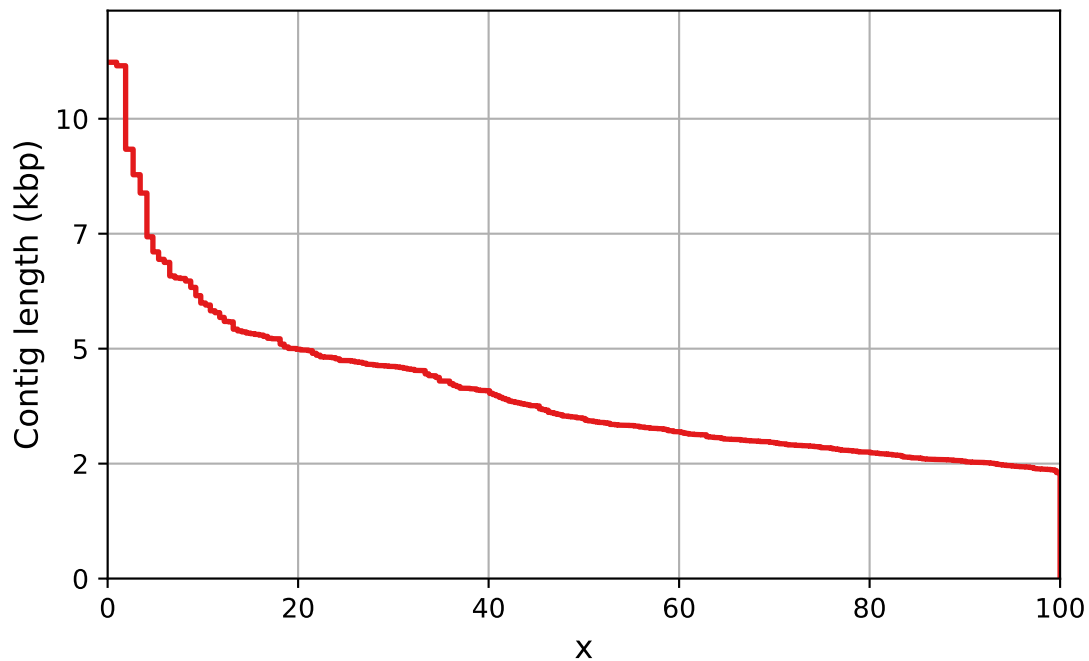


Report

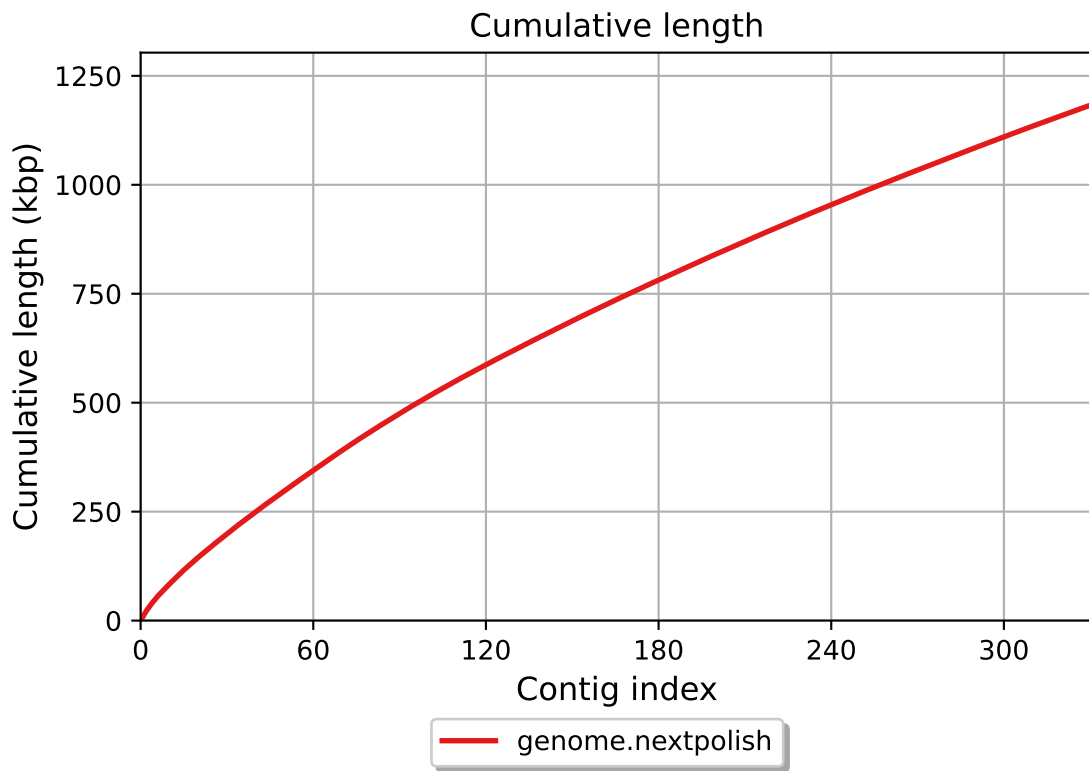
	genome.nextpolish
# contigs (>= 0 bp)	331
# contigs (>= 1000 bp)	331
# contigs (>= 5000 bp)	37
# contigs (>= 10000 bp)	2
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1184778
Total length (>= 1000 bp)	1184778
Total length (>= 5000 bp)	234757
Total length (>= 10000 bp)	22380
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	331
Largest contig	11228
Total length	1184778
GC (%)	62.03
N50	3481
N90	2553
auN	4056.3
L50	122
L90	283
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	8 + 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).

