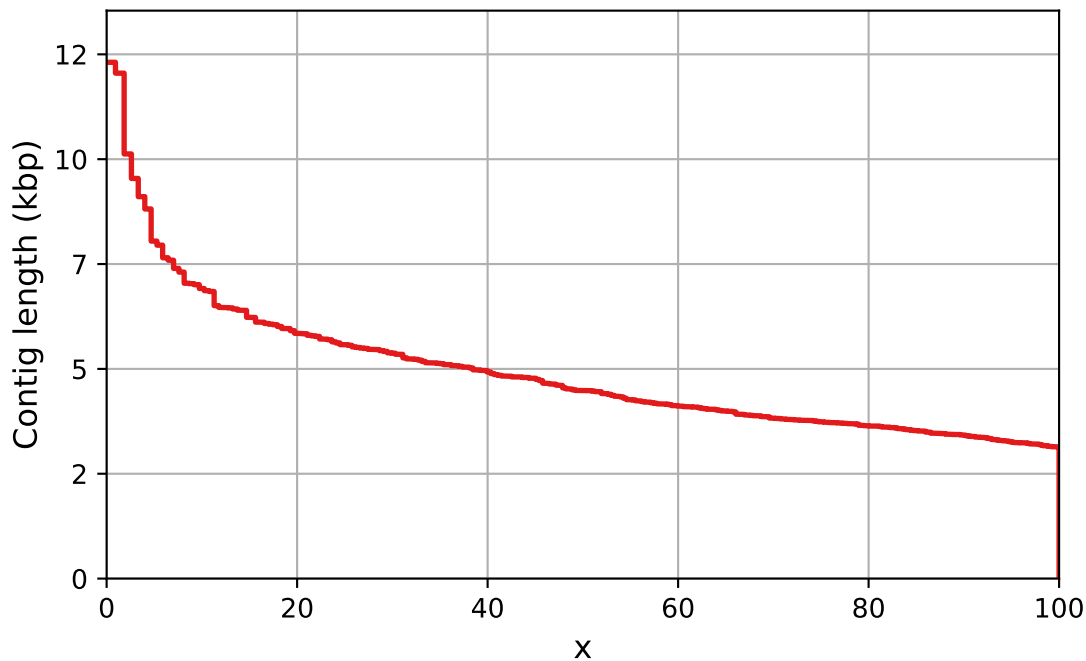


Report

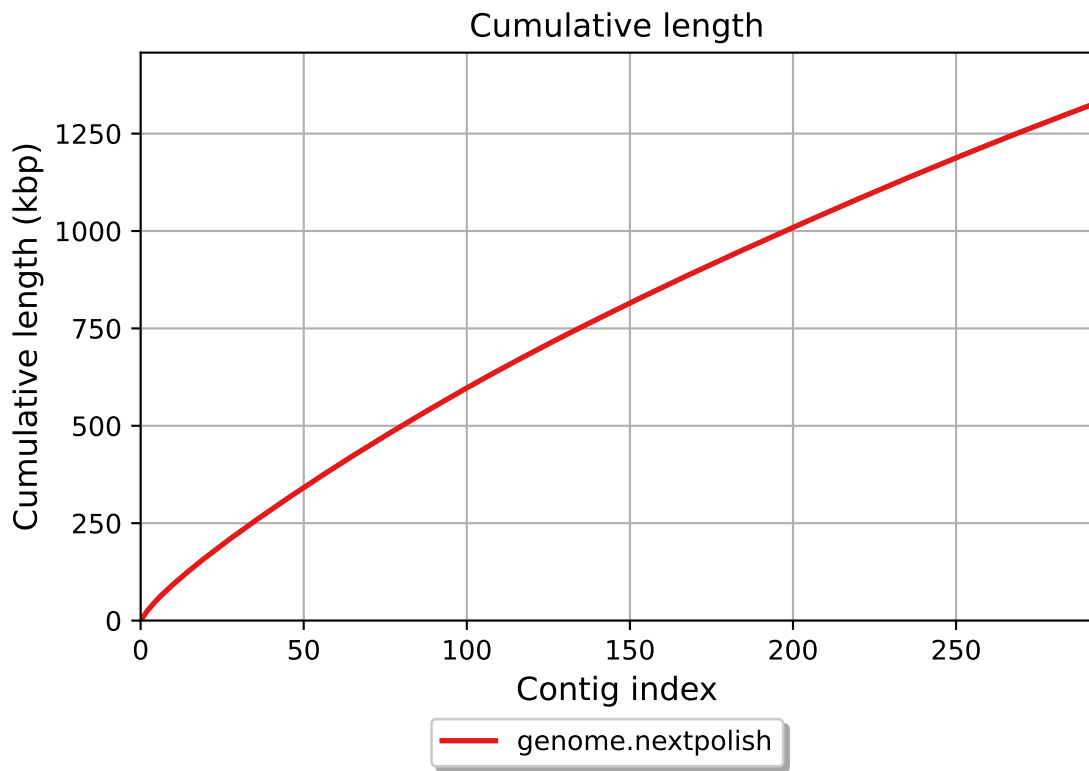
	genome.nextpolish
# contigs (>= 0 bp)	292
# contigs (>= 1000 bp)	292
# contigs (>= 5000 bp)	82
# contigs (>= 10000 bp)	3
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1325331
Total length (>= 1000 bp)	1325331
Total length (>= 5000 bp)	509717
Total length (>= 10000 bp)	34486
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	292
Largest contig	12310
Total length	1325331
GC (%)	33.56
N50	4481
N90	3421
auN	4943.3
L50	115
L90	252
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	0 + 2 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).

