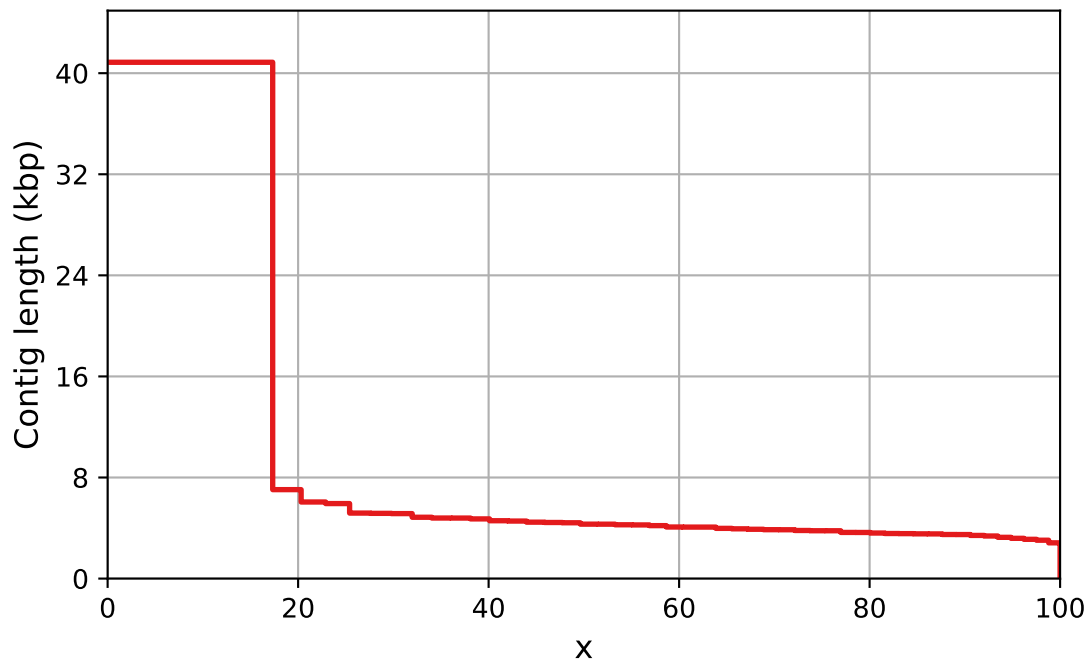


# Report

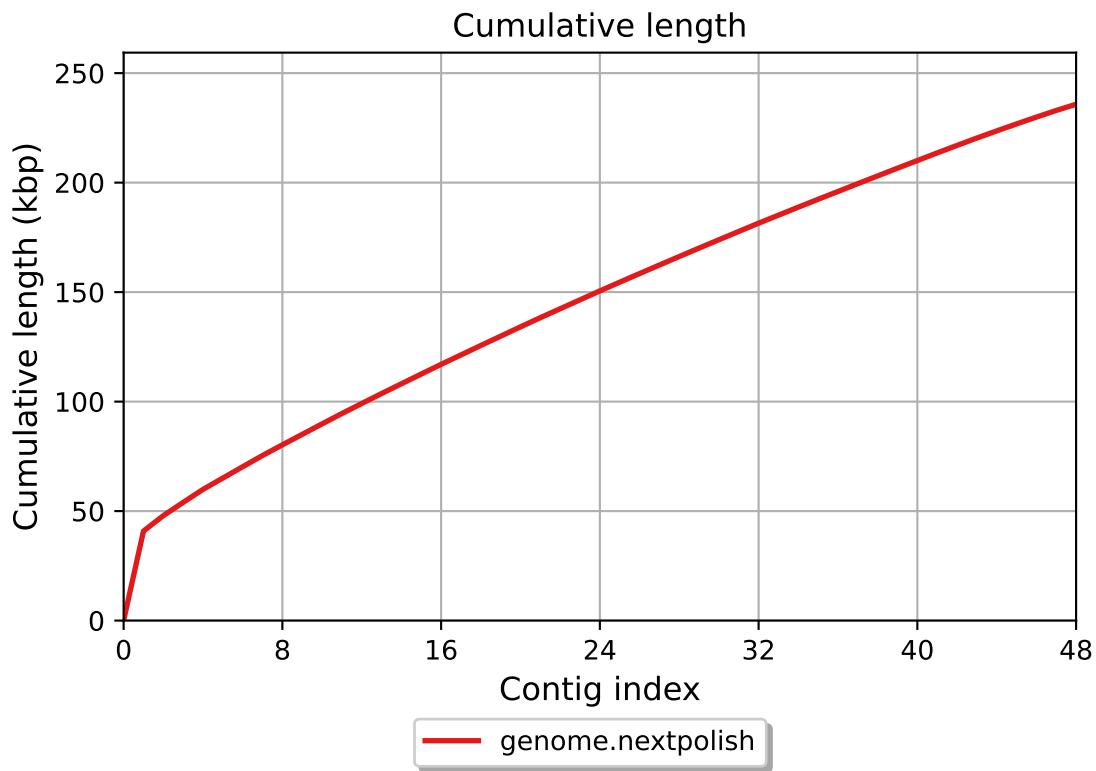
	genome.nextpolish
# contigs (>= 0 bp)	48
# contigs (>= 1000 bp)	48
# contigs (>= 5000 bp)	7
# contigs (>= 10000 bp)	1
# contigs (>= 25000 bp)	1
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	235782
Total length (>= 1000 bp)	235782
Total length (>= 5000 bp)	75392
Total length (>= 10000 bp)	40863
Total length (>= 25000 bp)	40863
Total length (>= 50000 bp)	0
# contigs	48
Largest contig	40863
Total length	235782
GC (%)	44.71
N50	4308
N90	3485
auN	10643.2
L50	17
L90	41
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	3 + 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).

