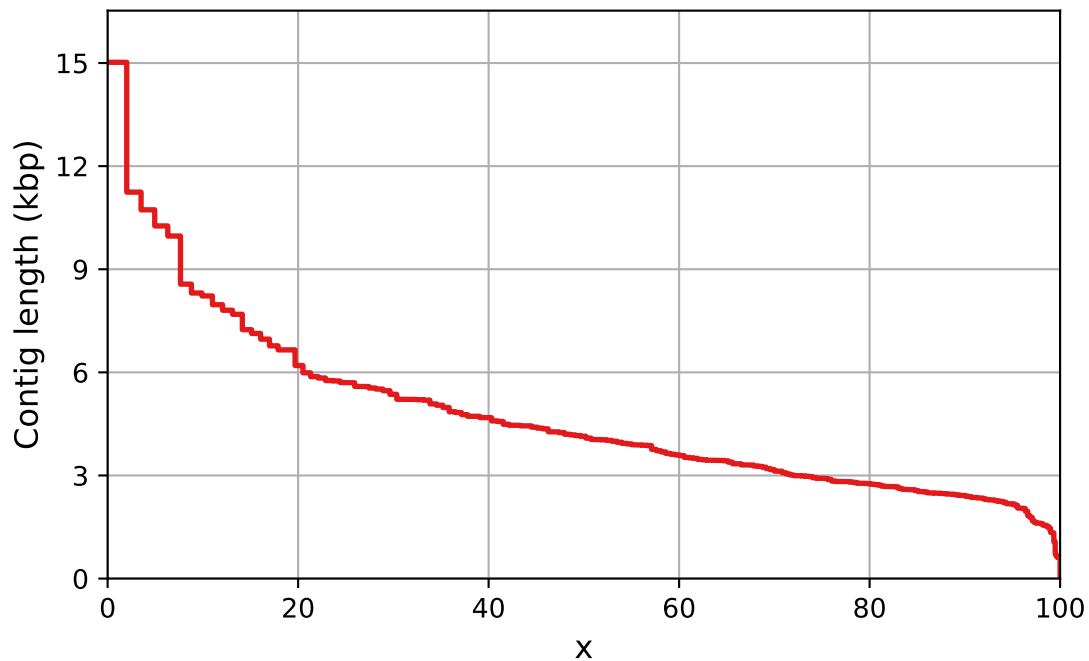


Report

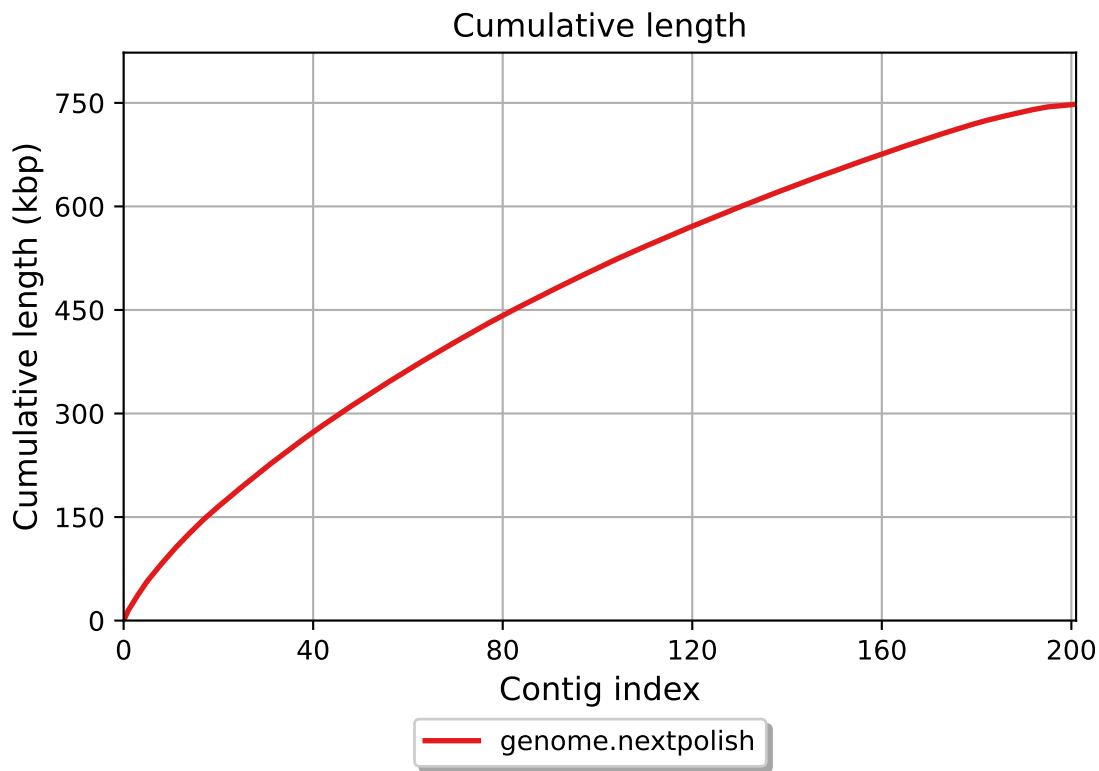
	genome.nextpolish
# contigs (>= 0 bp)	203
# contigs (>= 1000 bp)	195
# contigs (>= 5000 bp)	38
# contigs (>= 10000 bp)	4
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	748564
Total length (>= 1000 bp)	744157
Total length (>= 5000 bp)	263239
Total length (>= 10000 bp)	47253
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	201
Largest contig	15021
Total length	747986
GC (%)	37.87
N50	4141
N90	2417
auN	4806.5
L50	63
L90	159
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	0 + 0 part

