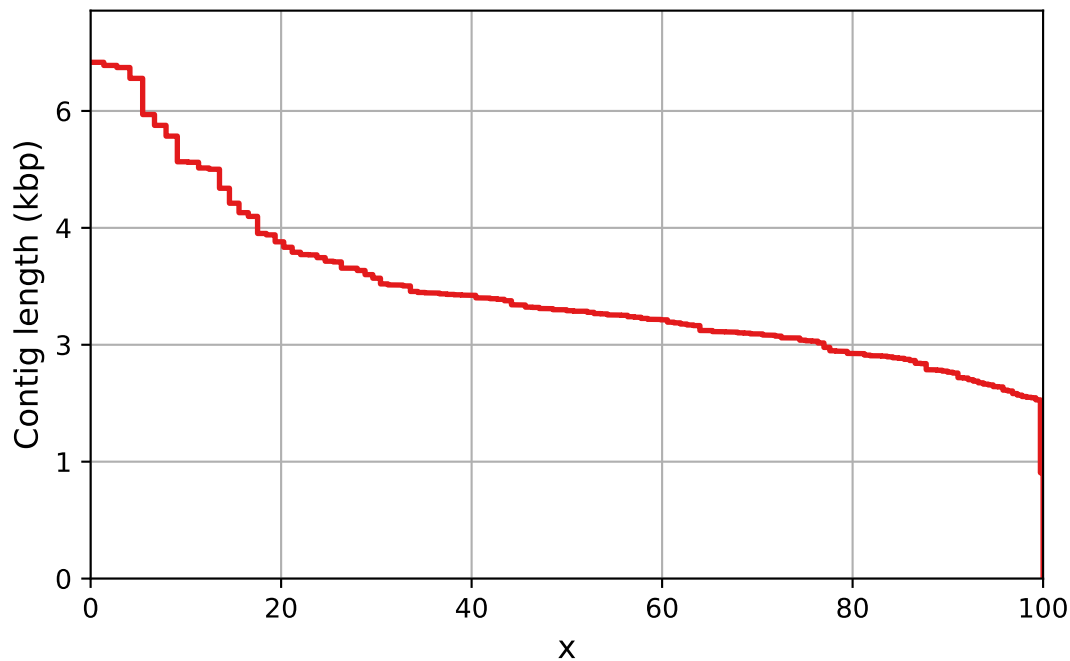


# Report

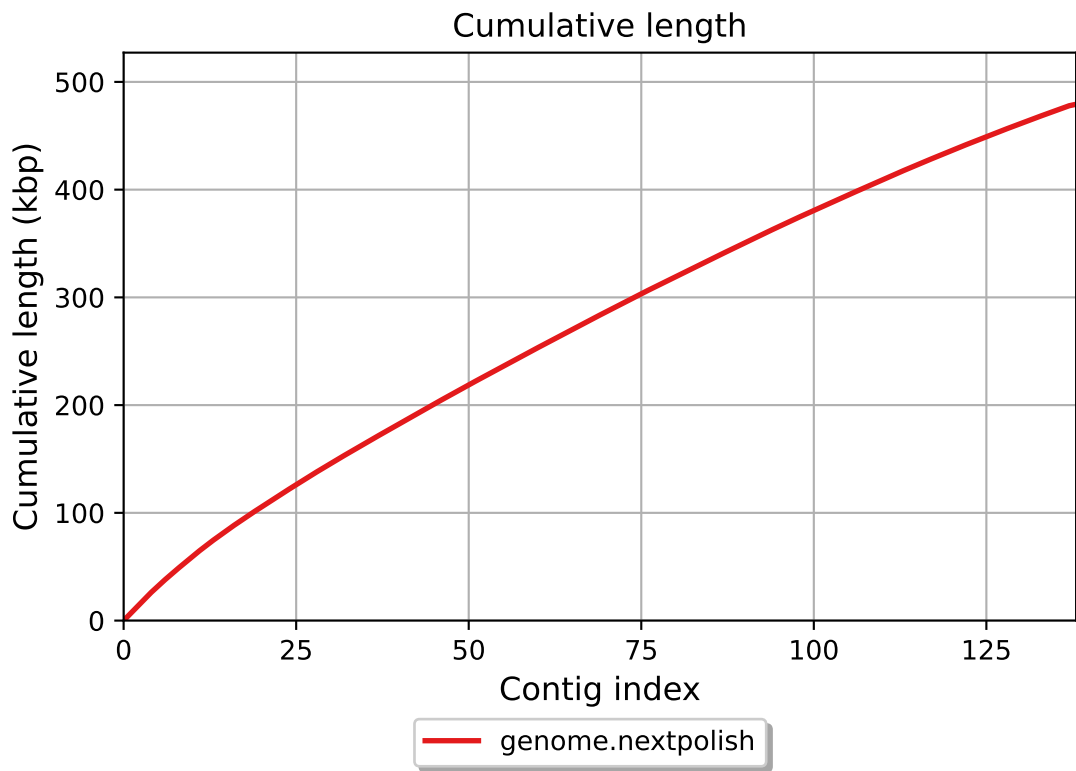
	genome.nextpolish
# contigs (>= 0 bp)	138
# contigs (>= 1000 bp)	138
# contigs (>= 5000 bp)	12
# contigs (>= 10000 bp)	0
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	479244
Total length (>= 1000 bp)	479244
Total length (>= 5000 bp)	69846
Total length (>= 10000 bp)	0
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	138
Largest contig	6624
Total length	479244
GC (%)	40.08
N50	3437
N90	2648
auN	3720.3
L50	57
L90	119
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	0 + 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).

