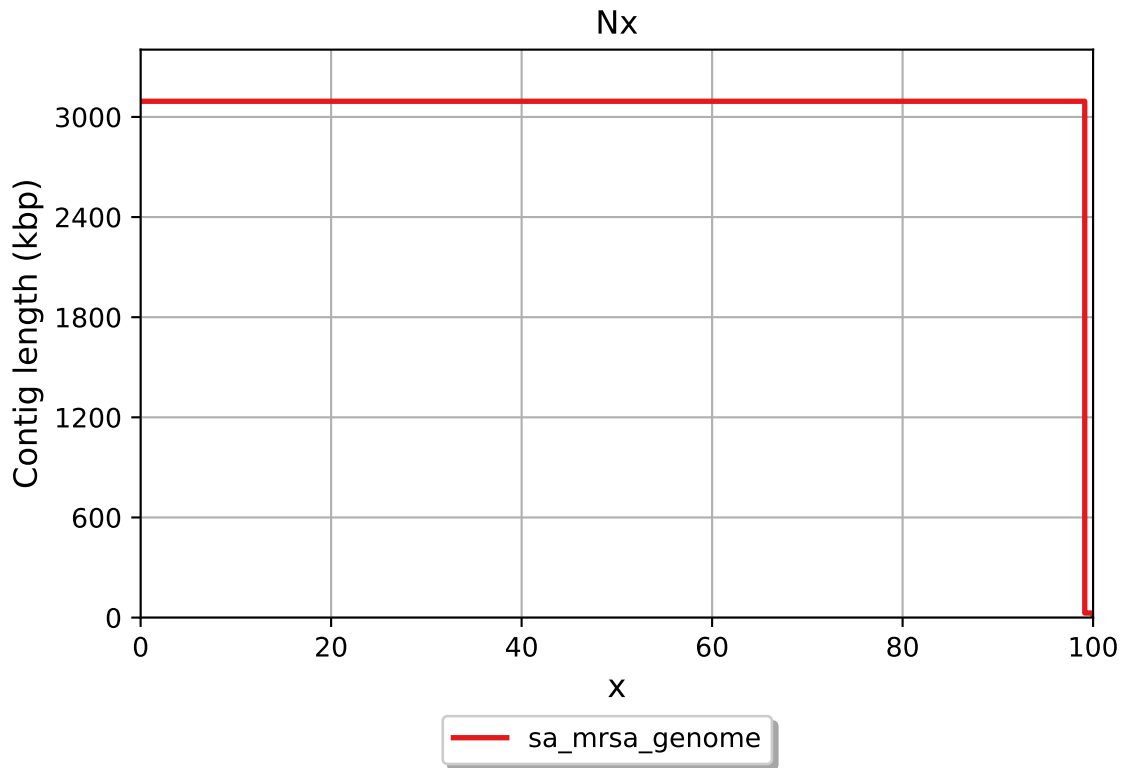


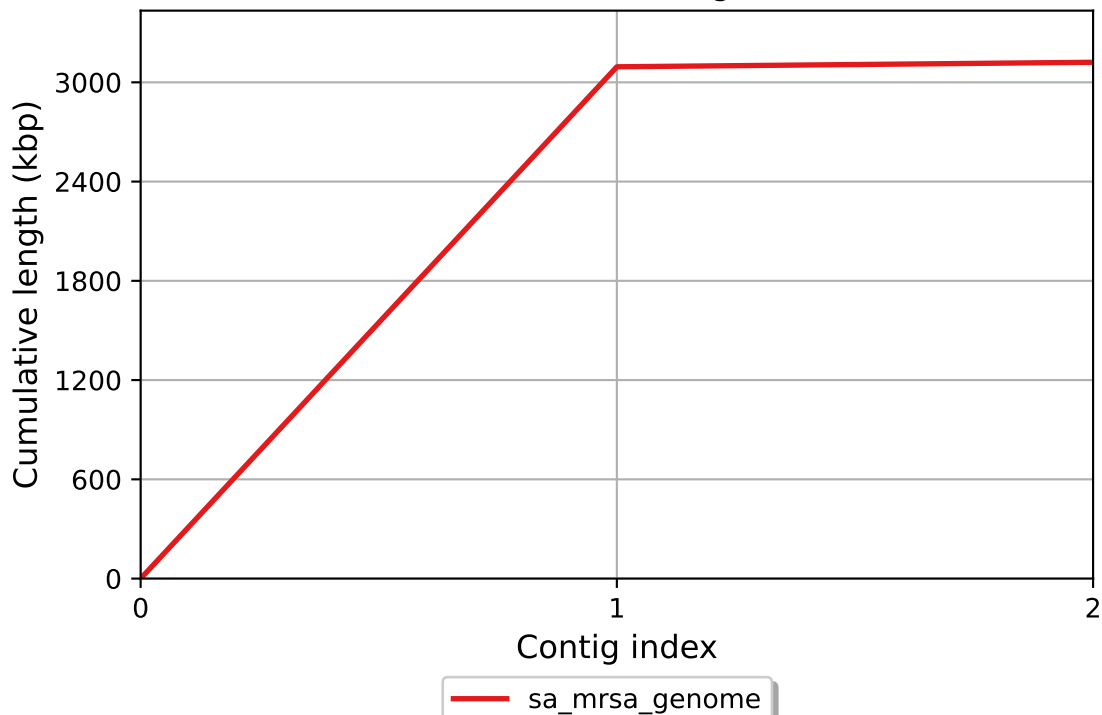
Report

	sa_mrsa_genome
# contigs (>= 0 bp)	2
# contigs (>= 1000 bp)	2
# contigs (>= 5000 bp)	2
# contigs (>= 10000 bp)	2
# contigs (>= 25000 bp)	2
# contigs (>= 50000 bp)	1
Total length (>= 0 bp)	3121265
Total length (>= 1000 bp)	3121265
Total length (>= 5000 bp)	3121265
Total length (>= 10000 bp)	3121265
Total length (>= 25000 bp)	3121265
Total length (>= 50000 bp)	3093943
# contigs	2
Largest contig	3093943
Total length	3121265
GC (%)	32.78
N50	3093943
N90	3093943
auN	3067099.3
L50	1
L90	1
# N's per 100 kbp	0.00

