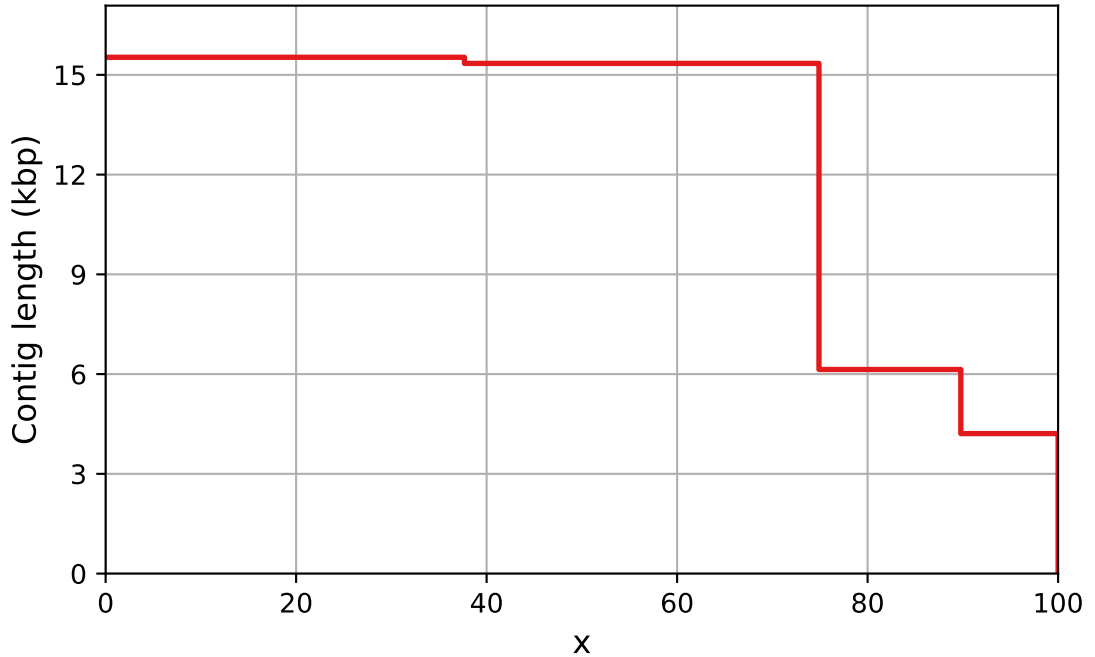


Report

	assembly_raven_lr-marine_sediment
# contigs (>= 0 bp)	4
# contigs (>= 1000 bp)	4
# contigs (>= 5000 bp)	3
# contigs (>= 10000 bp)	2
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	41222
Total length (>= 1000 bp)	41222
Total length (>= 5000 bp)	37011
Total length (>= 10000 bp)	30873
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	4
Largest contig	15529
Total length	41222
GC (%)	36.66
N50	15344
N90	4211
auN	12905.6
L50	2
L90	4
# N's per 100 kbp	0.00
# predicted genes (unique)	18
# predicted genes (>= 0 bp)	18 + 0 part
# predicted genes (>= 300 bp)	9 + 0 part
# predicted genes (>= 1500 bp)	2 + 0 part
# predicted genes (>= 3000 bp)	1 + 0 part

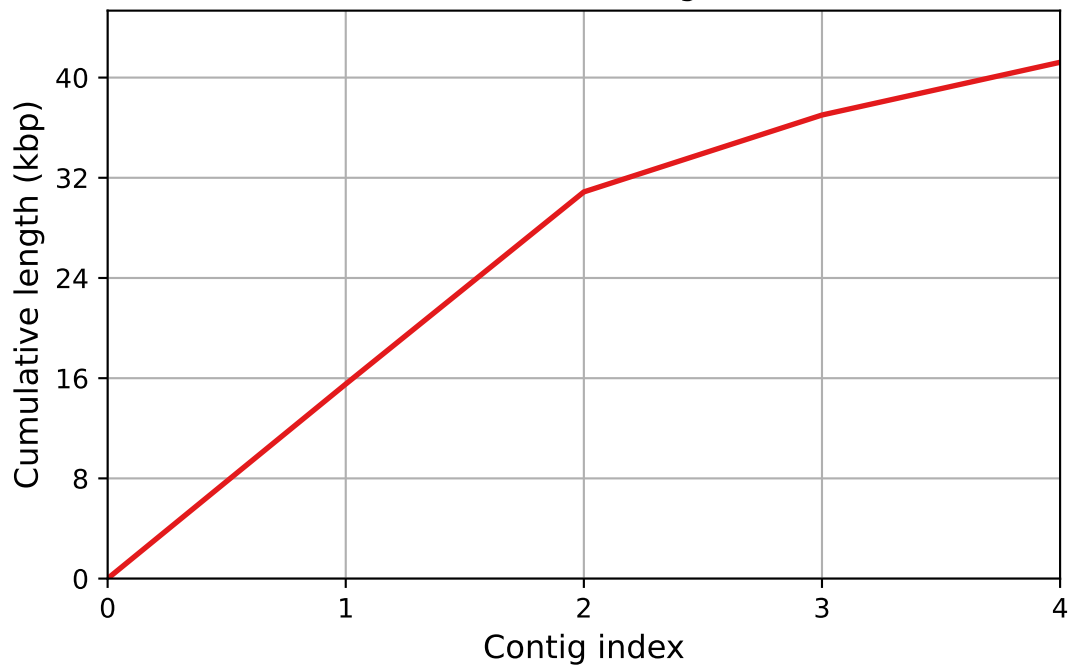
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx