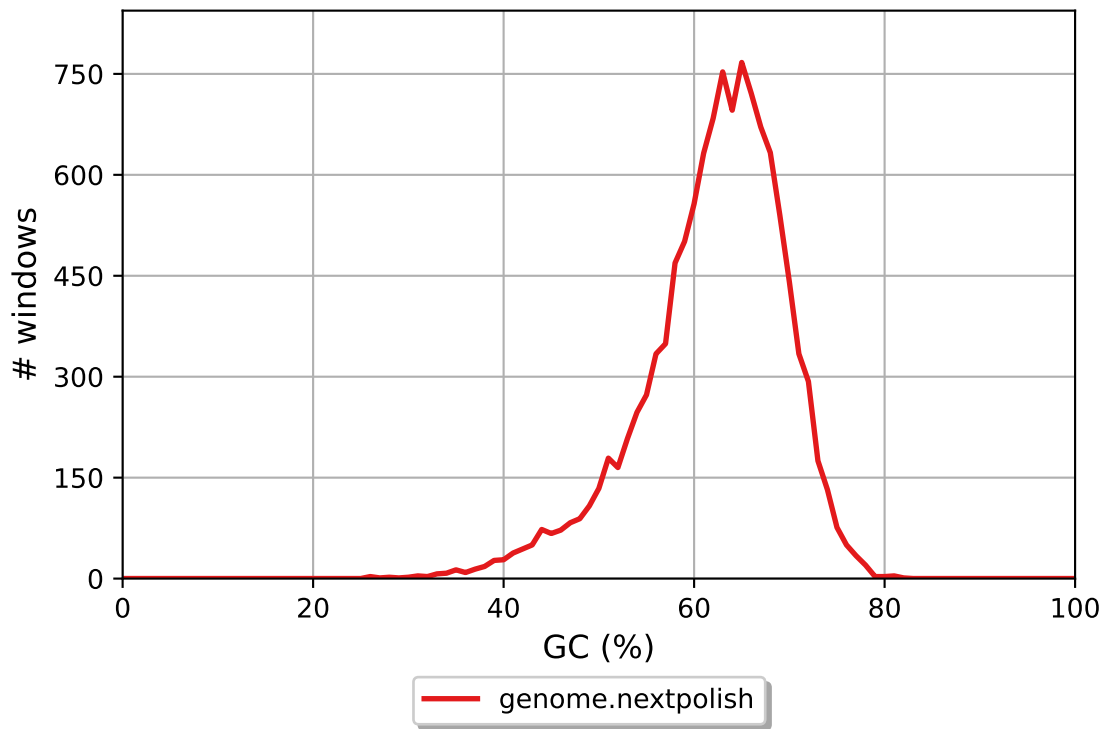
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

