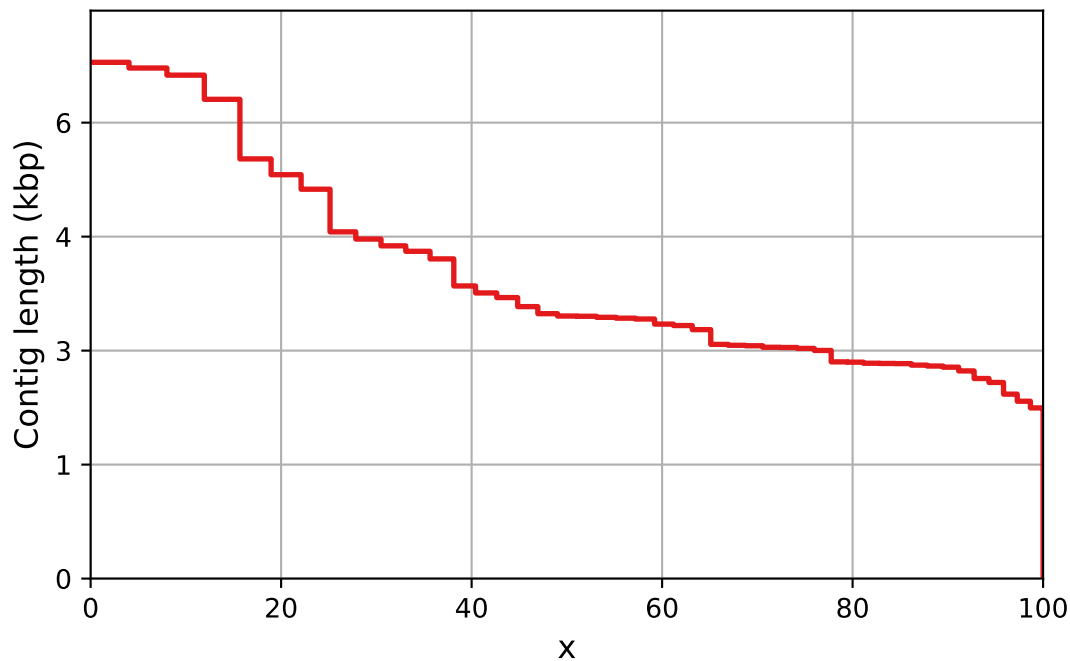


Report

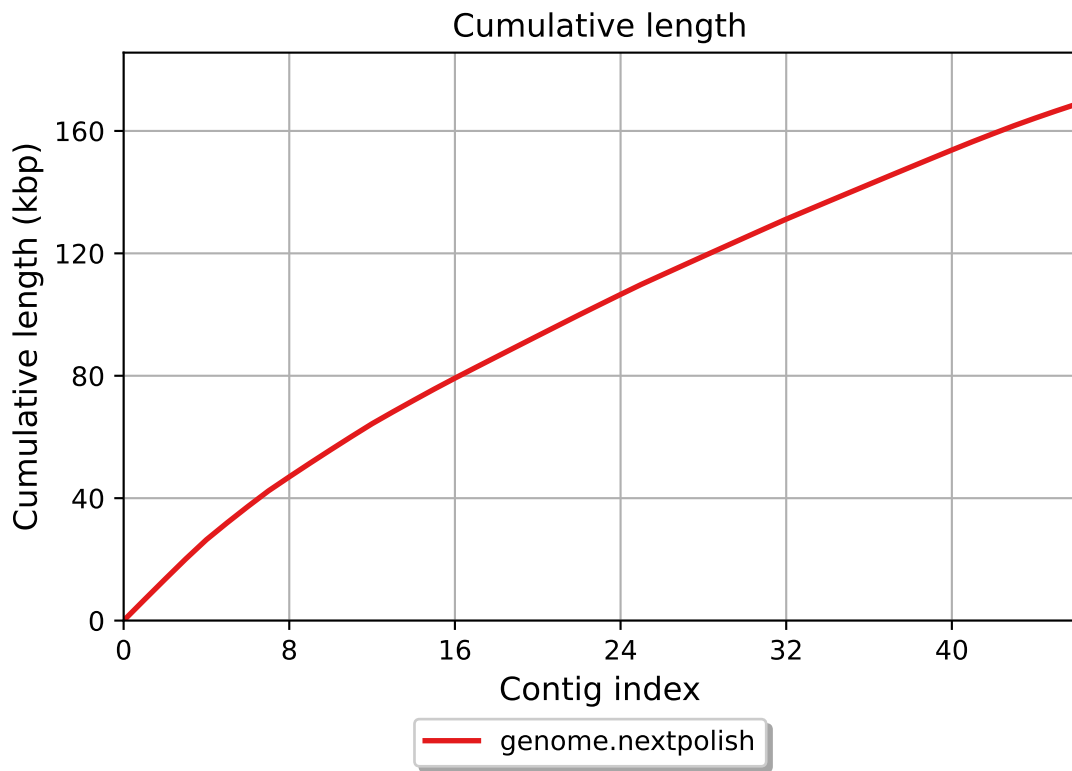
	genome.nextpolish
# contigs (>= 0 bp)	46
# contigs (>= 1000 bp)	46
# contigs (>= 5000 bp)	7
# contigs (>= 10000 bp)	0
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	168715
Total length (>= 1000 bp)	168715
Total length (>= 5000 bp)	42407
Total length (>= 10000 bp)	0
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	46
Largest contig	6794
Total length	168715
GC (%)	38.79
N50	3455
N90	2782
auN	4043.0
L50	18
L90	40
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	4 + 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).

