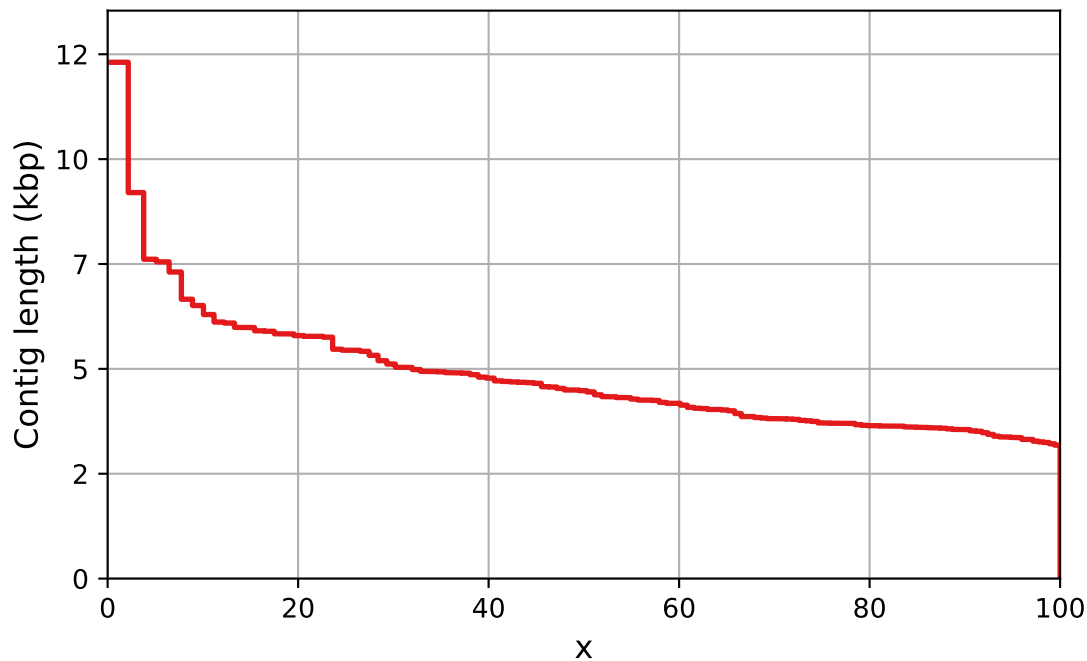


Report

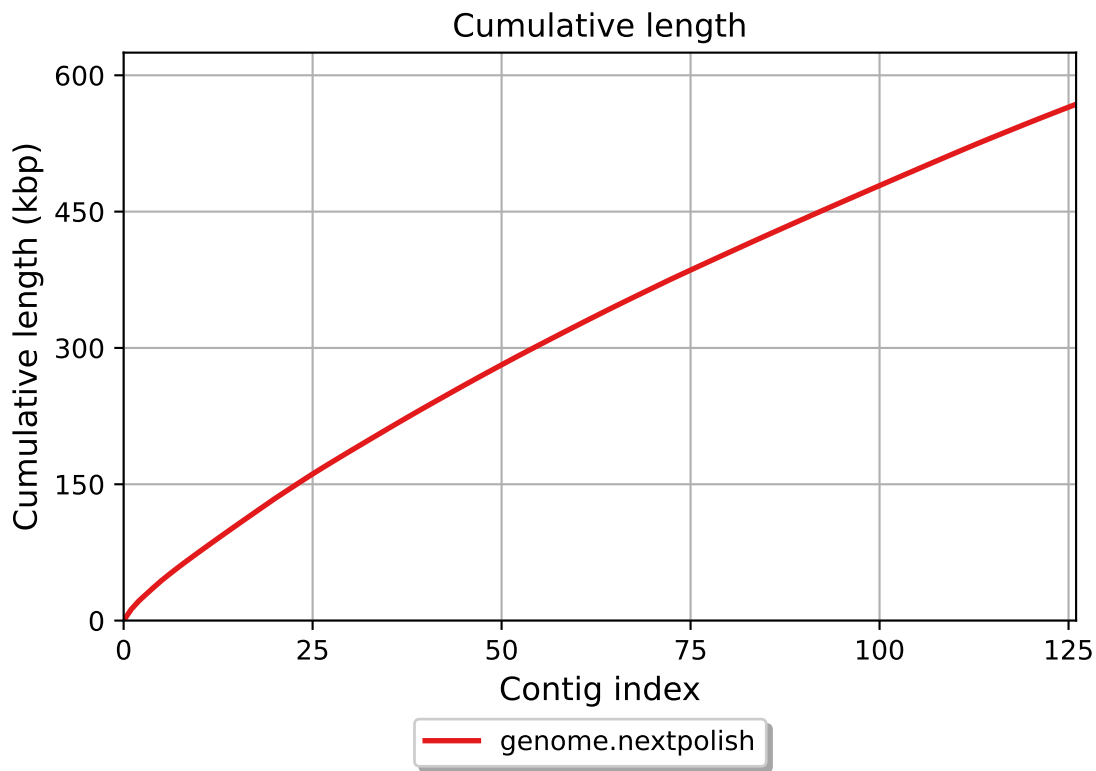
	genome.nextpolish
# contigs (>= 0 bp)	126
# contigs (>= 1000 bp)	126
# contigs (>= 5000 bp)	29
# contigs (>= 10000 bp)	1
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	568112
Total length (>= 1000 bp)	568112
Total length (>= 5000 bp)	181677
Total length (>= 10000 bp)	12309
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	126
Largest contig	12309
Total length	568112
GC (%)	37.20
N50	4480
N90	3551
auN	4863.3
L50	51
L90	110
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	2 + 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).

