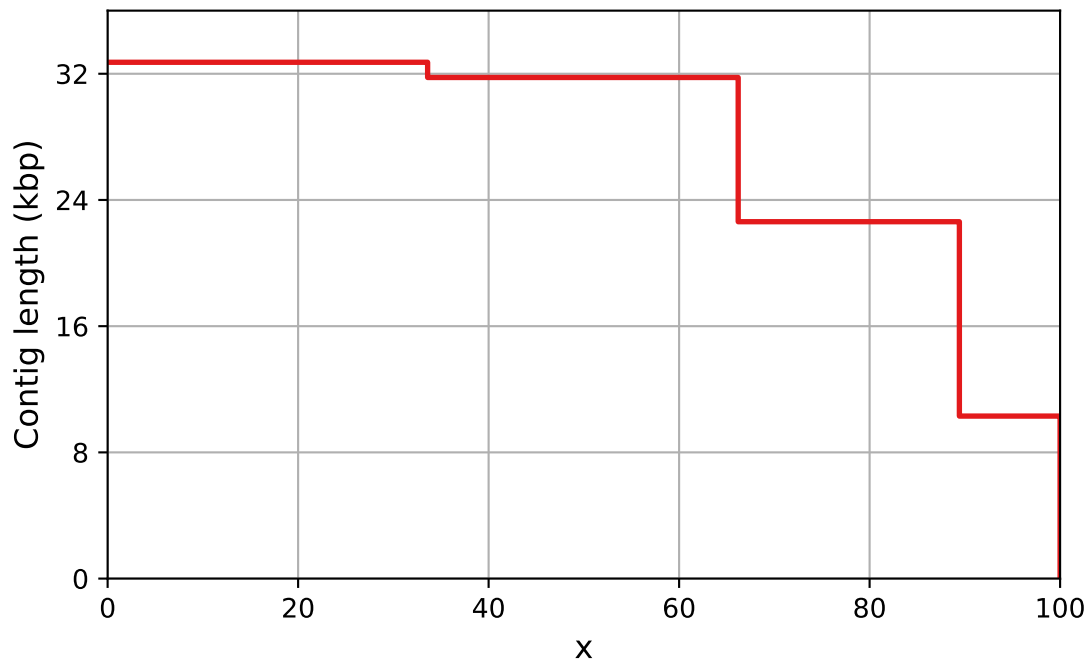


Report

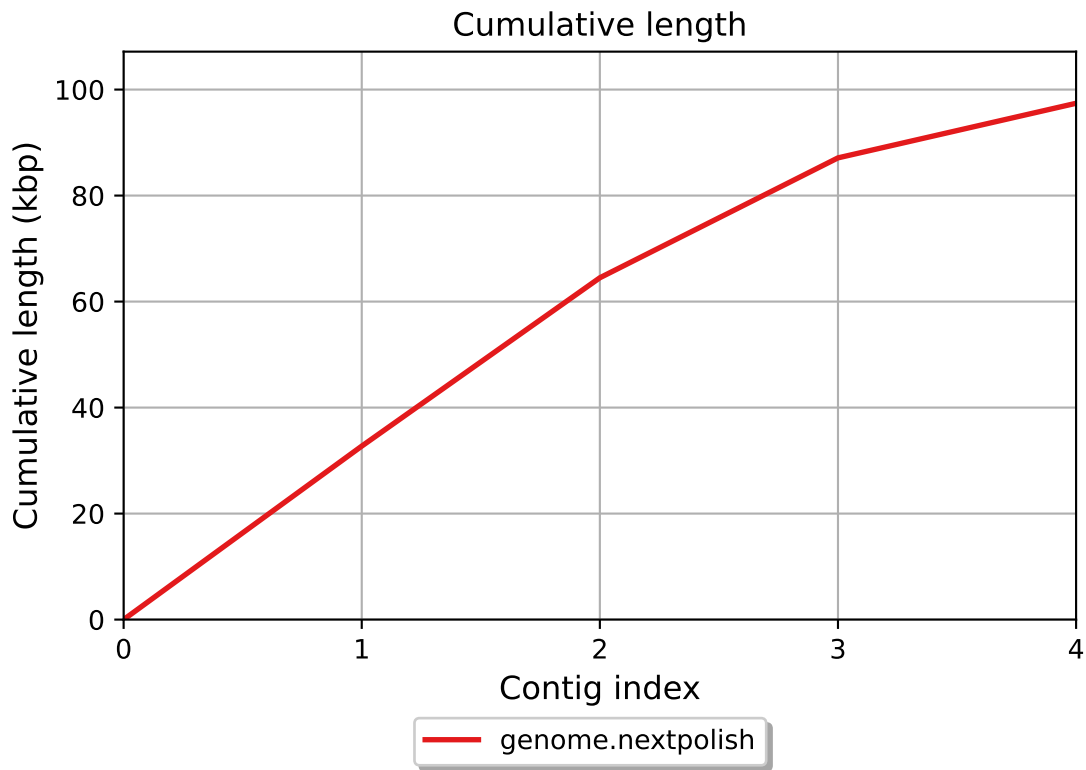
	genome.nextpolish
# contigs (>= 0 bp)	4
# contigs (>= 1000 bp)	4
# contigs (>= 5000 bp)	4
# contigs (>= 10000 bp)	4
# contigs (>= 25000 bp)	2
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	97415
Total length (>= 1000 bp)	97415
Total length (>= 5000 bp)	97415
Total length (>= 10000 bp)	97415
Total length (>= 25000 bp)	64492
Total length (>= 50000 bp)	0
# contigs	4
Largest contig	32728
Total length	97415
GC (%)	47.68
N50	31764
N90	10301
auN	27695.3
L50	2
L90	4
# 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).

