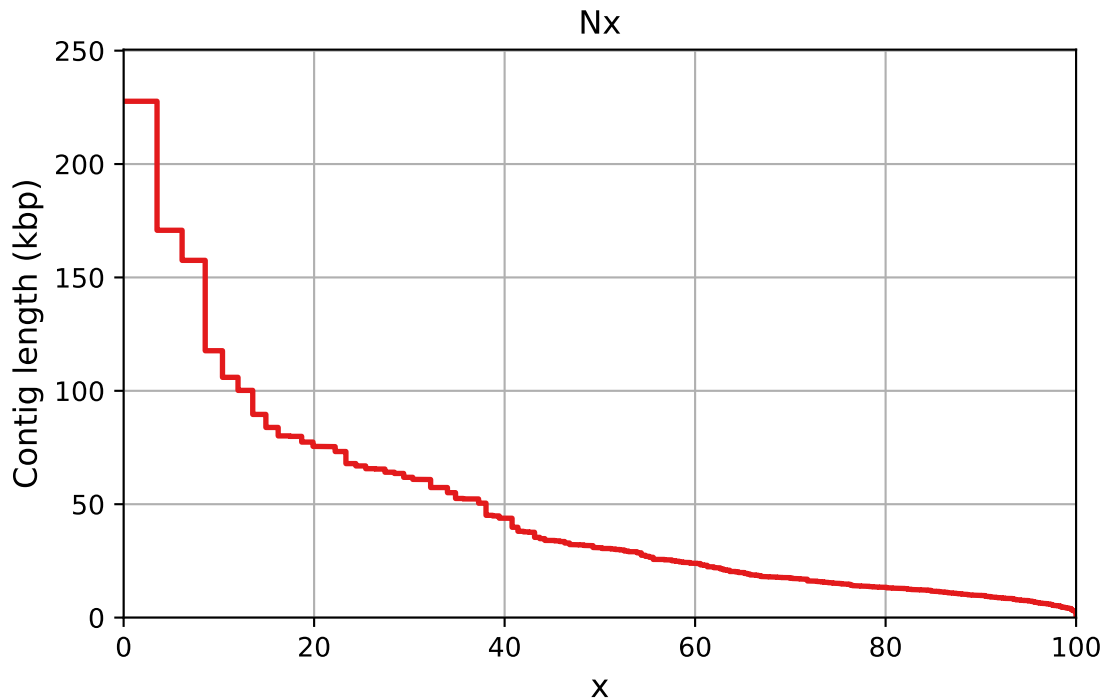


Report

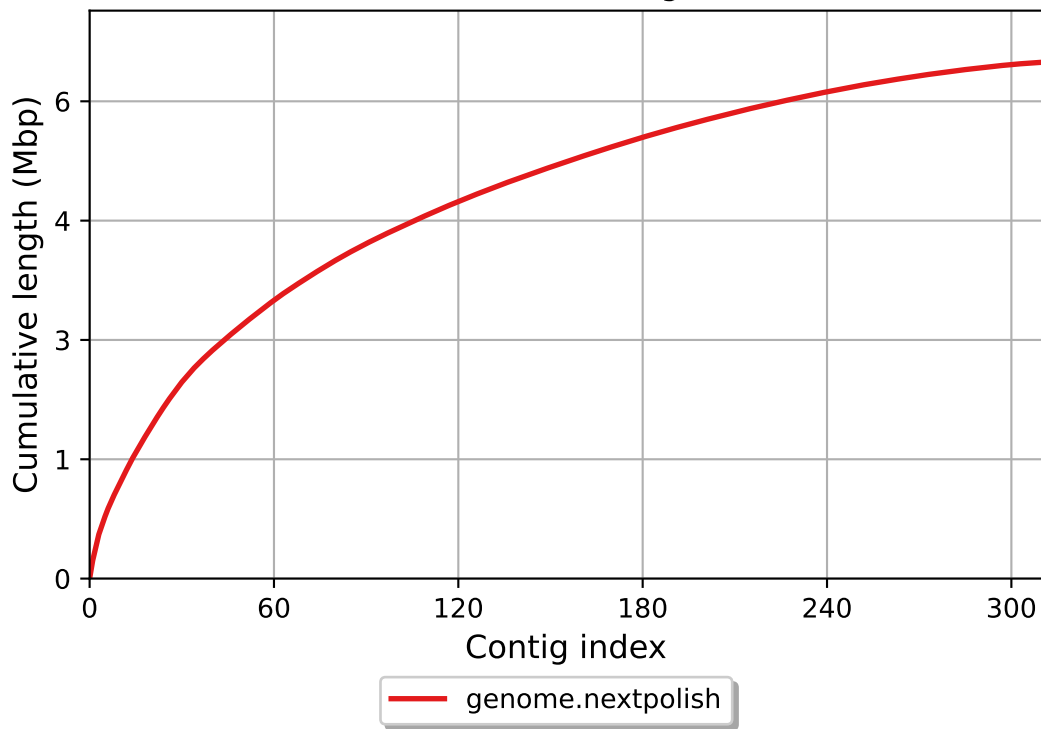
	genome.nextpolish
# contigs (>= 0 bp)	310
# contigs (>= 1000 bp)	310
# contigs (>= 5000 bp)	282
# contigs (>= 10000 bp)	201
# contigs (>= 25000 bp)	70
# contigs (>= 50000 bp)	30
Total length (>= 0 bp)	6490723
Total length (>= 1000 bp)	6490723
Total length (>= 5000 bp)	6384640
Total length (>= 10000 bp)	5772782
Total length (>= 25000 bp)	3761455
Total length (>= 50000 bp)	2469525
# contigs	310
Largest contig	227684
Total length	6490723
GC (%)	41.79
N50	30792
N90	9769
auN	50846.8
L50	52
L90	208
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	23 + 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).

