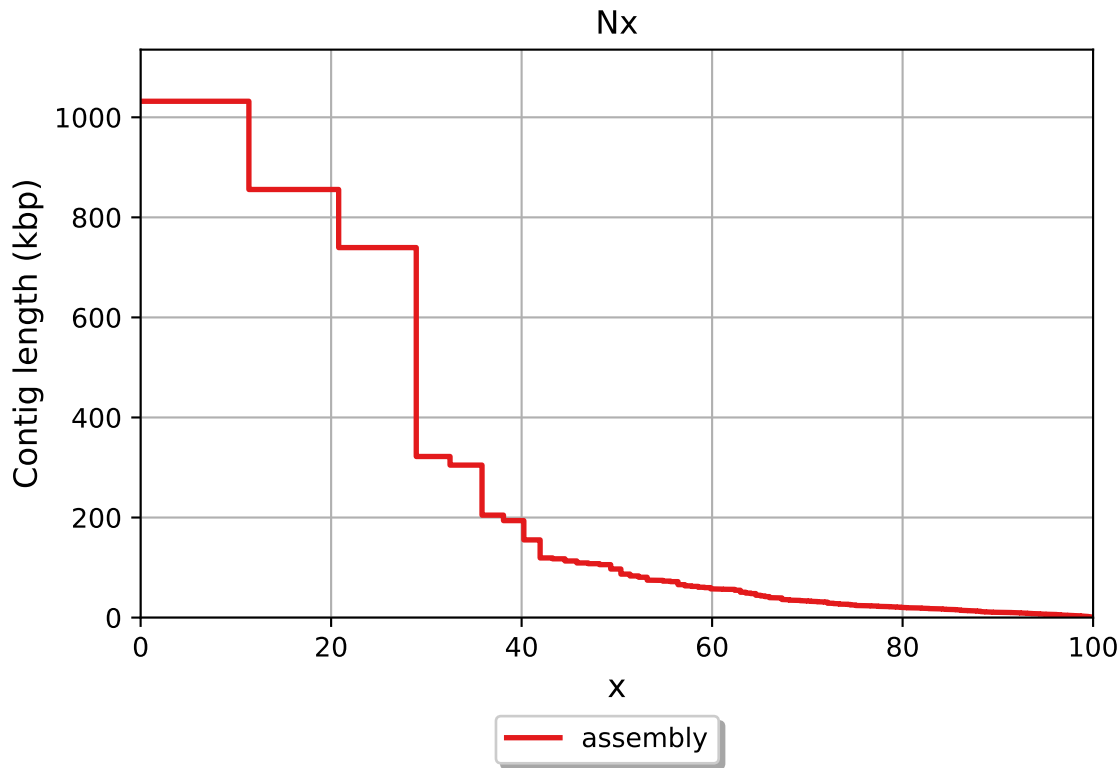


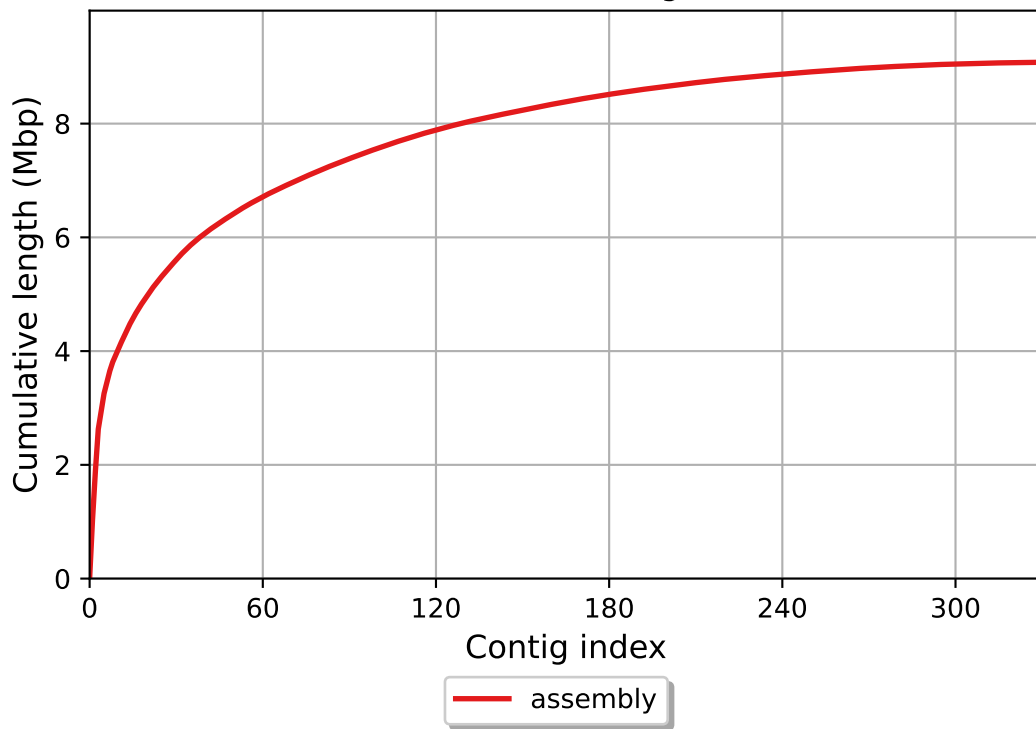
Report

	assembly
# contigs (≥ 0 bp)	331
# contigs (≥ 1000 bp)	318
# contigs (≥ 5000 bp)	222
# contigs (≥ 10000 bp)	157
# contigs (≥ 25000 bp)	64
# contigs (≥ 50000 bp)	33
Total length (≥ 0 bp)	9078912
Total length (≥ 1000 bp)	9070609
Total length (≥ 5000 bp)	8787775
Total length (≥ 10000 bp)	8308280
Total length (≥ 25000 bp)	6816522
Total length (≥ 50000 bp)	5767117
# contigs	330
Largest contig	1032034
Total length	9078758
GC (%)	56.60
