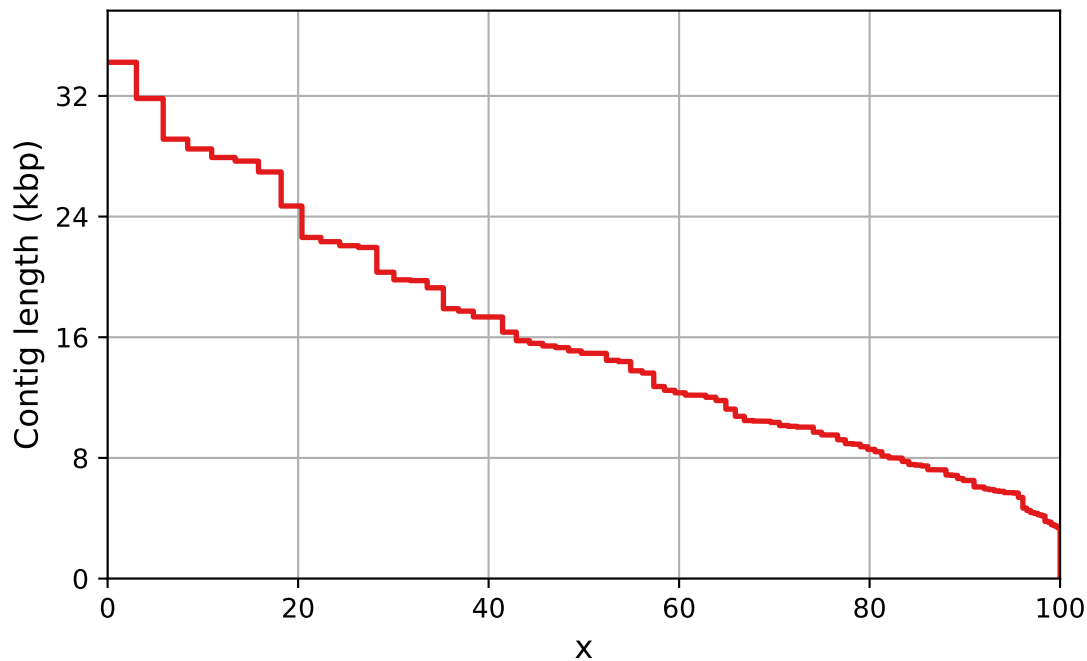


Report

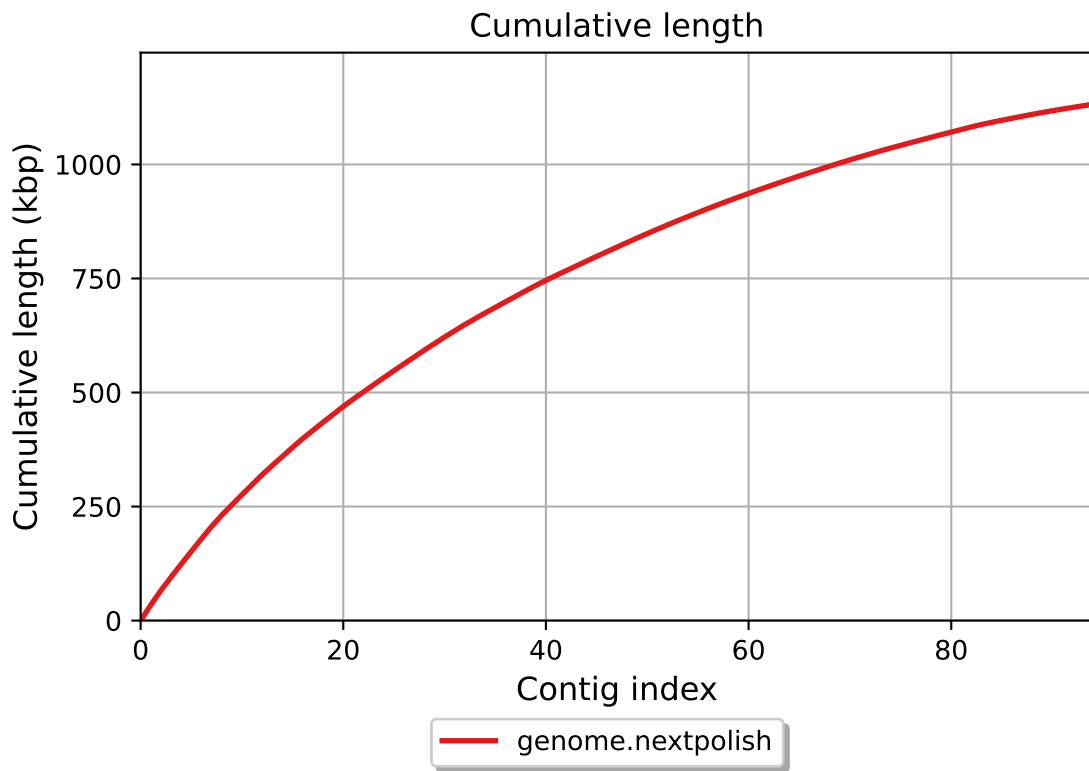
	genome.nextpolish
# contigs (>= 0 bp)	94
# contigs (>= 1000 bp)	94
# contigs (>= 5000 bp)	83
# contigs (>= 10000 bp)	49
# contigs (>= 25000 bp)	7
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1131931
Total length (>= 1000 bp)	1131931
Total length (>= 5000 bp)	1087695
Total length (>= 10000 bp)	838723
Total length (>= 25000 bp)	206228
Total length (>= 50000 bp)	0
# contigs	94
Largest contig	34230
Total length	1131931
GC (%)	42.06
N50	14937
N90	6504
auN	16253.7
L50	27
L90	72
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	3 + 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).

