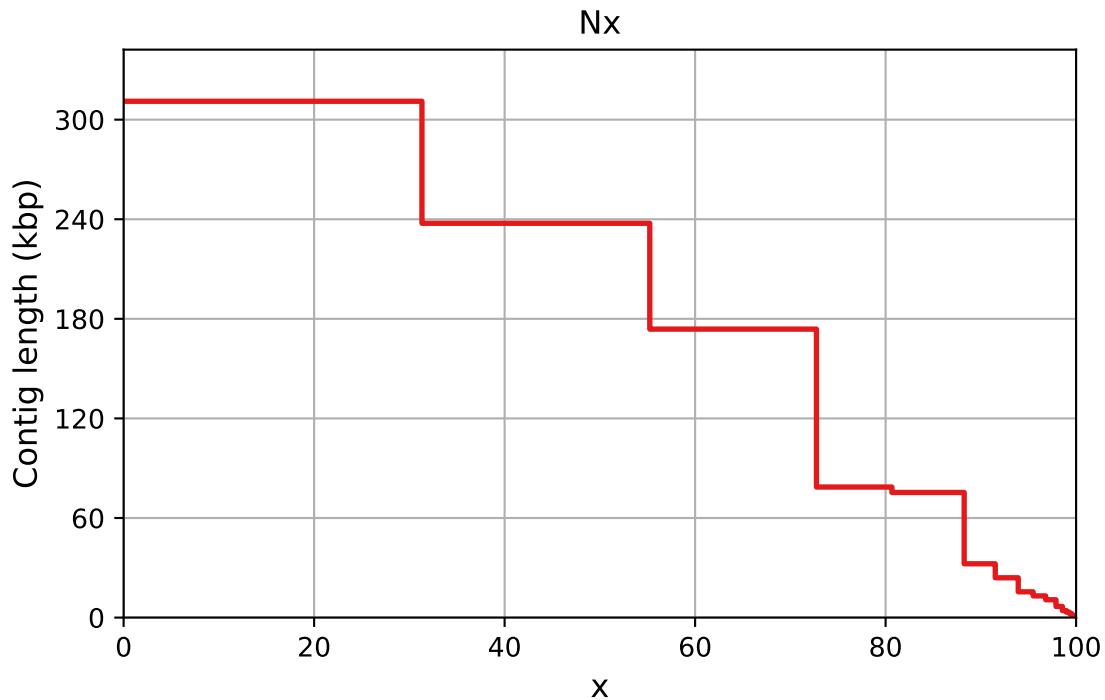
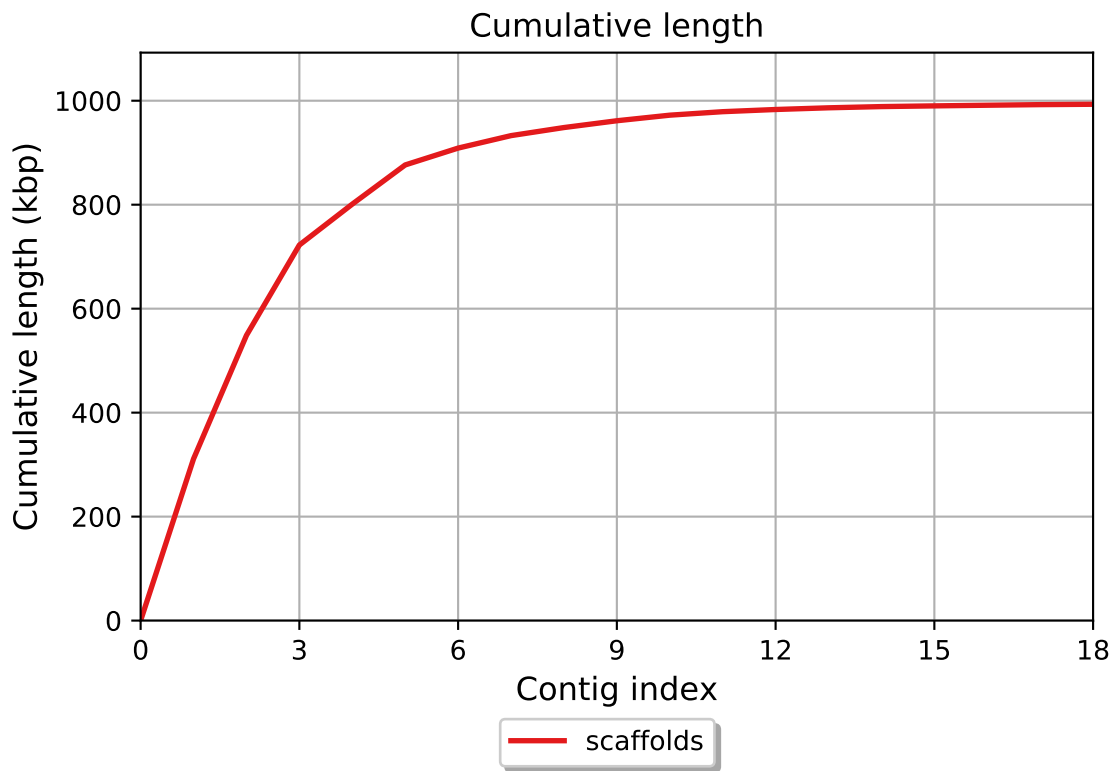