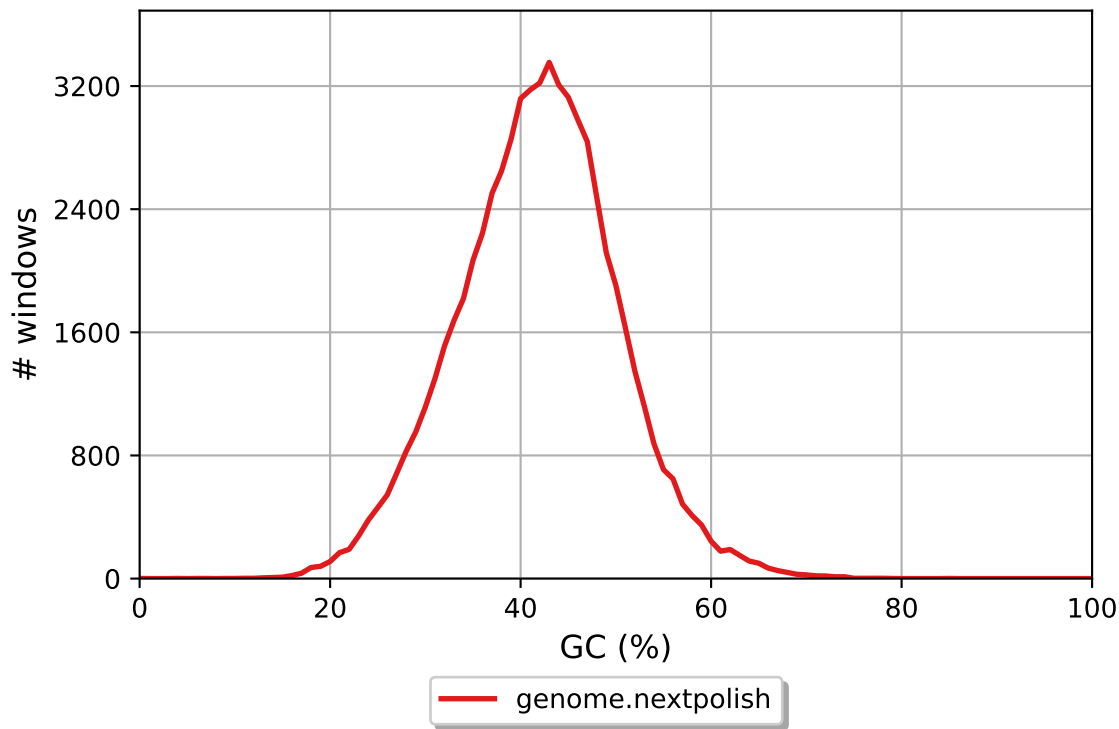


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

