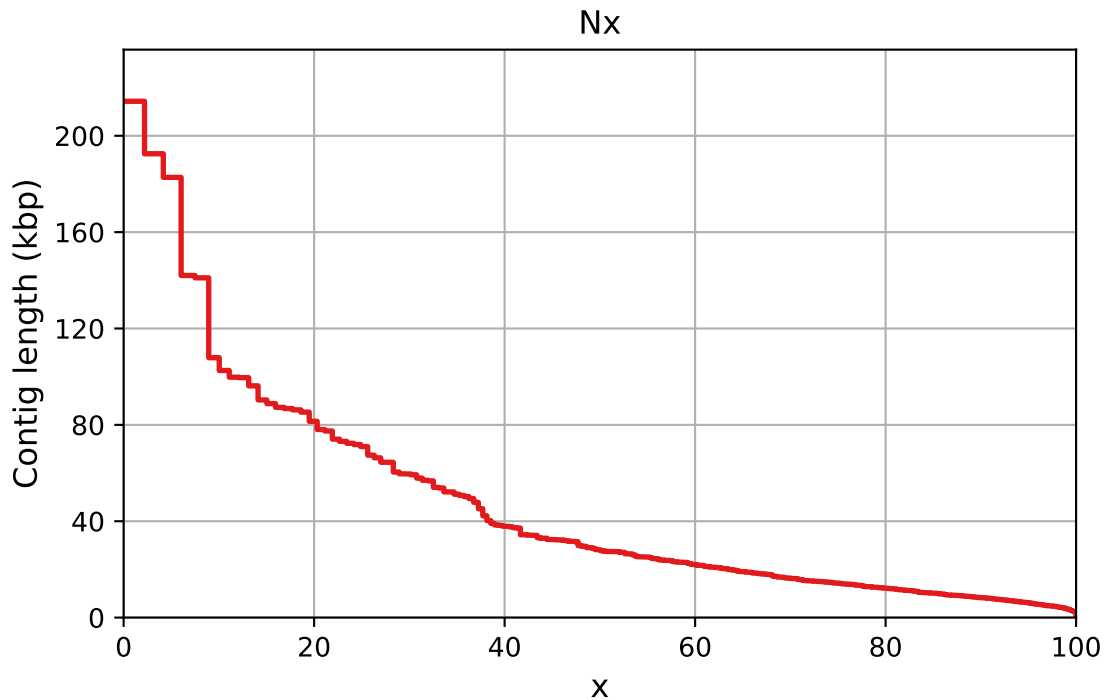


Report

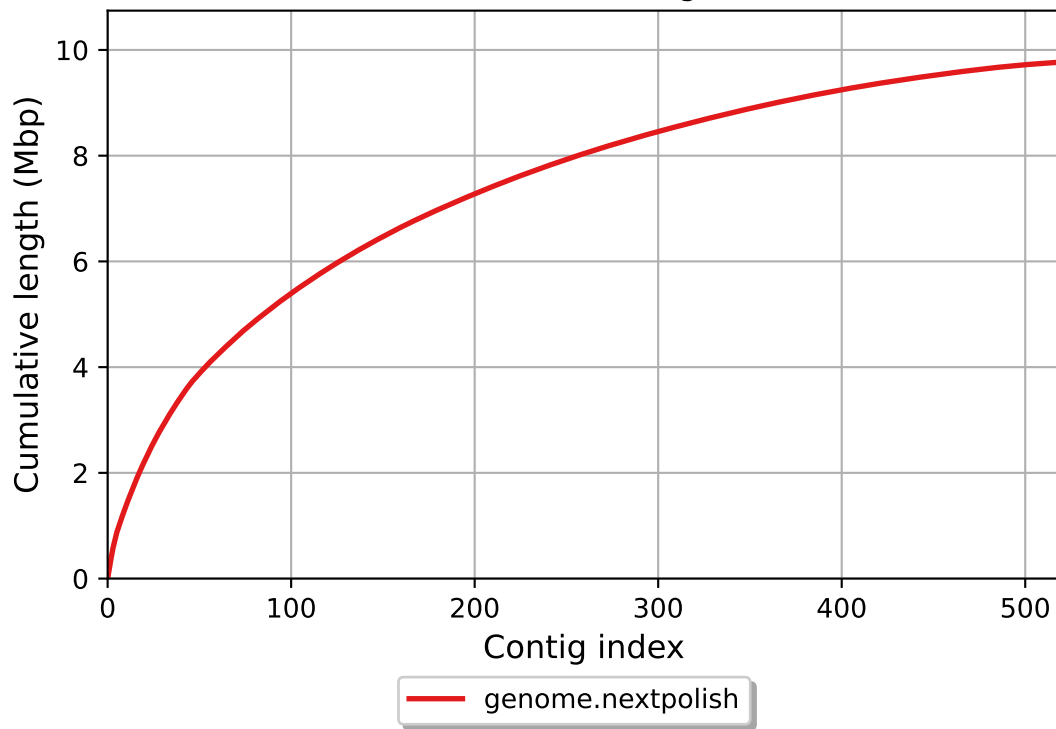
	genome.nextpolish
# contigs (>= 0 bp)	519
# contigs (>= 1000 bp)	519
# contigs (>= 5000 bp)	439
# contigs (>= 10000 bp)	288
# contigs (>= 25000 bp)	100
# contigs (>= 50000 bp)	42
Total length (>= 0 bp)	9767061
Total length (>= 1000 bp)	9767061
Total length (>= 5000 bp)	9466434
Total length (>= 10000 bp)	8340474
Total length (>= 25000 bp)	5395706
Total length (>= 50000 bp)	3540487
# contigs	519
Largest contig	214325
Total length	9767061
GC (%)	44.41
N50	28128
N90	8294
auN	49090.1
L50	81
L90	338
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	14 + 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).

