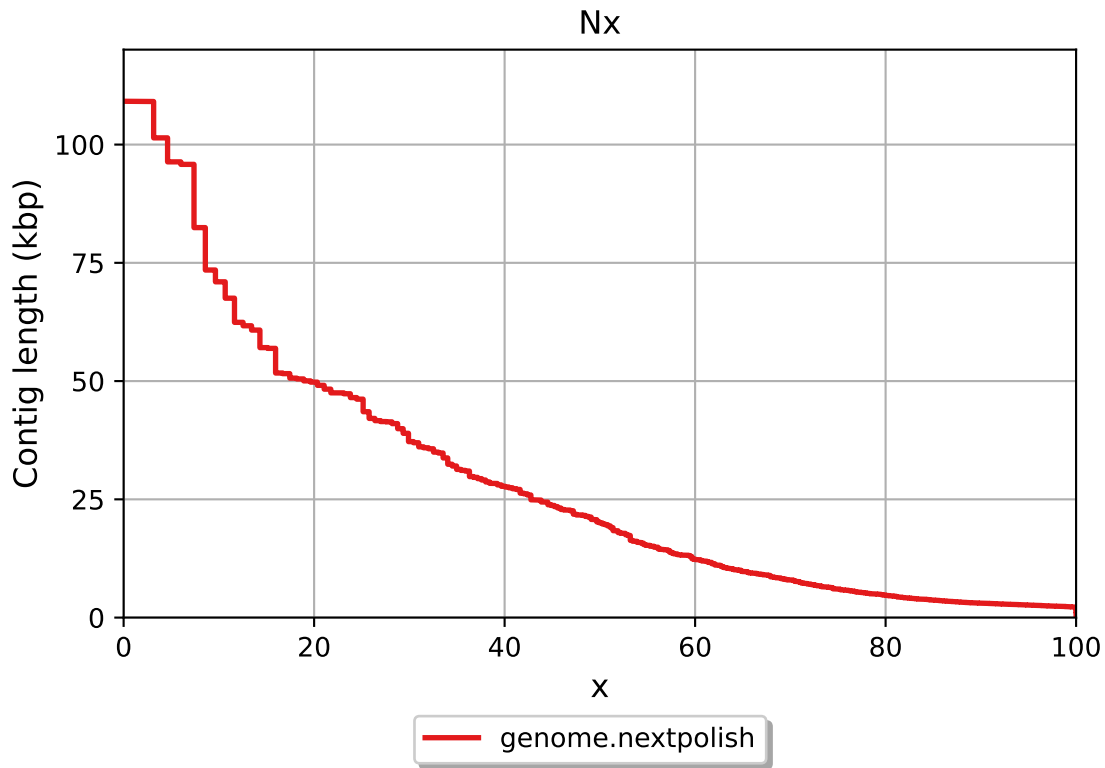


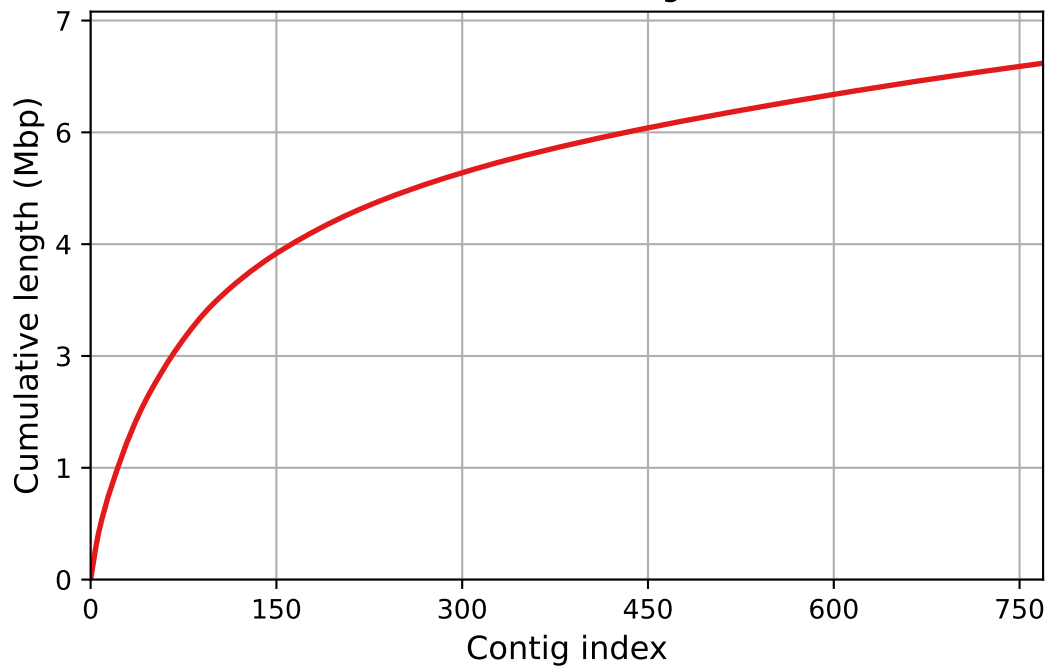
Report

	genome.nextpolish
# contigs (>= 0 bp)	769
# contigs (>= 1000 bp)	769
# contigs (>= 5000 bp)	297
# contigs (>= 10000 bp)	161
# contigs (>= 25000 bp)	64
# contigs (>= 50000 bp)	19
Total length (>= 0 bp)	6926827
Total length (>= 1000 bp)	6926827
Total length (>= 5000 bp)	5443161
Total length (>= 10000 bp)	4488864
