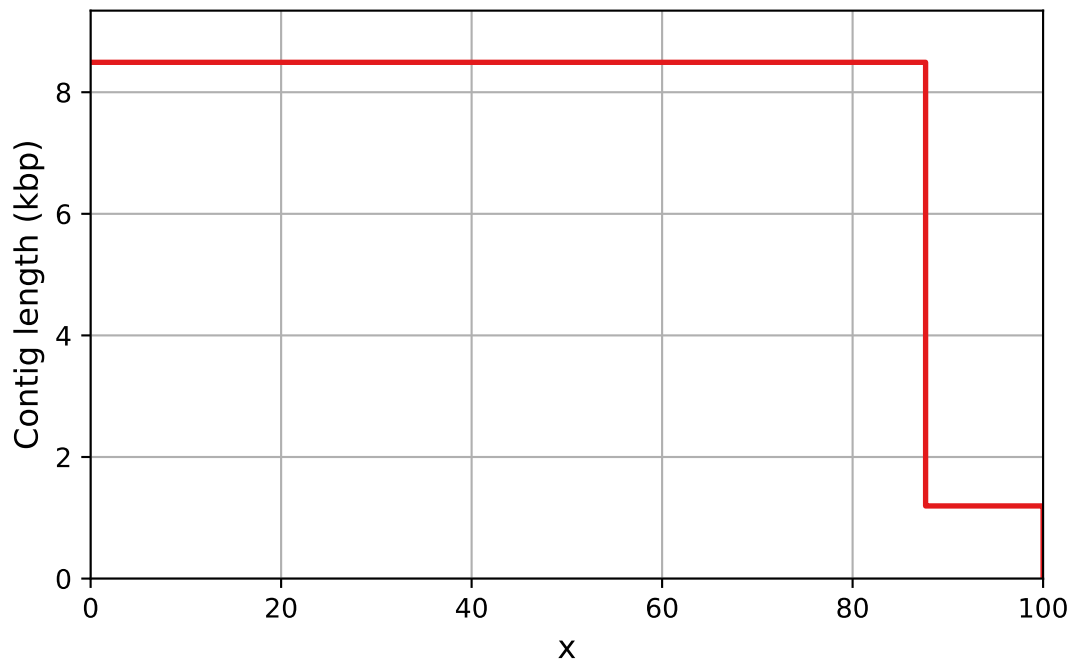


Report

	lr-ecoli.contigs
# contigs (>= 0 bp)	2
# contigs (>= 1000 bp)	2
# contigs (>= 5000 bp)	1
# contigs (>= 10000 bp)	0
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	9687
Total length (>= 1000 bp)	9687
Total length (>= 5000 bp)	8492
Total length (>= 10000 bp)	0
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	2
Largest contig	8492
Total length	9687
GC (%)	46.28
N50	8492
N90	1195
auN	7591.8
L50	1
L90	2
# N's per 100 kbp	0.00
# predicted genes (unique)	7
# predicted genes (>= 0 bp)	7 + 0 part
# predicted genes (>= 300 bp)	6 + 0 part
# predicted genes (>= 1500 bp)	1 + 0 part
# predicted genes (>= 3000 bp)	0 + 0 part

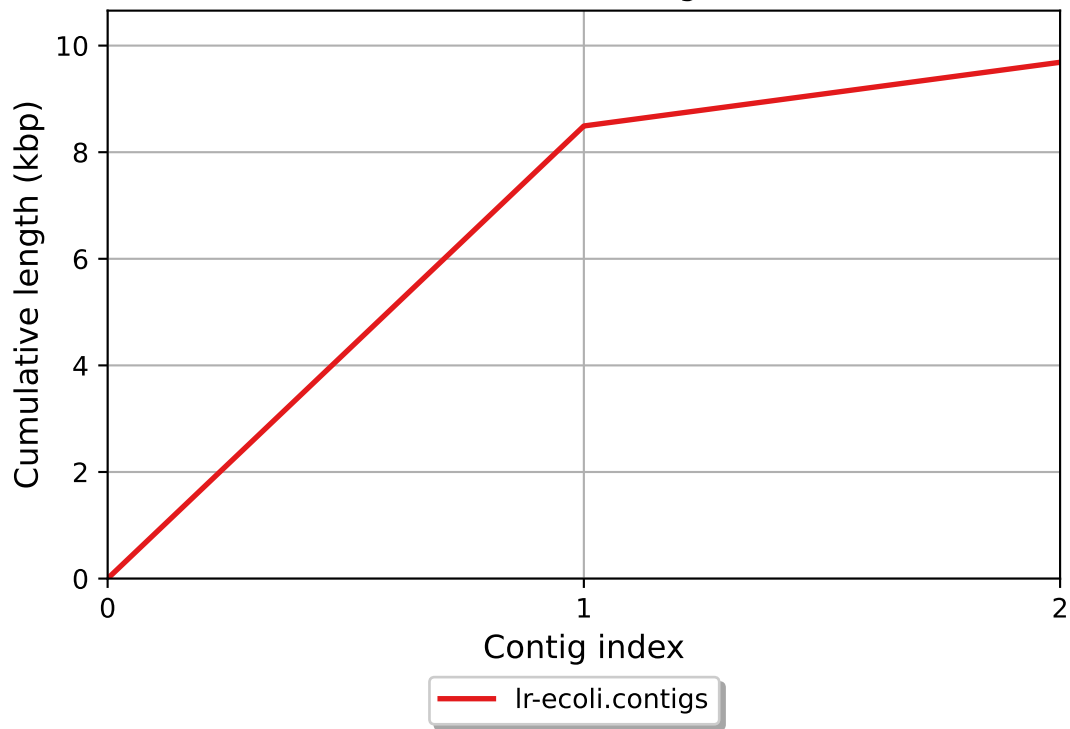
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

