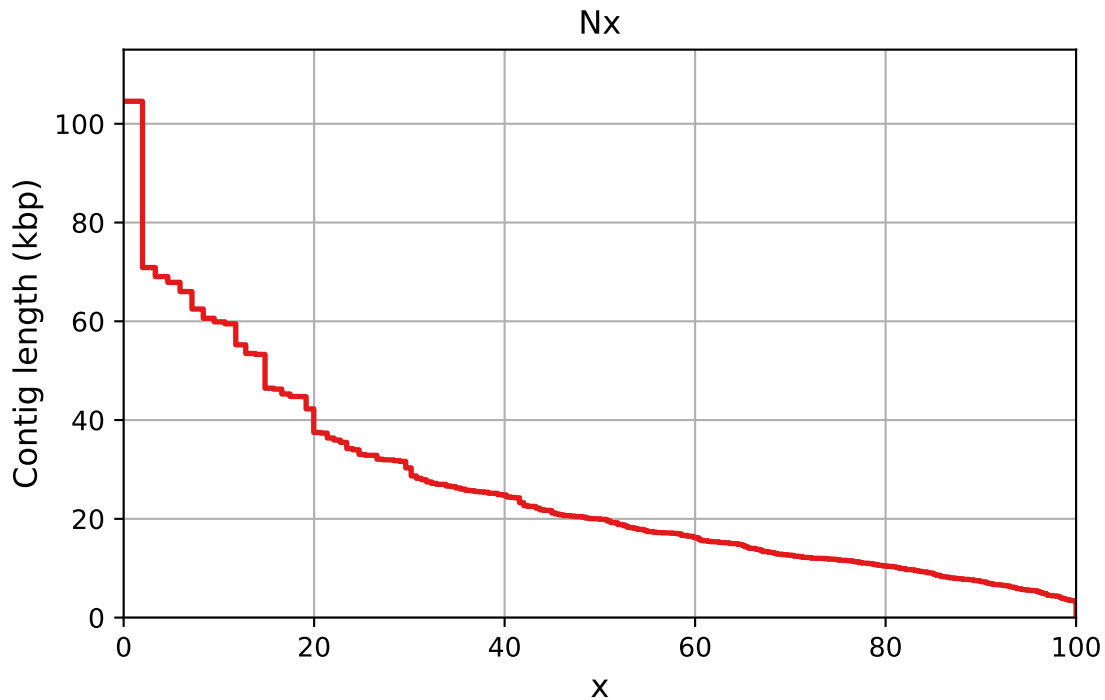


# Report

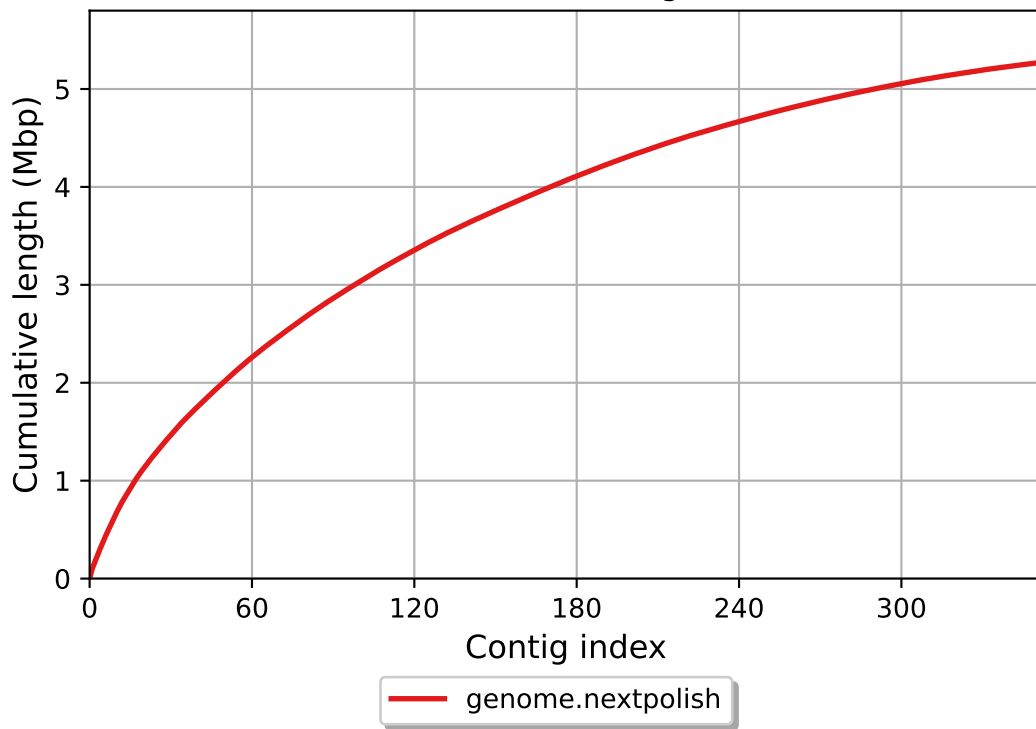
	genome.nextpolish
# contigs (>= 0 bp)	352
# contigs (>= 1000 bp)	352
# contigs (>= 5000 bp)	306
# contigs (>= 10000 bp)	198
# contigs (>= 25000 bp)	52
# contigs (>= 50000 bp)	12
Total length (>= 0 bp)	5273009
Total length (>= 1000 bp)	5273009
Total length (>= 5000 bp)	5086699
Total length (>= 10000 bp)	4300317
Total length (>= 25000 bp)	2069023
Total length (>= 50000 bp)	782898
# contigs	352
Largest contig	104557
Total length	5273009
GC (%)	39.45
N50	19915
N90	7282
auN	26421.9
L50	79
L90	251
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	9 + 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).

