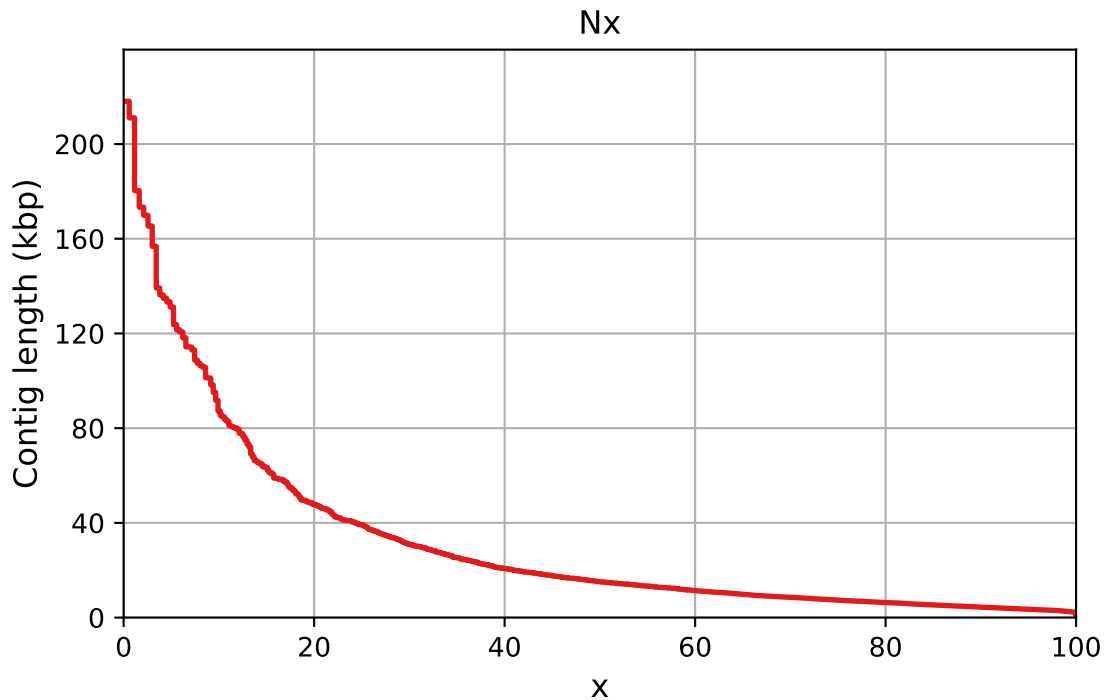


# Report

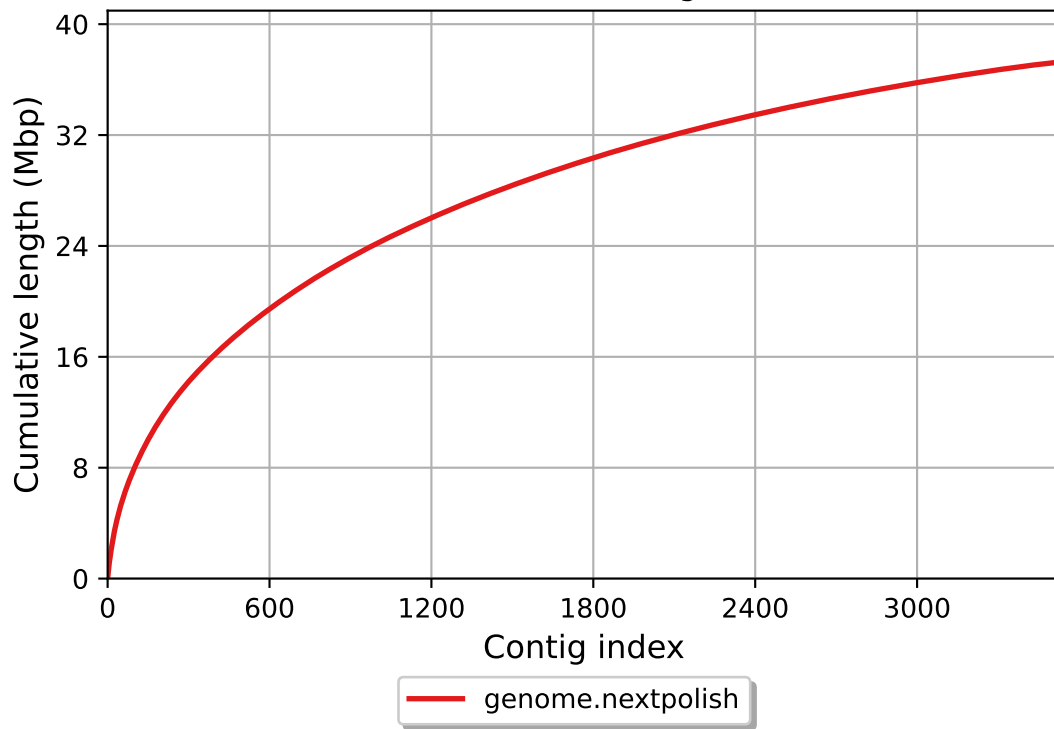
	genome.nextpolish
# contigs (>= 0 bp)	3530
# contigs (>= 1000 bp)	3520
# contigs (>= 5000 bp)	2162
# contigs (>= 10000 bp)	986
# contigs (>= 25000 bp)	255
# contigs (>= 50000 bp)	77
Total length (>= 0 bp)	37251381
Total length (>= 1000 bp)	37244593
Total length (>= 5000 bp)	32343378
Total length (>= 10000 bp)	24062483
Total length (>= 25000 bp)	13141394
Total length (>= 50000 bp)	6928436
# contigs	3530
Largest contig	218063
Total length	37251381
GC (%)	38.87
N50	15124
N90	4420
auN	32646.8
L50	544
L90	2414
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	7 + 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).

