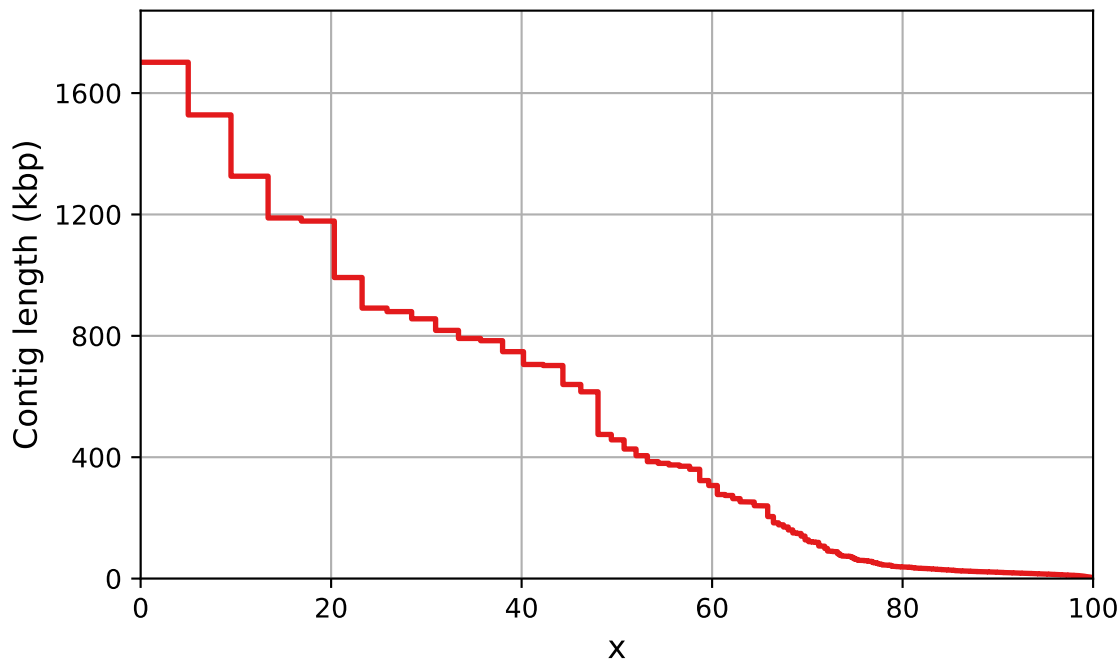


# Report

	assembly
# contigs (>= 0 bp)	512
# contigs (>= 1000 bp)	507
# contigs (>= 5000 bp)	471
# contigs (>= 10000 bp)	406
# contigs (>= 25000 bp)	168
# contigs (>= 50000 bp)	76
Total length (>= 0 bp)	34039534
Total length (>= 1000 bp)	34035824
Total length (>= 5000 bp)	33932280
Total length (>= 10000 bp)	33441531
Total length (>= 25000 bp)	29423238
Total length (>= 50000 bp)	26312434
# contigs	512
Largest contig	1701574
Total length	34039534
GC (%)	30.99
N50	457292
N75	64519
L50	19
L75	63
# N's per 100 kbp	0.00
# predicted genes (unique)	4704
# predicted genes (>= 0 bp)	30526 + 0 part
# predicted genes (>= 300 bp)	3570 + 0 part
# predicted genes (>= 1500 bp)	197 + 0 part
# predicted genes (>= 3000 bp)	20 + 0 part