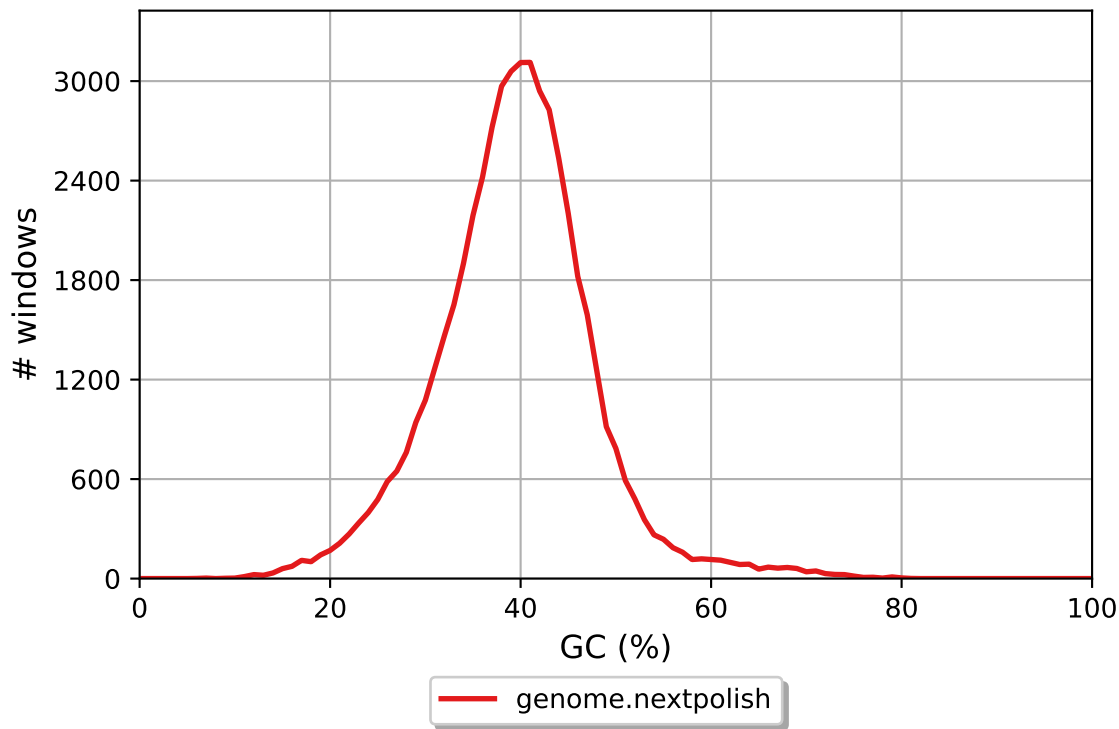


— genome.nextpolish

Cumulative length



## GC content



genome.nextpolish GC content