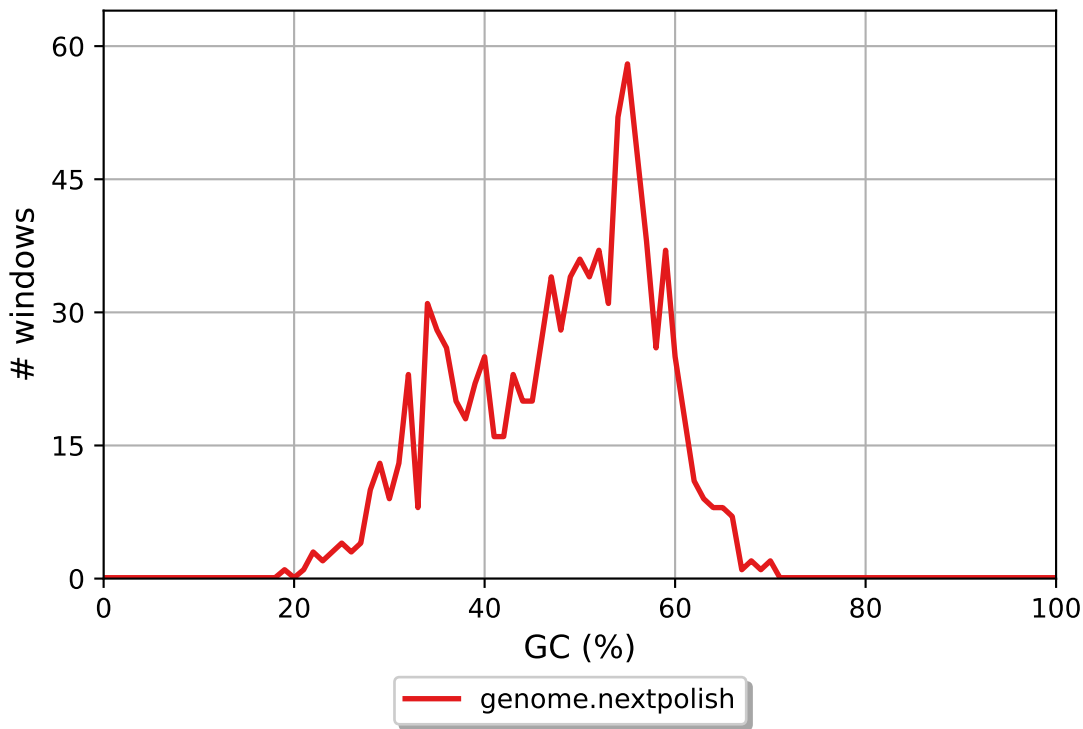
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