Report

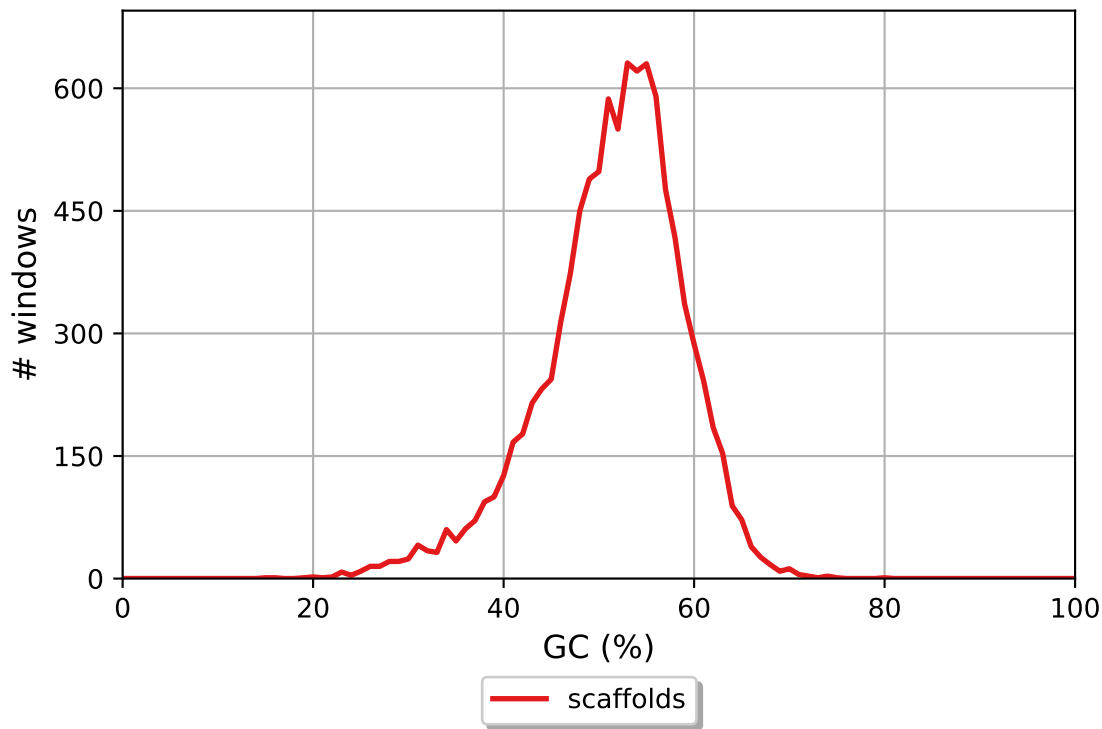
	scaffolds
# contigs (>= 0 bp)	29
# contigs (>= 1000 bp)	17
# contigs (>= 5000 bp)	11
# contigs (>= 10000 bp)	10
# contigs (>= 25000 bp)	6
# contigs (>= 50000 bp)	5
Total length (>= 0 bp)	995374
Total length (>= 1000 bp)	992468
Total length (>= 5000 bp)	978907
Total length (>= 10000 bp)	972202
Total length (>= 25000 bp)	908881
Total length (>= 50000 bp)	876464
# contigs	18
Largest contig	311072
Total length	993171
GC (%)	51.42
N50	237562
N75	78653
L50	2
L75	4
# N's per 100 kbp	10.07
# predicted genes (unique)	914
# predicted genes (>= 0 bp)	910 + 4 part
# predicted genes (>= 300 bp)	813 + 1 part
# predicted genes (>= 1500 bp)	129 + 0 part
# predicted genes (>= 3000 bp)	11 + 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).

