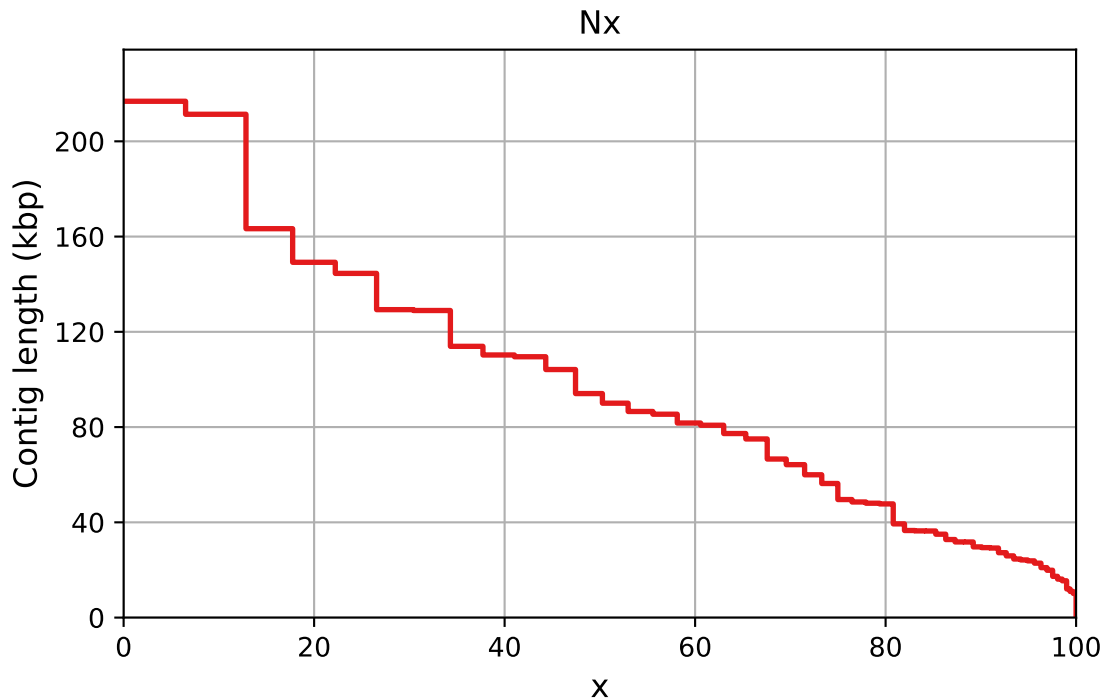


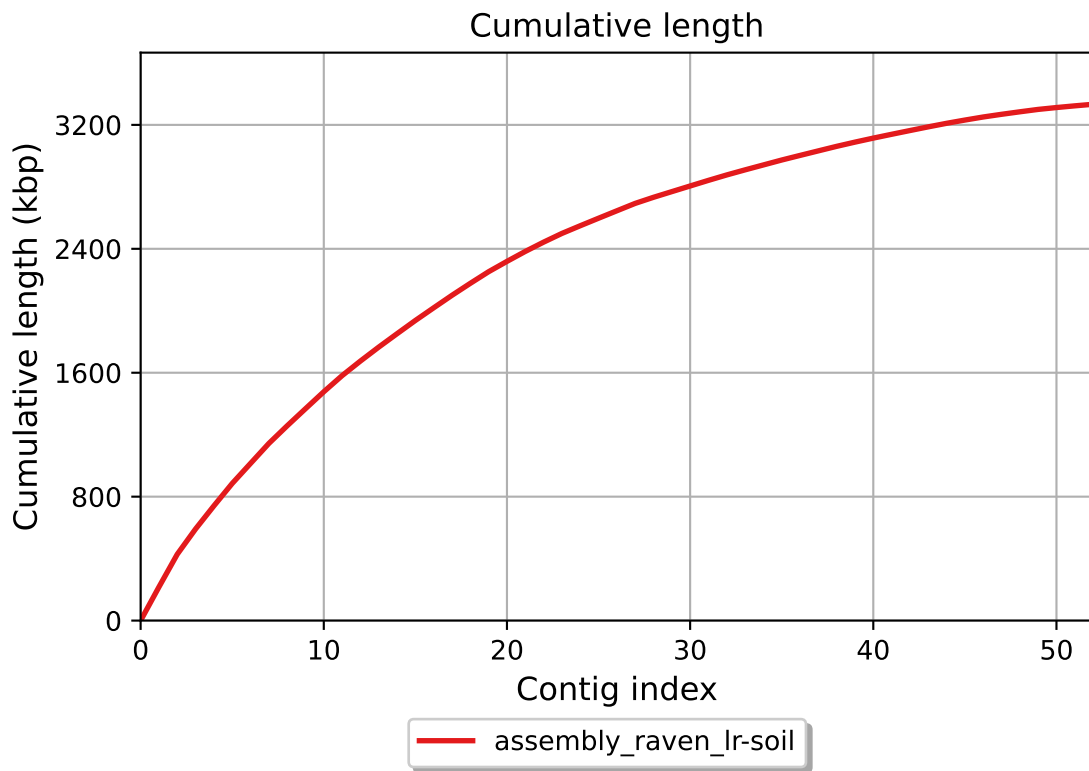
Report

	assembly_raven_lr-soil
# contigs (>= 0 bp)	52
# contigs (>= 1000 bp)	52
# contigs (>= 5000 bp)	52
# contigs (>= 10000 bp)	52
# contigs (>= 25000 bp)	40
# contigs (>= 50000 bp)	23
Total length (>= 0 bp)	3333078
Total length (>= 1000 bp)	3333078
Total length (>= 5000 bp)	3333078
Total length (>= 10000 bp)	3333078
Total length (>= 25000 bp)	3114641
Total length (>= 50000 bp)	2499308
# contigs	52
Largest contig	216819
Total length	3333078
GC (%)	66.23
N50	94074
N90	29693
auN	102684.2
L50	12
L90	36
# N's per 100 kbp	0.00
# predicted genes (unique)	3978
# predicted genes (>= 0 bp)	3960 + 19 part
# predicted genes (>= 300 bp)	3324 + 17 part
# predicted genes (>= 1500 bp)	206 + 1 part
# predicted genes (>= 3000 bp)	4 + 0 part

