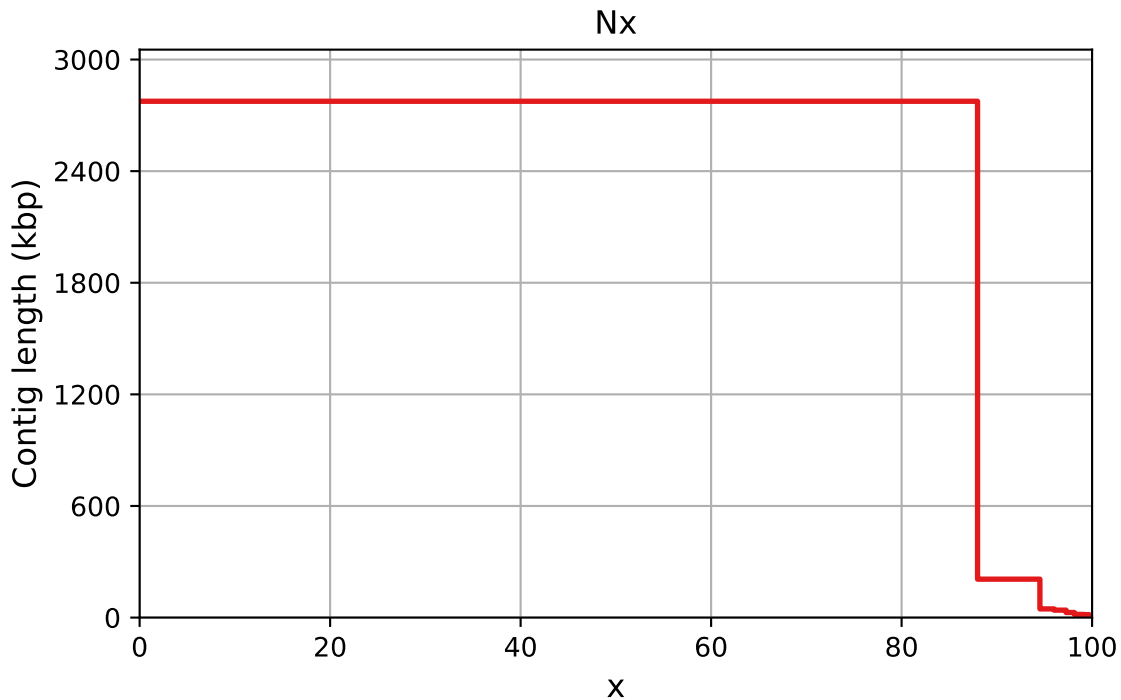


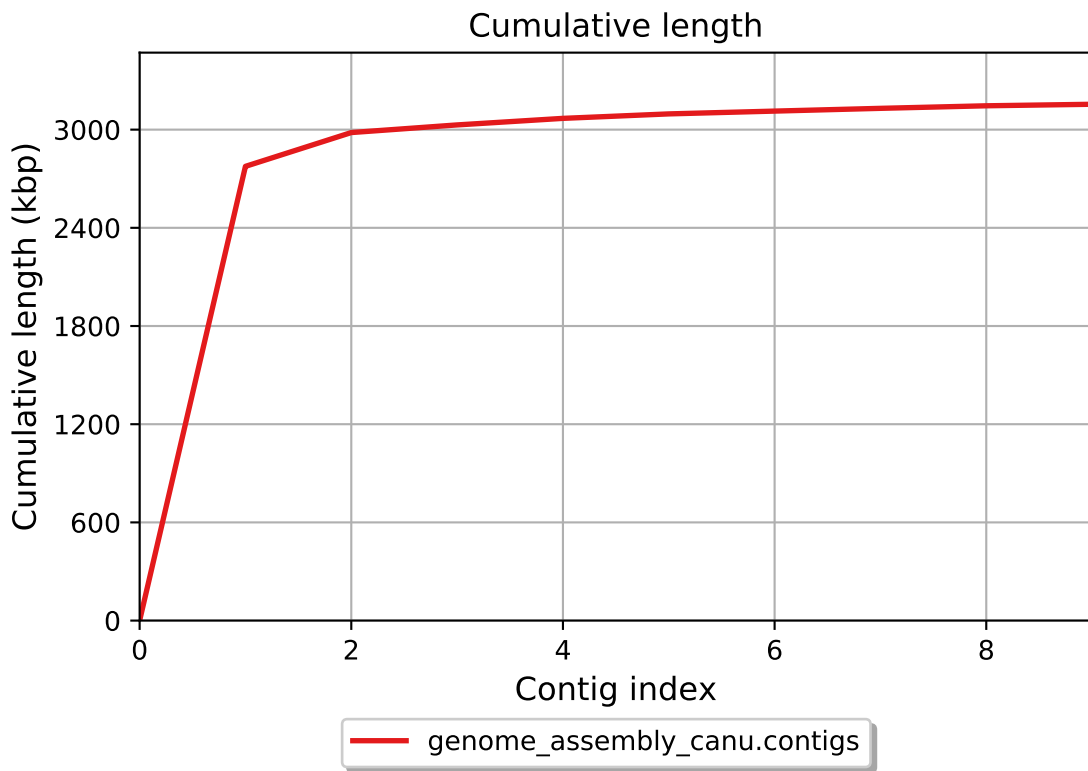
Report

	genome_assembly_canu.contigs
# contigs (>= 0 bp)	9
# contigs (>= 1000 bp)	9
# contigs (>= 5000 bp)	9
# contigs (>= 10000 bp)	8
# contigs (>= 25000 bp)	5
# contigs (>= 50000 bp)	2
Total length (>= 0 bp)	3154995
Total length (>= 1000 bp)	3154995
Total length (>= 5000 bp)	3154995
Total length (>= 10000 bp)	3145708
Total length (>= 25000 bp)	3096150
Total length (>= 50000 bp)	2982128
# contigs	9
Largest contig	2775620
Total length	3154995
GC (%)	37.75
N50	2775620
N75	2775620
L50	1
L75	1
# N's per 100 kbp	0.00