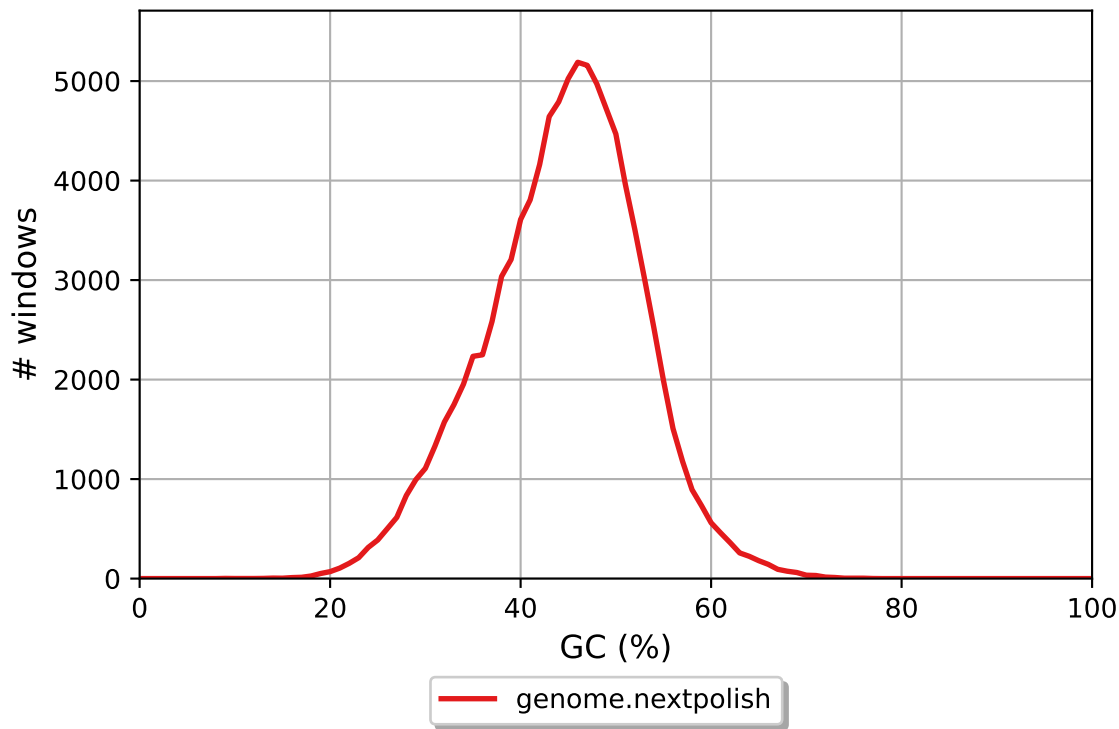


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

