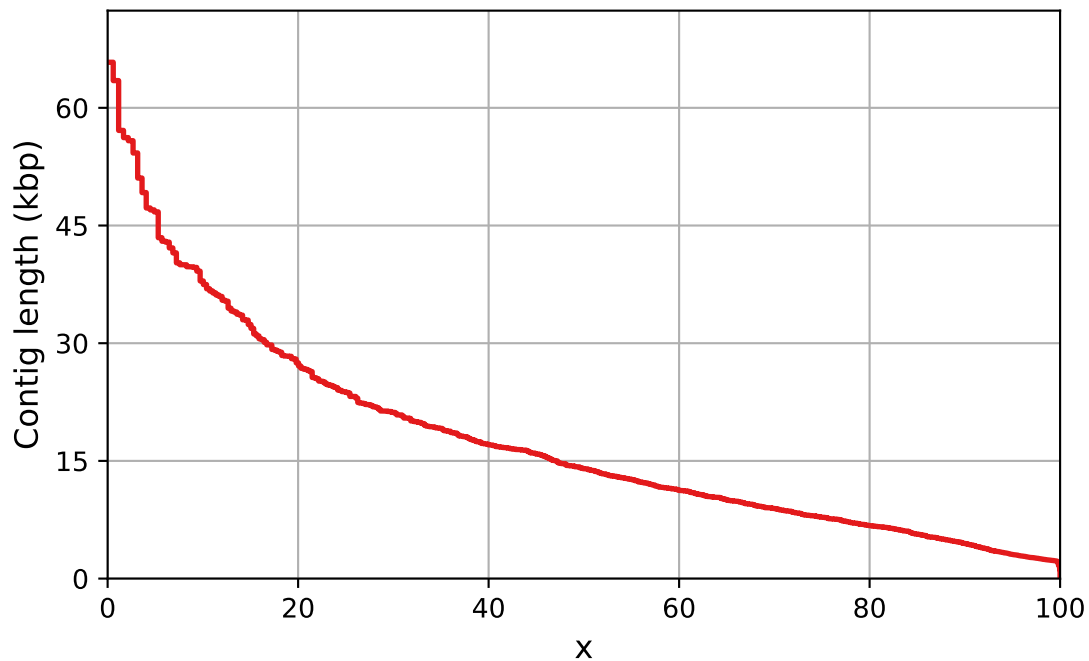


Report

	assembly
# contigs (≥ 0 bp)	1153
# contigs (≥ 1000 bp)	1152
# contigs (≥ 5000 bp)	731
# contigs (≥ 10000 bp)	379
# contigs (≥ 25000 bp)	71
# contigs (≥ 50000 bp)	7
Total length (≥ 0 bp)	11175781
Total length (≥ 1000 bp)	11174829
Total length (≥ 5000 bp)	9824715
Total length (≥ 10000 bp)	7273864
Total length (≥ 25000 bp)	2551938
Total length (≥ 50000 bp)	403714
# contigs	1153
Largest contig	65807
Total length	11175781
GC (%)	38.77
N50	14009
N90	4474
auN	17827.4
L50	237
L90	781
# N's per 100 kbp	0.00
# predicted genes (unique)	4694
# predicted genes (≥ 0 bp)	4679 + 15 part
# predicted genes (≥ 300 bp)	3022 + 9 part
# predicted genes (≥ 1500 bp)	284 + 0 part
# predicted genes (≥ 3000 bp)	29 + 0 part