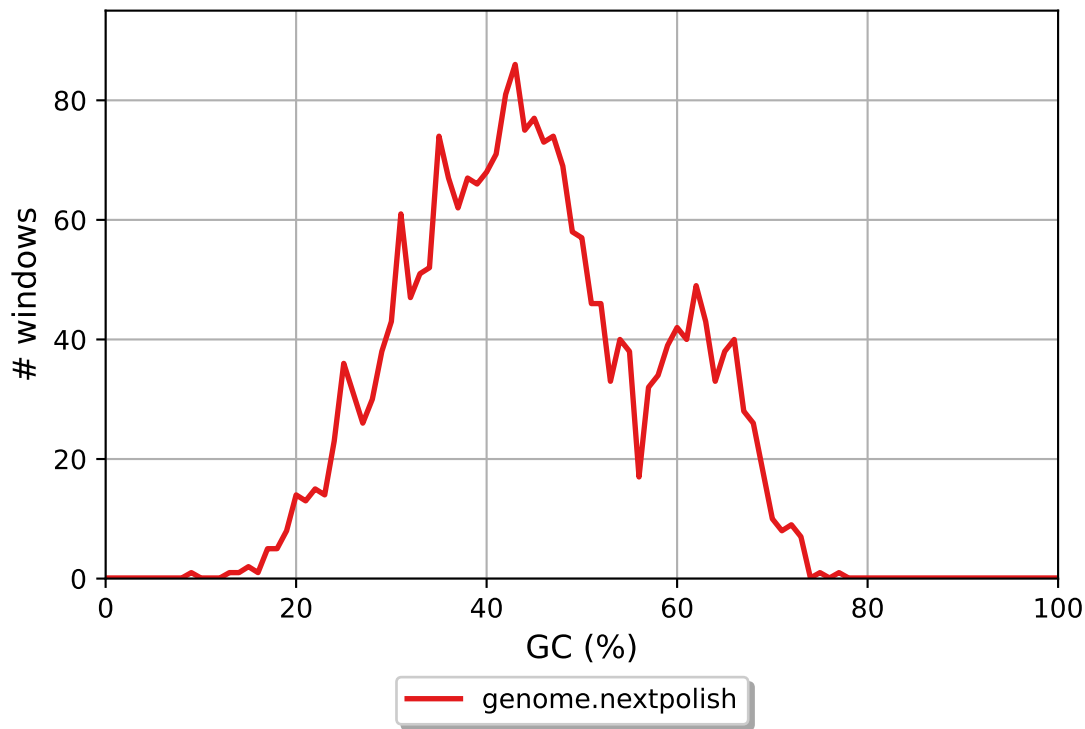
Nx



— genome.nextpolish



## GC content



genome.nextpolish GC content

