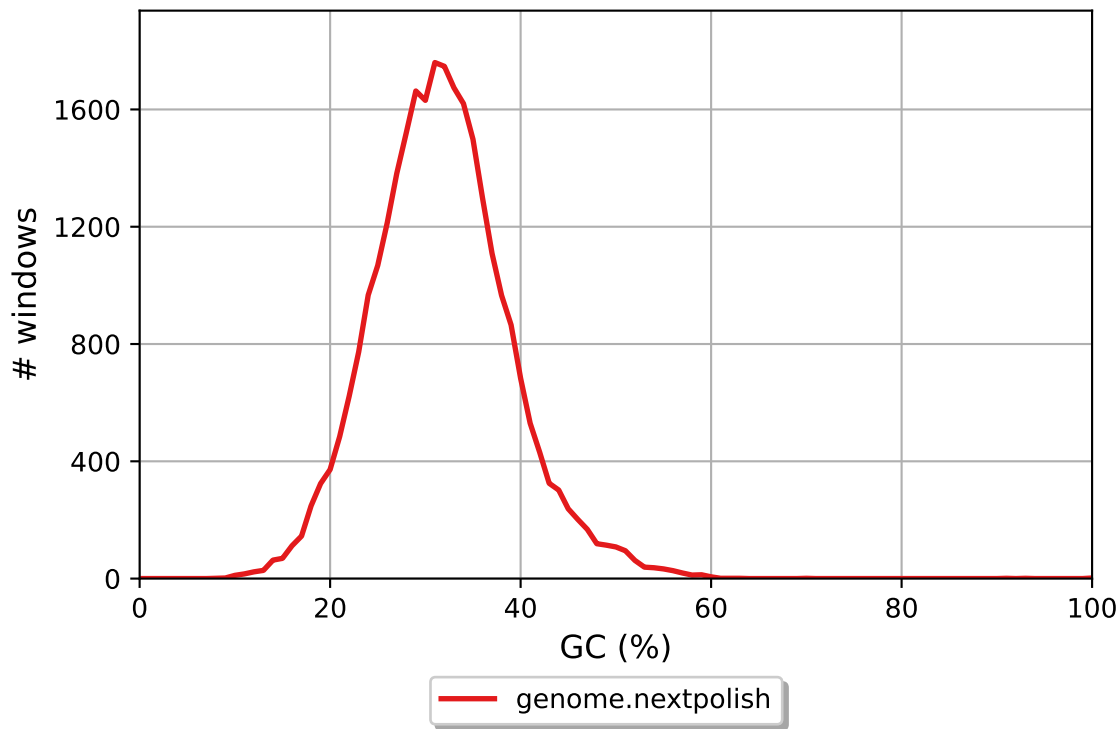
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



— genome.nextpolish



GC content



genome.nextpolish GC content

