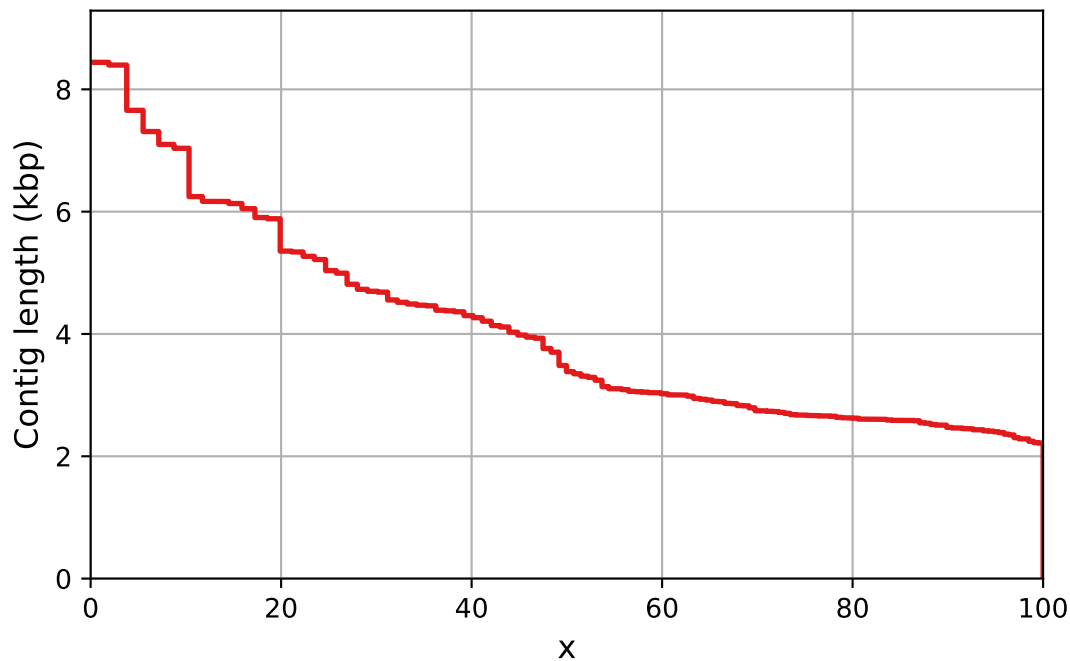


# Report

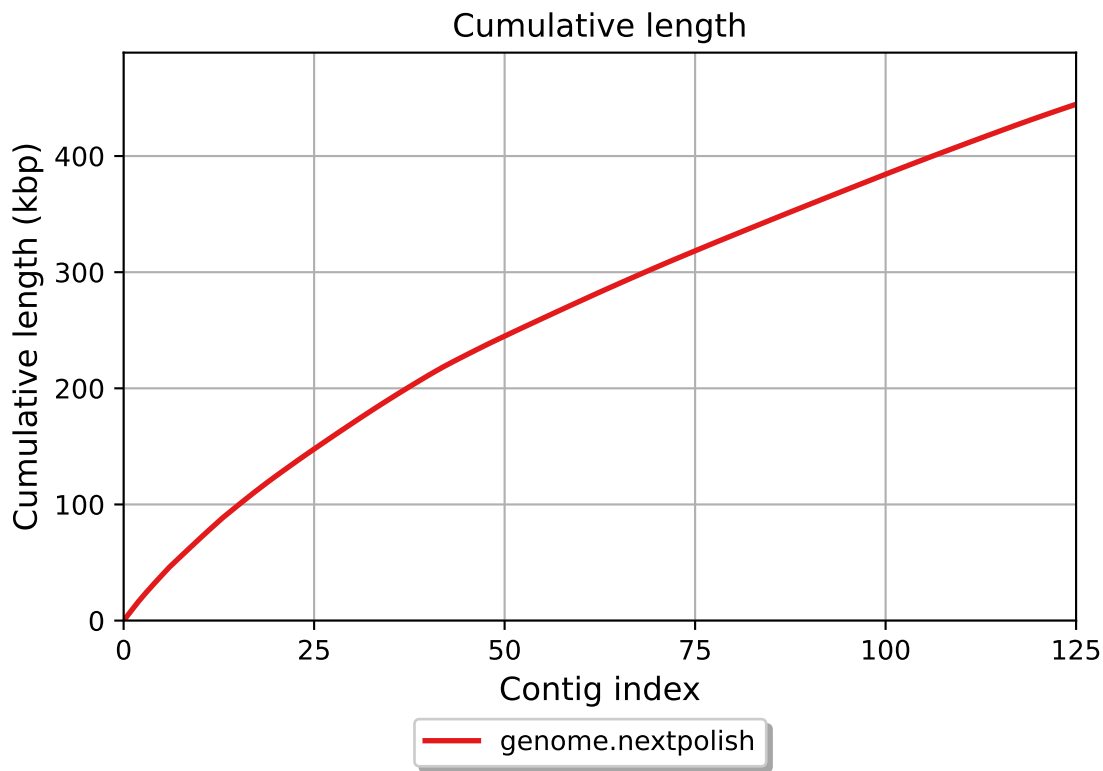
	genome.nextpolish
# contigs (>= 0 bp)	125
# contigs (>= 1000 bp)	125
# contigs (>= 5000 bp)	18
# contigs (>= 10000 bp)	0
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	444589
Total length (>= 1000 bp)	444589
Total length (>= 5000 bp)	114709
Total length (>= 10000 bp)	0
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	125
Largest contig	8442
Total length	444589
GC (%)	38.57
N50	3384
N90	2472
auN	4104.5
L50	44
L90	107
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	3 + 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).

