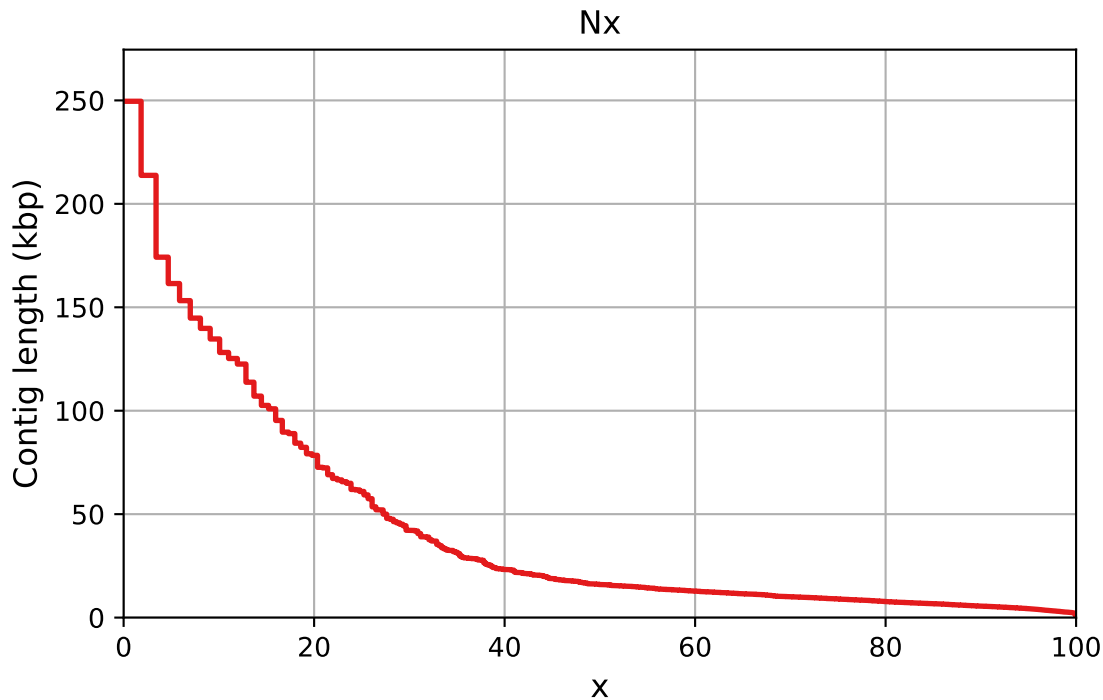


Report

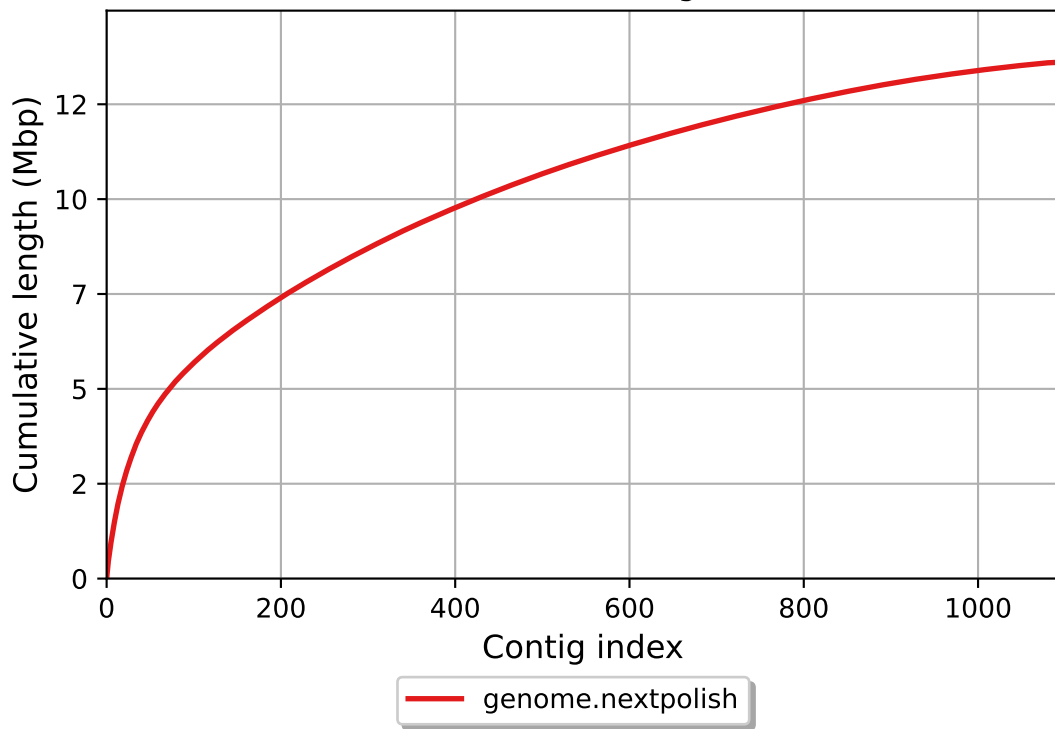
	genome.nextpolish
# contigs (>= 0 bp)	1093
# contigs (>= 1000 bp)	1087
# contigs (>= 5000 bp)	798
# contigs (>= 10000 bp)	382
# contigs (>= 25000 bp)	81
# contigs (>= 50000 bp)	38
Total length (>= 0 bp)	13606780
Total length (>= 1000 bp)	13602786
Total length (>= 5000 bp)	12589795
Total length (>= 10000 bp)	9596238
Total length (>= 25000 bp)	5246712
Total length (>= 50000 bp)	3757392
# contigs	1093
Largest contig	249583
Total length	13606780
GC (%)	41.31
N50	15990
N90	5536
auN	43974.7
L50	161
L90	733
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	38 + 6 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).