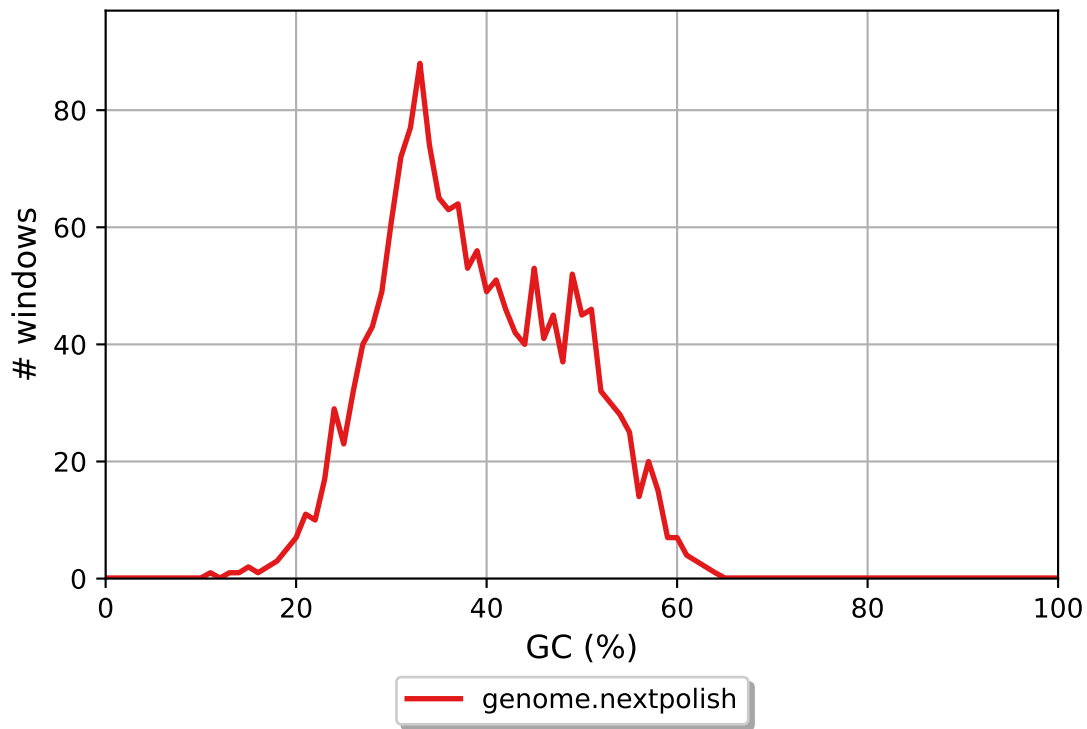
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

