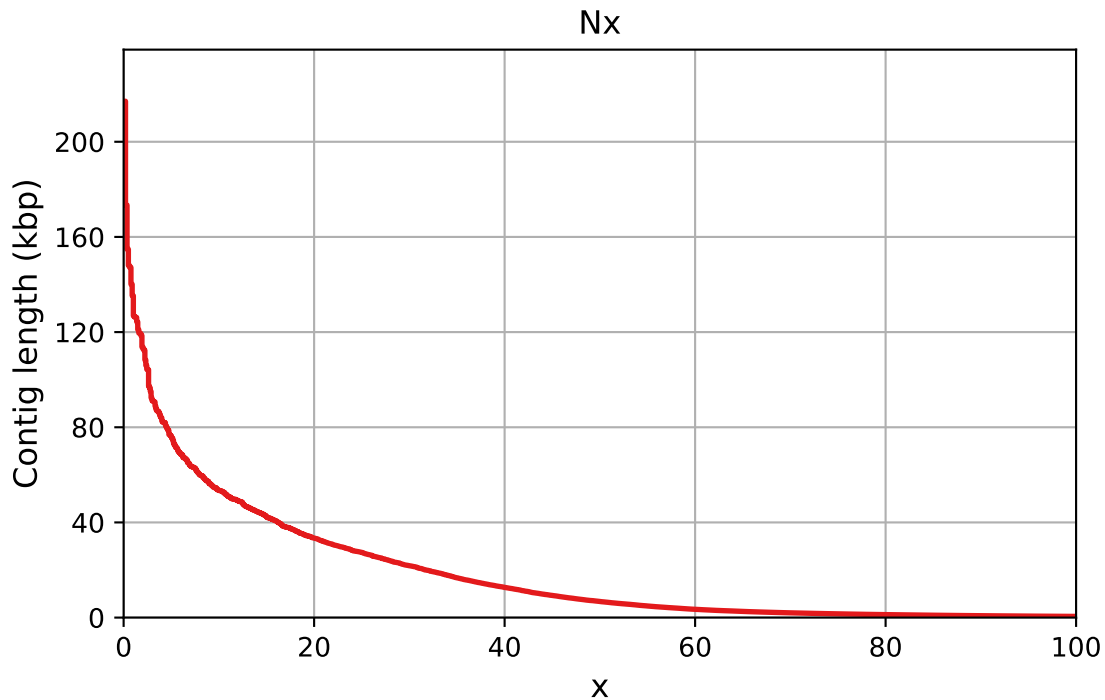


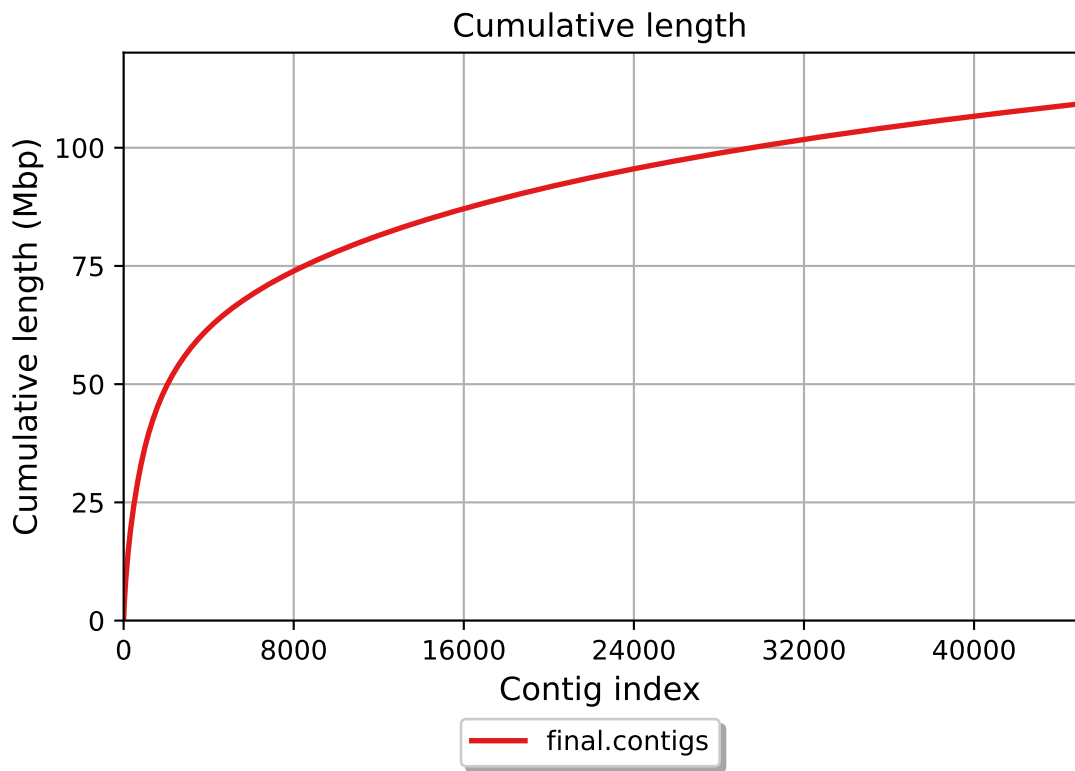
# Report

	final.contigs
# contigs ( $\geq 0$ bp)	62587
# contigs ( $\geq 1000$ bp)	21011
# contigs ( $\geq 5000$ bp)	3535
# contigs ( $\geq 10000$ bp)	1834
# contigs ( $\geq 25000$ bp)	669
# contigs ( $\geq 50000$ bp)	169
Total length ( $\geq 0$ bp)	115981271
Total length ( $\geq 1000$ bp)	92727758
Total length ( $\geq 5000$ bp)	59668293
Total length ( $\geq 10000$ bp)	47818131
Total length ( $\geq 25000$ bp)	29564231
Total length ( $\geq 50000$ bp)	12411536
# contigs	44799
Largest contig	217021
Total length	109191644