Total length (>= 25000 bp)	2961920
Total length (>= 50000 bp)	1359563
# contigs	769
Largest contig	109156
Total length	6926827
GC (%)	44.41
N50	20020
N90	3050
auN	29264.6
L50	87
L90	505
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	15 + 2 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).

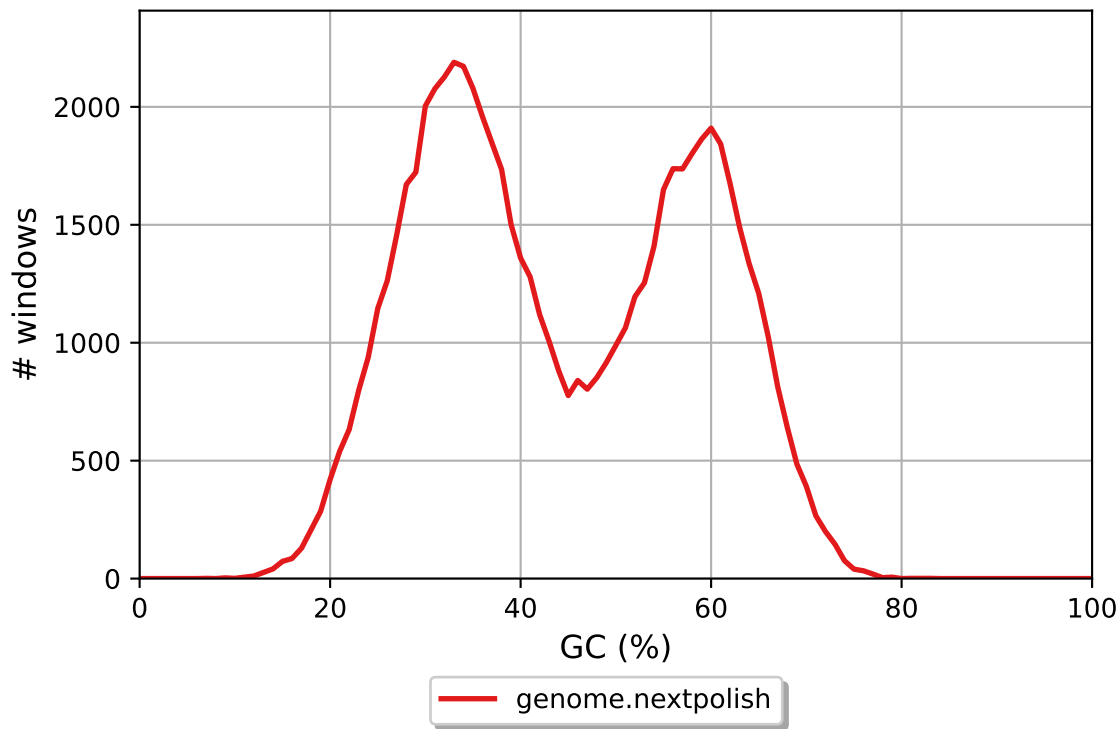


Cumulative length



— genome.nextpolish

GC content



genome.nextpolish GC content