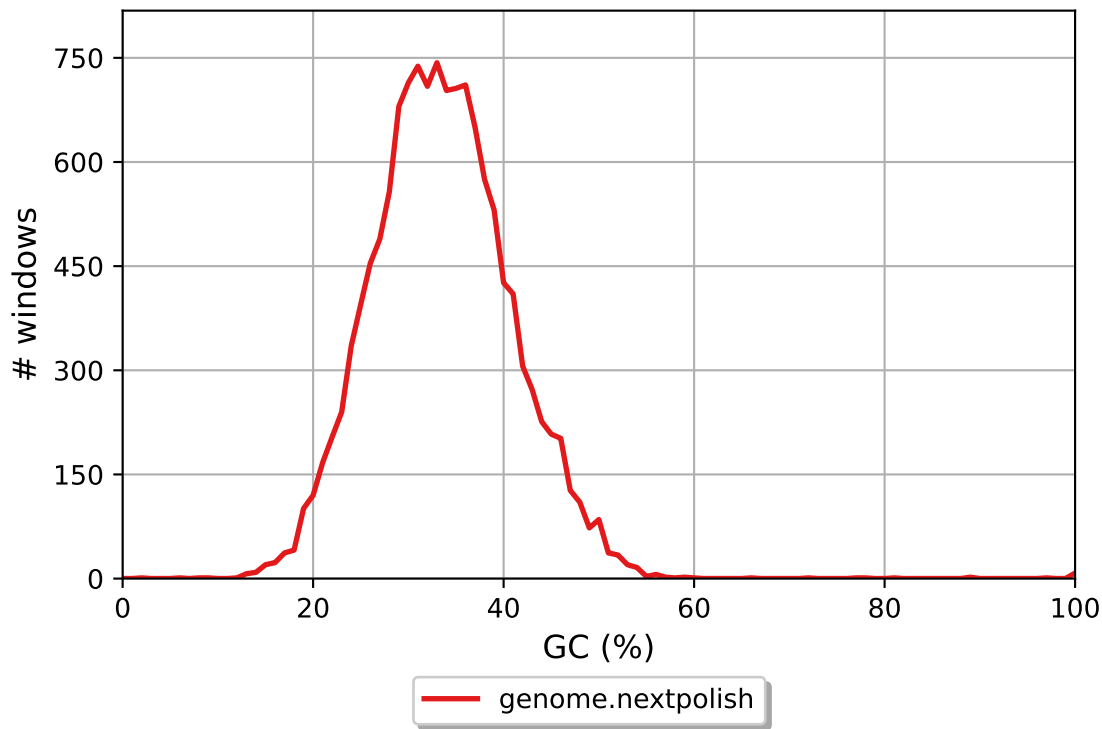
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

