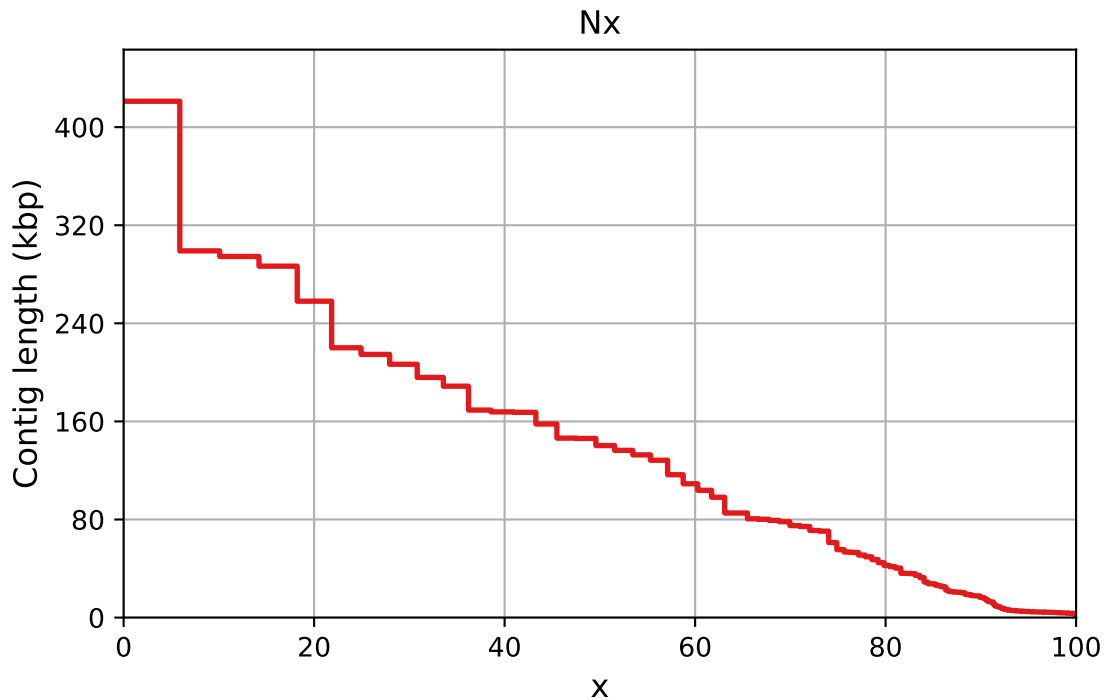


Report

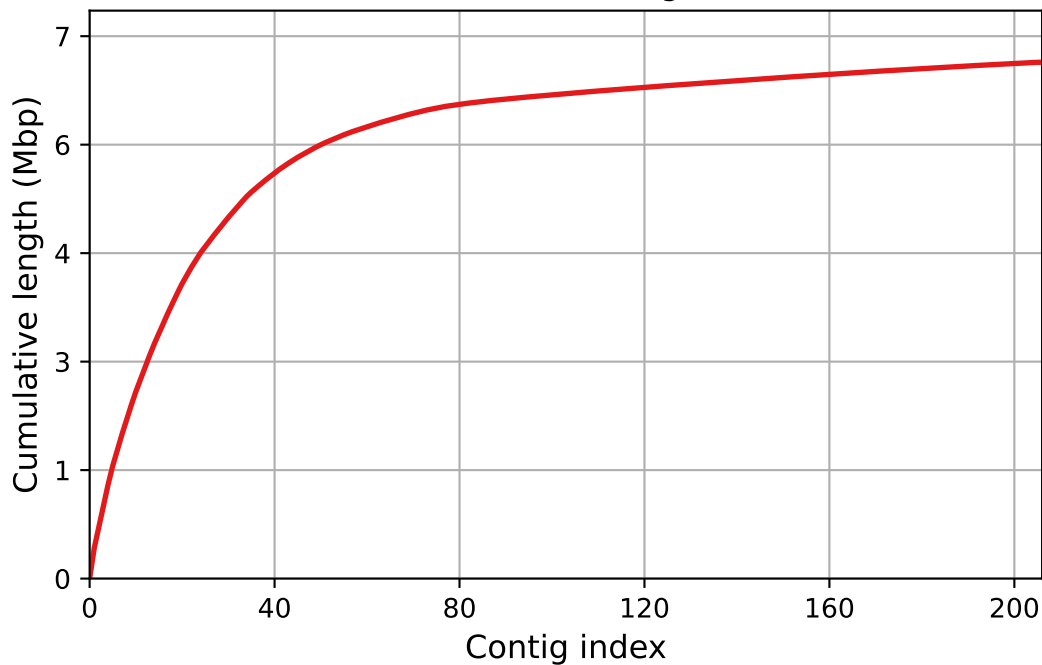
	genome.nextpolish
# contigs (>= 0 bp)	206
# contigs (>= 1000 bp)	205
# contigs (>= 5000 bp)	115
# contigs (>= 10000 bp)	77
# contigs (>= 25000 bp)	56
# contigs (>= 50000 bp)	39
Total length (>= 0 bp)	7139379
Total length (>= 1000 bp)	7138737
Total length (>= 5000 bp)	6767996
Total length (>= 10000 bp)	6529789
Total length (>= 25000 bp)	6161603
Total length (>= 50000 bp)	5559271
# contigs	206
Largest contig	421107
Total length	7139379
GC (%)	50.85
N50	140291
N90	16512
auN	153020.9
L50	17
L90	70
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	18 + 3 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).