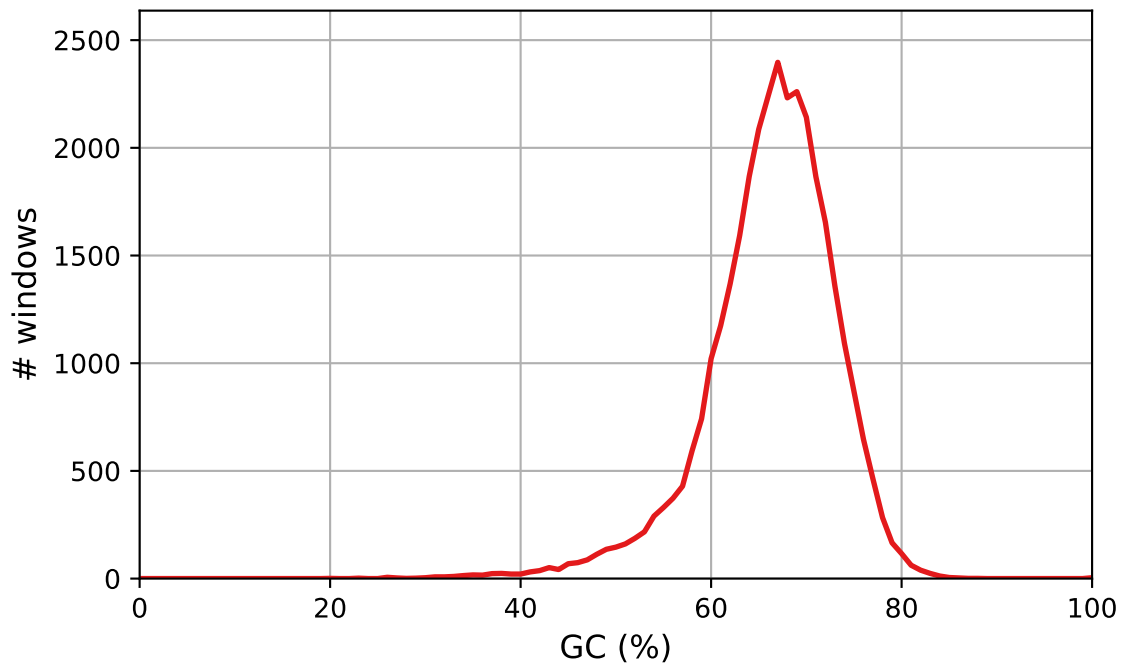
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



— assembly_raven_lr-soil

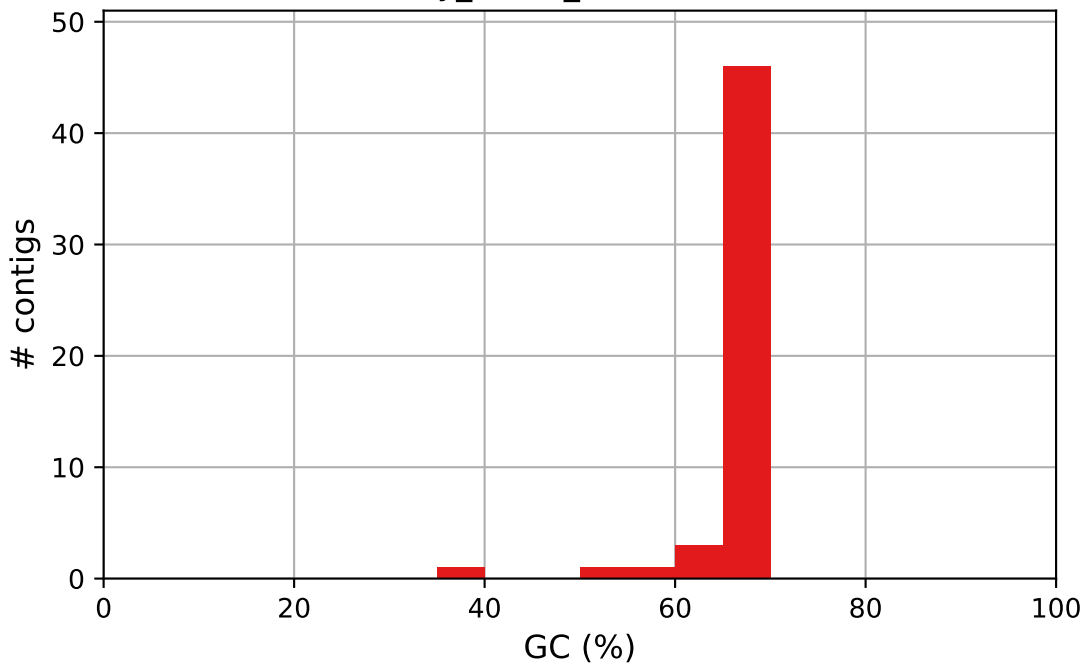


GC content



assembly_raven_lr-soil

assembly_raven_lr-soil GC content



assembly_raven_lr-soil