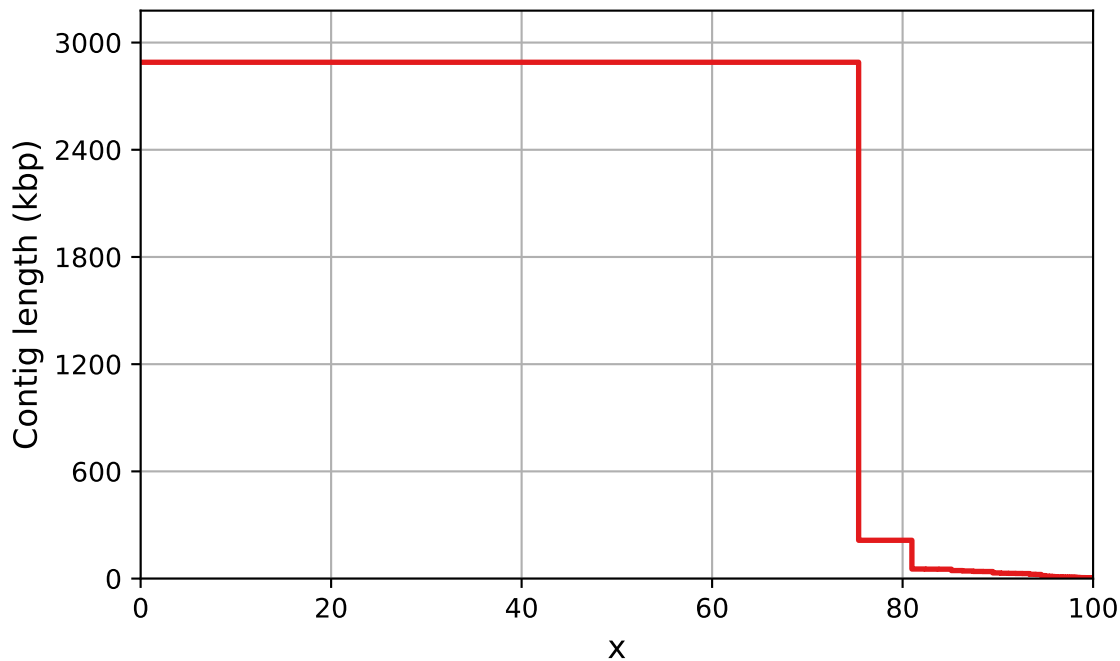


Report

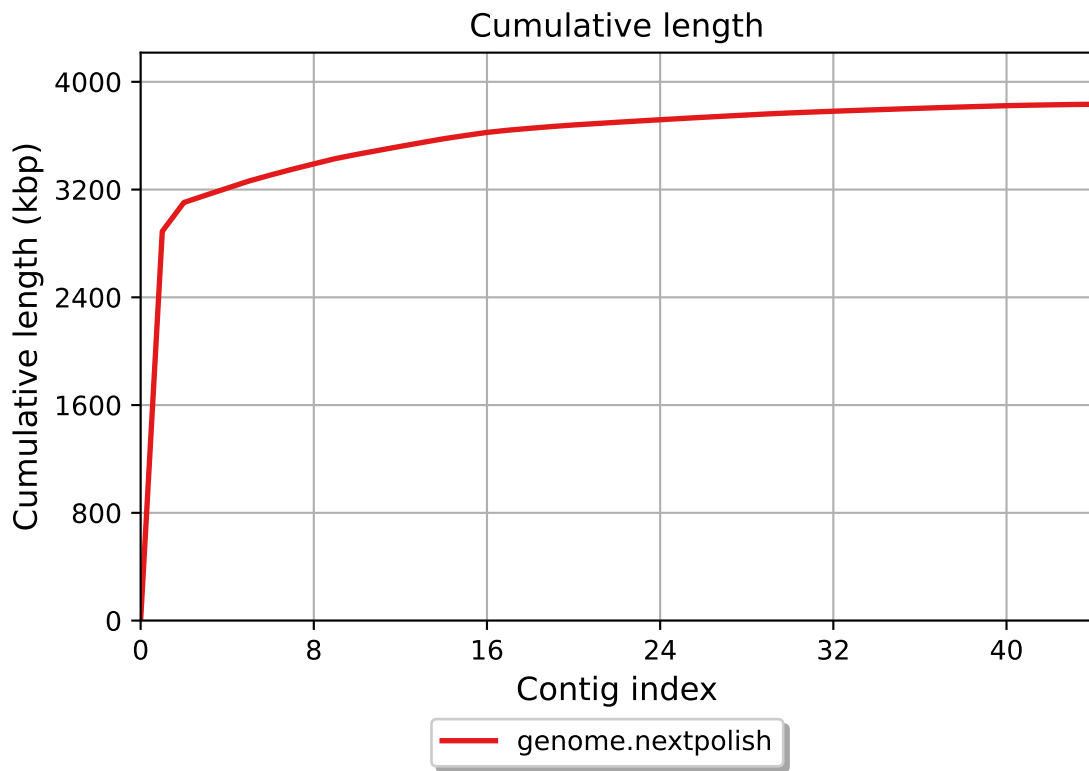
	genome.nextpolish
# contigs (>= 0 bp)	44
# contigs (>= 1000 bp)	44
# contigs (>= 5000 bp)	37
# contigs (>= 10000 bp)	21
# contigs (>= 25000 bp)	14
# contigs (>= 50000 bp)	5
Total length (>= 0 bp)	3833316
Total length (>= 1000 bp)	3833316
Total length (>= 5000 bp)	3808674
Total length (>= 10000 bp)	3689641
Total length (>= 25000 bp)	3577340
Total length (>= 50000 bp)	3263075
# contigs	44
Largest contig	2889715
Total length	3833316
GC (%)	37.73
N50	2889715
N90	31692
auN	2196317.2
L50	1
L90	10
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	21 + 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).

