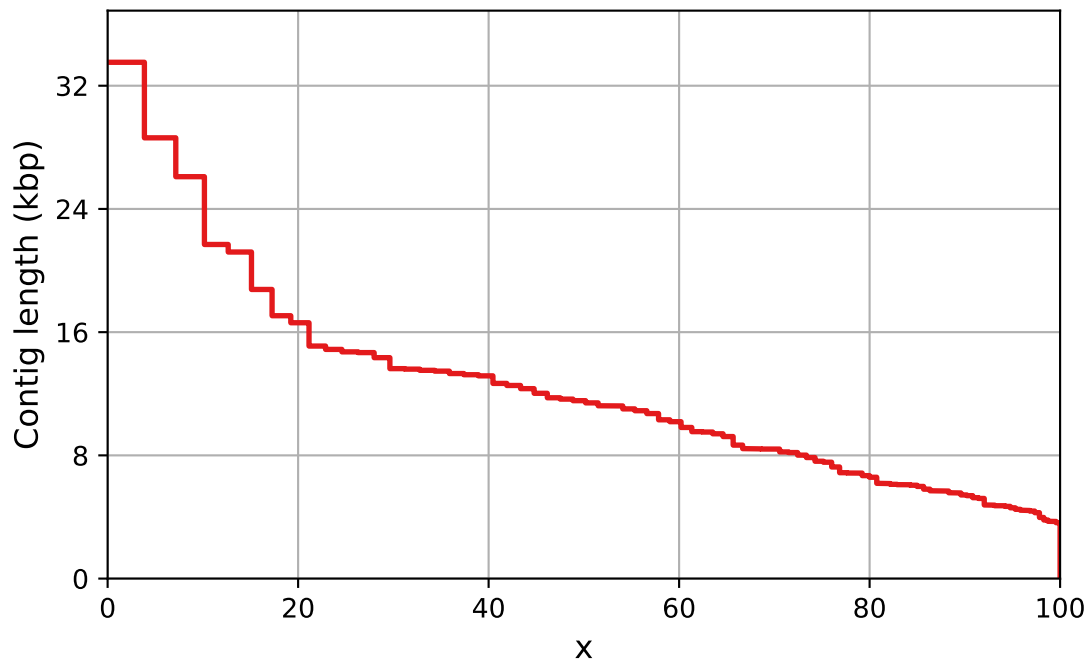


Report

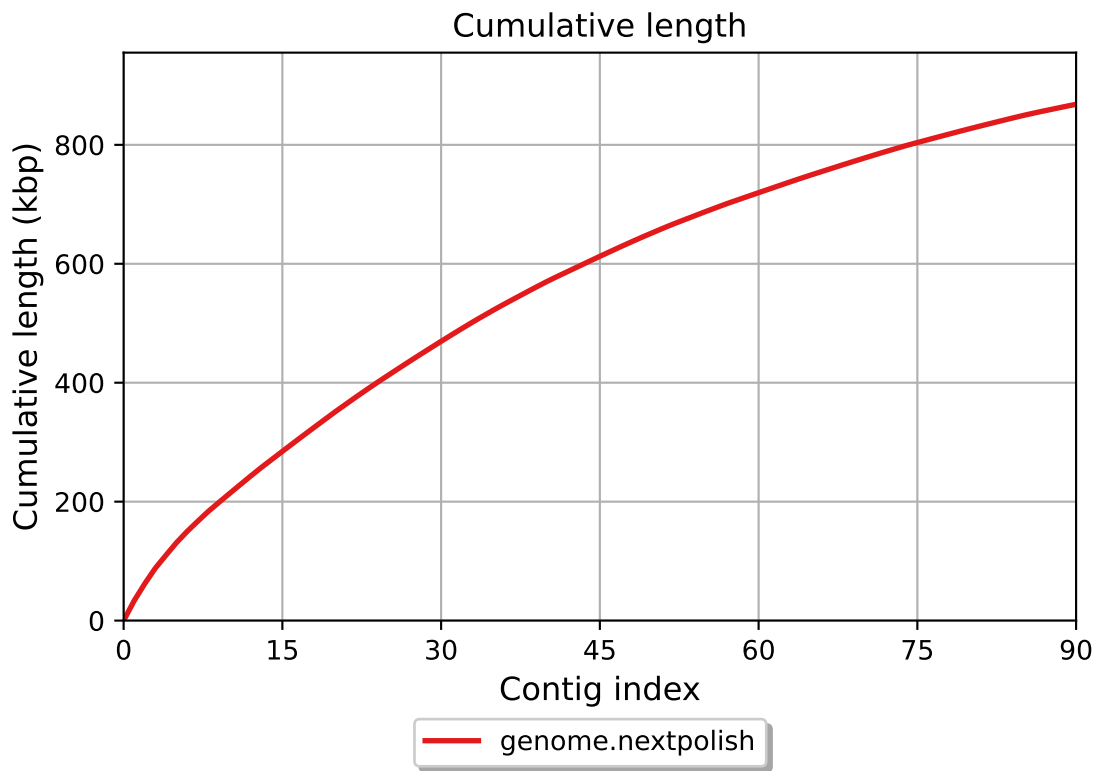
	genome.nextpolish
# contigs (>= 0 bp)	90
# contigs (>= 1000 bp)	90
# contigs (>= 5000 bp)	74
# contigs (>= 10000 bp)	35
# contigs (>= 25000 bp)	3
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	868225
Total length (>= 1000 bp)	868225
Total length (>= 5000 bp)	799084
Total length (>= 10000 bp)	522707
Total length (>= 25000 bp)	88233
Total length (>= 50000 bp)	0
# contigs	90
Largest contig	33522
Total length	868225
GC (%)	41.87
N50	11551
N90	5430
auN	12801.7
L50	27
L90	71
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	2 + 2 part