GC (%)	43.35
N50	6772
N90	791
auN	19447.6
L50	2663
L90	27257
# N's per 100 kbp	0.00
# predicted genes (unique)	75591
# predicted genes ( $\geq 0$ bp)	69452 + 6157 part
# predicted genes ( $\geq 300$ bp)	62140 + 5763 part
# predicted genes ( $\geq 1500$ bp)	12059 + 428 part
# predicted genes ( $\geq 3000$ bp)	2454 + 64 part

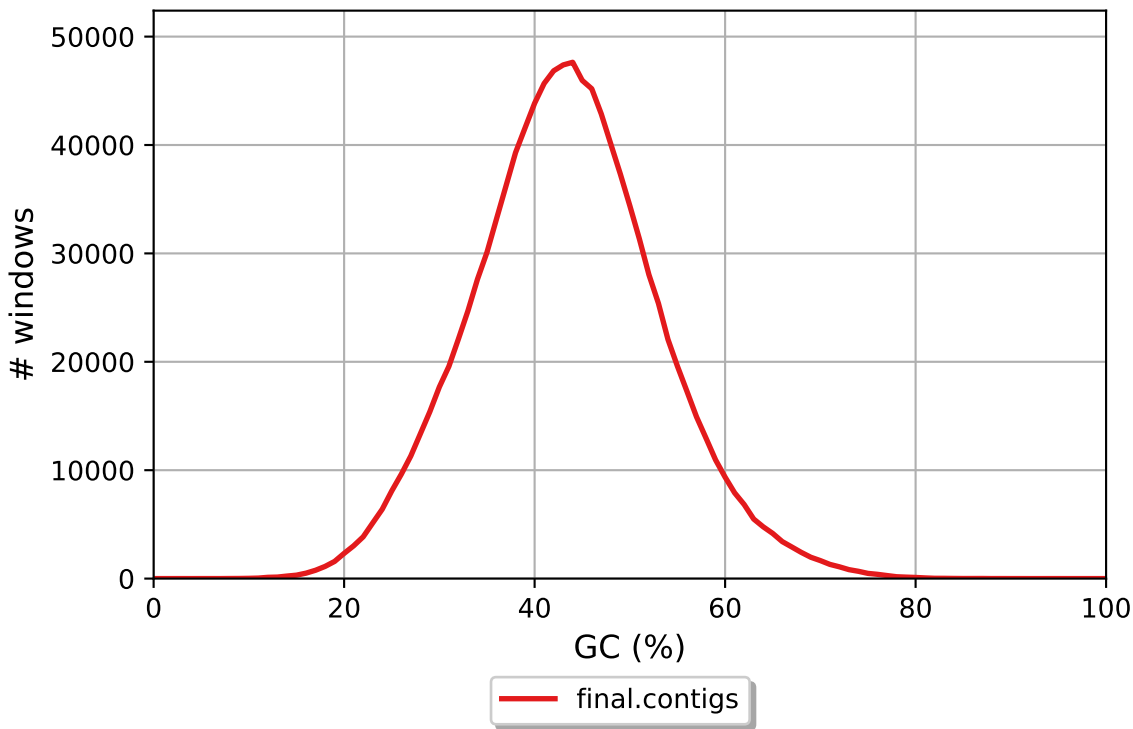
All statistics are based on contigs of size  $\geq 500$  bp, unless otherwise noted (e.g., "# contigs ( $\geq 0$  bp)" and "Total length ( $\geq 0$  bp)" include all contigs).



— final.contigs



GC content



final.contigs GC content