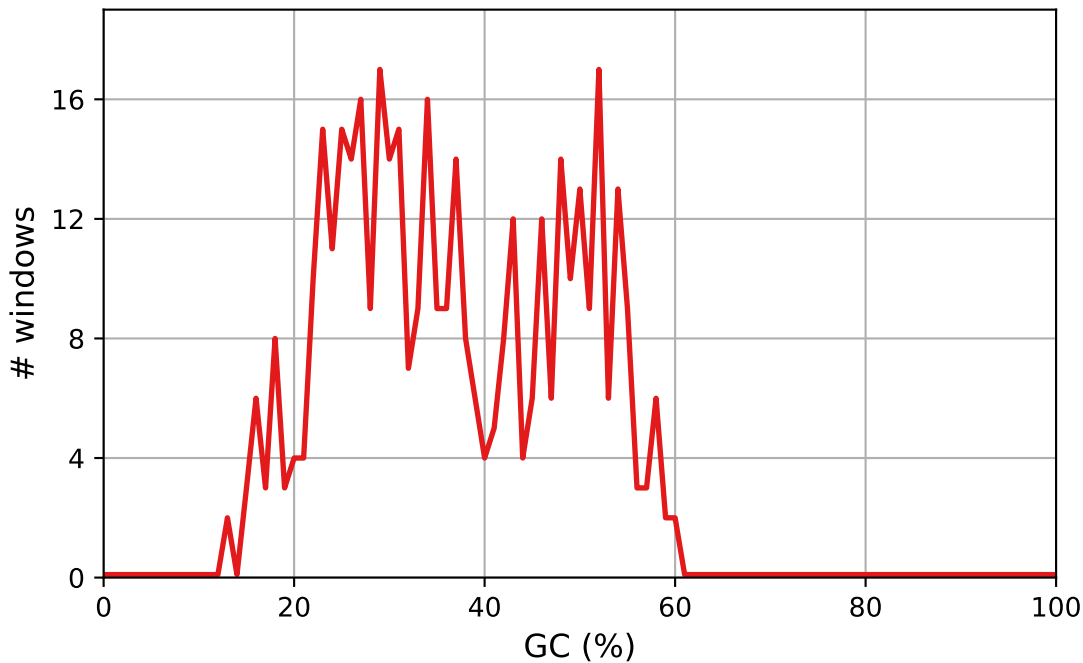
— assembly_raven_Ir-marine_sediment

Cumulative length



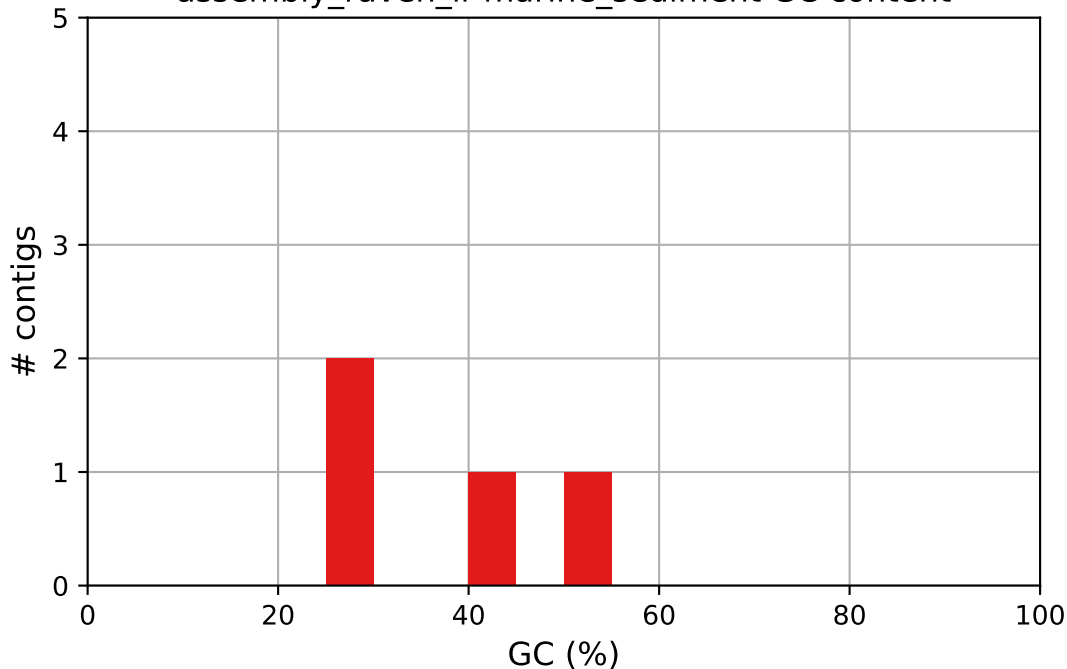
— assembly_raven_lr-marine_sediment

GC content



— assembly_raven_Ir-marine_sediment

assembly_raven_lr-marine_sediment GC content



assembly_raven_lr-marine_sediment