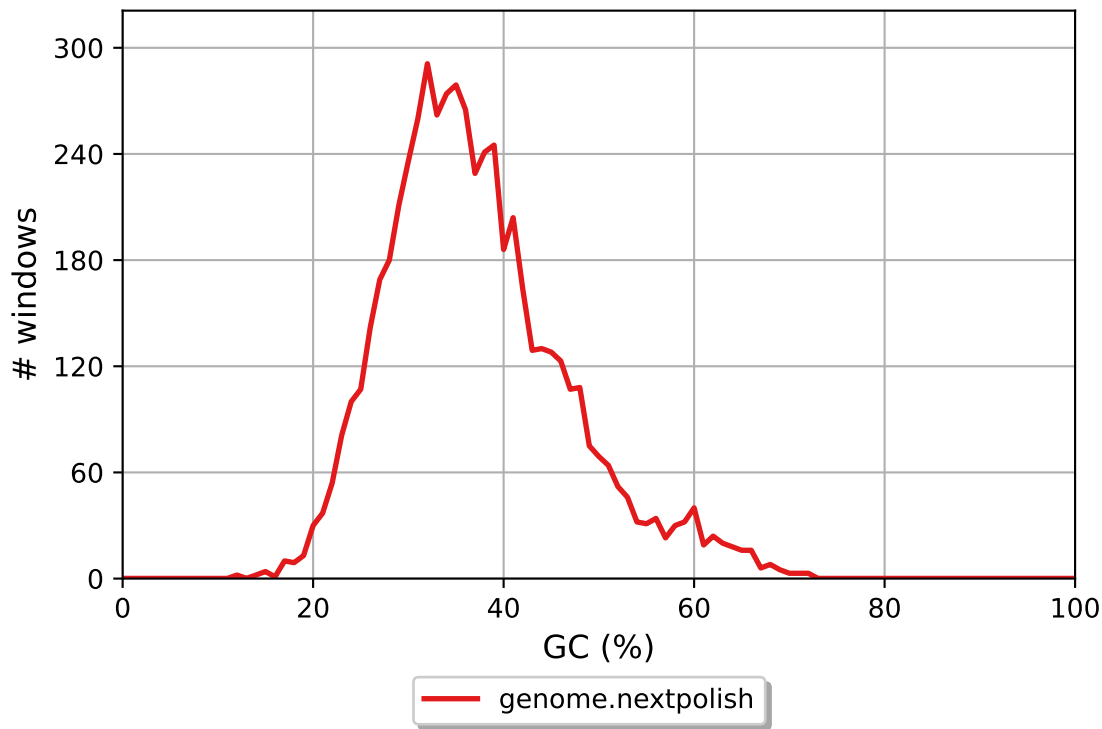
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

