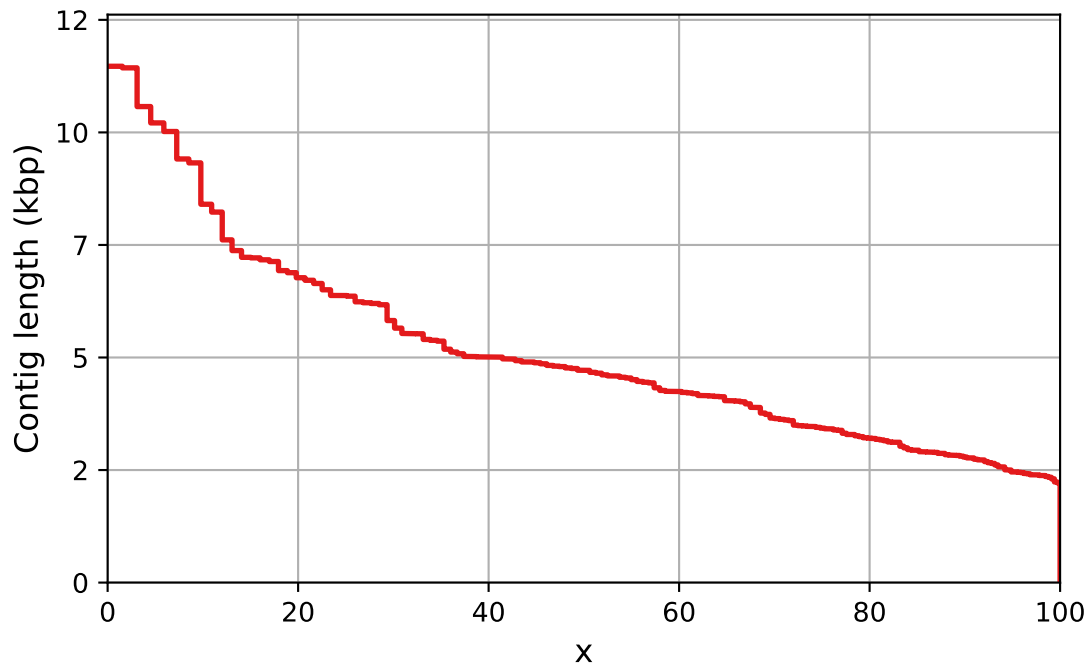


# Report

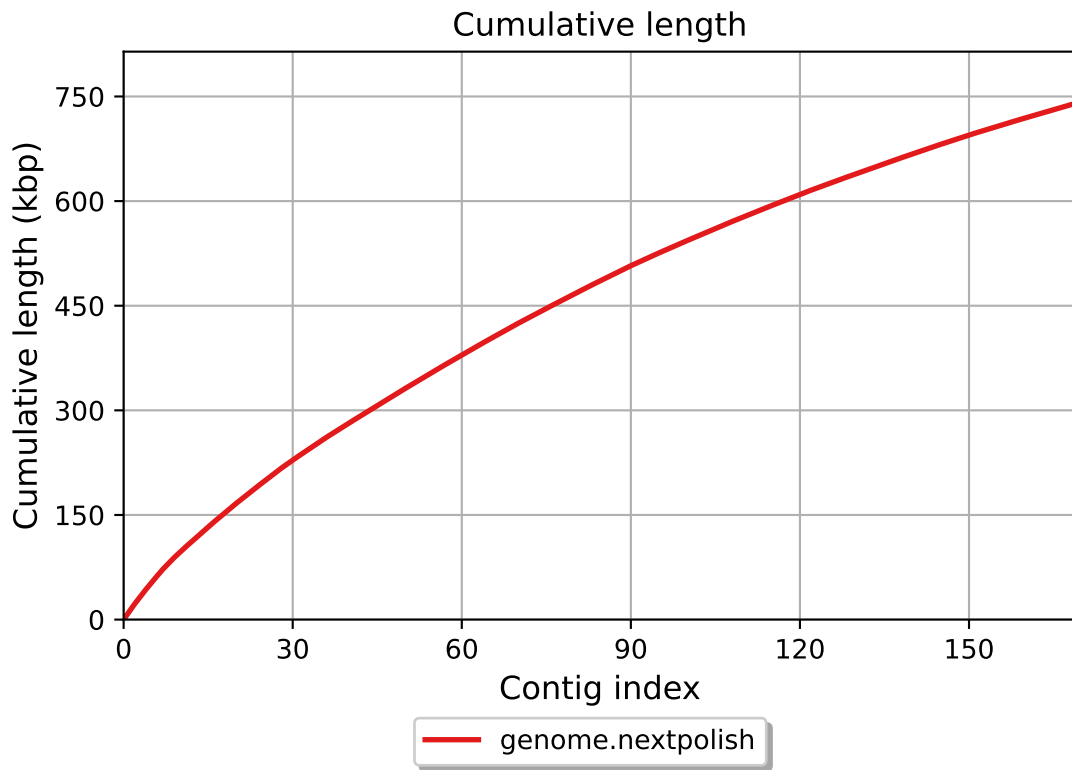
	genome.nextpolish
# contigs (>= 0 bp)	169
# contigs (>= 1000 bp)	169
# contigs (>= 5000 bp)	45
# contigs (>= 10000 bp)	5
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	740263
Total length (>= 1000 bp)	740263
Total length (>= 5000 bp)	306878
Total length (>= 10000 bp)	53708
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	169
Largest contig	11469
Total length	740263
GC (%)	34.96
N50	4714
N90	2789
auN	5177.7
L50	59
L90	140
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	5 + 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).