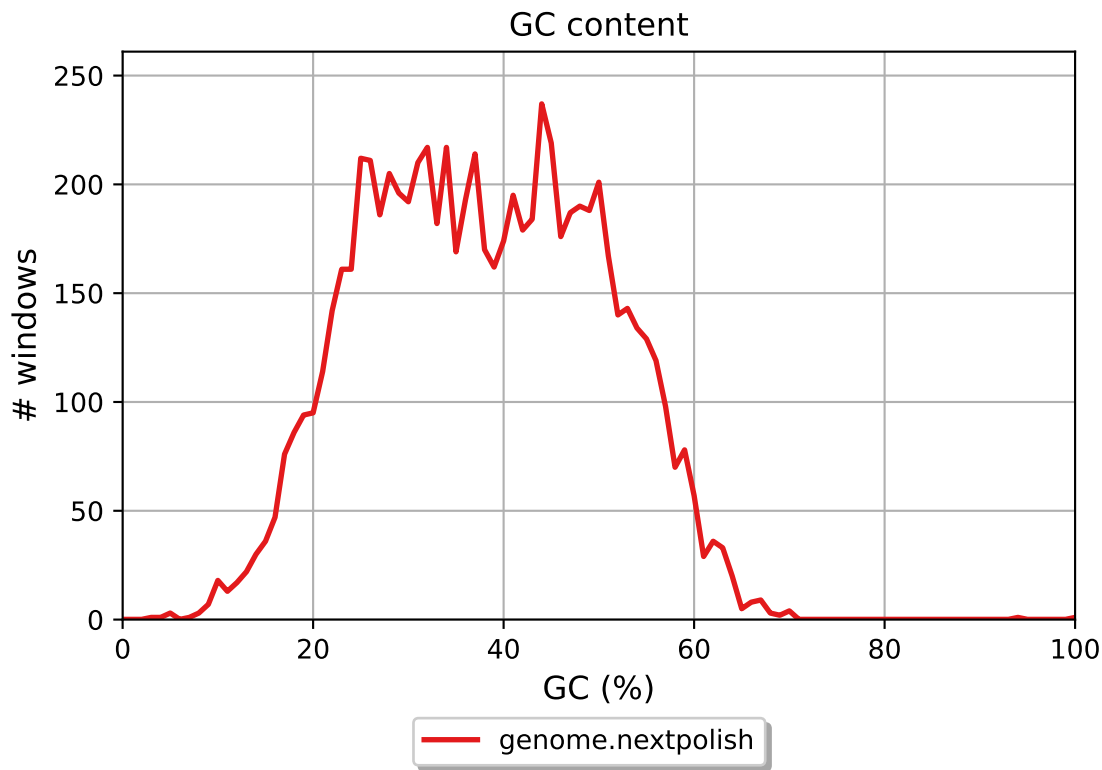
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx



— genome.nextpolish





genome.nextpolish GC content