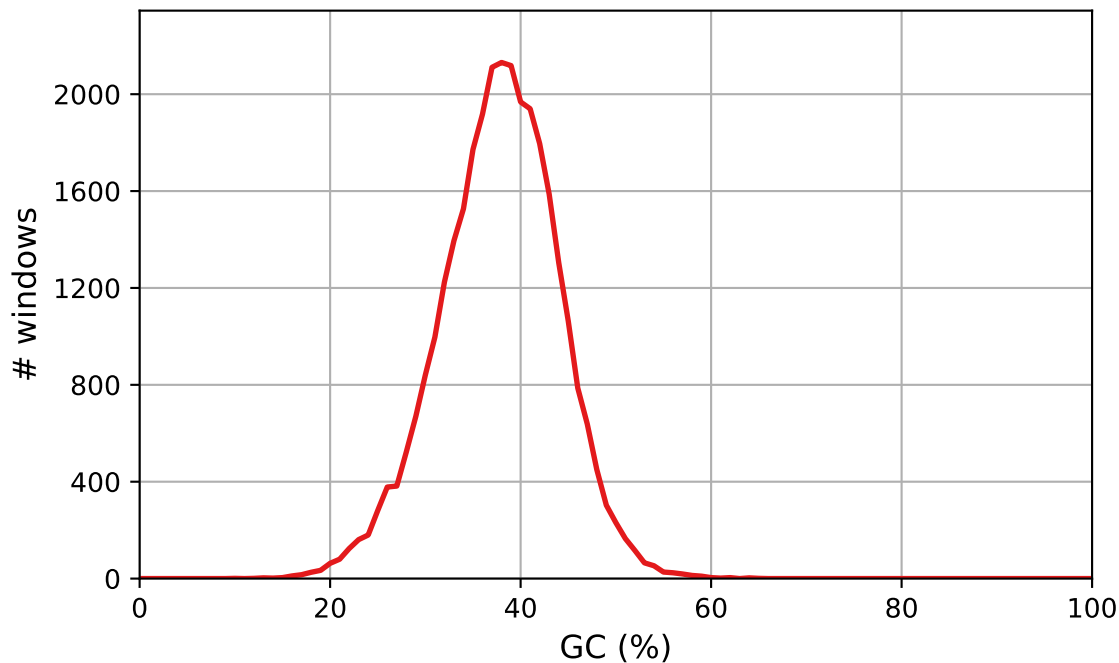
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



— genome_assembly_canu.contigs

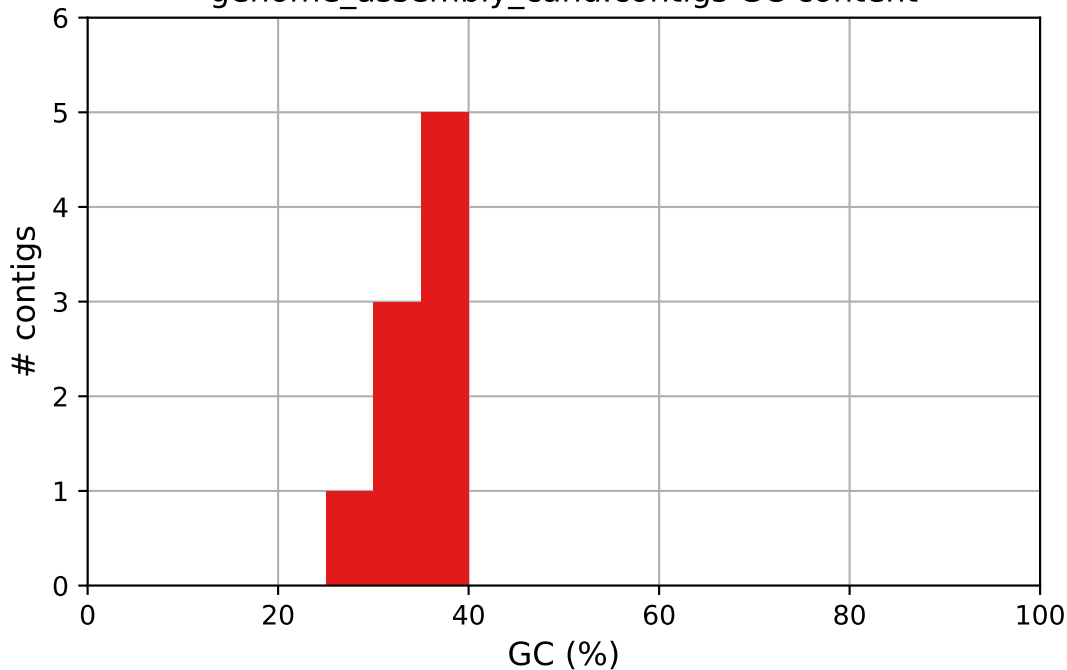


GC content



— genome_assembly_canu.contigs

genome_assembly_canu.contigs GC content



genome_assembly_canu.contigs