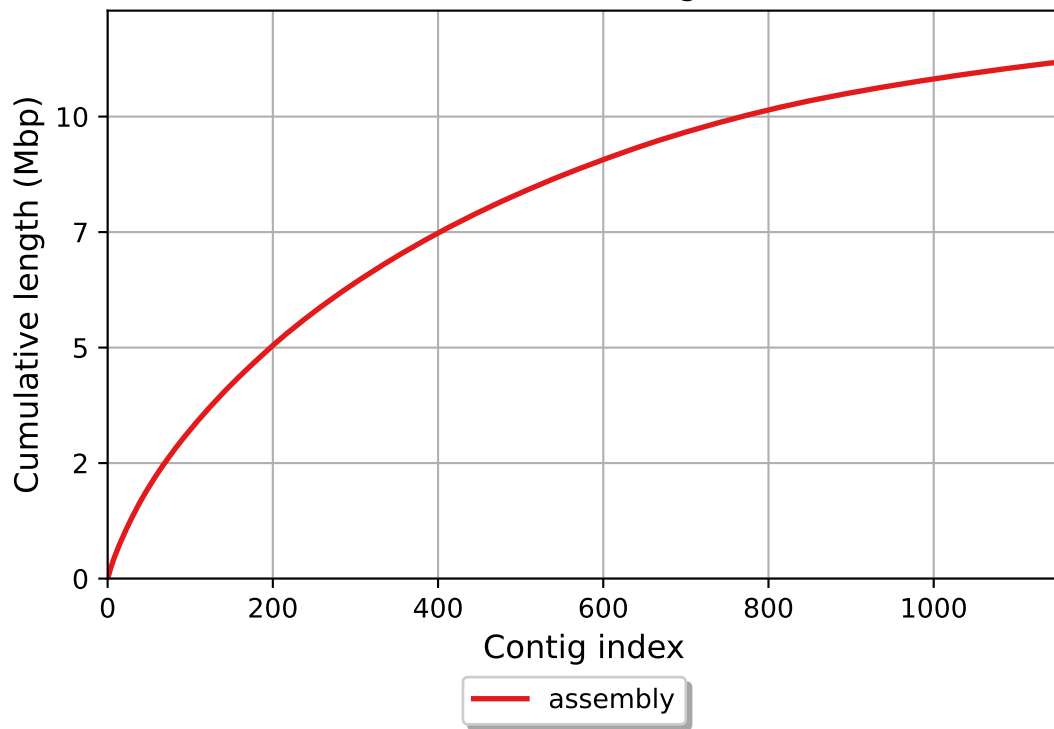
All statistics are based on contigs of size ≥ 500 bp, unless otherwise noted (e.g., "# contigs (≥ 0 bp)" and "Total length (≥ 0 bp)" include all contigs).

Nx

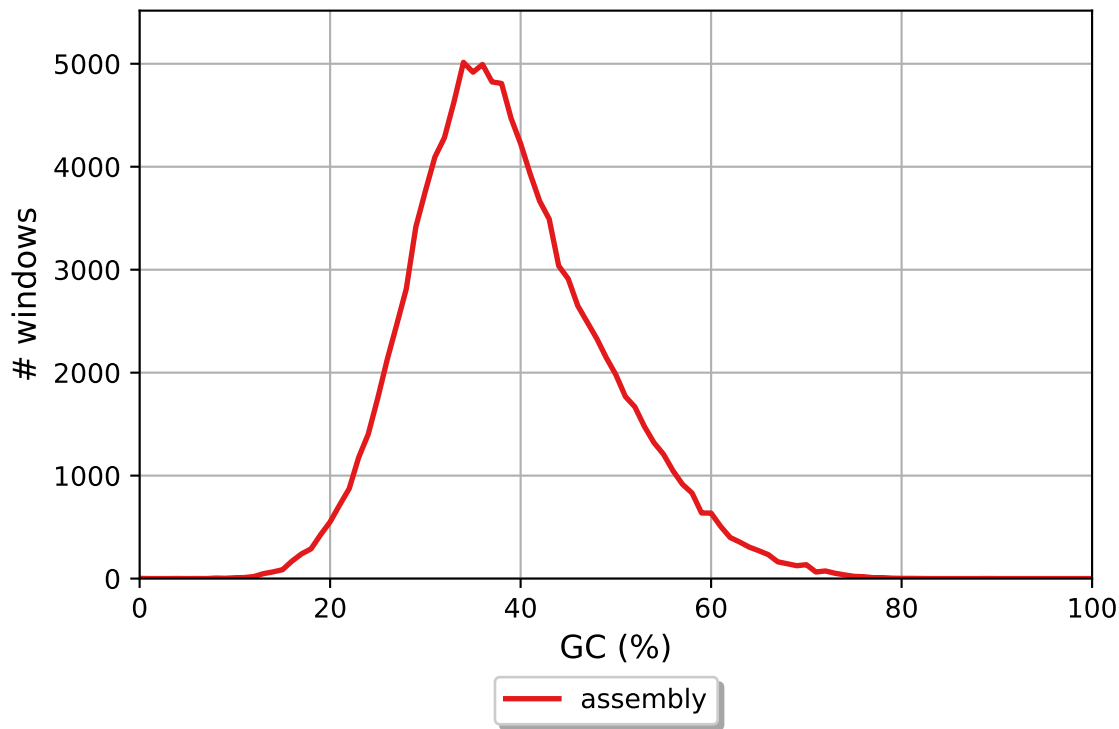


— assembly

Cumulative length



GC content



assembly GC content

