Report

	assembly	medaka_assembly_1_consensus	medaka_assembly_2_consensus	medaka_assembly_3_consensus
# contigs (>= 0 bp)	315	243	241	238
# contigs (>= 1000 bp)	307	241	239	236
# contigs (>= 5000 bp)	251	228	226	223
# contigs (>= 10000 bp)	199	192	189	186
# contigs (>= 25000 bp)	140	140	141	141
# contigs (>= 50000 bp)	120	121	120	120
Total length (>= 0 bp)	32329712	32101582	32137739	32134007
Total length (>= 1000 bp)	32324184	32100326	32136481	32132751
Total length (>= 5000 bp)	32144434	32049278	32085562	32081830
Total length (>= 10000 bp)	31758269	31780740	31802978	31798405
Total length (>= 25000 bp)	30873848	30994456	31080126	31115178
Total length (>= 50000 bp)	30132304	30298266	30308400	30344019
# contigs	315	243	241	238
Largest contig	825024	828718	828729	828750
Total length	32329712	32101582	32137739	32134007
GC (%)	28.65	28.52	28.52	28.52
N50	277586	280068	280441	280848
N75	202230	204655	207158	208682
L50	43	42	42	42
L75	77	76	76	76
# N's per 100 kbp	0.31	0.00	0.00	0.00