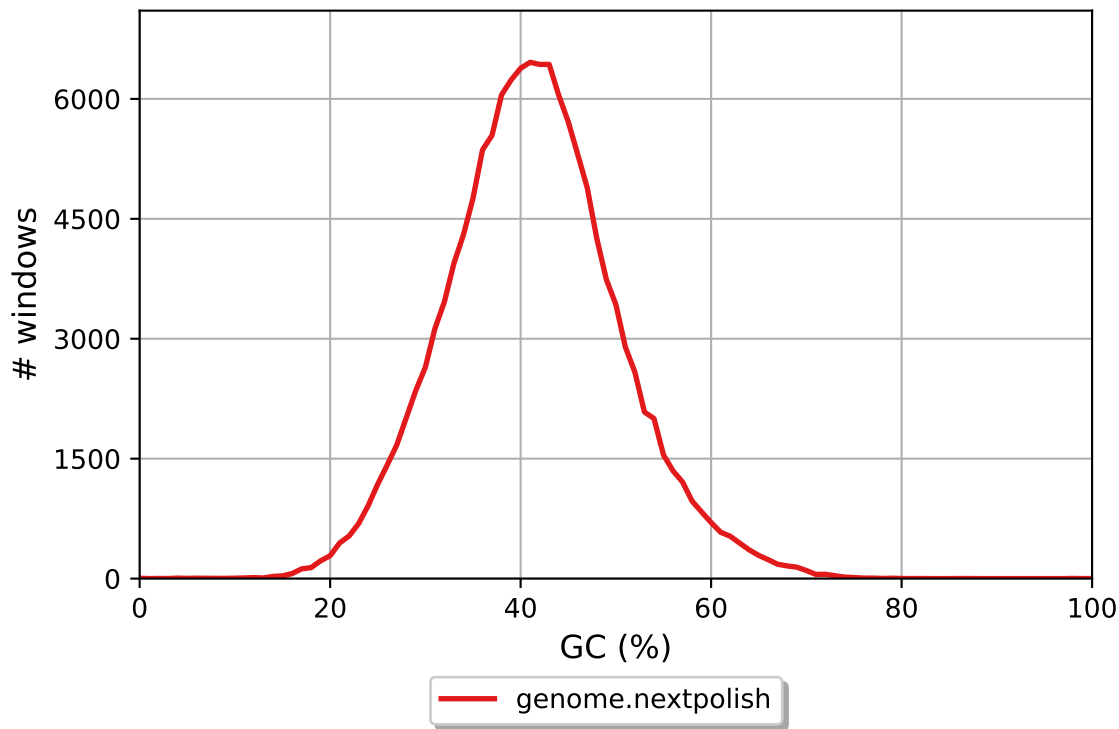


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

