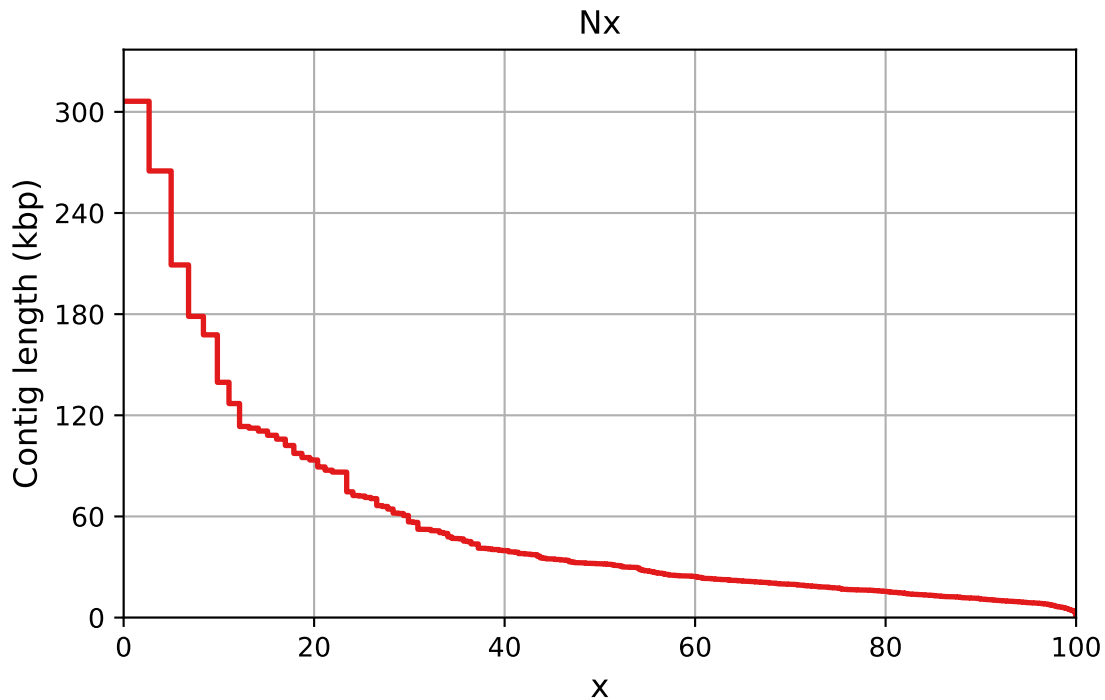


Report

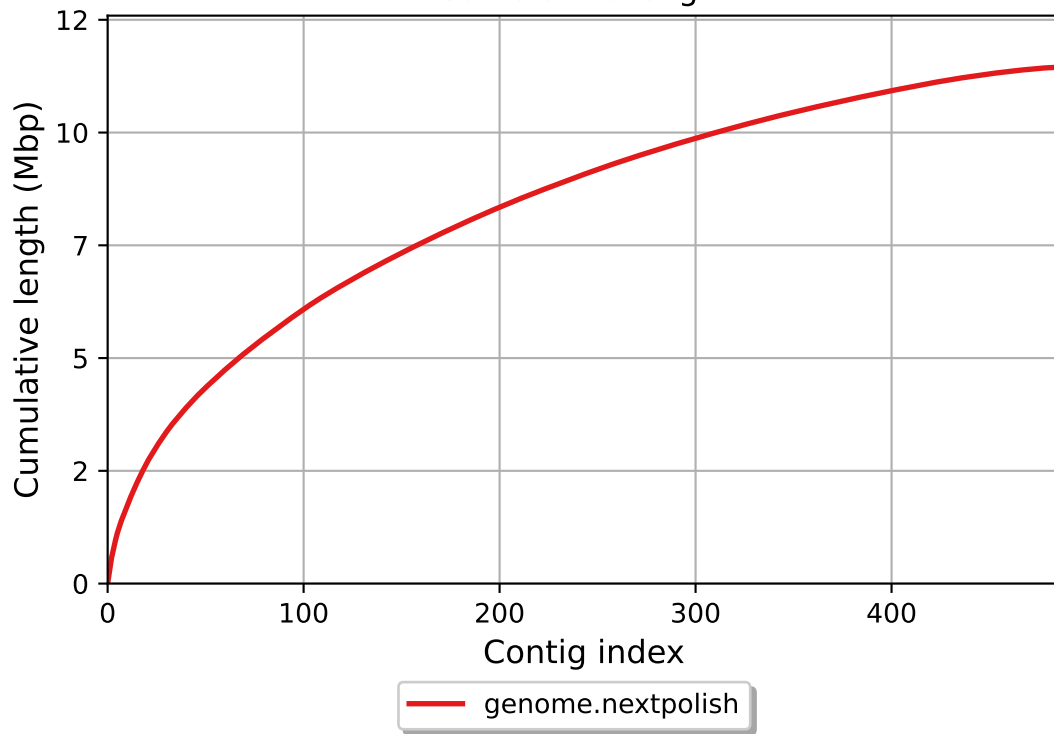
	genome.nextpolish
# contigs (>= 0 bp)	486
# contigs (>= 1000 bp)	483
# contigs (>= 5000 bp)	460
# contigs (>= 10000 bp)	361
# contigs (>= 25000 bp)	120
# contigs (>= 50000 bp)	39
Total length (>= 0 bp)	11447541
Total length (>= 1000 bp)	11445221
Total length (>= 5000 bp)	11356327
Total length (>= 10000 bp)	10563563
Total length (>= 25000 bp)	6625962
Total length (>= 50000 bp)	3846675
# contigs	486
Largest contig	306285
Total length	11447541
GC (%)	46.64
N50	31893
N90	10970
auN	59313.9
L50	89
L90	337
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	43 + 5 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).

