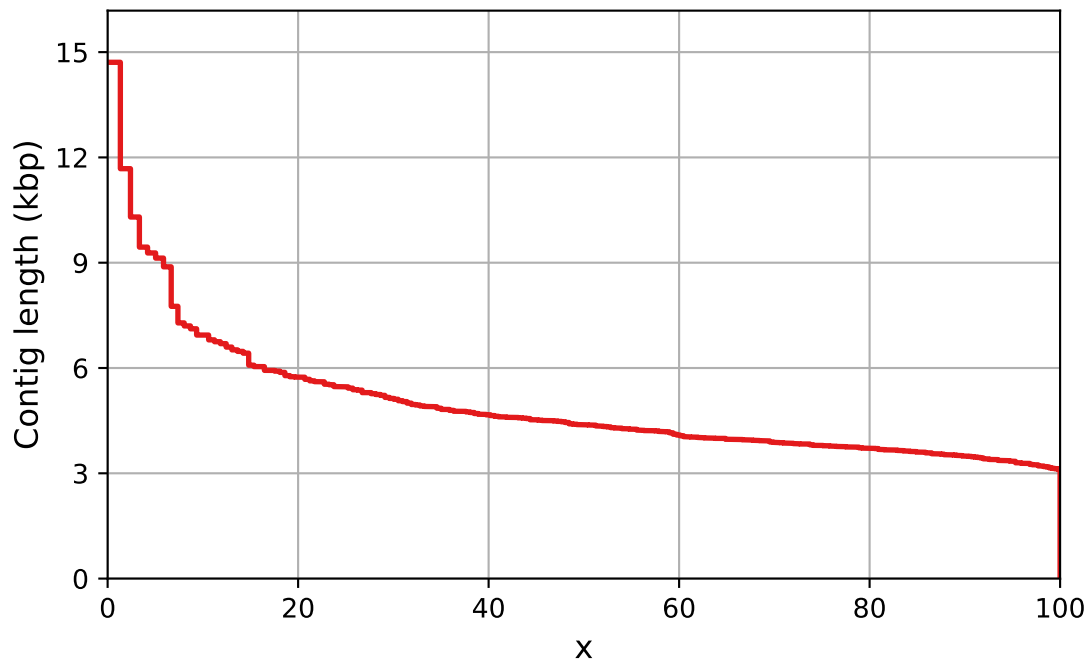


Report

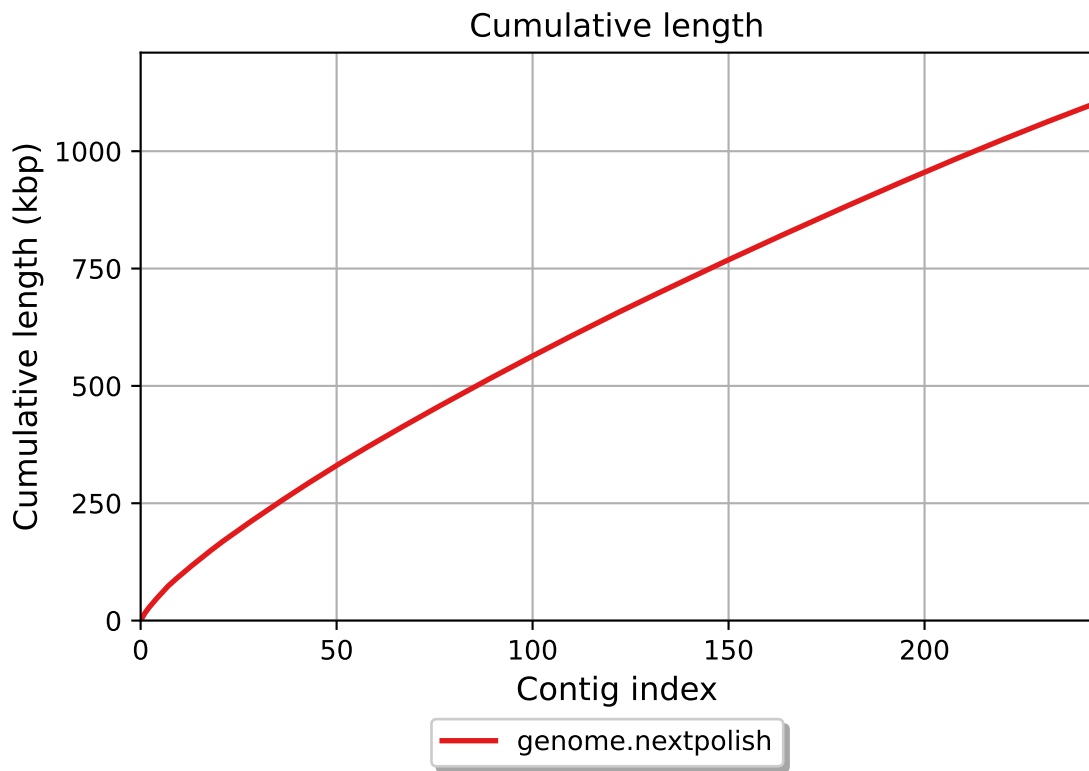
	genome.nextpolish
# contigs (>= 0 bp)	243
# contigs (>= 1000 bp)	243
# contigs (>= 5000 bp)	53
# contigs (>= 10000 bp)	3
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1099999
Total length (>= 1000 bp)	1099999
Total length (>= 5000 bp)	345749
Total length (>= 10000 bp)	36691
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	243
Largest contig	14710
Total length	1099999
GC (%)	35.82
N50	4387
N90	3492
auN	4971.2
L50	97
L90	210
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	3 + 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).

