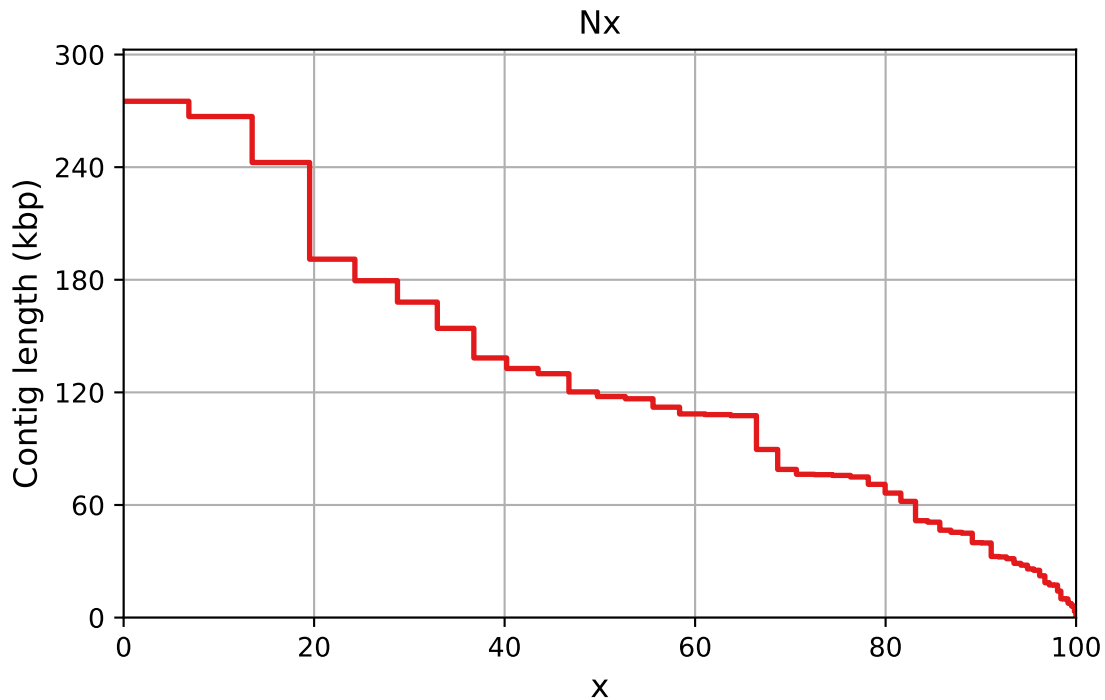


Report

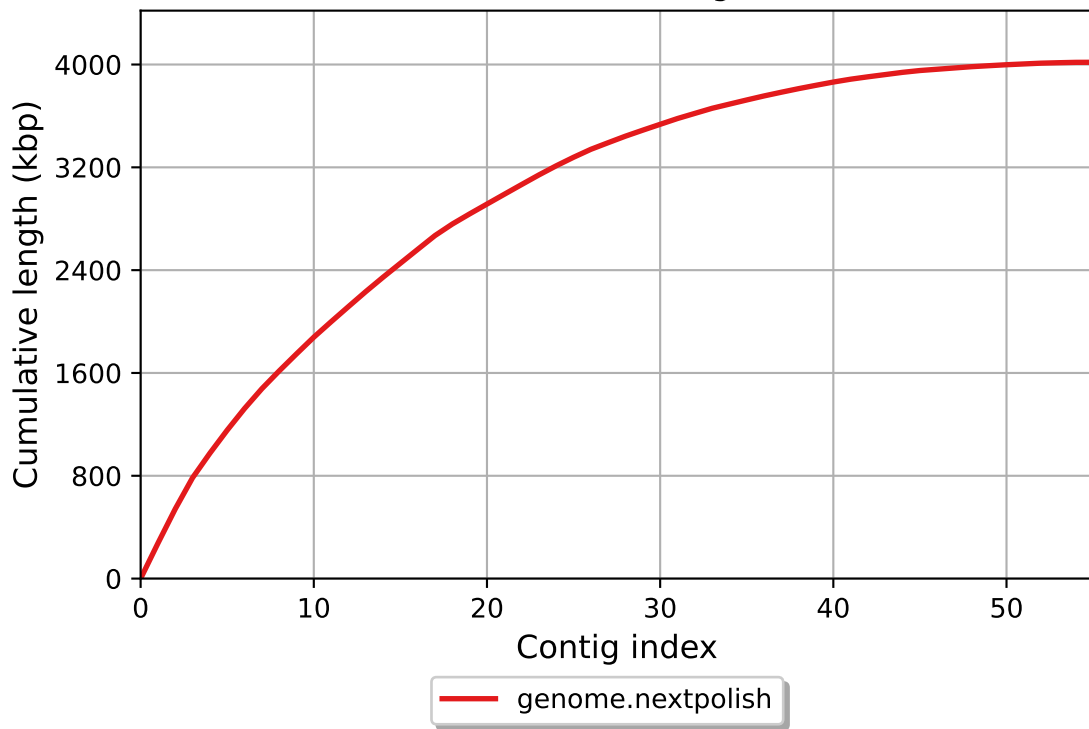
	genome.nextpolish
# contigs (>= 0 bp)	55
# contigs (>= 1000 bp)	54
# contigs (>= 5000 bp)	52
# contigs (>= 10000 bp)	46
# contigs (>= 25000 bp)	40
# contigs (>= 50000 bp)	28
Total length (>= 0 bp)	4017540
Total length (>= 1000 bp)	4016950
Total length (>= 5000 bp)	4010591
Total length (>= 10000 bp)	3963337
Total length (>= 25000 bp)	3863615
Total length (>= 50000 bp)	3442736
# contigs	55
Largest contig	275113
Total length	4017540
GC (%)	39.09
N50	117784
N90	39928
auN	135160.3
L50	12
L90	32
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	17 + 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).

