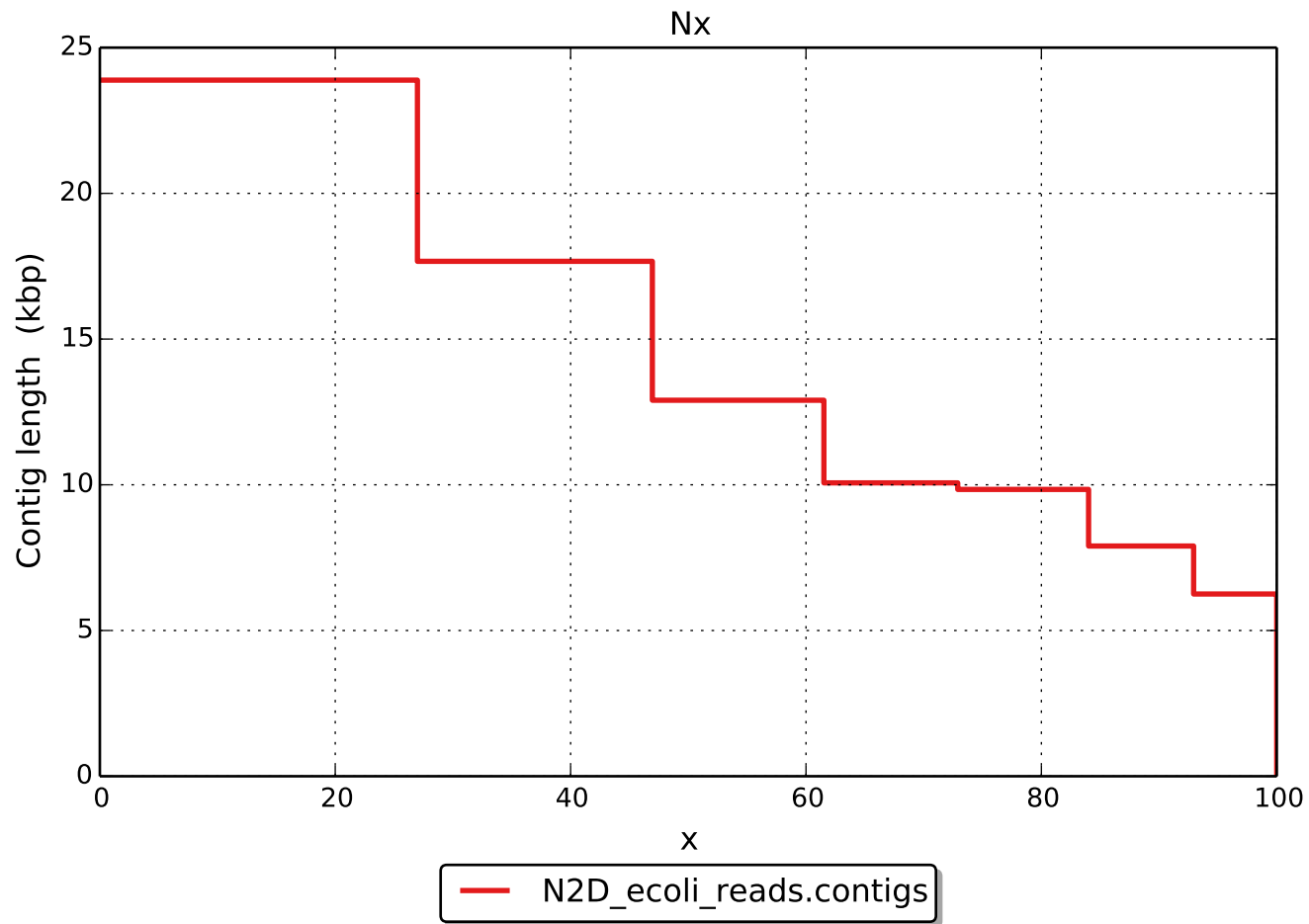
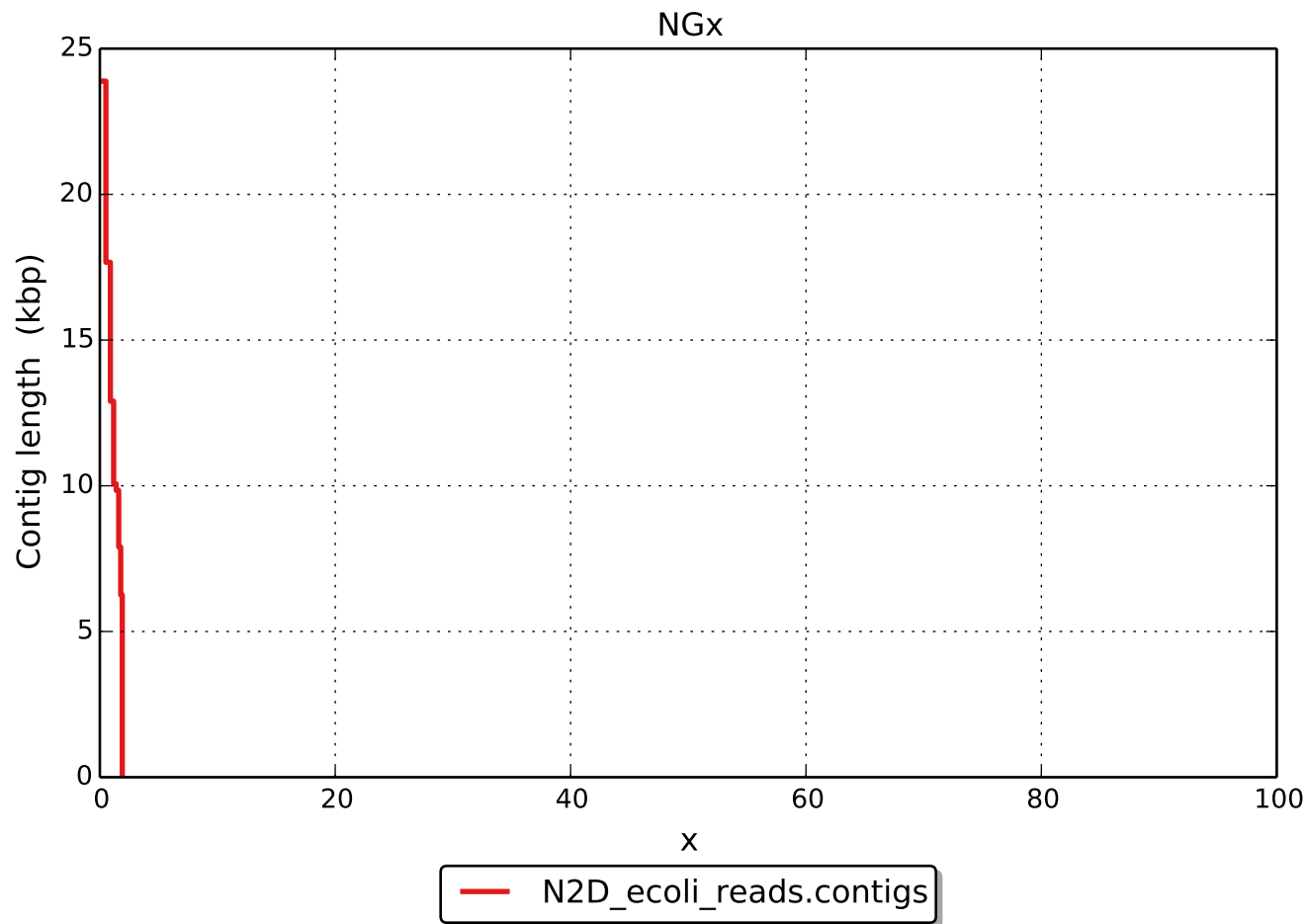


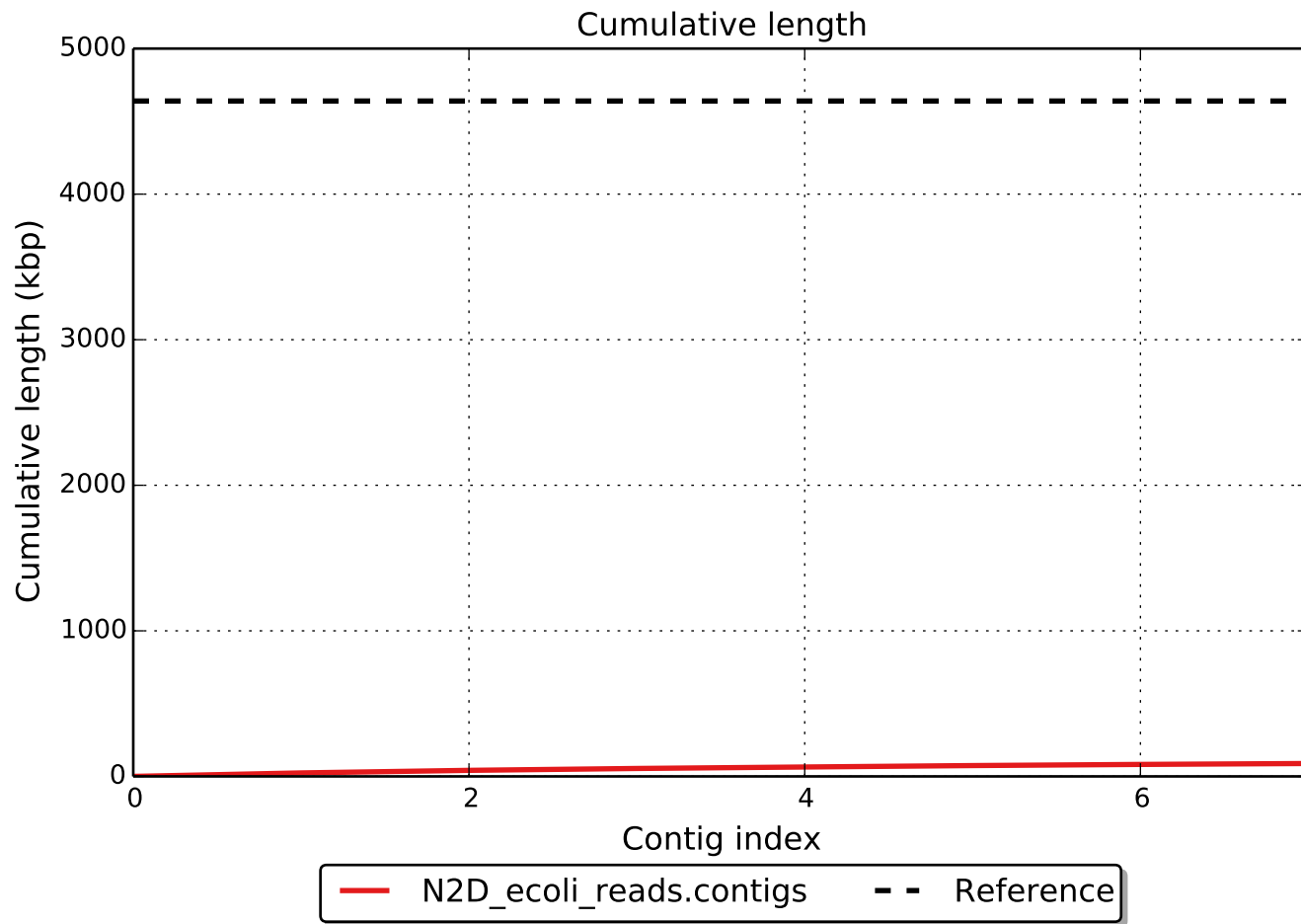
## Report

	N2D_ecoli_reads.contigs
# contigs ( $\geq 0$ bp)	7
# contigs ( $\geq 1000$ bp)	7
# contigs ( $\geq 5000$ bp)	7
# contigs ( $\geq 10000$ bp)	4
# contigs ( $\geq 25000$ bp)	0
# contigs ( $\geq 50000$ bp)	0
Total length ( $\geq 0$ bp)	88513
Total length ( $\geq 1000$ bp)	88513
Total length ( $\geq 5000$ bp)	88513
Total length ( $\geq 10000$ bp)	64520
Total length ( $\geq 25000$ bp)	0
Total length ( $\geq 50000$ bp)	0
# contigs	7
Largest contig	23885
Total length	88513
Reference length	4639675
GC (%)	47.43
Reference GC (%)	50.79
N50	12901
N75	9841
L50	3
L75	5
# unaligned contigs	7 + 0 part
Unaligned length	88513
# N's per 100 kbp	0.00
NGA50	-

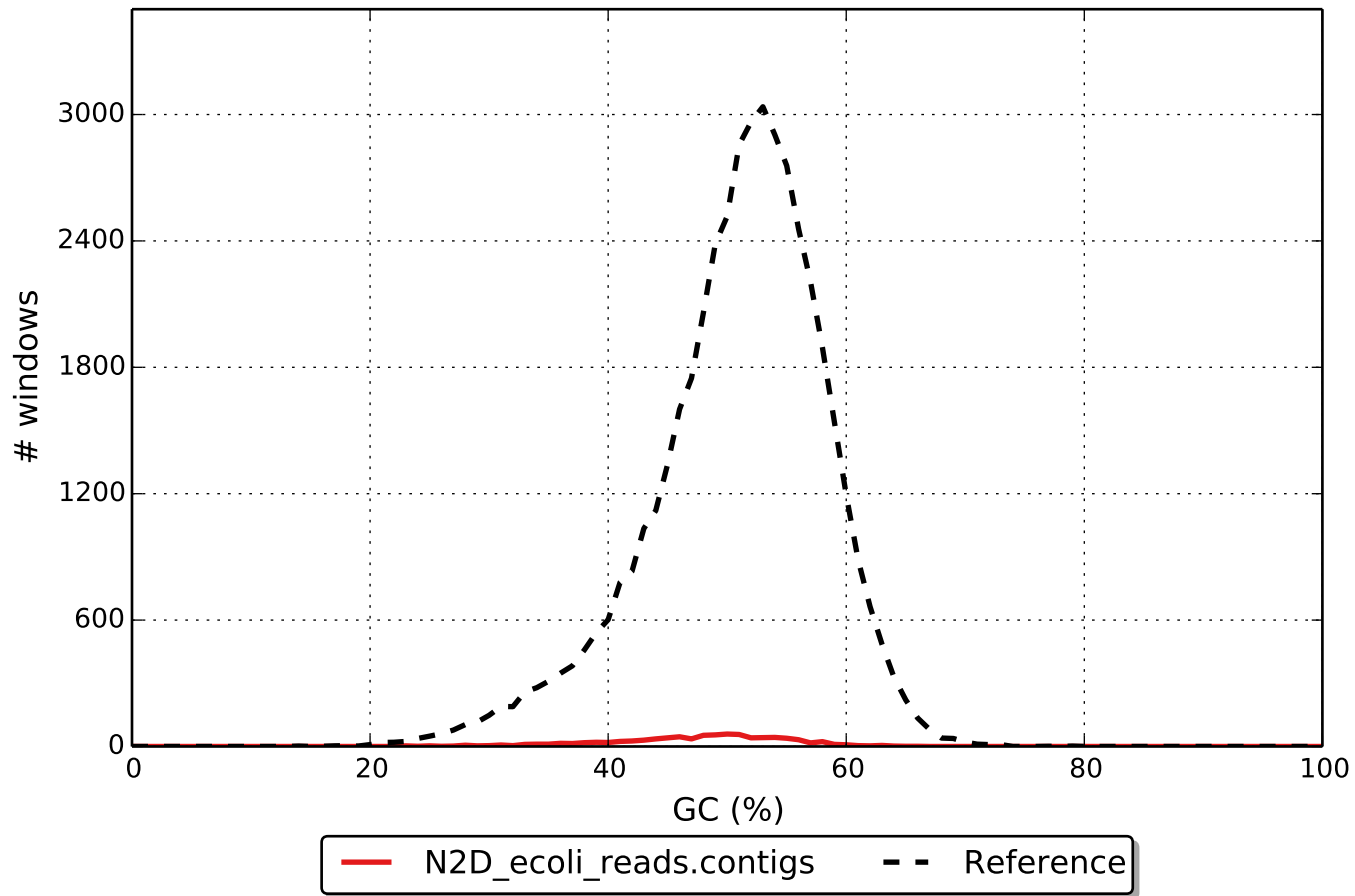
All statistics are based on contigs of size  $\geq 500$  bp, unless otherwise noted (e.g., "# contigs ( $\geq 0$  bp)" and "Total length ( $\geq 0$  bp)" include all contigs).







# GC content



# Misassemblies

