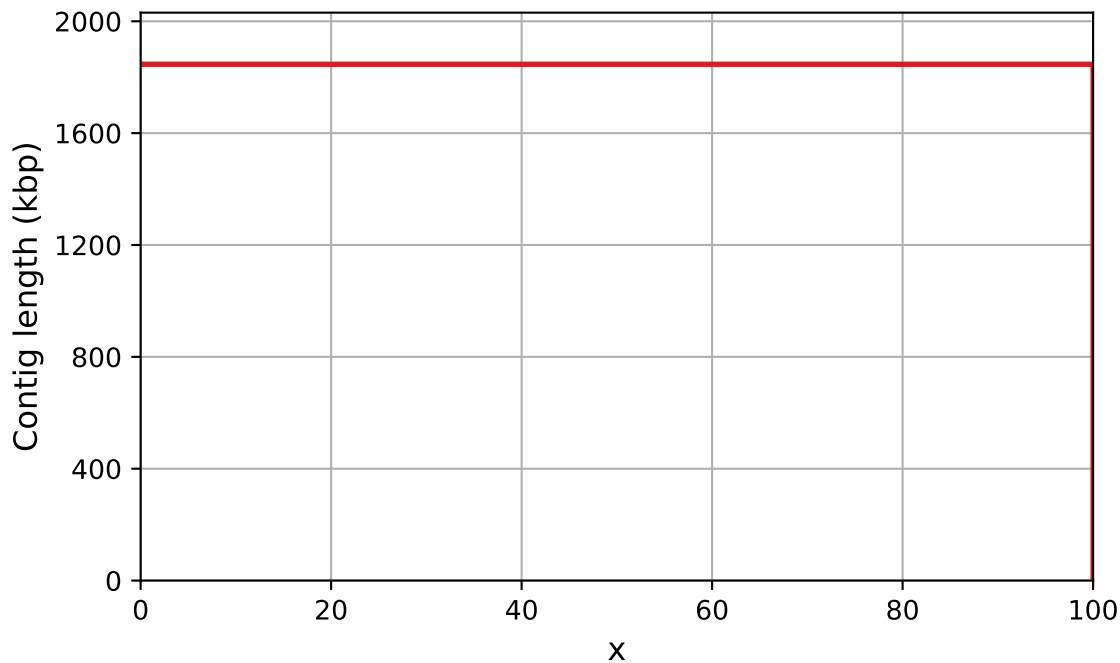


# Report

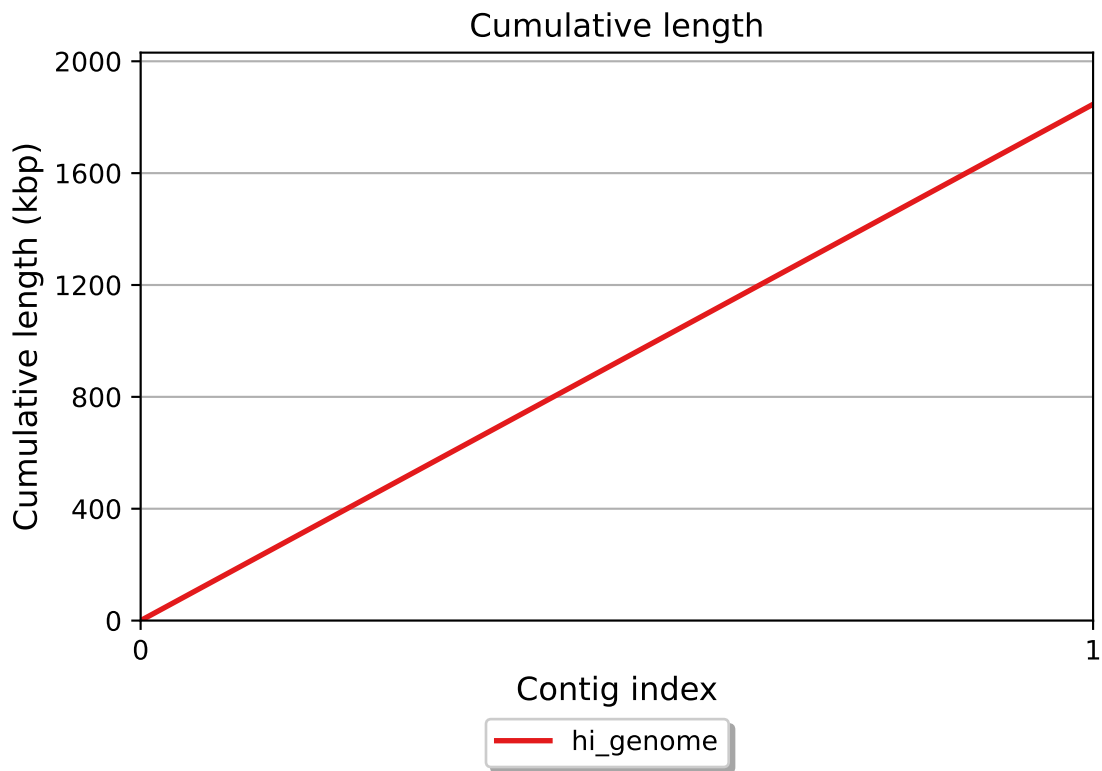
	hi_genome
# contigs ( $\geq 0$ bp)	1
# contigs ( $\geq 1000$ bp)	1
# contigs ( $\geq 5000$ bp)	1
# contigs ( $\geq 10000$ bp)	1
# contigs ( $\geq 25000$ bp)	1
# contigs ( $\geq 50000$ bp)	1
Total length ( $\geq 0$ bp)	1846259
Total length ( $\geq 1000$ bp)	1846259
Total length ( $\geq 5000$ bp)	1846259
Total length ( $\geq 10000$ bp)	1846259
Total length ( $\geq 25000$ bp)	1846259
Total length ( $\geq 50000$ bp)	1846259
# contigs	1
Largest contig	1846259
Total length	1846259
GC (%)	38.16
N50	1846259
N90	1846259
auN	1846259.0
L50	1
L90	1
# N's per 100 kbp	0.00