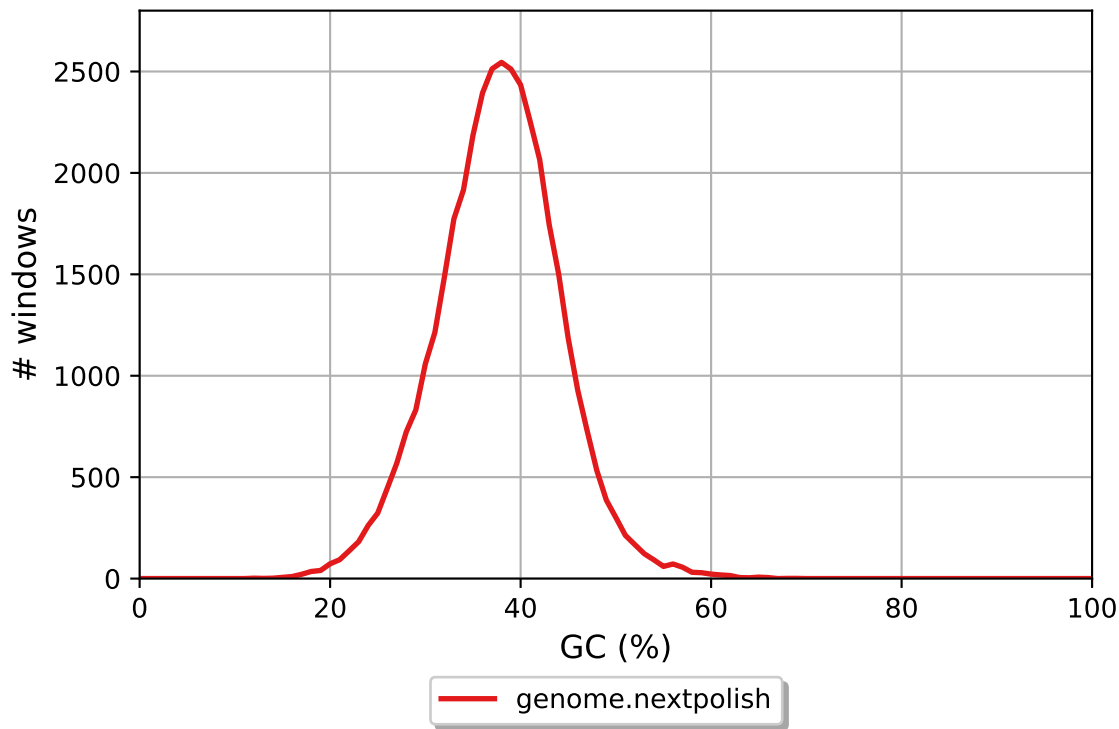
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

