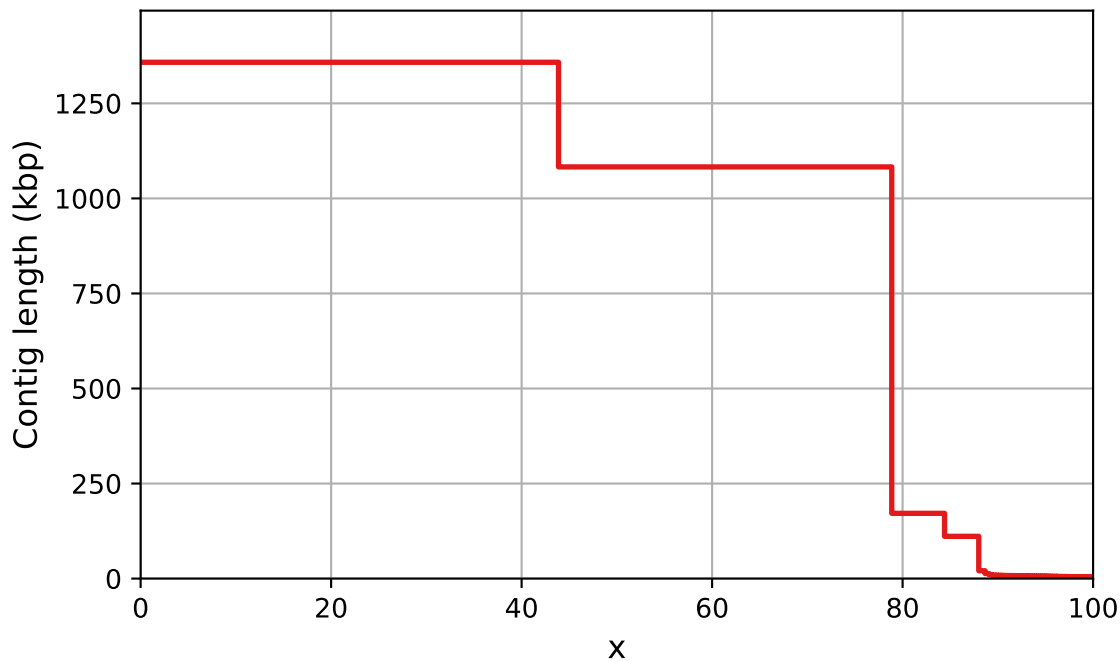


Report

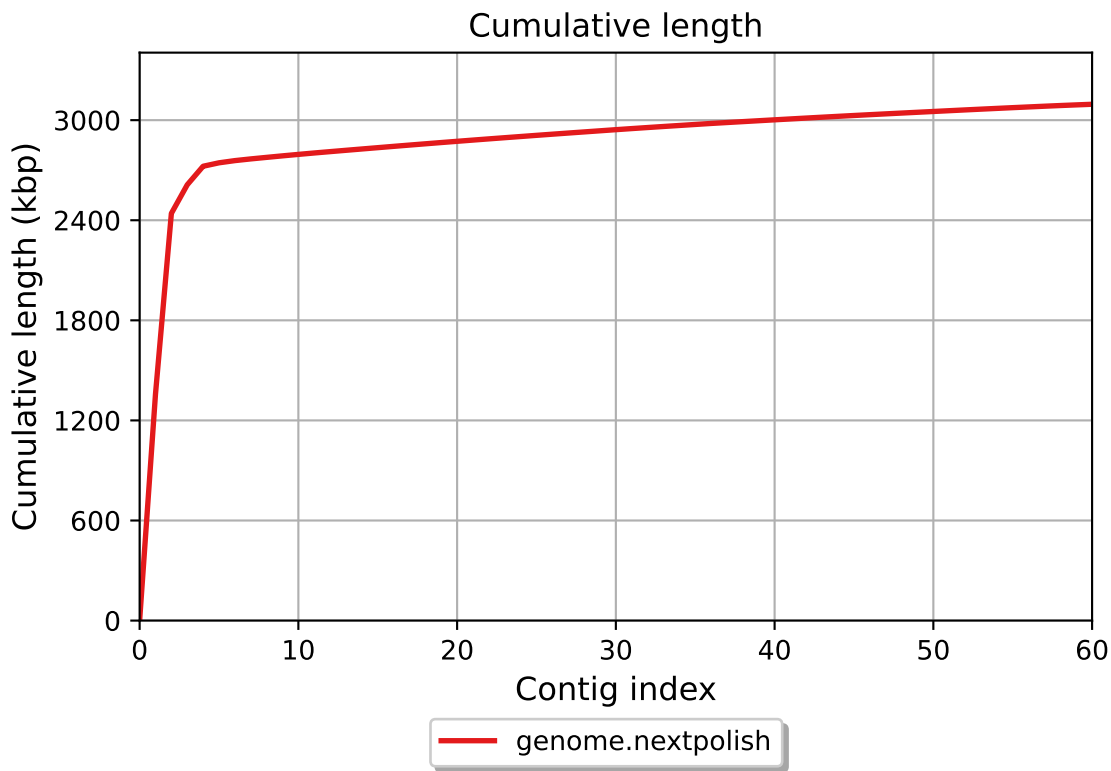
	genome.nextpolish
# contigs (>= 0 bp)	60
# contigs (>= 1000 bp)	60
# contigs (>= 5000 bp)	45
# contigs (>= 10000 bp)	7
# contigs (>= 25000 bp)	4
# contigs (>= 50000 bp)	4
Total length (>= 0 bp)	3095258
Total length (>= 1000 bp)	3095258
Total length (>= 5000 bp)	3027965
Total length (>= 10000 bp)	2767691
Total length (>= 25000 bp)	2723697
Total length (>= 50000 bp)	2723697
# contigs	60
Largest contig	1358099
Total length	3095258
GC (%)	34.97
N50	1082833
N90	9094
auN	989127.7
L50	2
L90	9
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	10 + 9 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).

