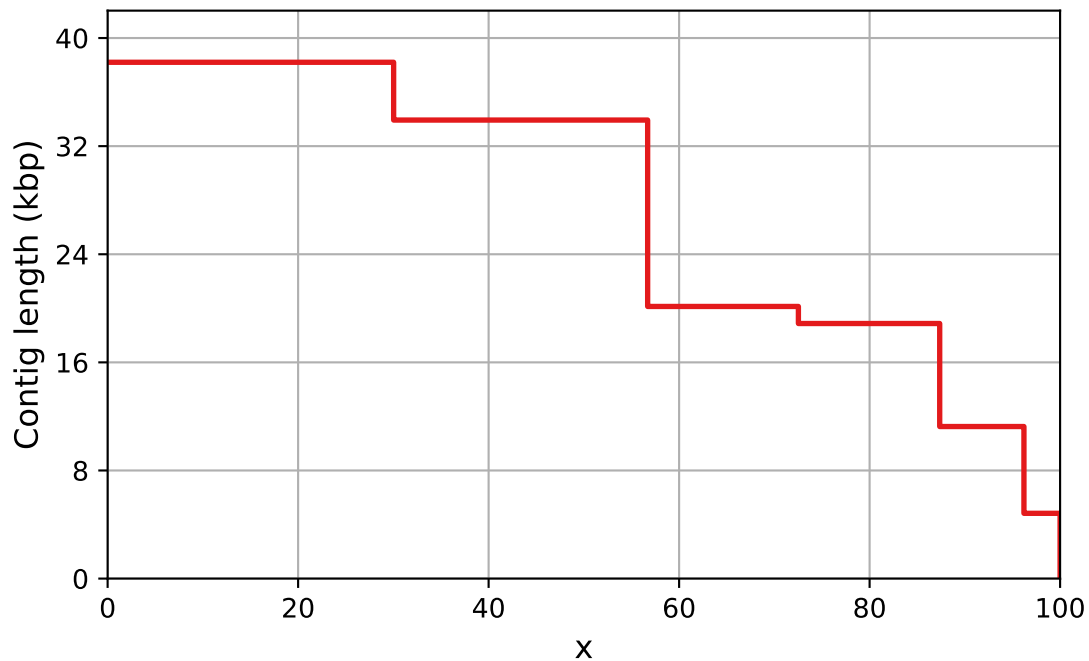


Report

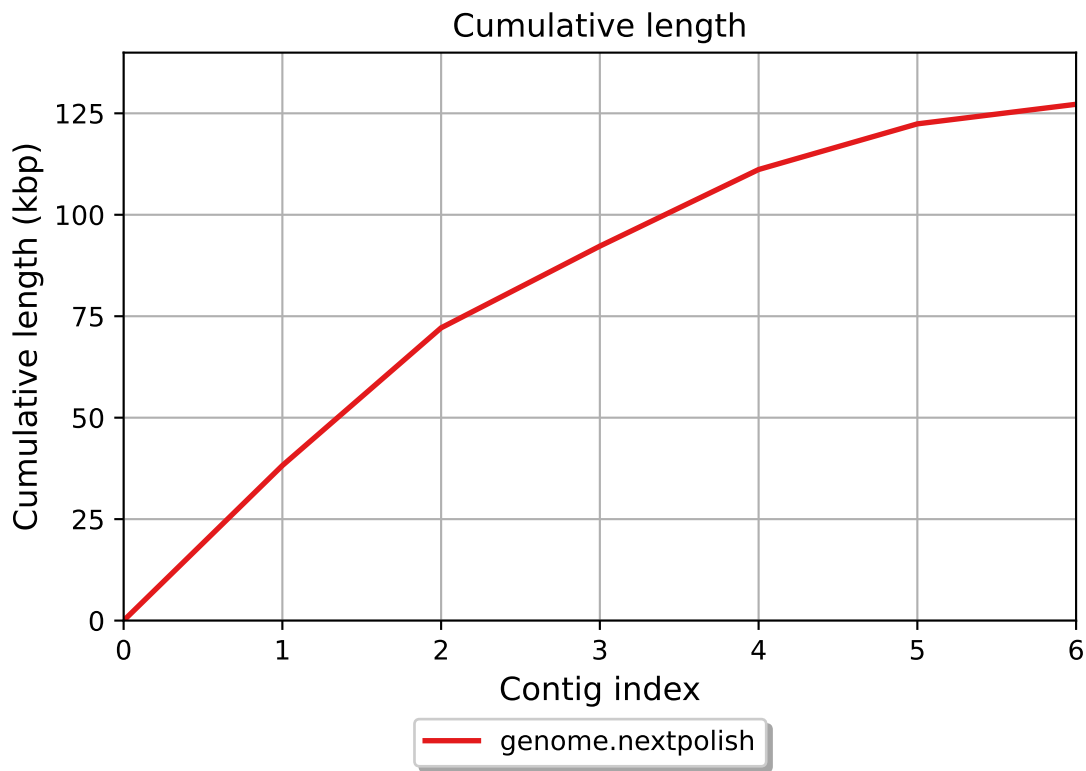
	genome.nextpolish
# contigs (>= 0 bp)	6
# contigs (>= 1000 bp)	6
# contigs (>= 5000 bp)	5
# contigs (>= 10000 bp)	5
# contigs (>= 25000 bp)	2
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	127230
Total length (>= 1000 bp)	127230
Total length (>= 5000 bp)	122400
Total length (>= 10000 bp)	122400
Total length (>= 25000 bp)	72136
Total length (>= 50000 bp)	0
# contigs	6
Largest contig	38206
Total length	127230
GC (%)	39.42
N50	33930
N90	11252
auN	27687.2
L50	2
L90	5
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	2 + 1 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).

