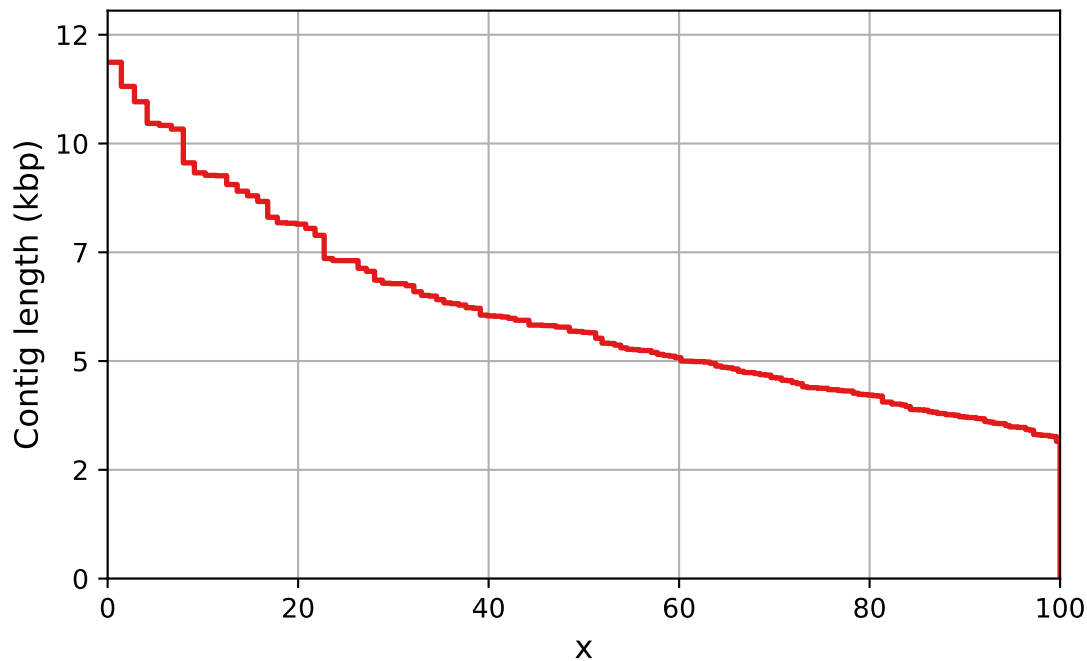


# Report

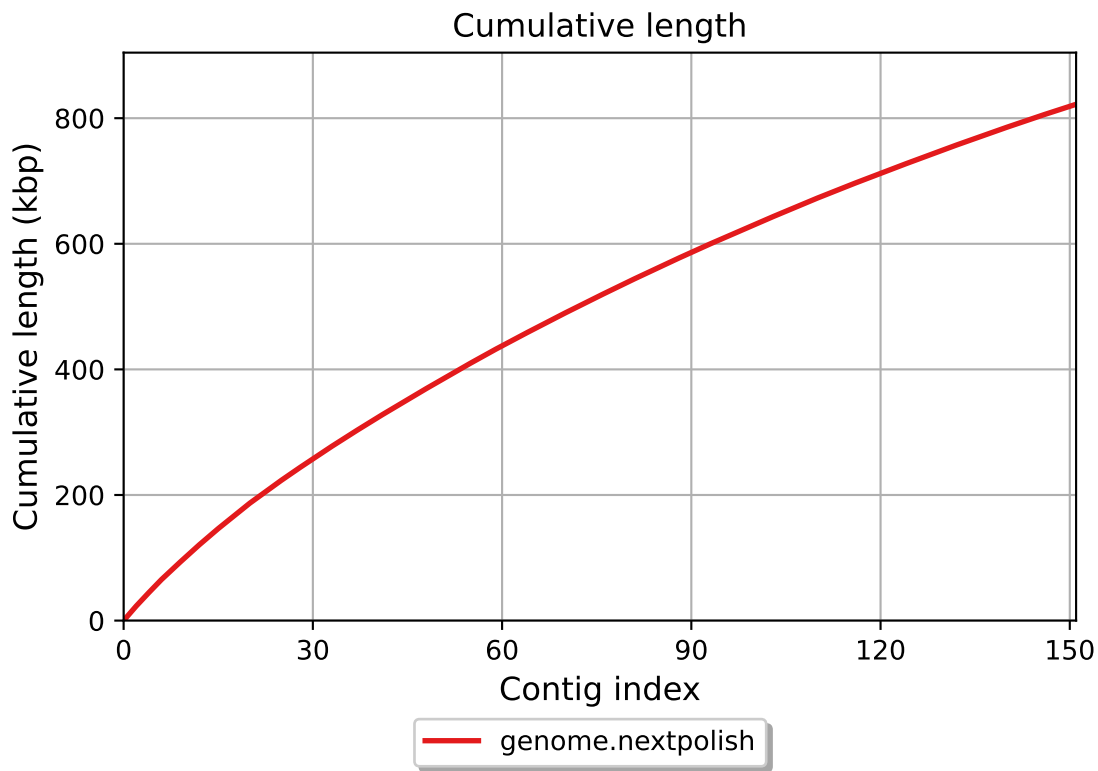
	genome.nextpolish
# contigs (>= 0 bp)	151
# contigs (>= 1000 bp)	151
# contigs (>= 5000 bp)	72
# contigs (>= 10000 bp)	6
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	822163
Total length (>= 1000 bp)	822163
Total length (>= 5000 bp)	499924
Total length (>= 10000 bp)	65348
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	151
Largest contig	11867
Total length	822163
GC (%)	50.87
N50	5659
N90	3714
auN	6101.5
L50	56
L90	128
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	4 + 1 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).