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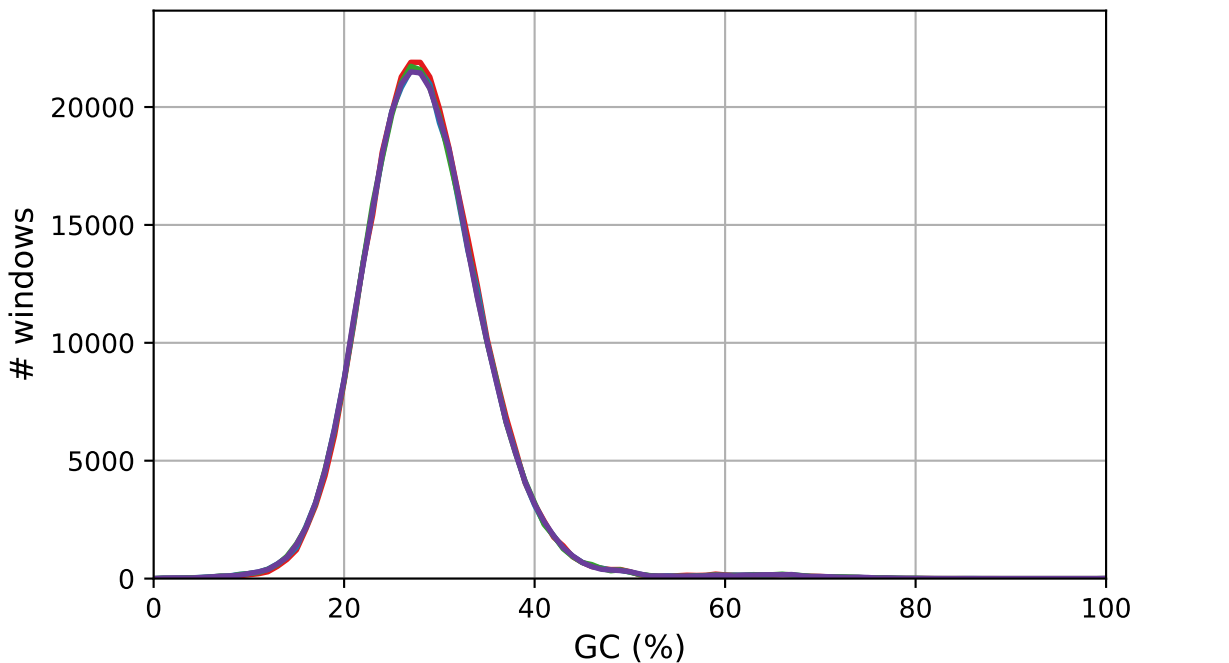(e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).
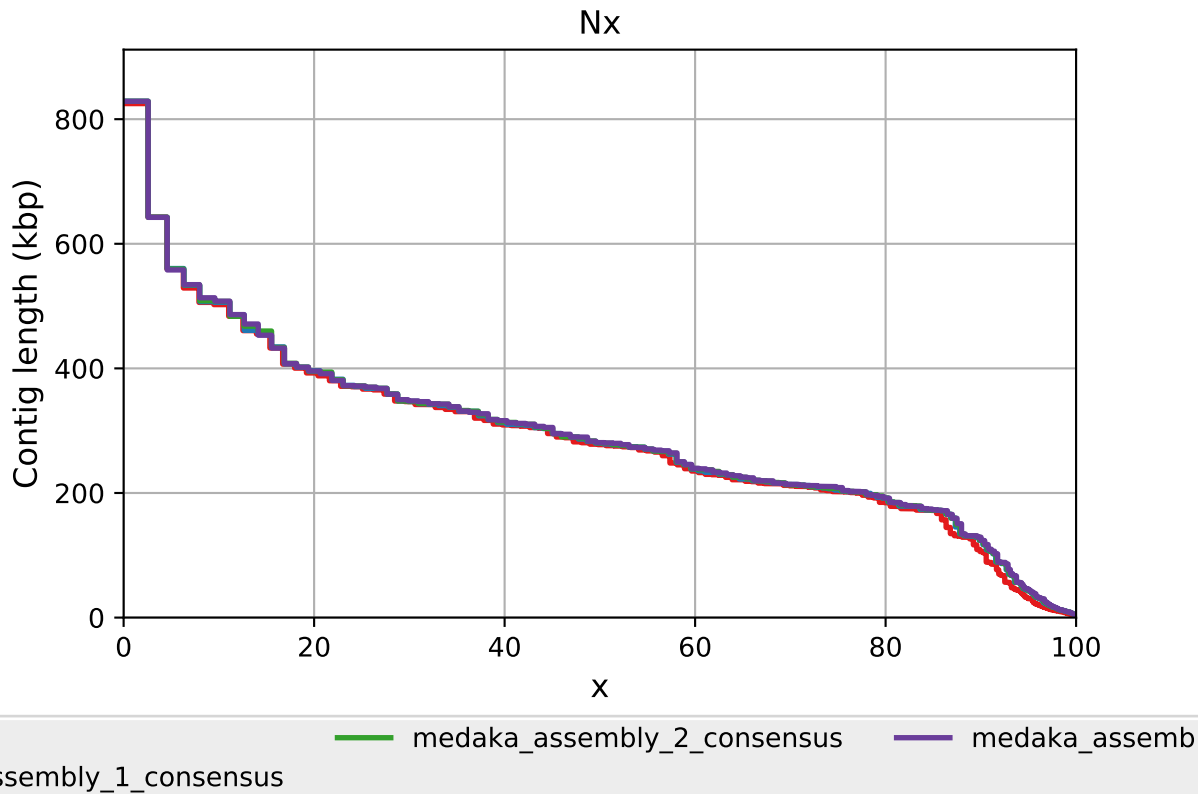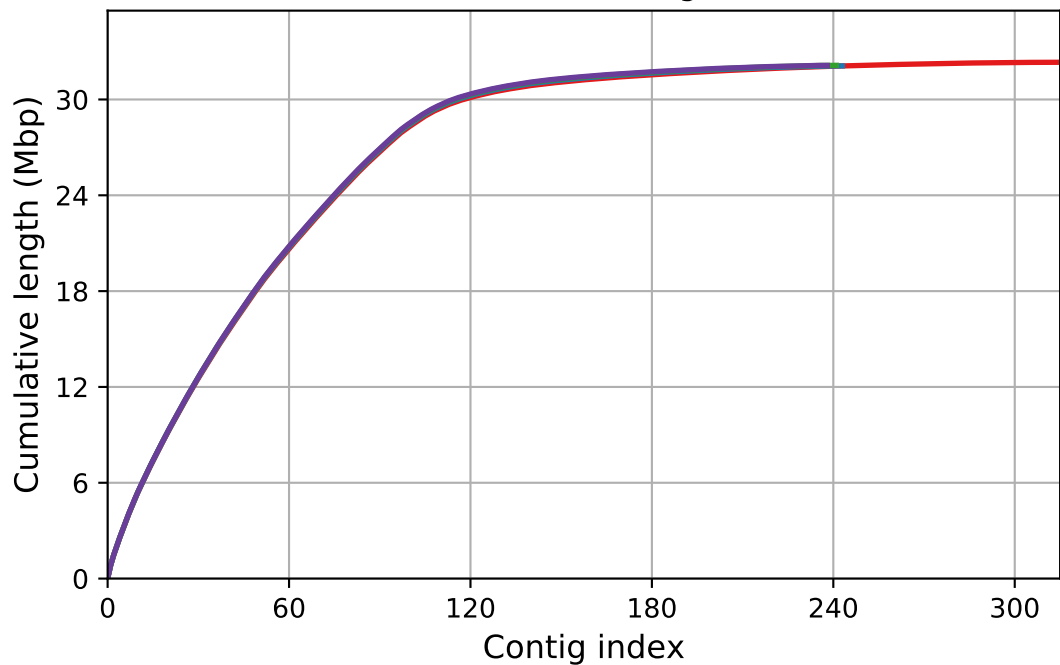
Nx

