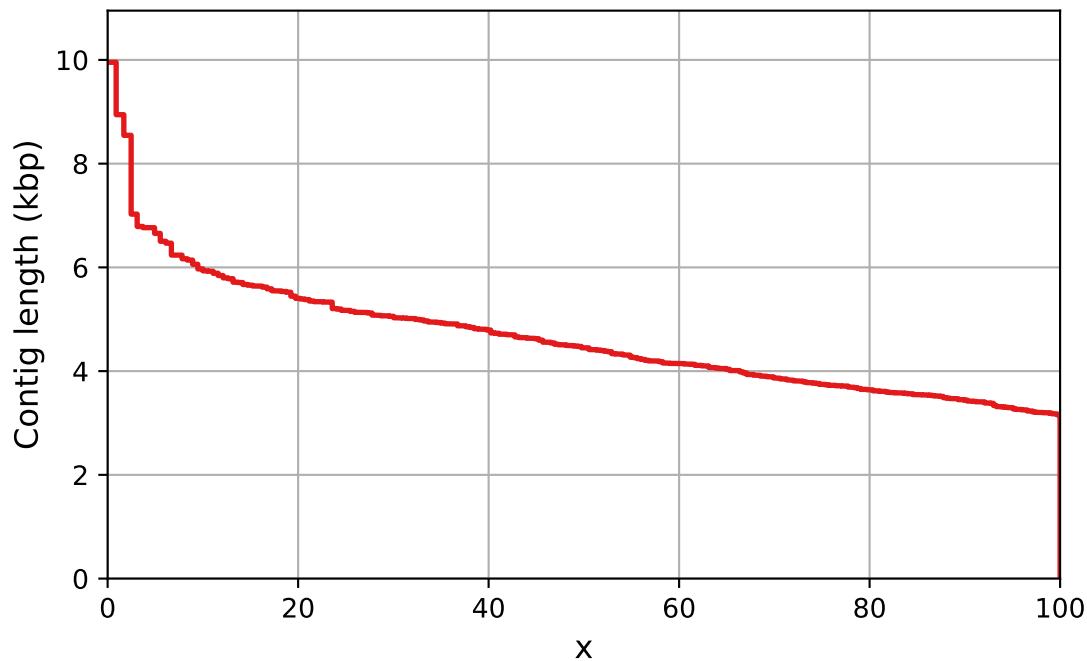


Report

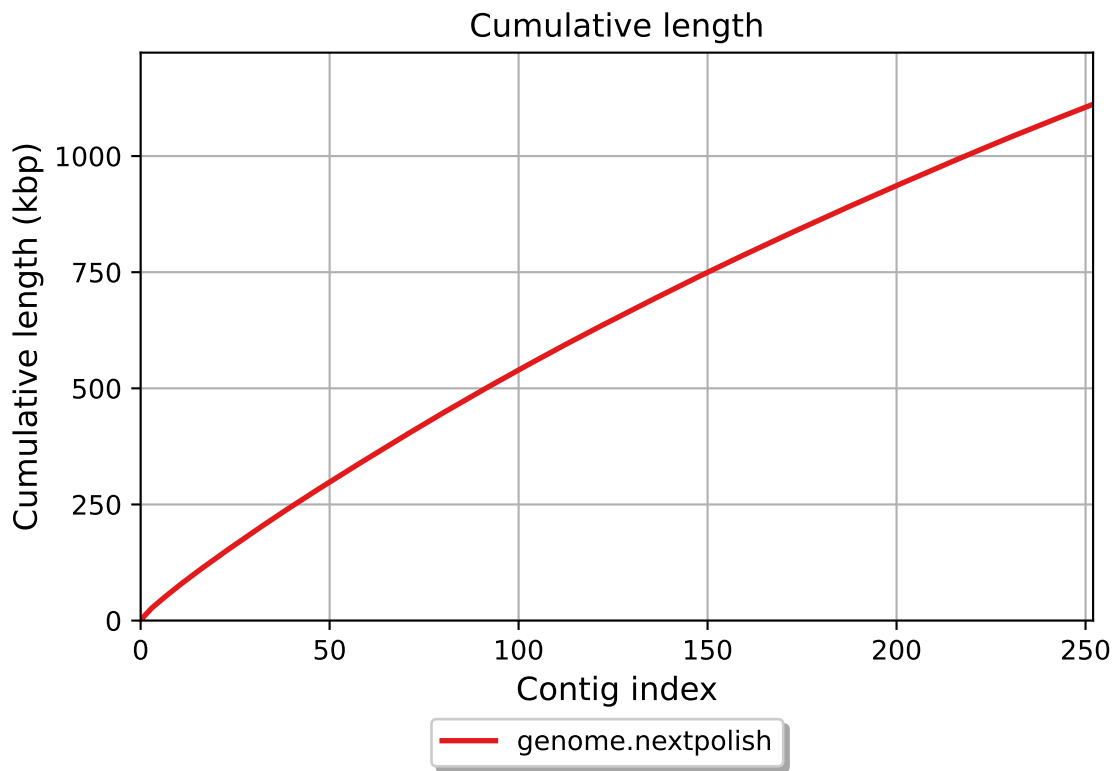
	genome.nextpolish
# contigs (>= 0 bp)	252
# contigs (>= 1000 bp)	252
# contigs (>= 5000 bp)	62
# contigs (>= 10000 bp)	0
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1111537
Total length (>= 1000 bp)	1111537
Total length (>= 5000 bp)	358948
Total length (>= 10000 bp)	0
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	252
Largest contig	9954
Total length	1111537
GC (%)	48.41
N50	4450
N90	3441
auN	4645.2
L50	104
L90	219
# 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).