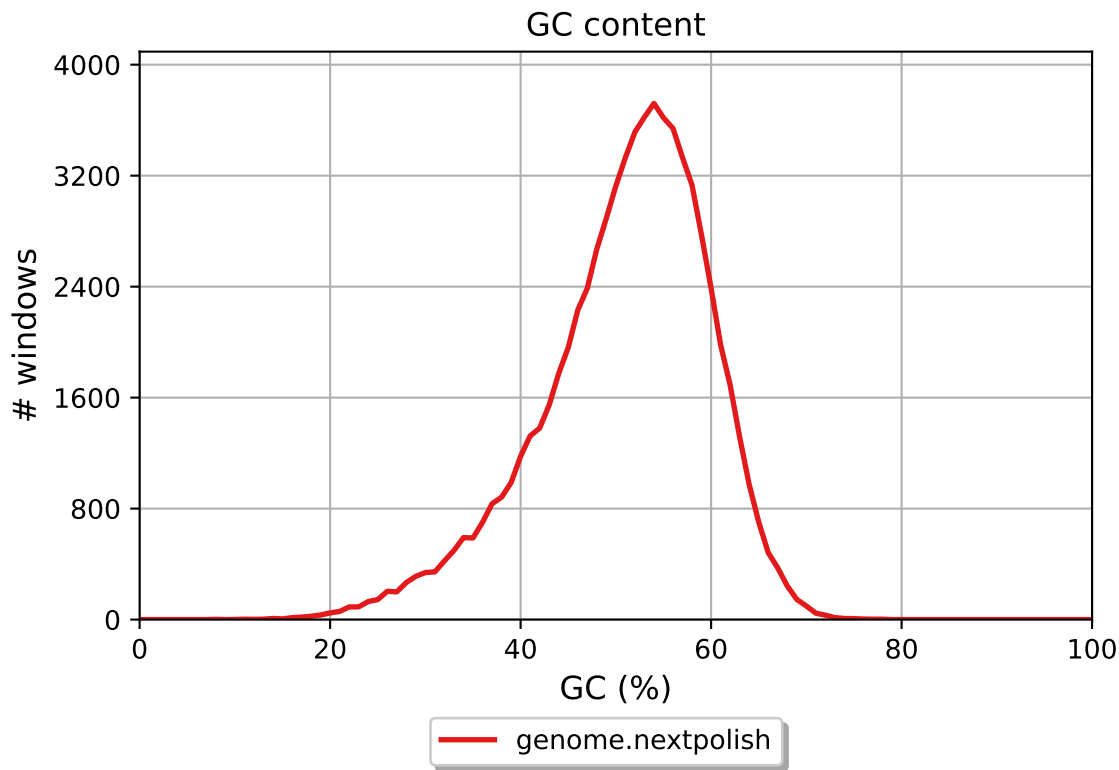


— genome.nextpolish

Cumulative length



— genome.nextpolish



genome.nextpolish GC content

