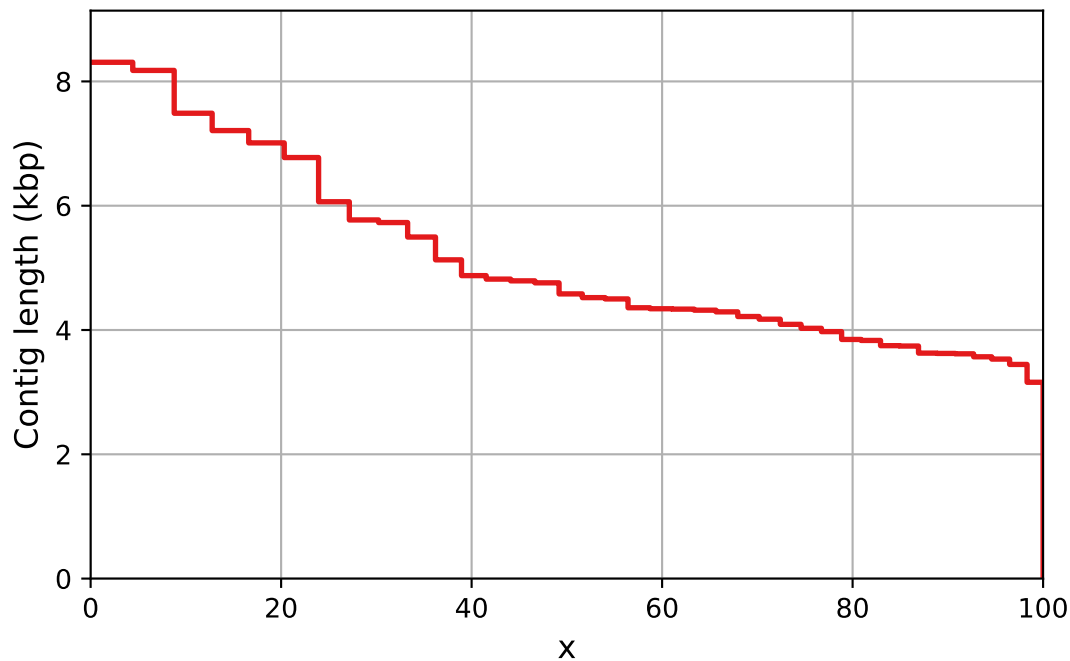


# Report

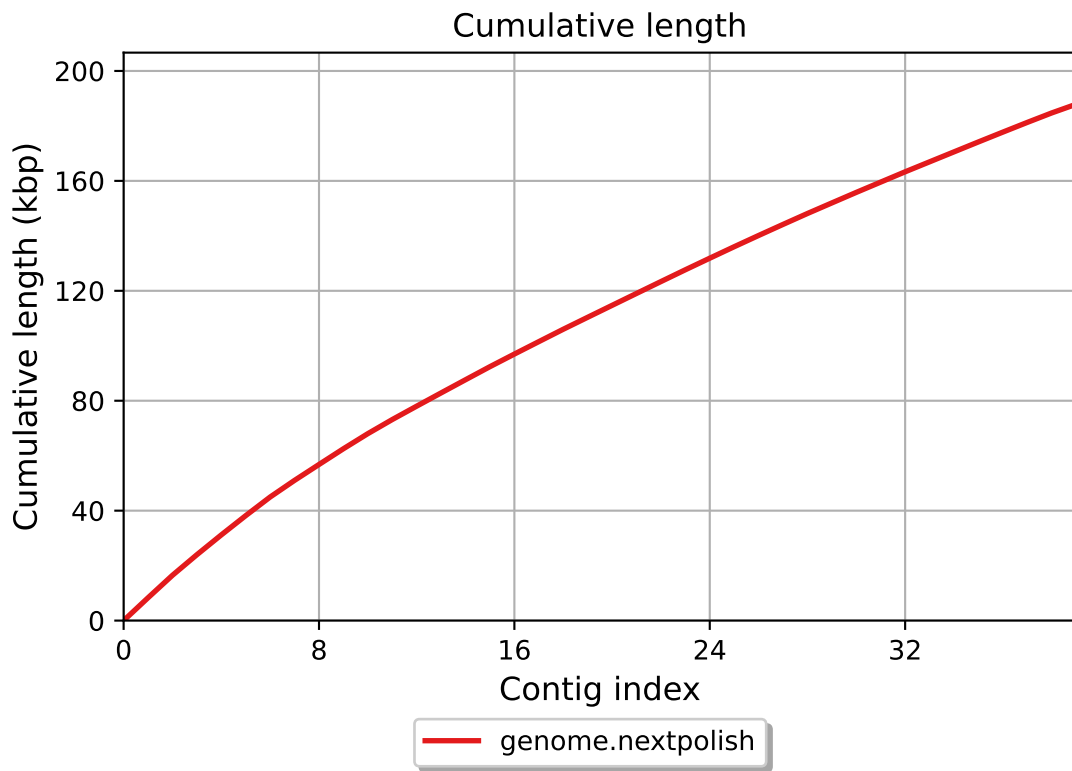
	genome.nextpolish
# contigs (>= 0 bp)	39
# contigs (>= 1000 bp)	39
# contigs (>= 5000 bp)	11
# contigs (>= 10000 bp)	0
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	187877
Total length (>= 1000 bp)	187877
Total length (>= 5000 bp)	73157
Total length (>= 10000 bp)	0
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	39
Largest contig	8308
Total length	187877
GC (%)	49.21
N50	4581
N90	3623
auN	5185.9
L50	16
L90	34
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	2 + 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).