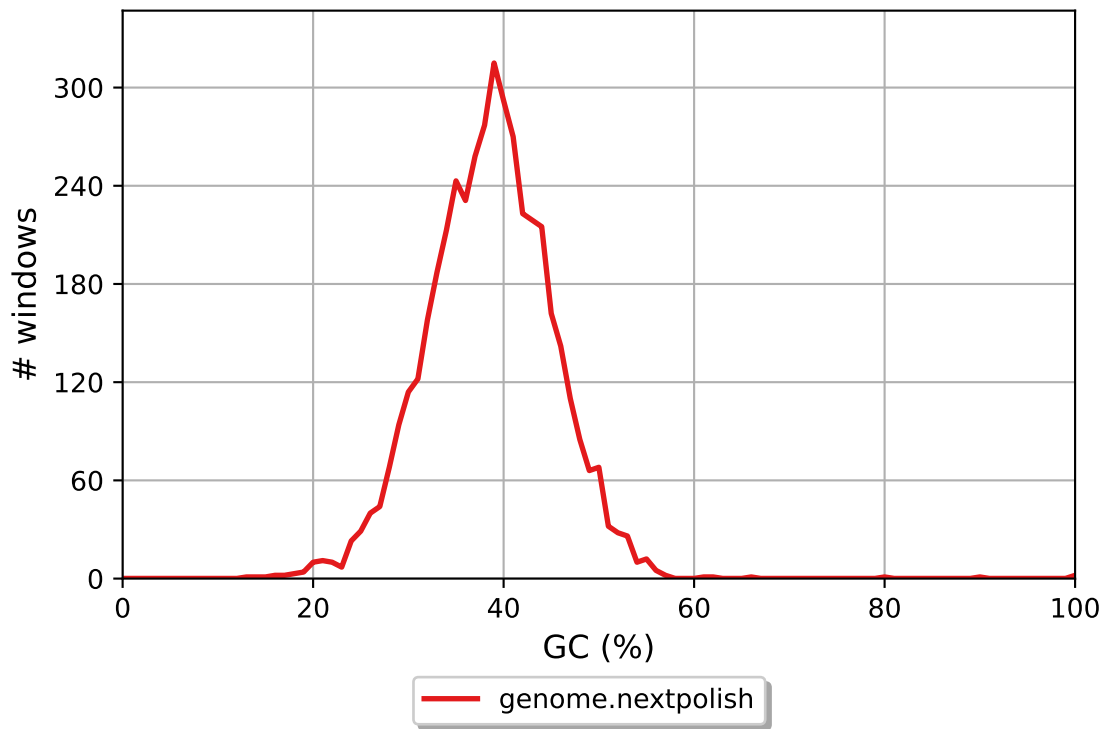
Nx



— genome.nextpolish



## GC content



genome.nextpolish GC content