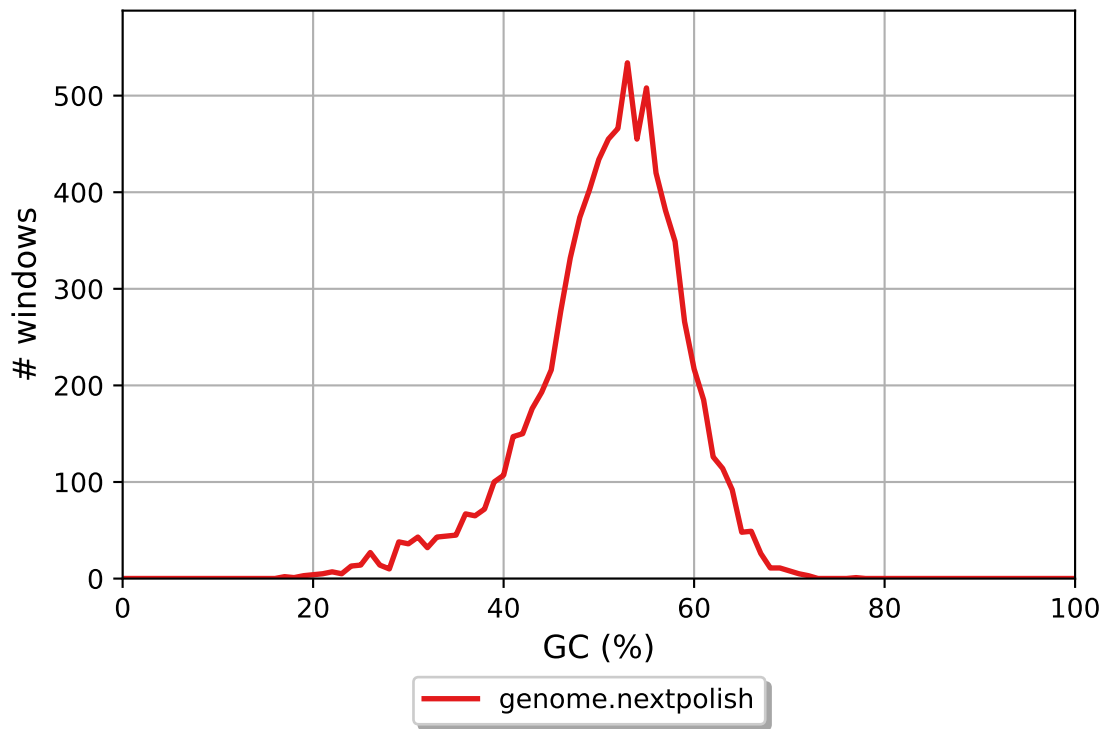
Nx



— genome.nextpolish



## GC content



genome.nextpolish GC content