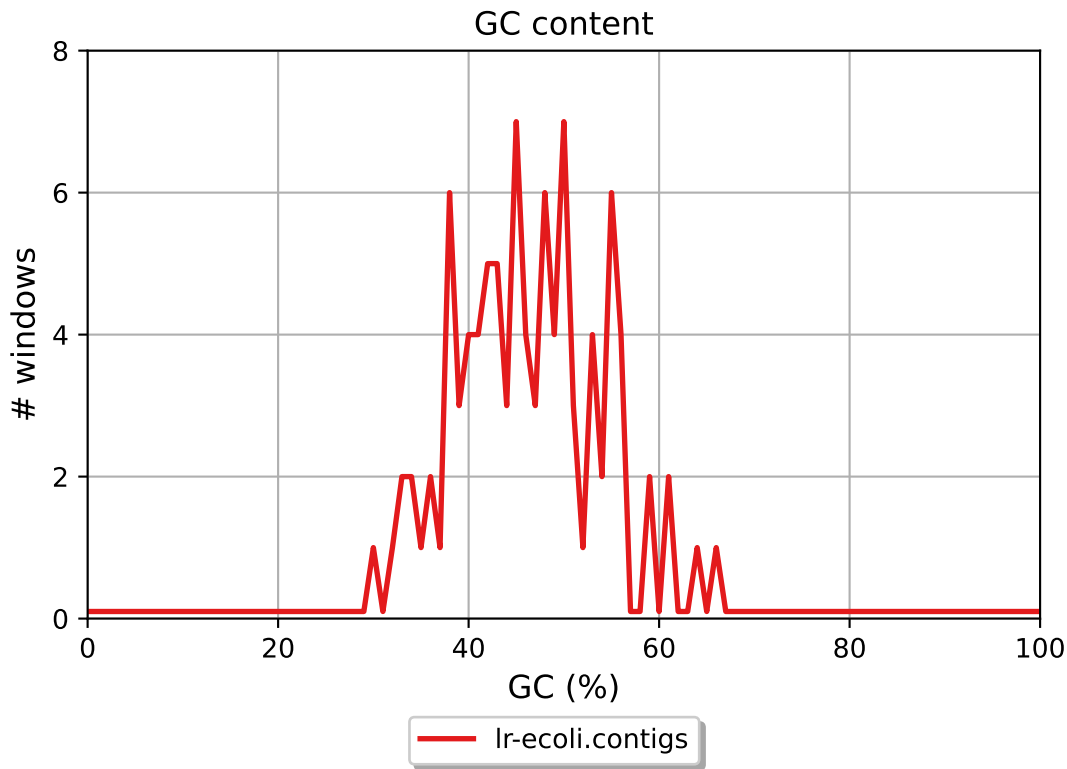
Nx



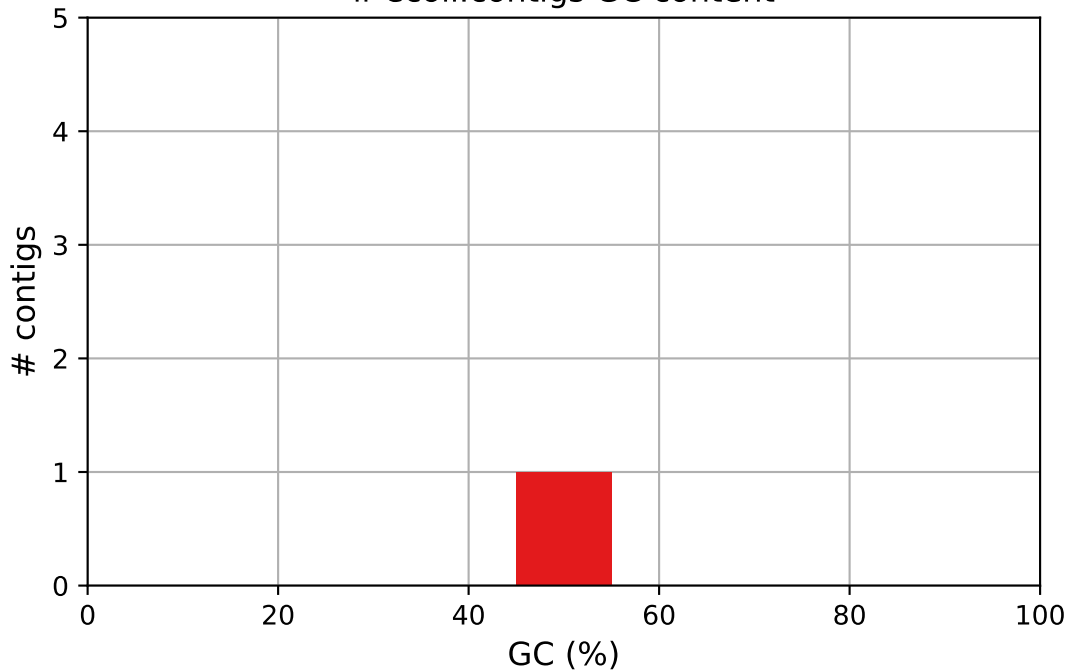
Ir-ecoli.contigs

Cumulative length





Ir-ecoli.contigs GC content



Ir-ecoli.contigs