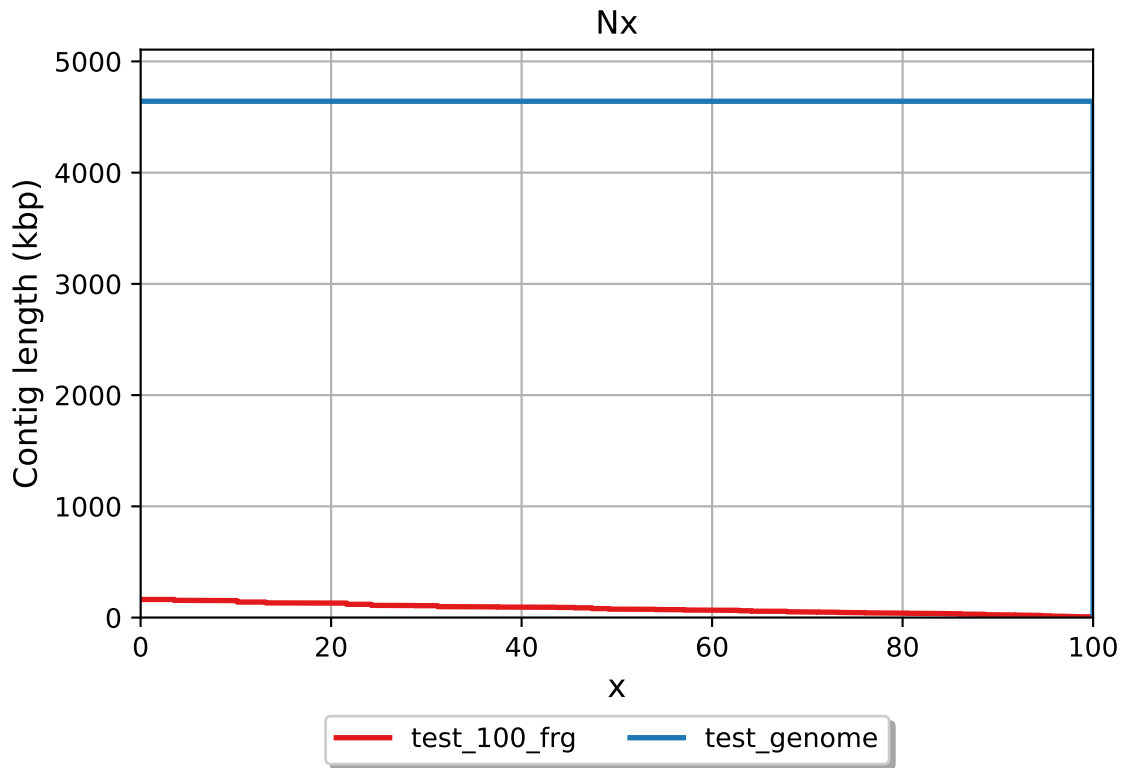
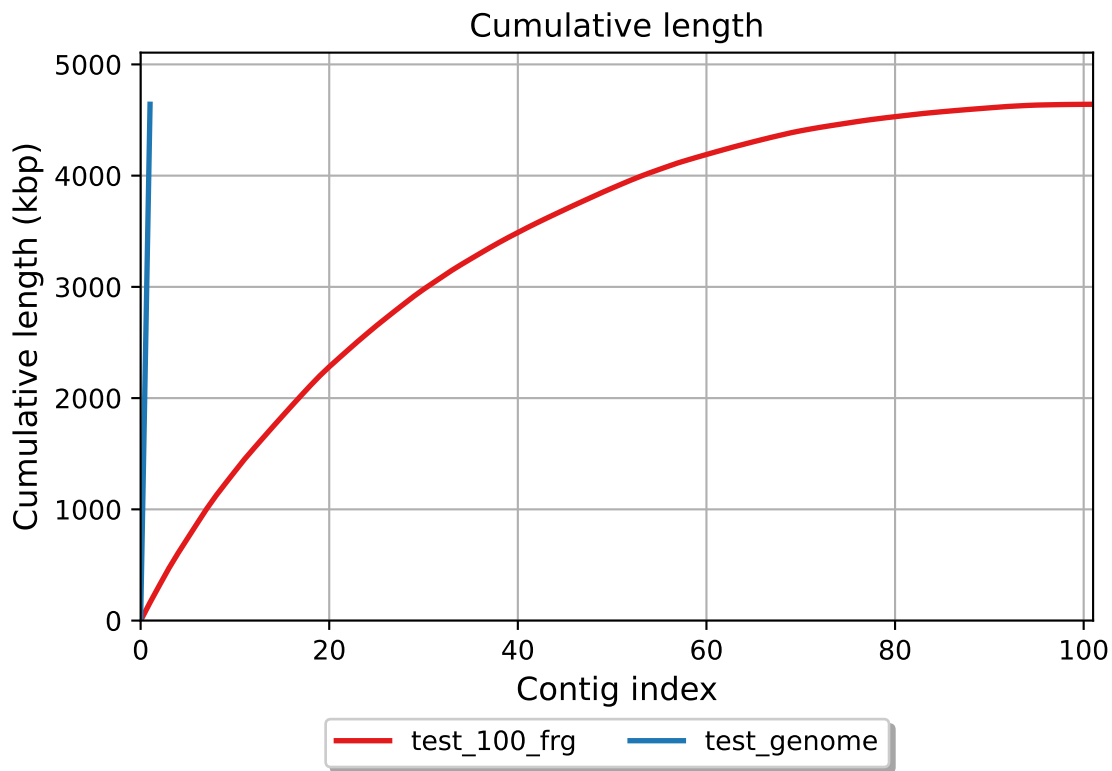