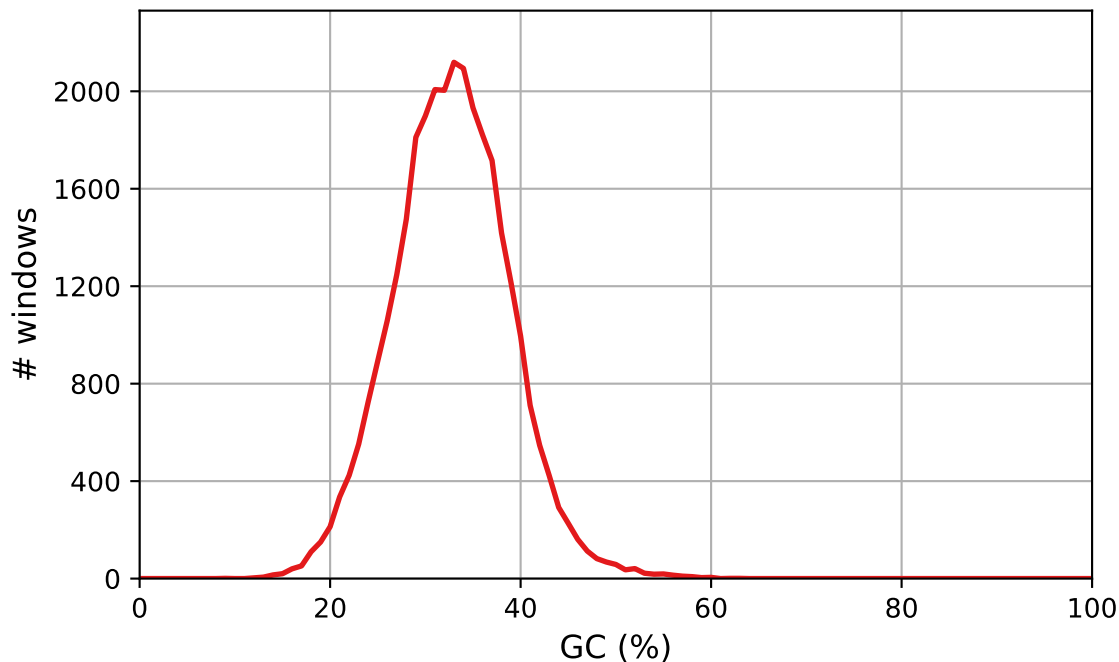
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length

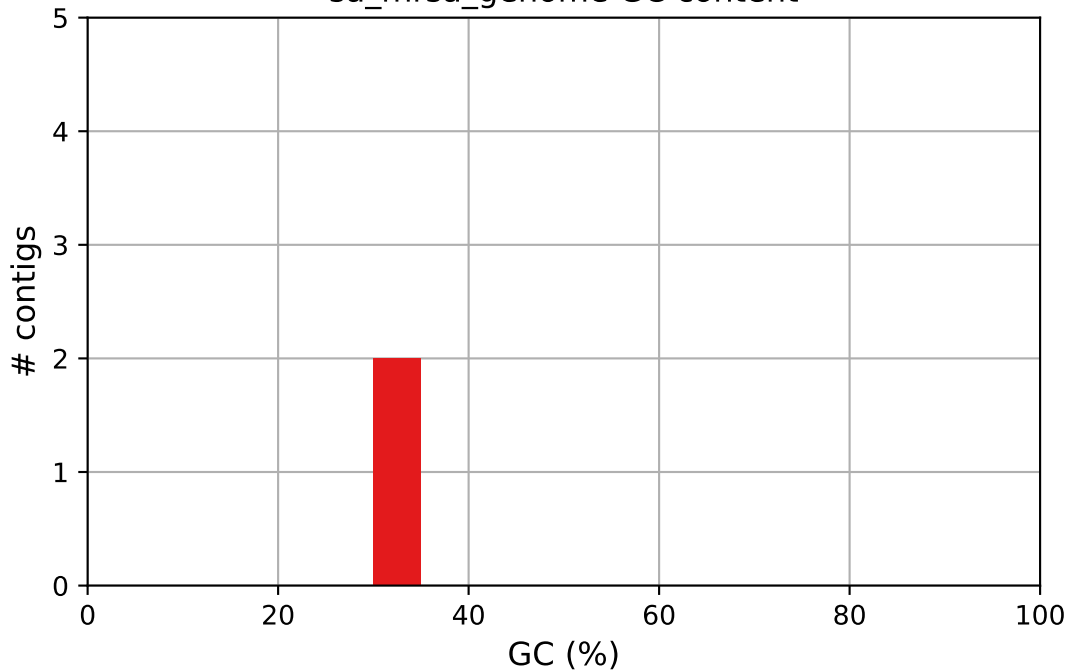


GC content



sa_mrsa_genome

sa_mrsa_genome GC content



sa_mrsa_genome