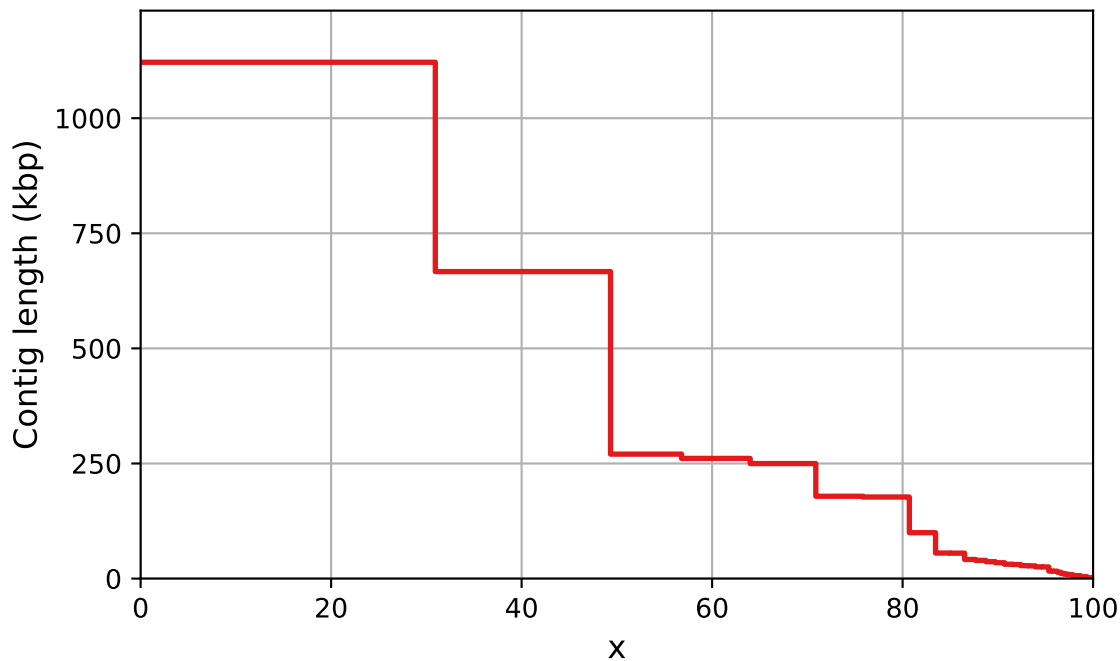


Report

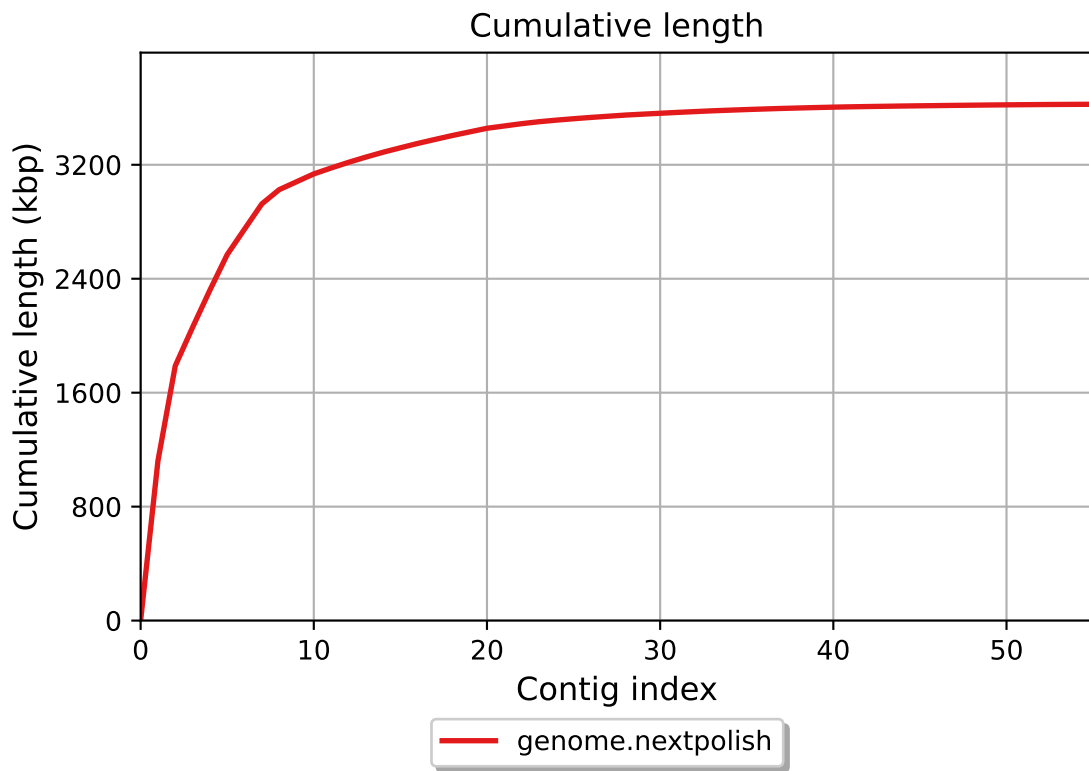
	genome.nextpolish
# contigs (>= 0 bp)	55
# contigs (>= 1000 bp)	51
# contigs (>= 5000 bp)	33
# contigs (>= 10000 bp)	24
# contigs (>= 25000 bp)	20
# contigs (>= 50000 bp)	10
Total length (>= 0 bp)	3624855
Total length (>= 1000 bp)	3621876
Total length (>= 5000 bp)	3578845
Total length (>= 10000 bp)	3513375
Total length (>= 25000 bp)	3456261
Total length (>= 50000 bp)	3135940
# contigs	55
Largest contig	1121485
Total length	3624855
GC (%)	37.54
N50	270359
N90	34439
auN	551060.3
L50	3
L90	14
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	18 + 0 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).

