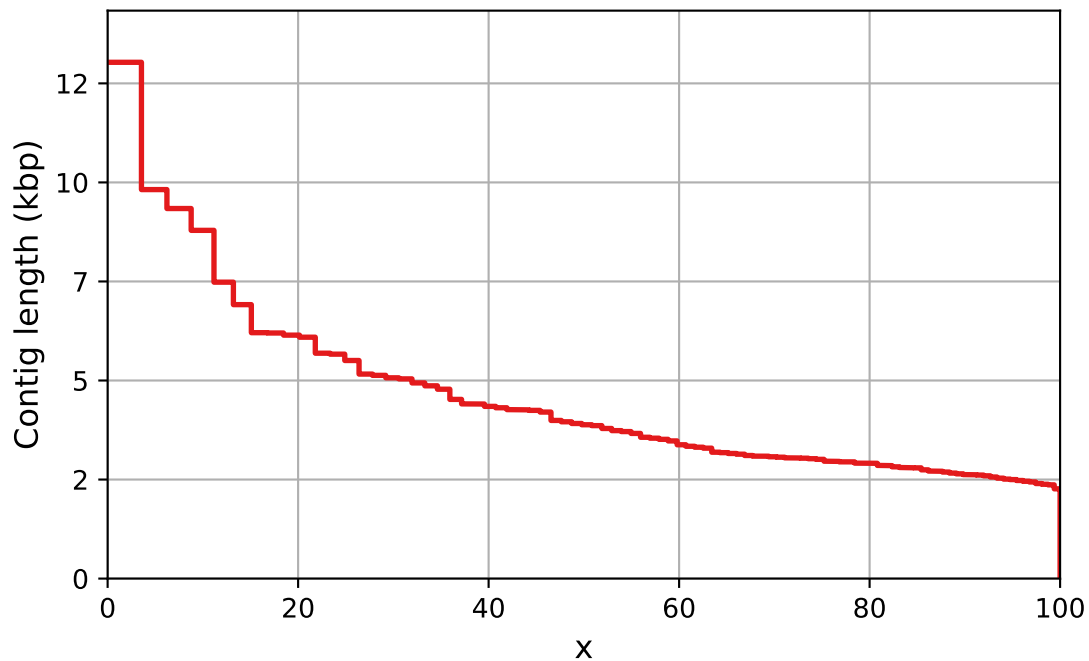


Report

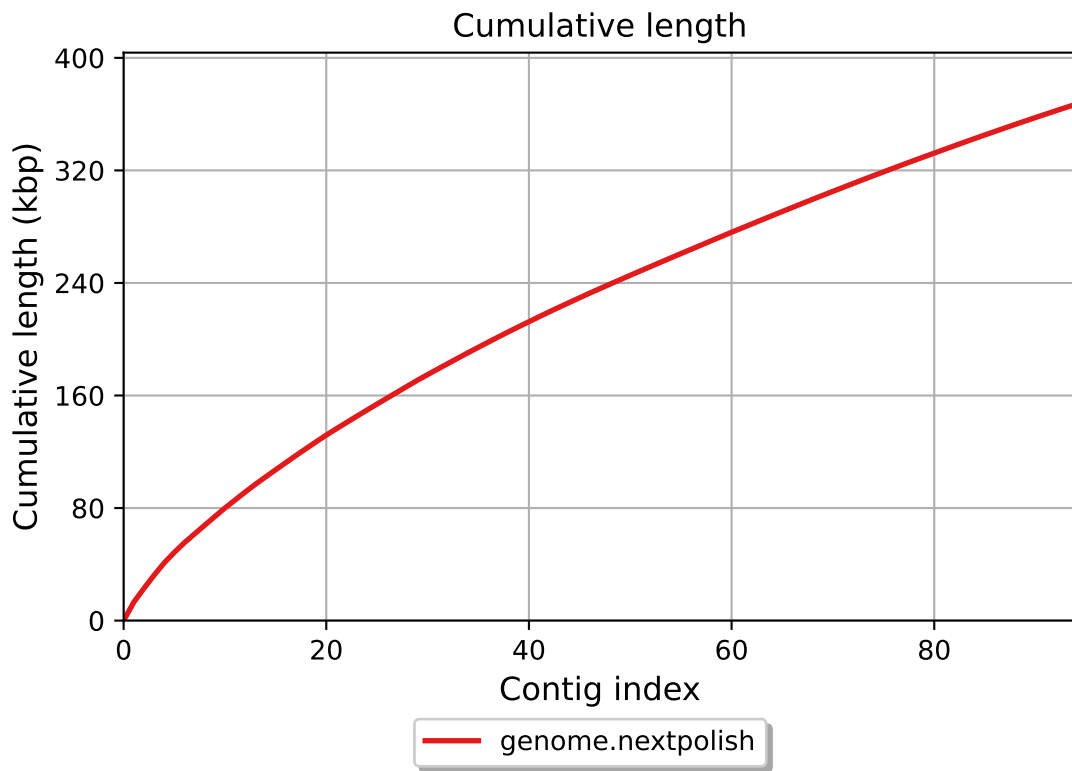
	genome.nextpolish
# contigs (>= 0 bp)	94
# contigs (>= 1000 bp)	94
# contigs (>= 5000 bp)	17
# contigs (>= 10000 bp)	1
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	367016
Total length (>= 1000 bp)	367016
Total length (>= 5000 bp)	117283
Total length (>= 10000 bp)	13032
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	94
Largest contig	13032
Total length	367016
GC (%)	34.61
N50	3883
N90	2625
auN	4713.6
L50	33
L90	80
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	1 + 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).