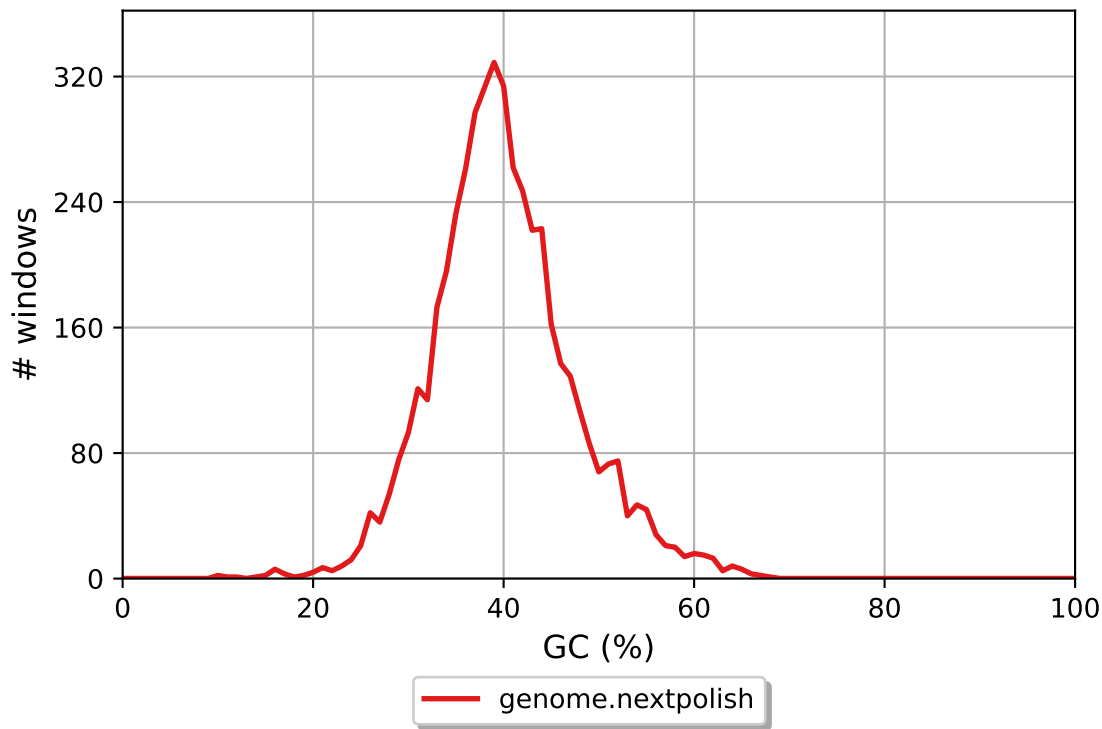
Nx



— genome.nextpolish



## GC content



genome.nextpolish GC content

