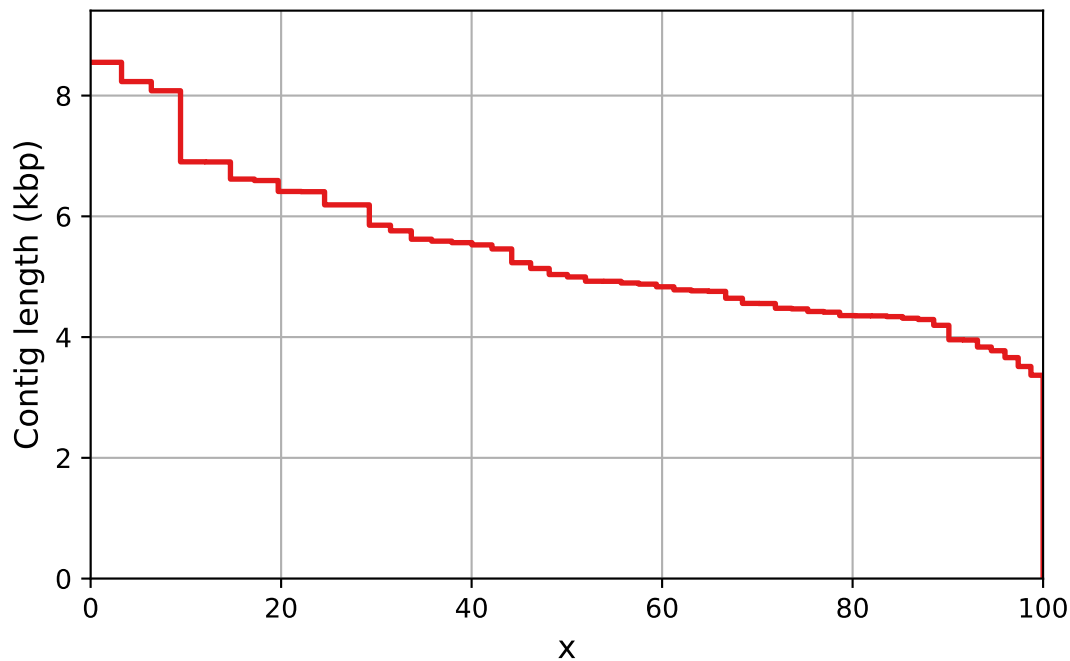


Report

	genome.nextpolish
# contigs (>= 0 bp)	51
# contigs (>= 1000 bp)	51
# contigs (>= 5000 bp)	21
# contigs (>= 10000 bp)	0
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	263401
Total length (>= 1000 bp)	263401
Total length (>= 5000 bp)	131855
Total length (>= 10000 bp)	0
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	51
Largest contig	8551
Total length	263401
GC (%)	43.84
N50	5036
N90	4195
auN	5431.6
L50	21
L90	44
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	1 + 0 part

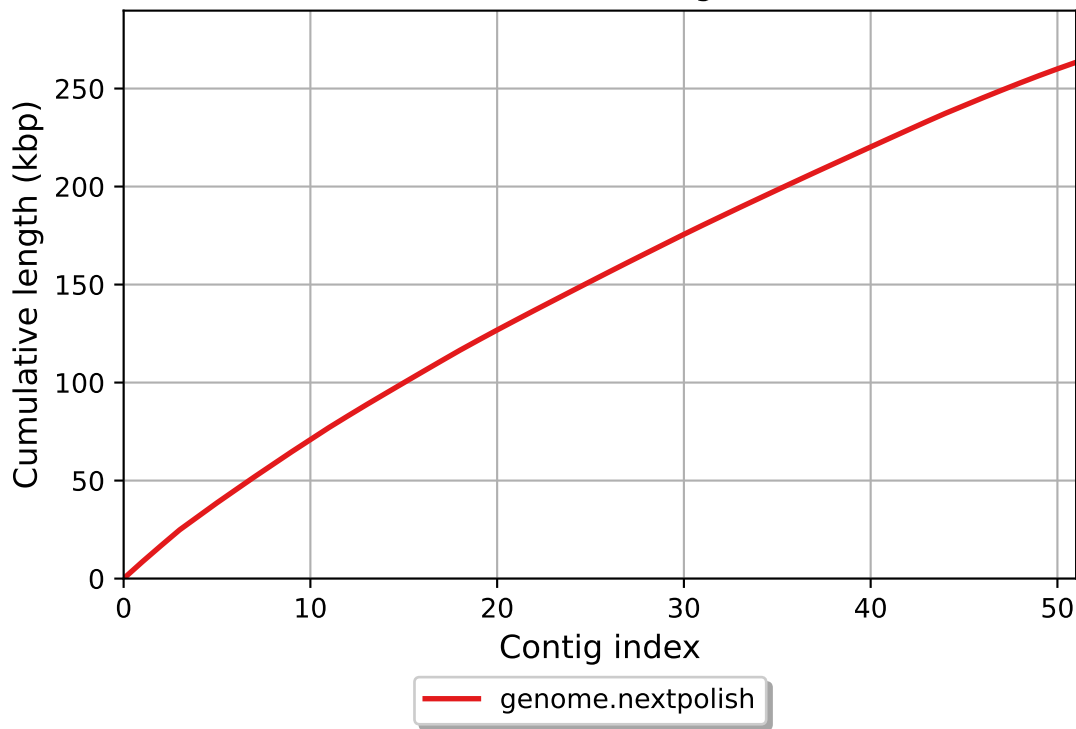
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