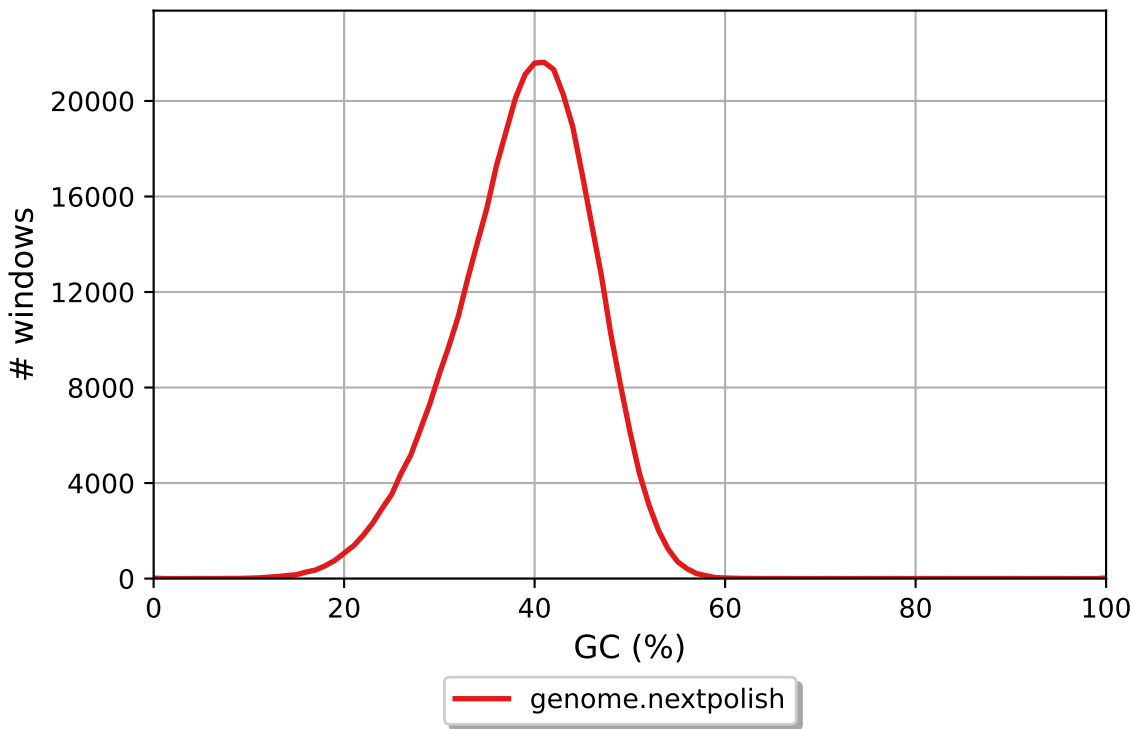


— genome.nextpolish

Cumulative length



# GC content



genome.nextpolish GC content