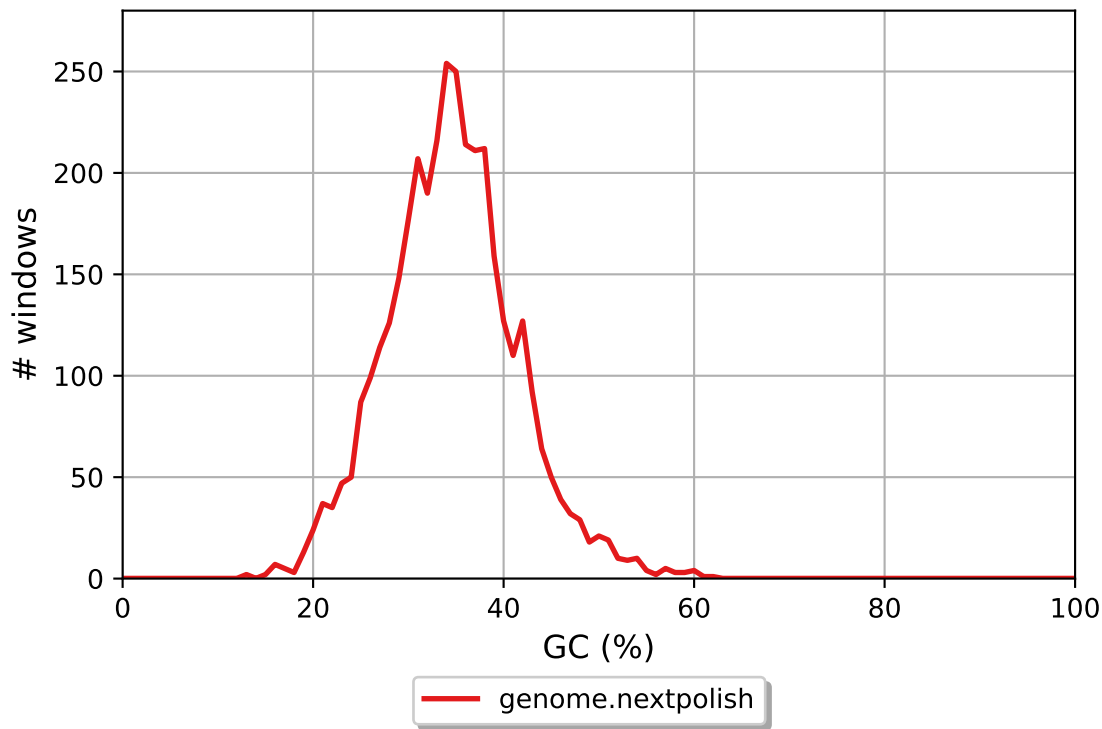
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

