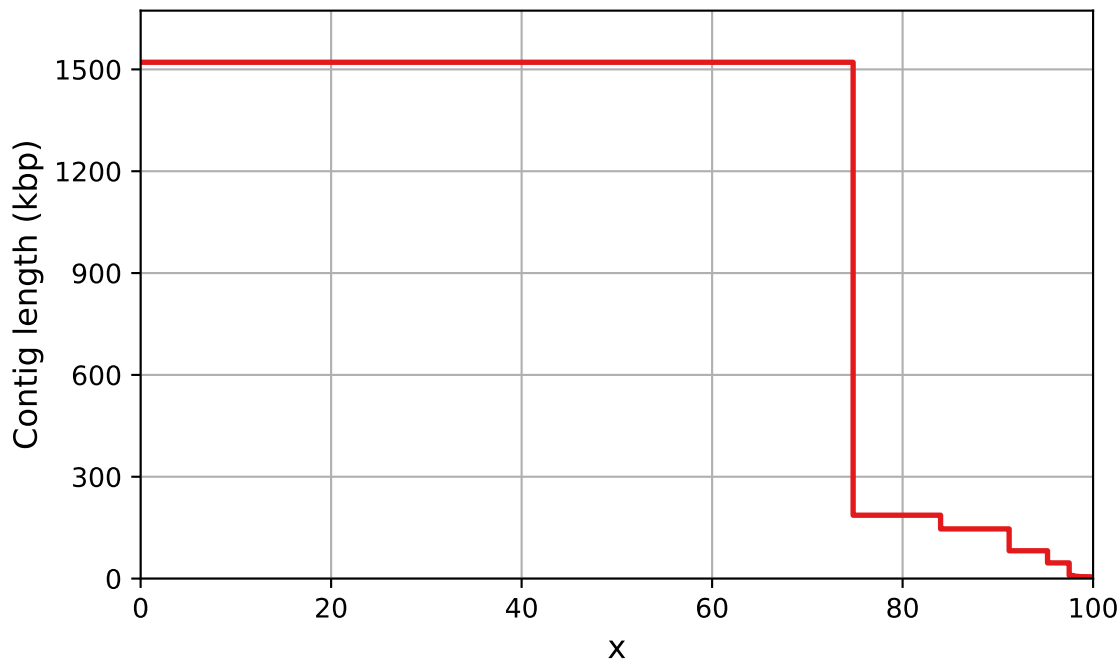


Report

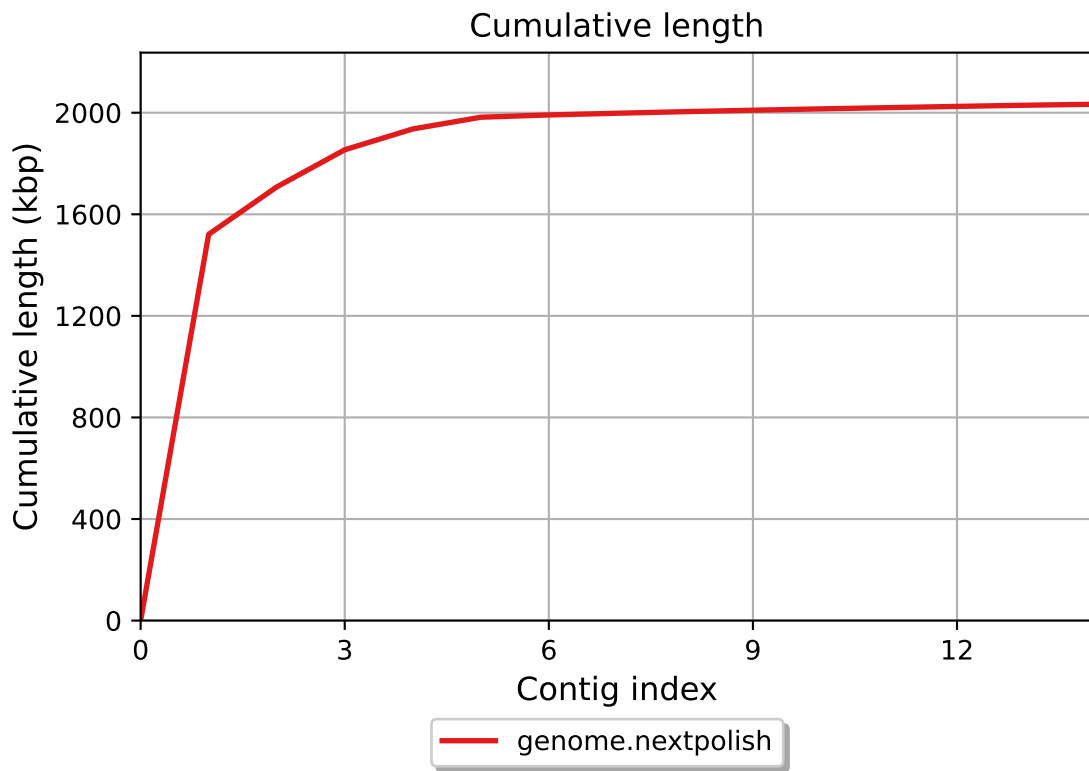
	genome.nextpolish
# contigs (>= 0 bp)	15
# contigs (>= 1000 bp)	14
# contigs (>= 5000 bp)	10
# contigs (>= 10000 bp)	5
# contigs (>= 25000 bp)	5
# contigs (>= 50000 bp)	4
Total length (>= 0 bp)	2033337
Total length (>= 1000 bp)	2033266
Total length (>= 5000 bp)	2015164
Total length (>= 10000 bp)	1982208
Total length (>= 25000 bp)	1982208
Total length (>= 50000 bp)	1935992
# contigs	14
Largest contig	1520982
Total length	2033266
GC (%)	41.45
N50	1520982
N90	146363
auN	1169953.3
L50	1
L90	3
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	13 + 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).