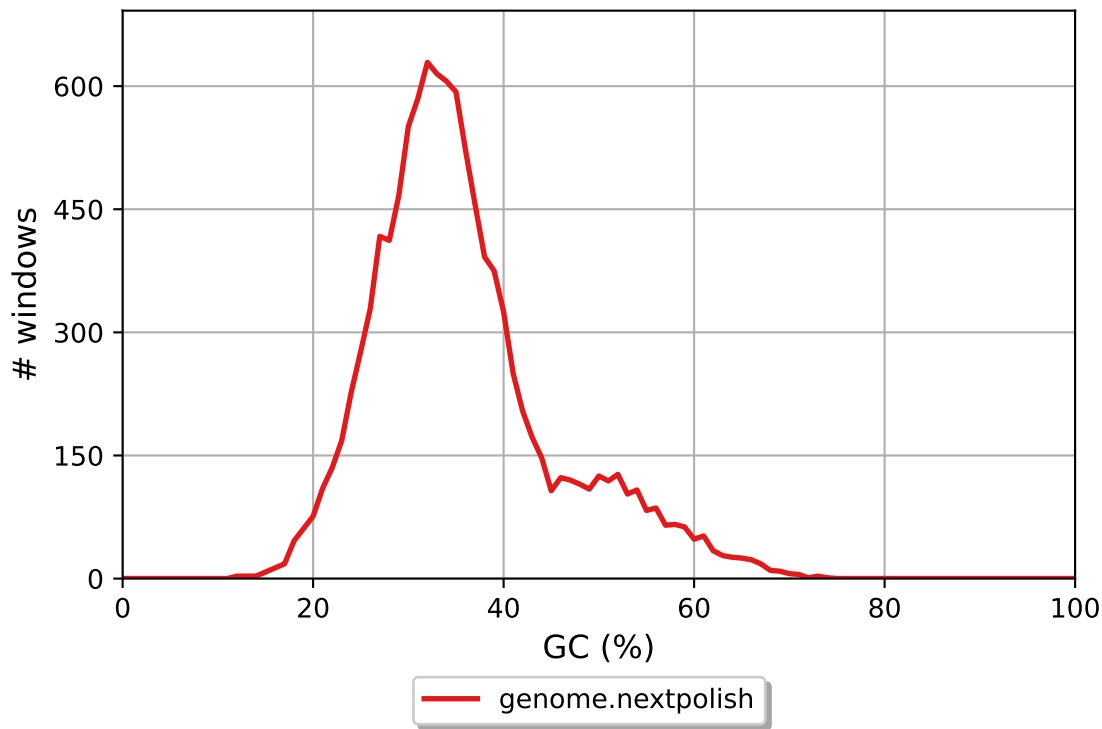
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