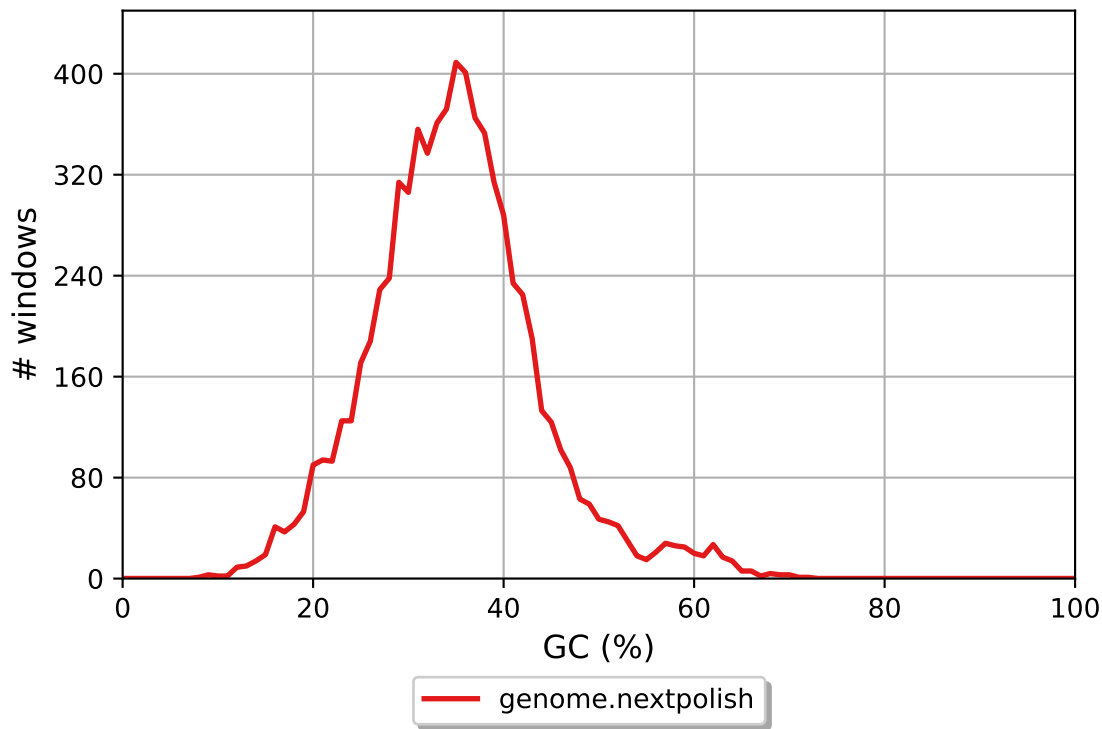
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