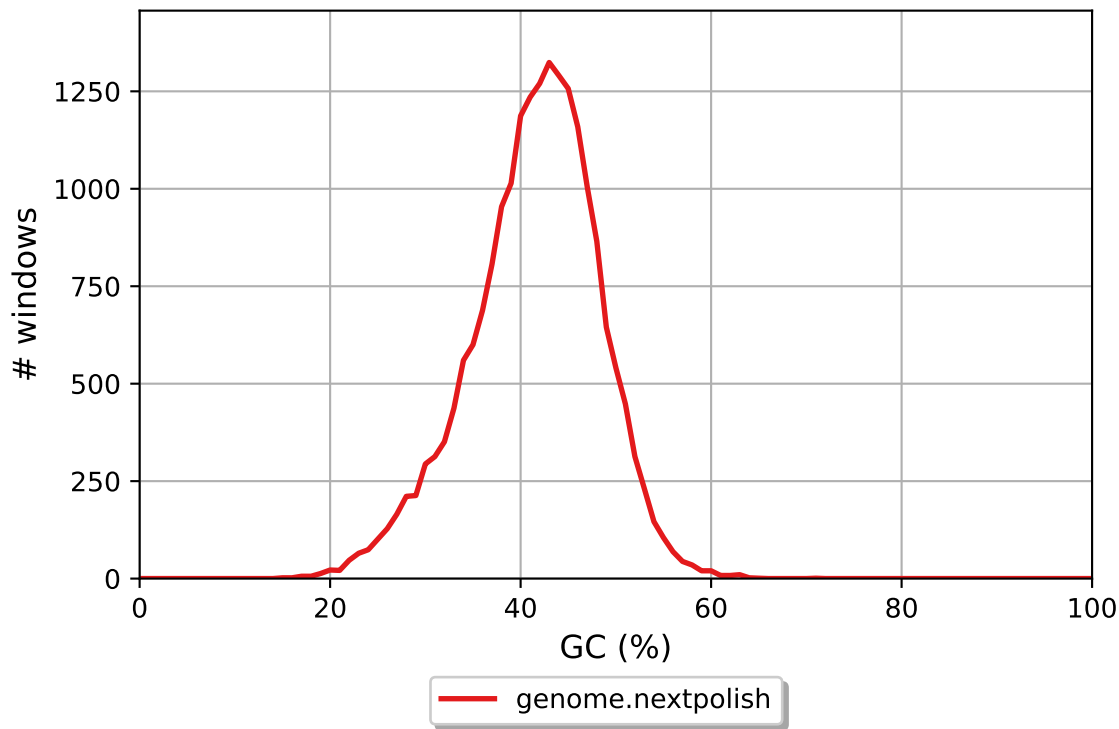
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