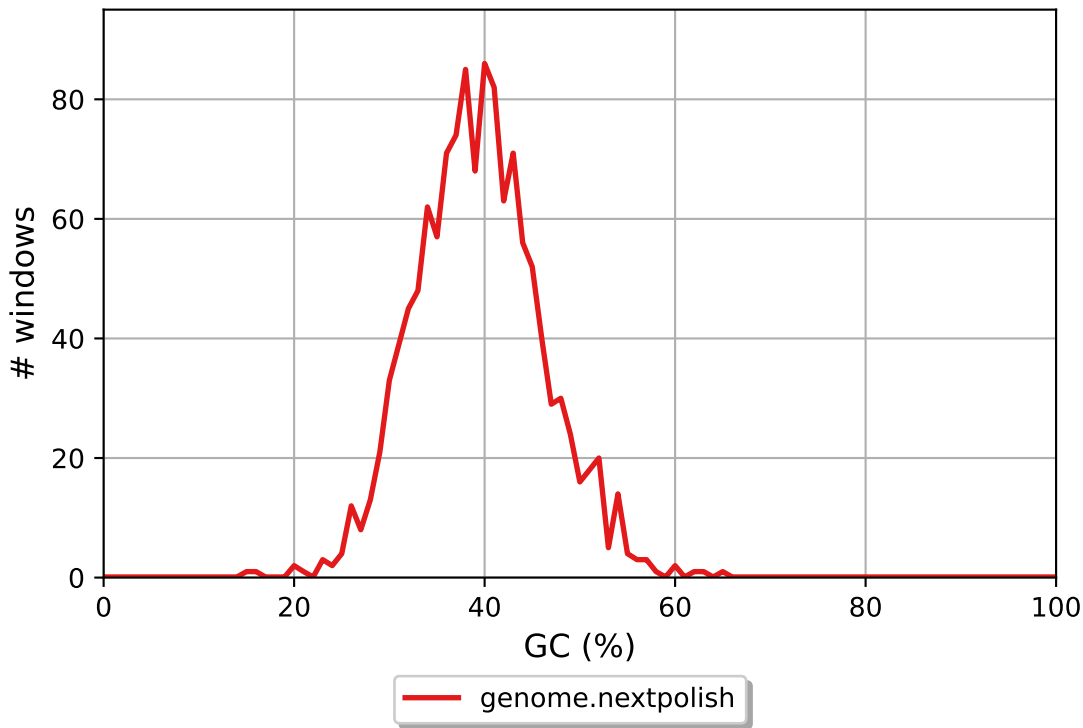
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