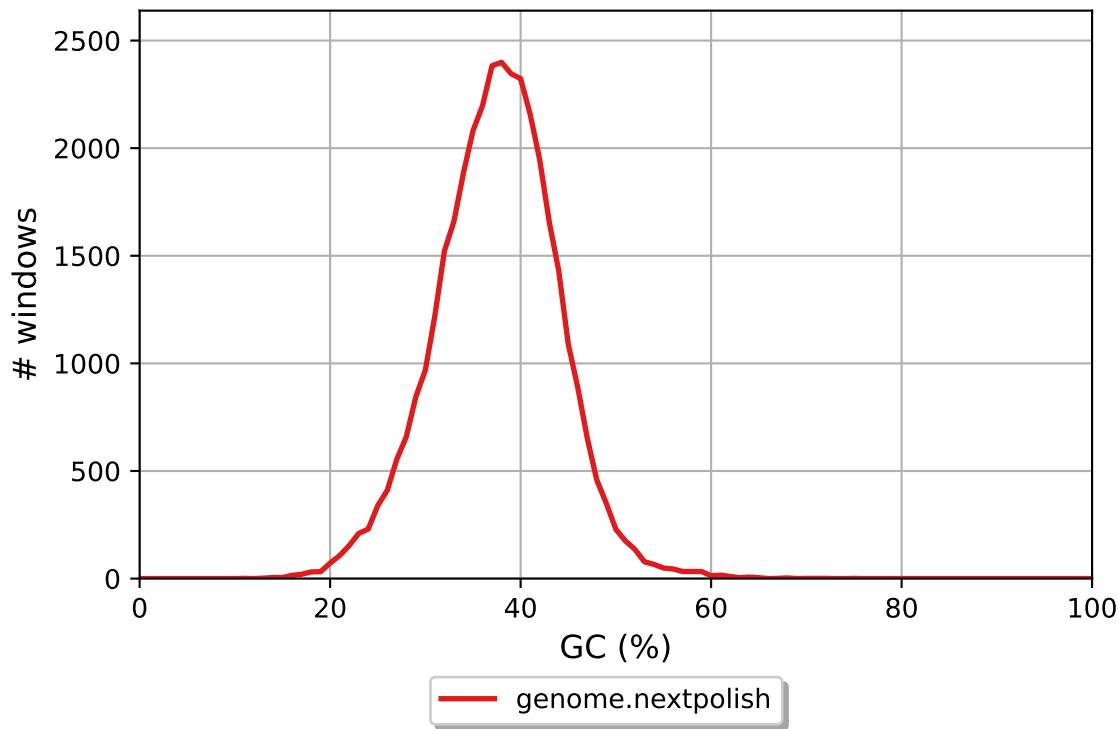
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