Contig length (kbp) vs x

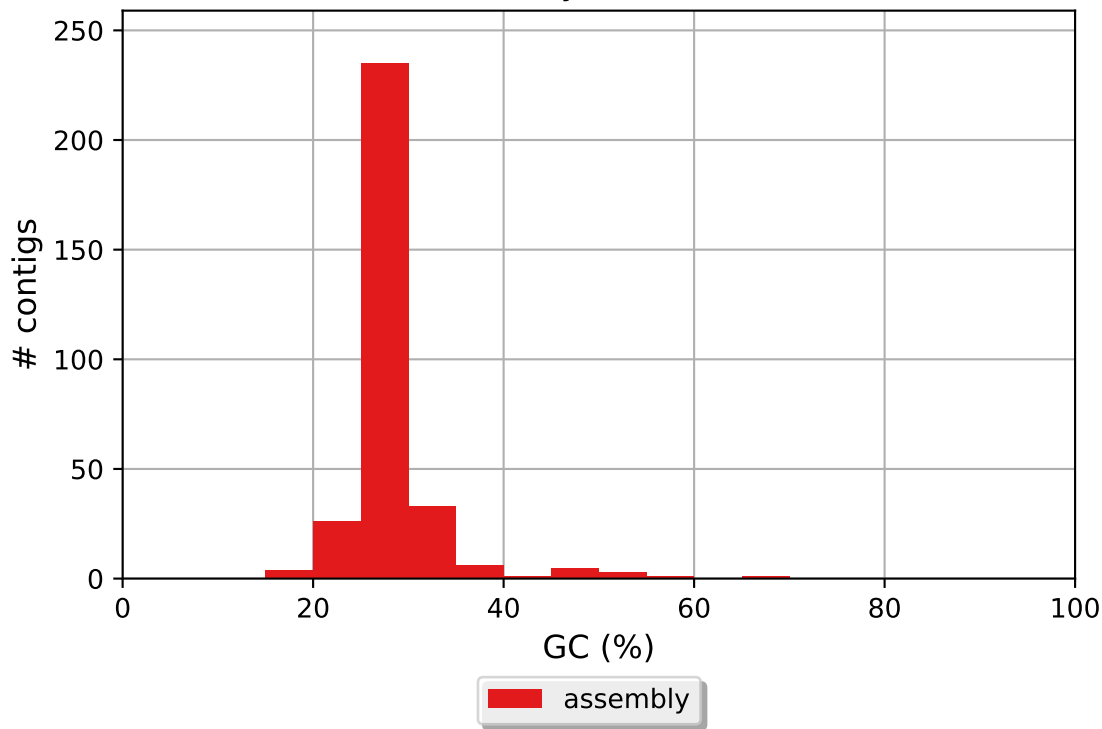medaka_assembly_1_consensus — medaka_assembly_2_consensus — medaka_assembl...