# Report

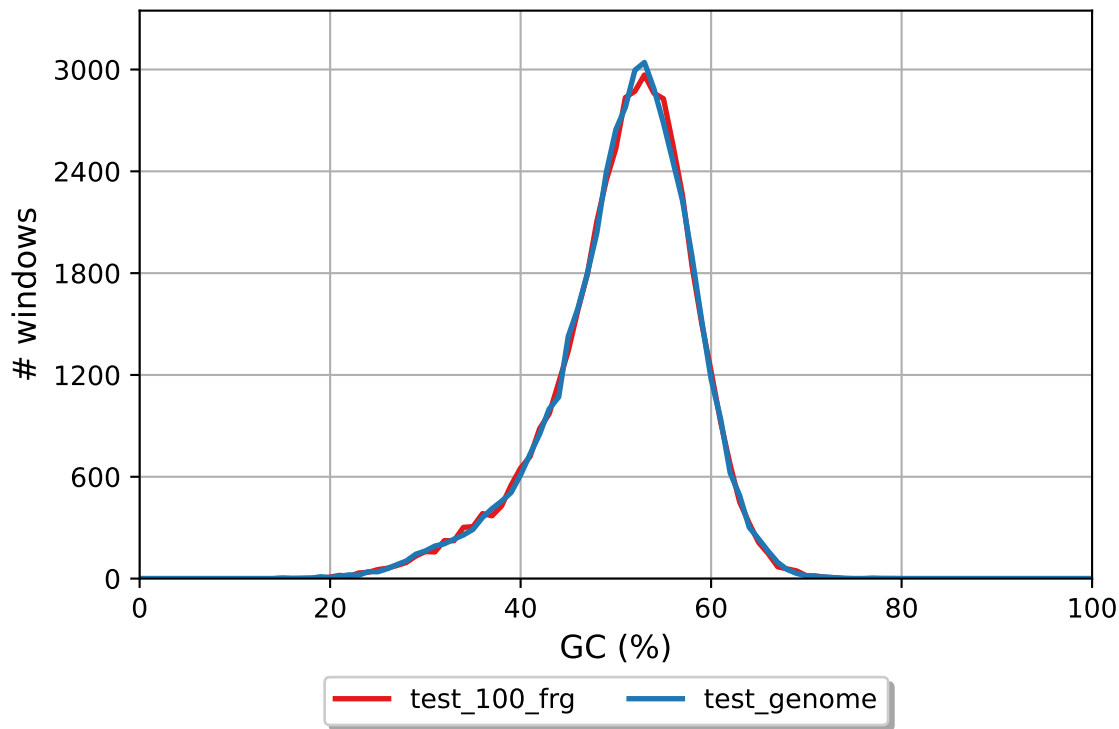
	test_100_frg	test_genome
# contigs (>= 0 bp)	1	1
# contigs (>= 1000 bp)	1	1
# contigs (>= 5000 bp)	1	1
# contigs (>= 10000 bp)	1	1
# contigs (>= 25000 bp)	1	1
# contigs (>= 50000 bp)	1	1
Total length (>= 0 bp)	74586	4641652
Total length (>= 1000 bp)	74586	4641652
Total length (>= 5000 bp)	74586	4641652
Total length (>= 10000 bp)	74586	4641652
Total length (>= 25000 bp)	74586	4641652
Total length (>= 50000 bp)	74586	4641652
# contigs	101	1
Largest contig	162841	4641652
Total length	4641652	4641652
GC (%)	50.79	50.79
N50	76028	4641652
N90	24233	4641652
auN	81690.5	4641652.0
L50	21	1
L90	60	1
# N's per 100 kbp	0.00	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).