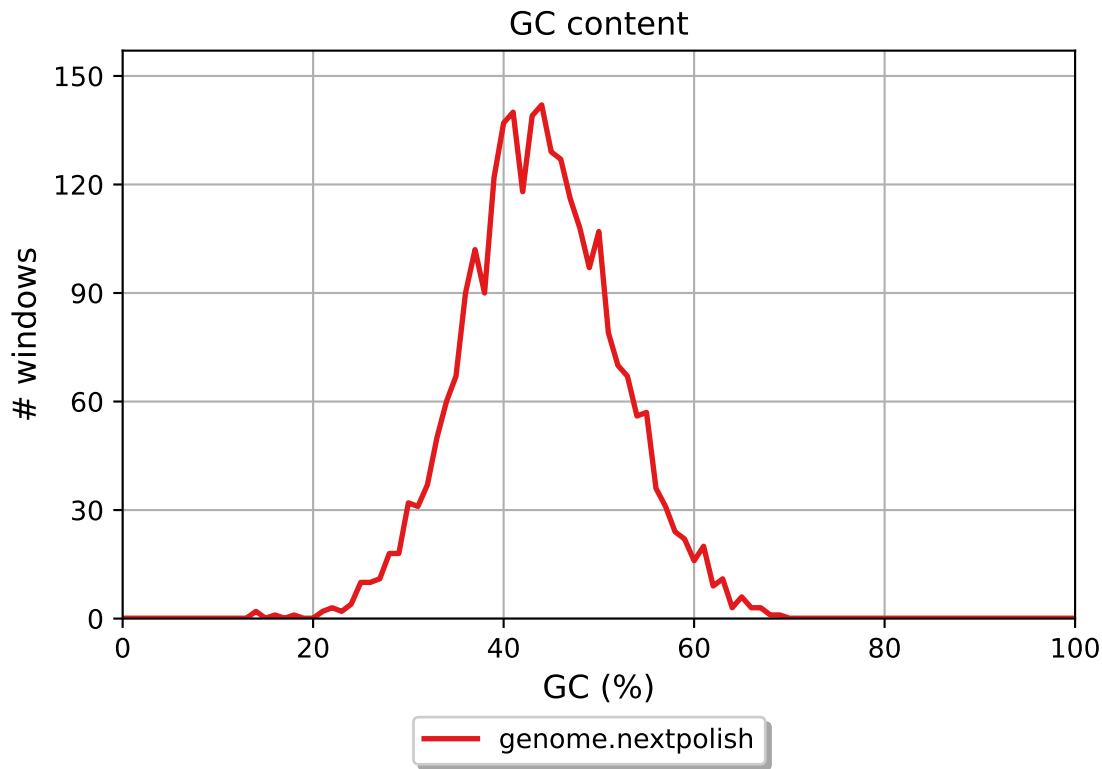
Nx



— genome.nextpolish

Cumulative length





genome.nextpolish GC content