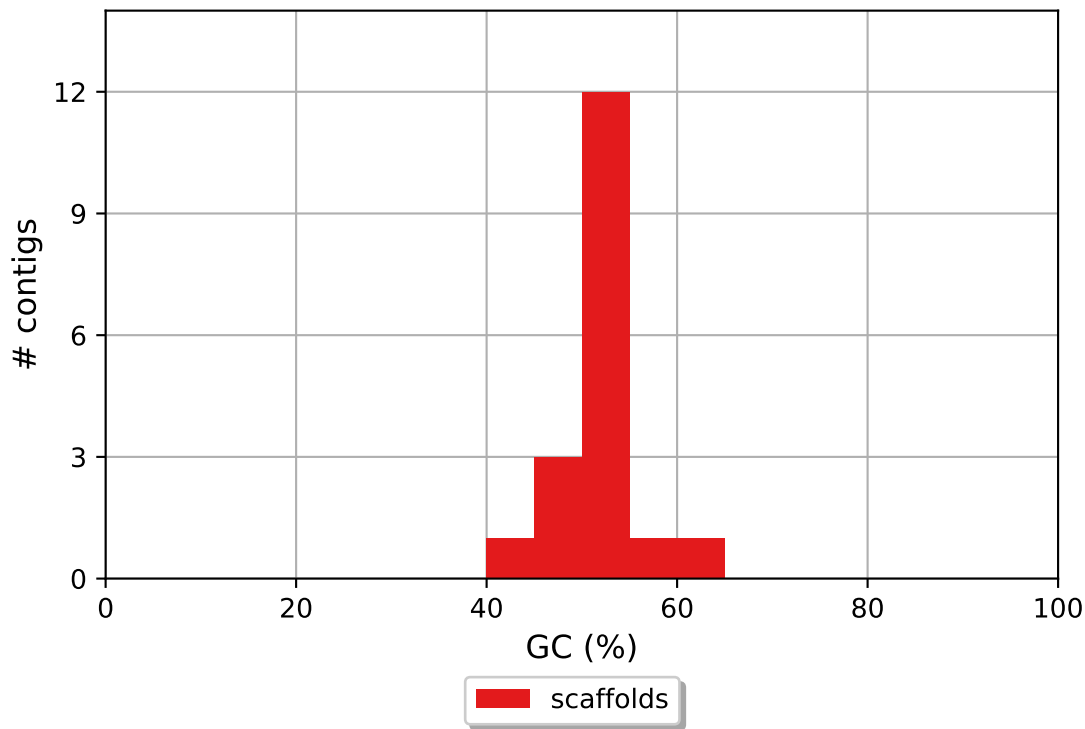




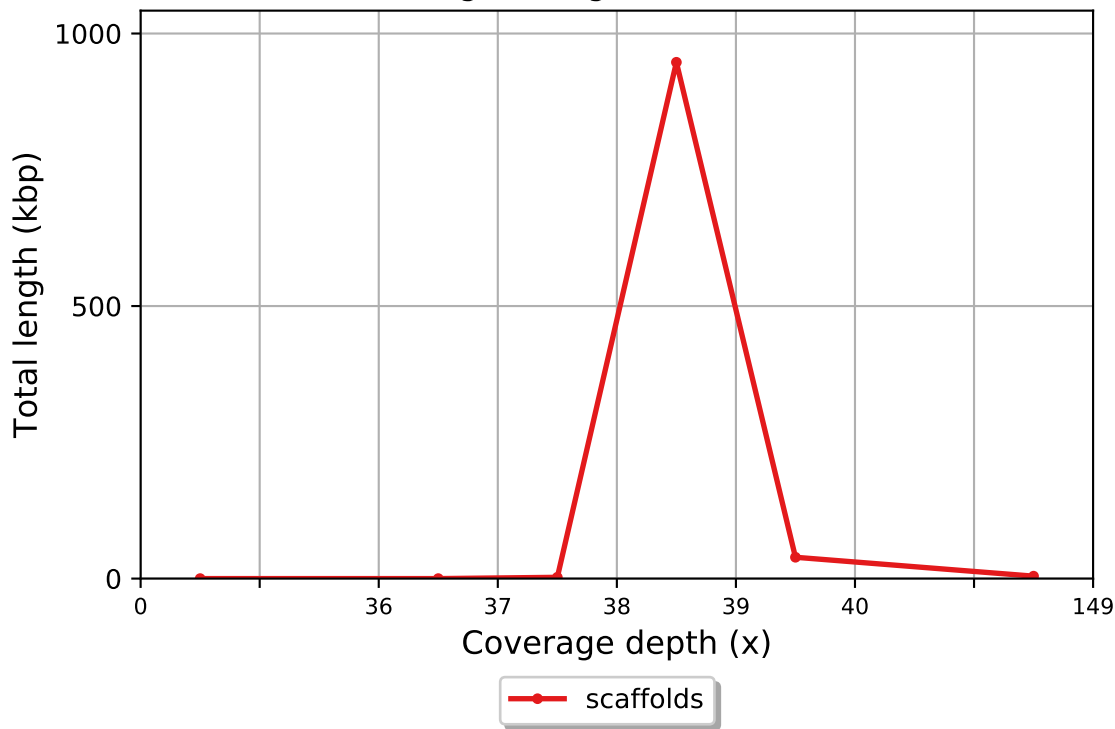
GC content



scaffolds GC content



Coverage histogram (bin size: 1x)



scaffolds coverage histogram (bin size: 1x)