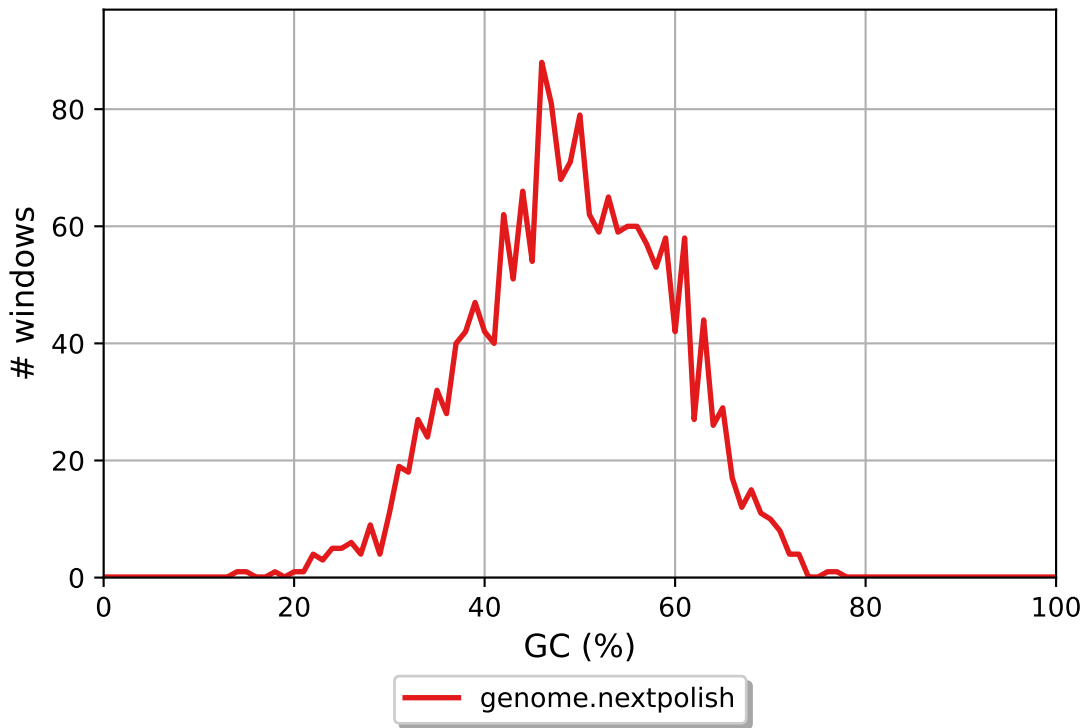
Nx



— genome.nextpolish



## GC content



genome.nextpolish GC content