N50	97090
N90	10518
auN	316925.2
L50	15
L90	144
# N's per 100 kbp	0.00
# predicted genes (unique)	8041
# predicted genes (≥ 0 bp)	7968 + 73 part
# predicted genes (≥ 300 bp)	6671 + 66 part
# predicted genes (≥ 1500 bp)	539 + 2 part
# predicted genes (≥ 3000 bp)	27 + 0 part

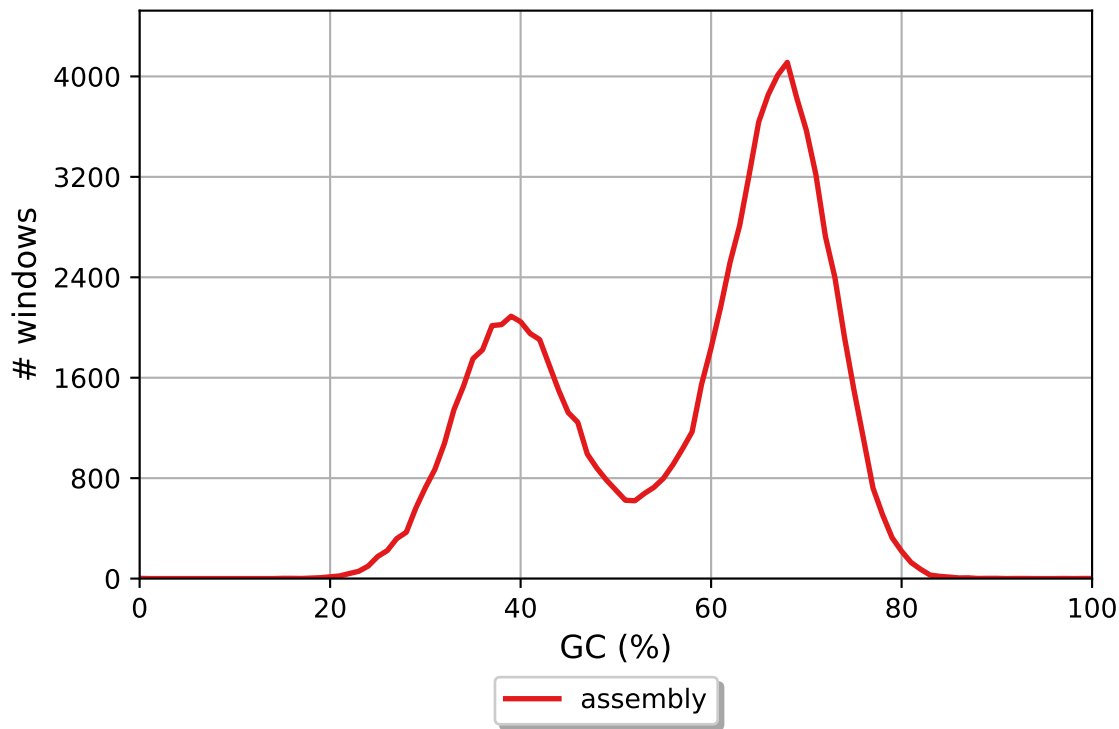
All statistics are based on contigs of size ≥ 500 bp, unless otherwise noted (e.g., "# contigs (≥ 0 bp)" and "Total length (≥ 0 bp)" include all contigs).



Cumulative length



GC content



assembly GC content