Cumulative length

Cumulative length (Mbp) vs Contig index

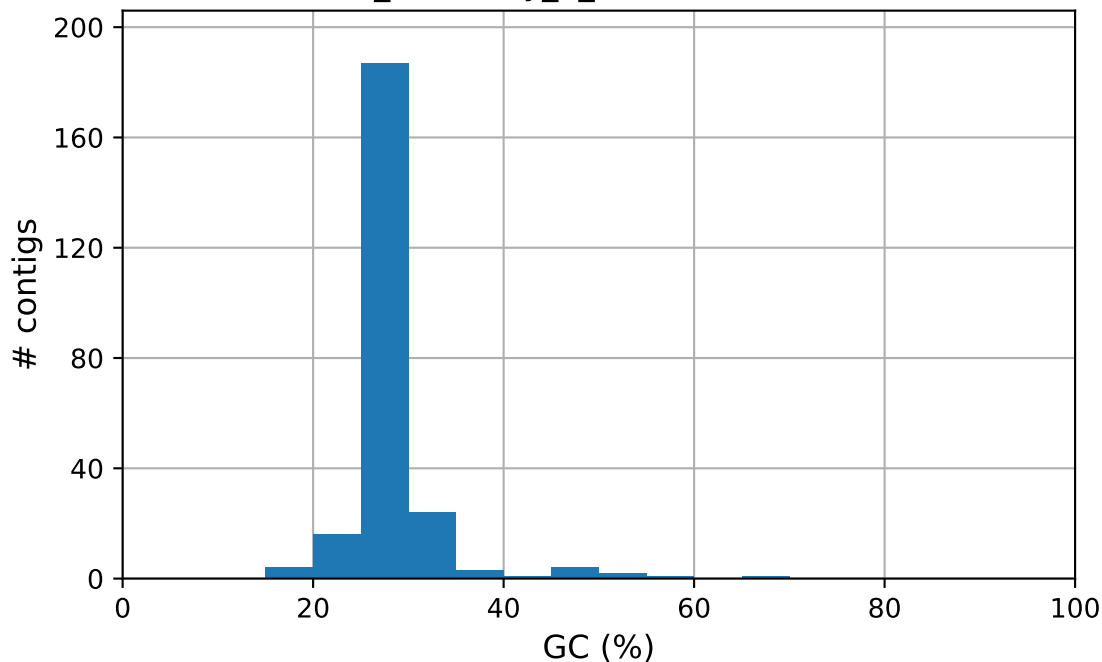Legend:
- medaka_assembly_2_consensus
- medaka_assembl...
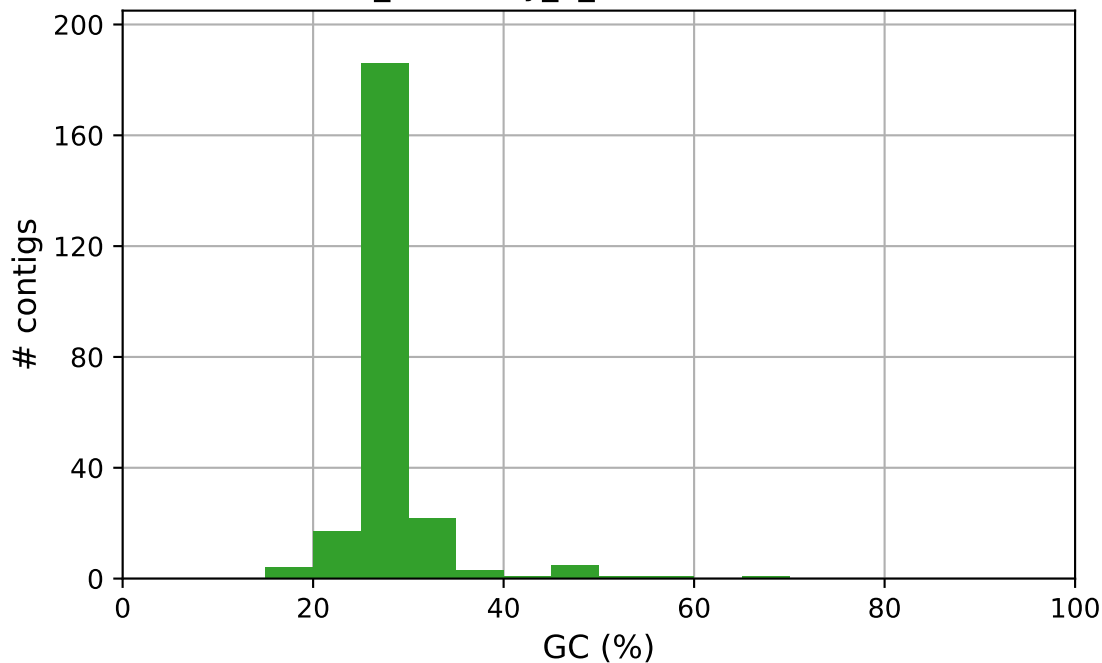- _assembly_1_consensus

GC content

assembly GC content

medaka_assembly_1_consensus GC content

medaka_assembly_2_consensus GC content

medaka_assembly_3_consensus GC content