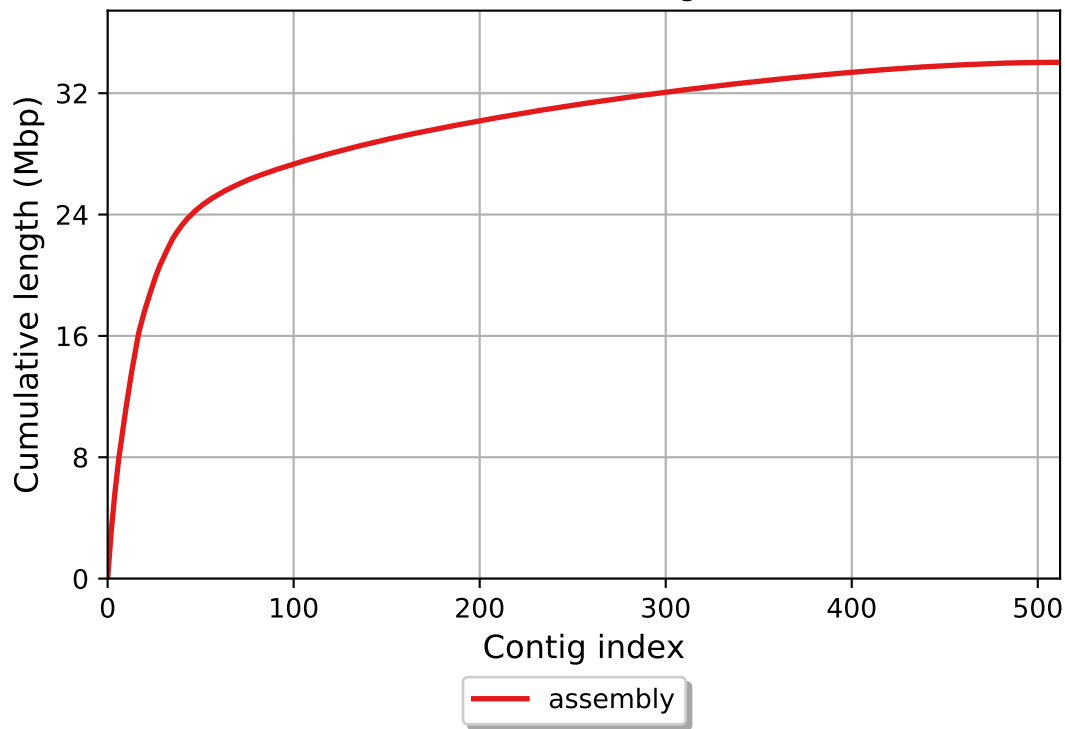
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

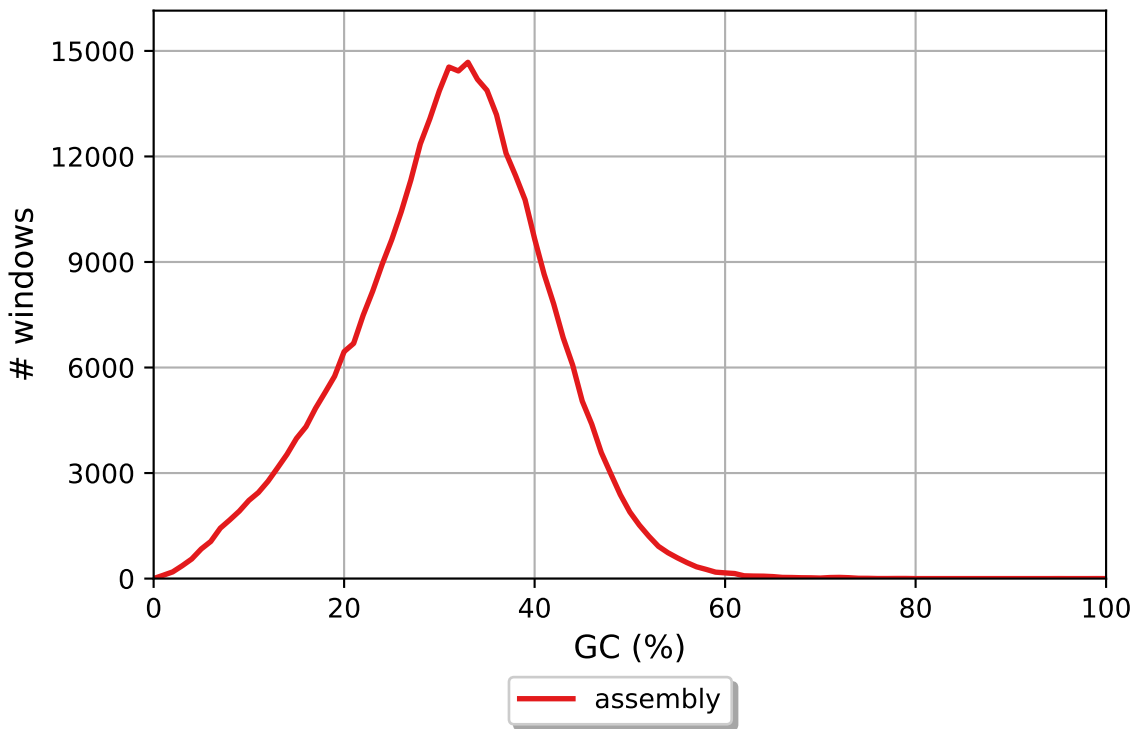


assembly

Cumulative length



GC content



assembly GC content