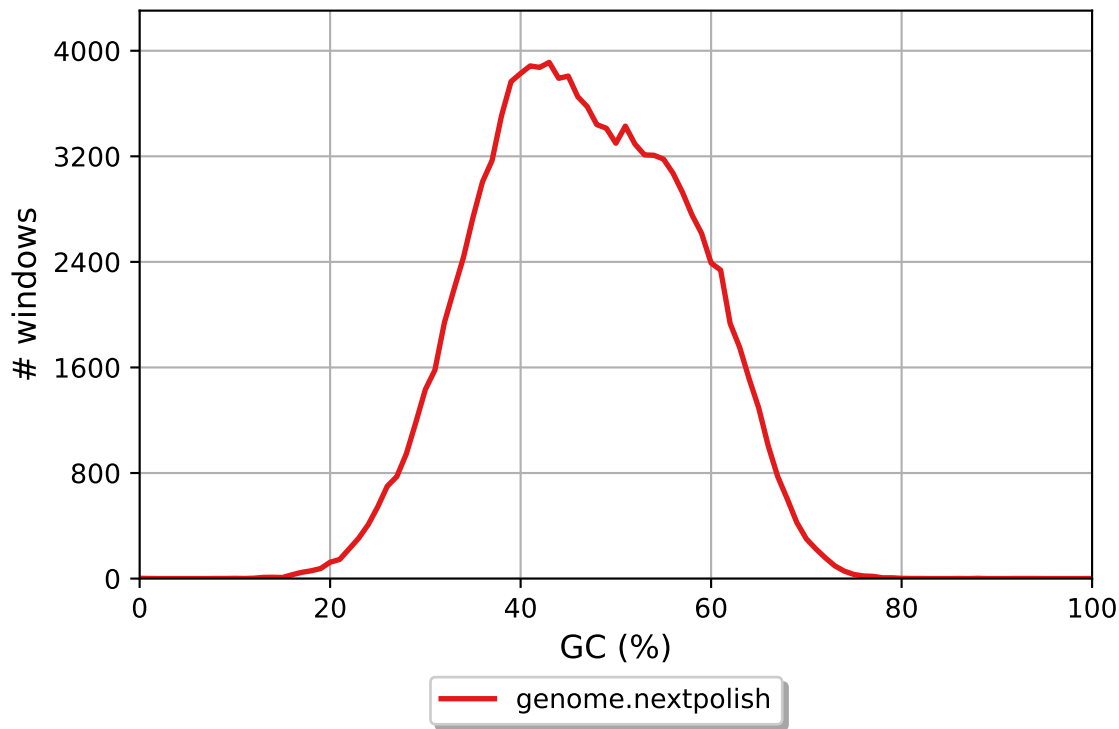


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