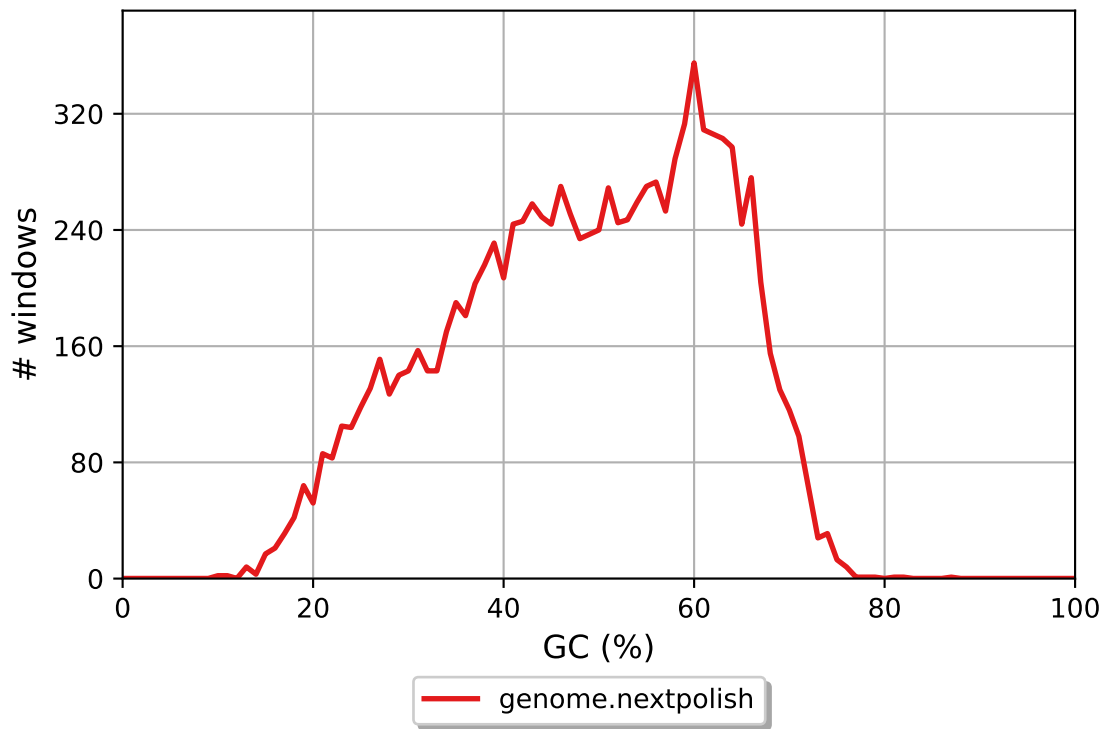
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