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

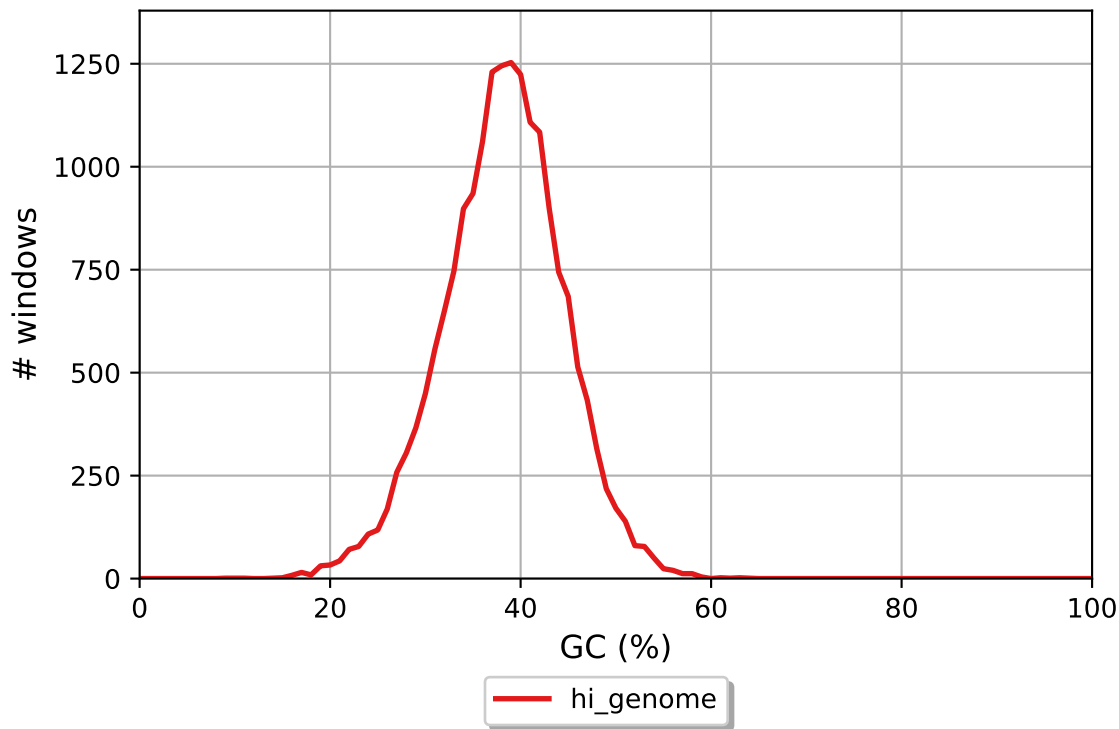
Nx



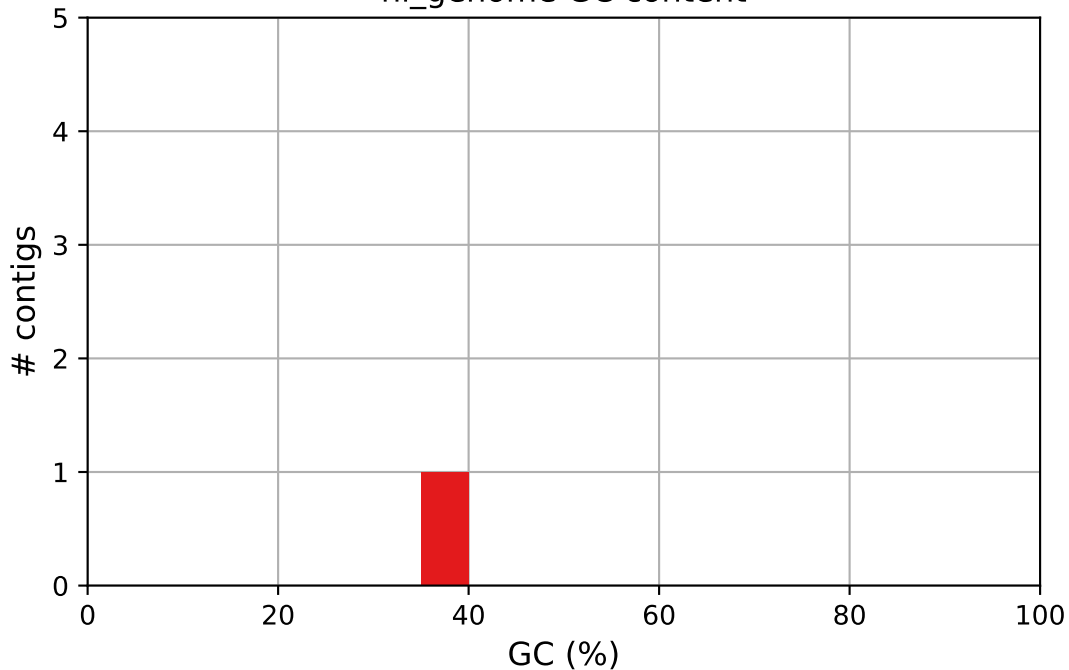
— hi\_genome



GC content



hi\_genome GC content



hi\_genome