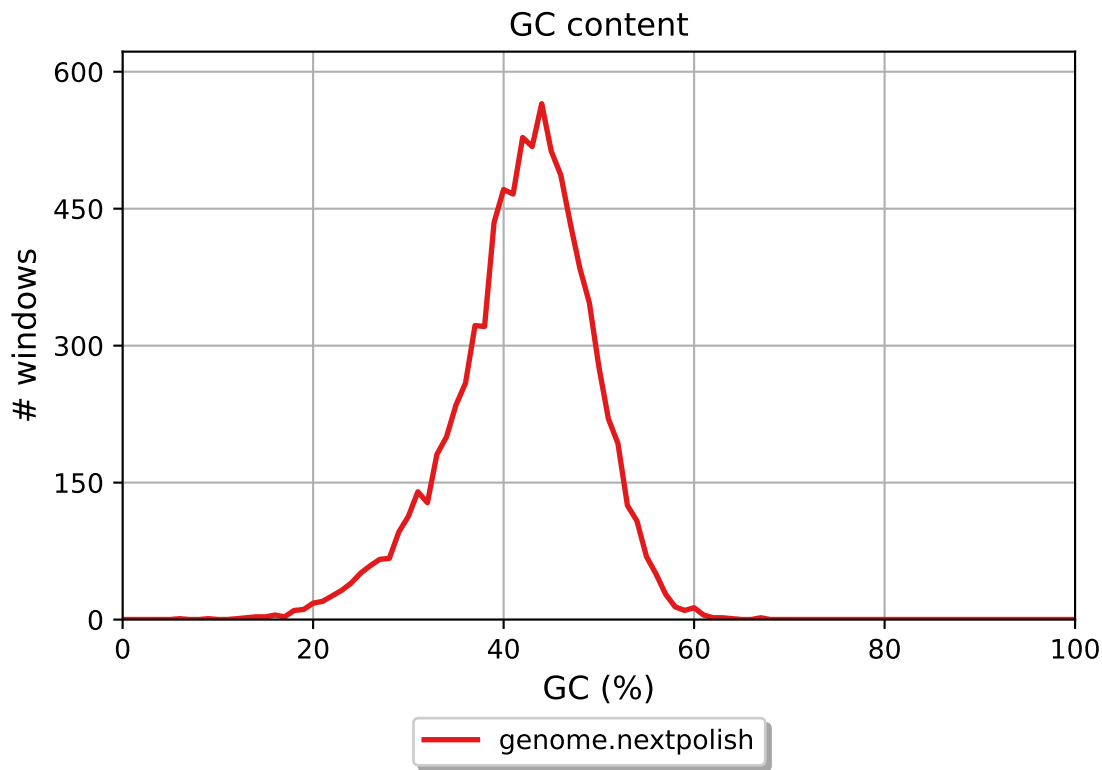
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx



— genome.nextpolish





genome.nextpolish GC content