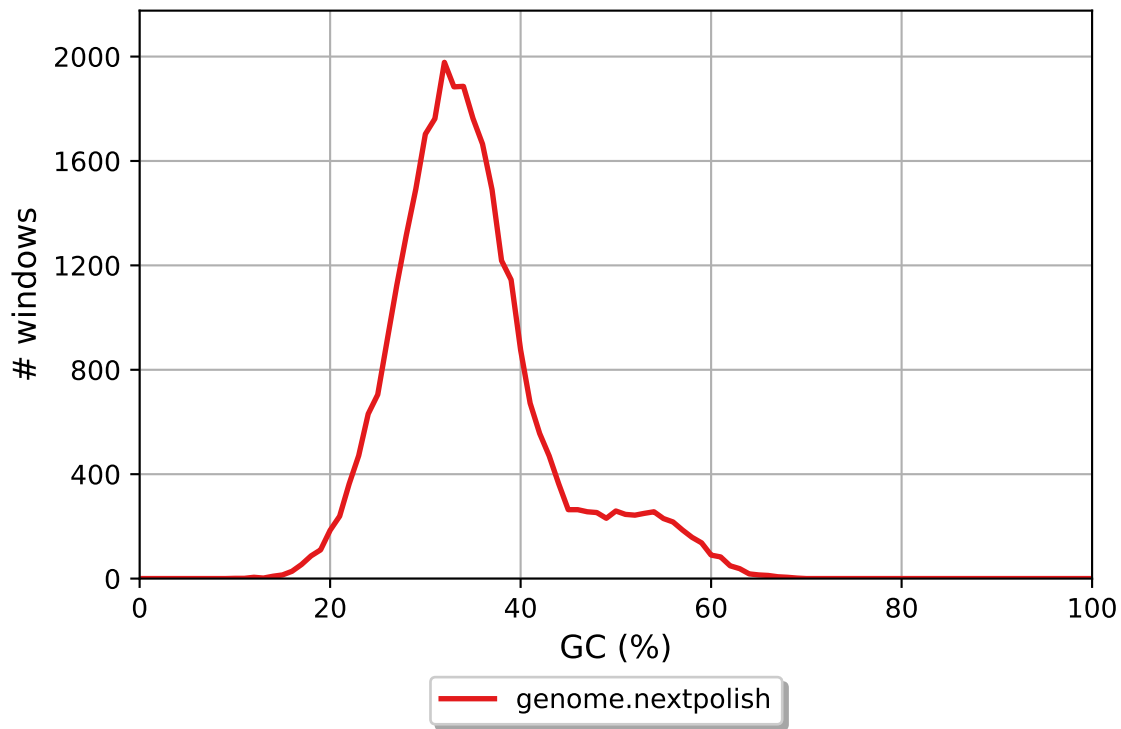
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