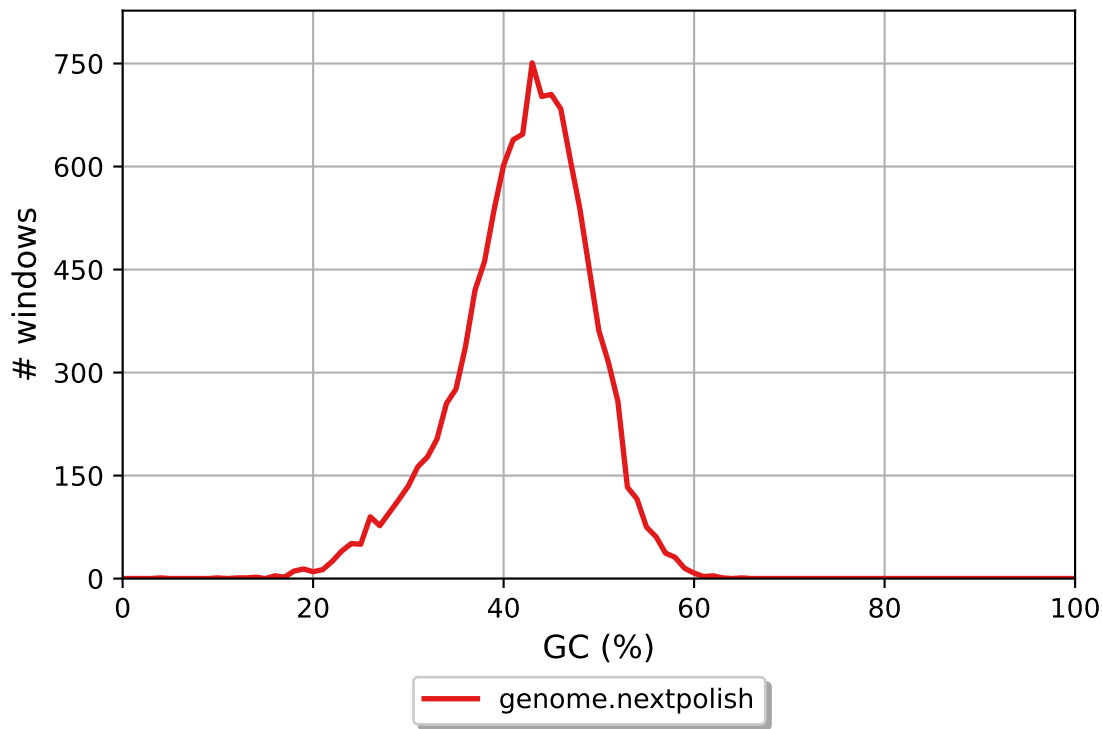
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