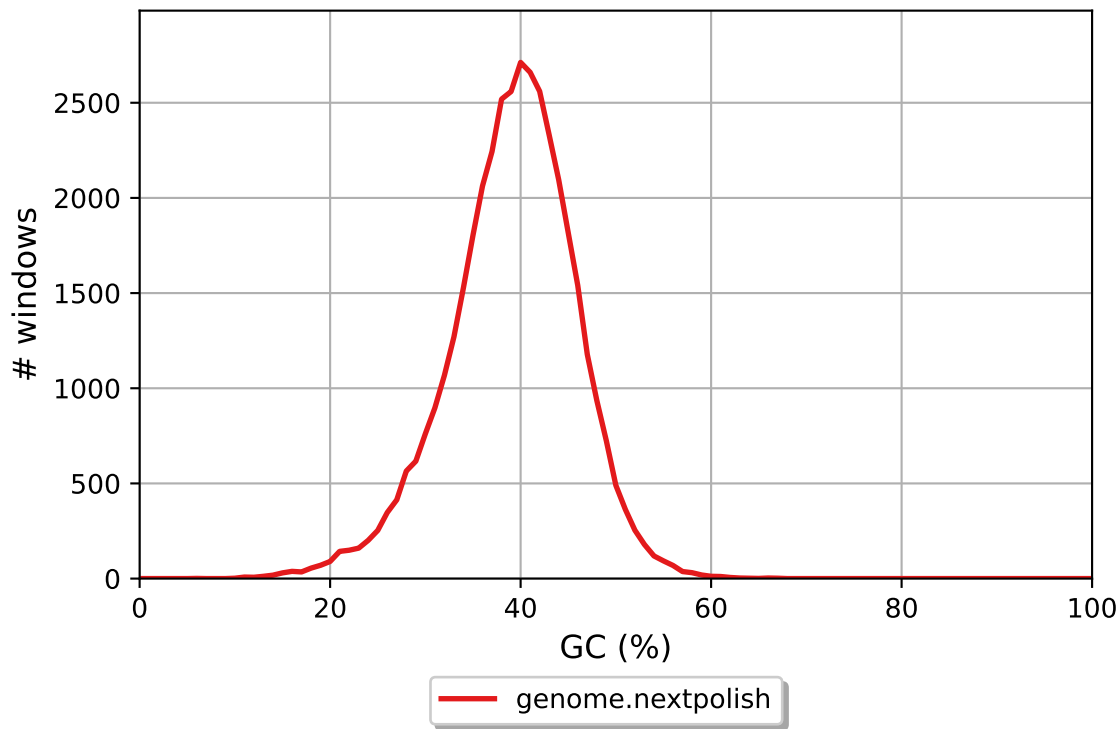


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