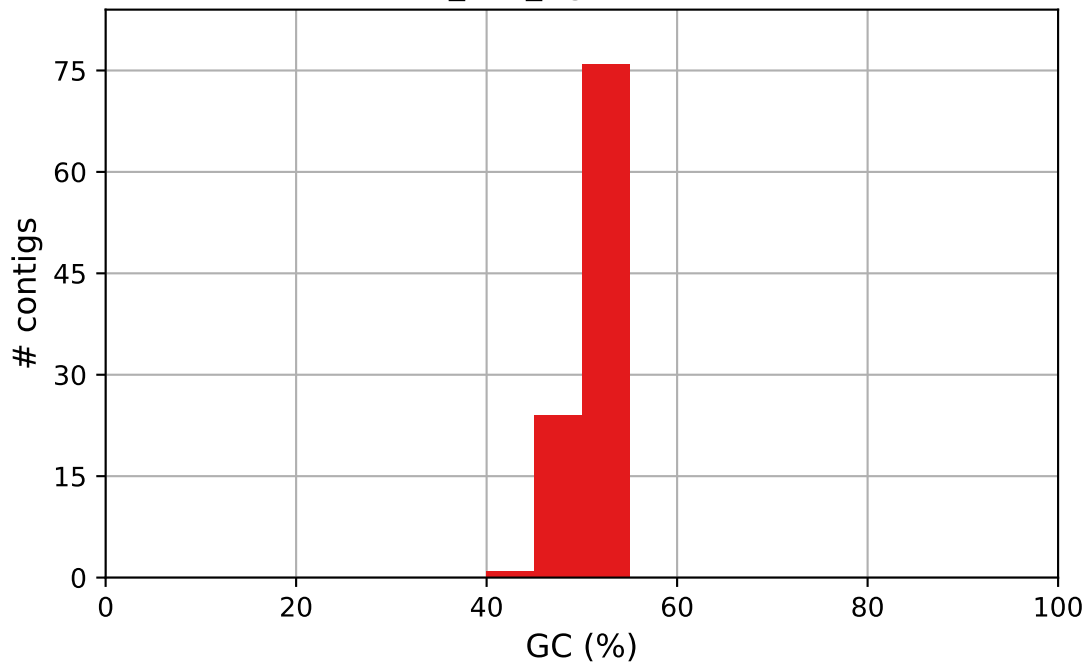




GC content

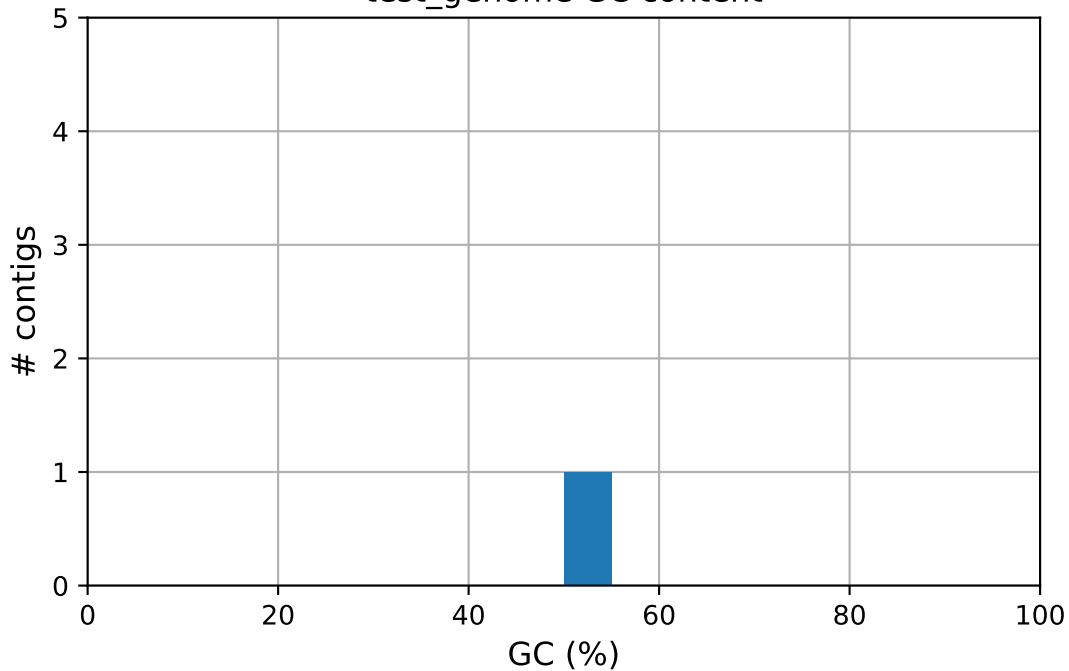


test\_100\_frg GC content



test\_100\_frg

test\_genome GC content



test\_genome