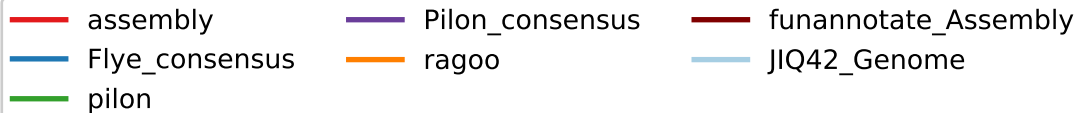
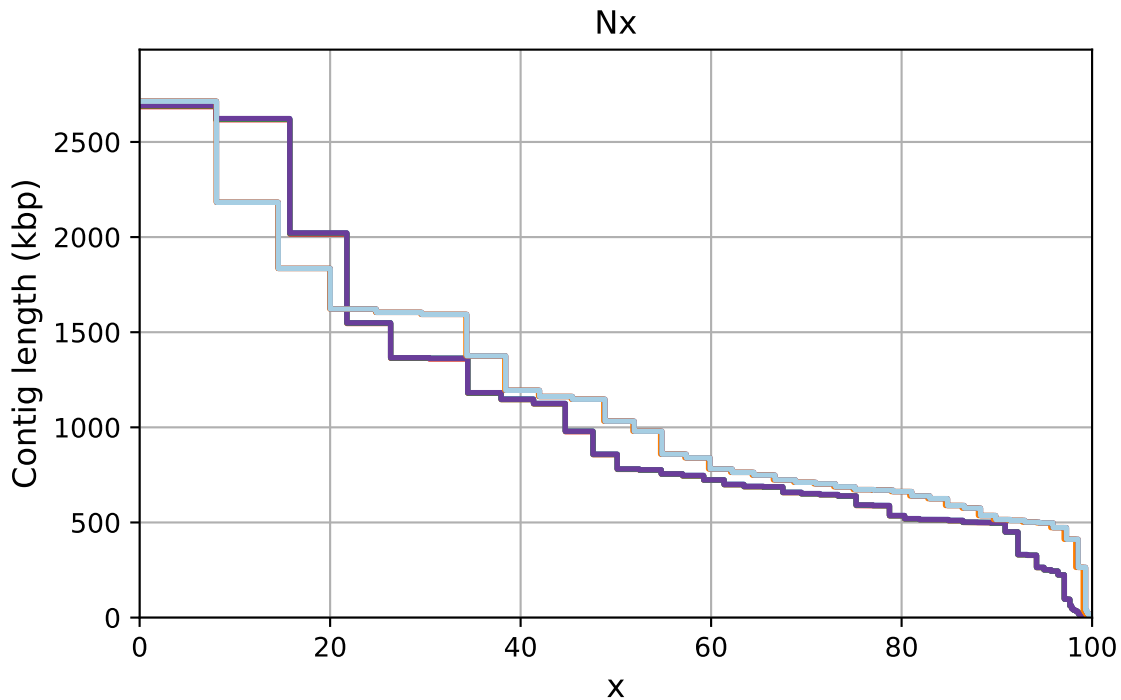


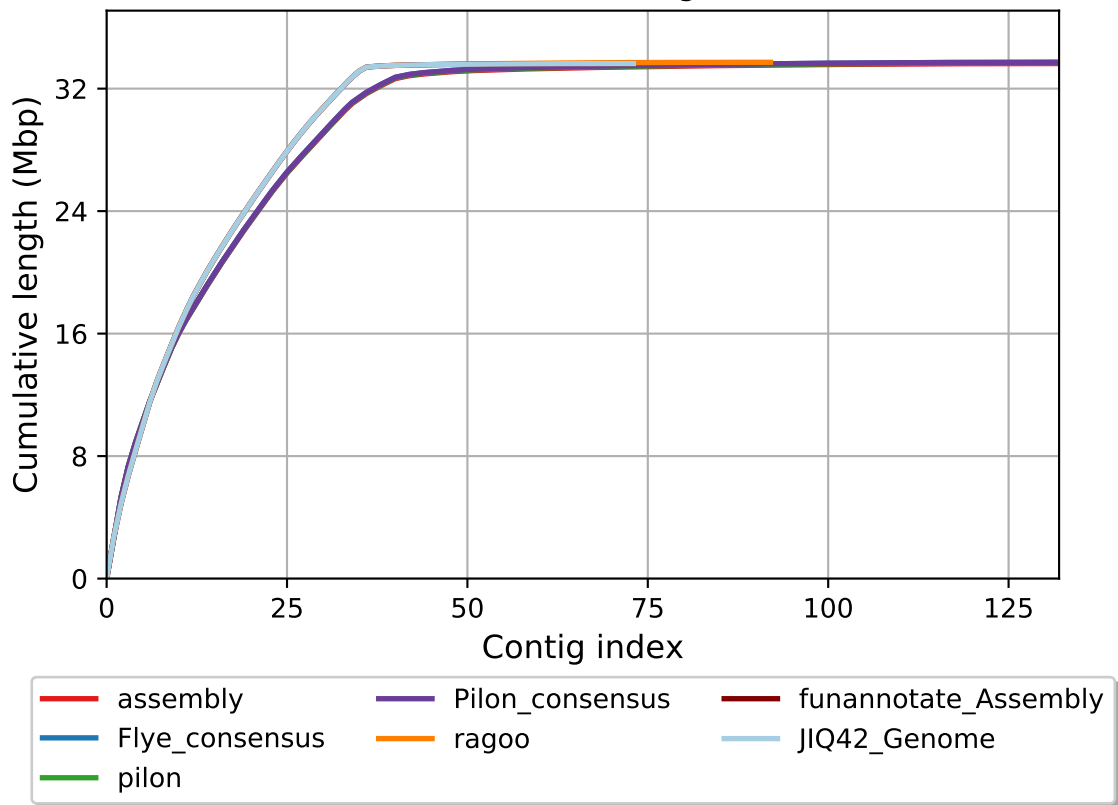
## Report

	assembly	Flye_consensus	pilon	Pilon_consensus	ragoo	funannotate_Assembly	JIQ42_Genome
# contigs (>= 0 bp)	132	132	132	132	92	73	73
# contigs (>= 1000 bp)	121	121	121	121	81	63	63
# contigs (>= 5000 bp)	89	89	89	89	54	47	47
# contigs (>= 10000 bp)	62	62	62	62	43	42	42
# contigs (>= 25000 bp)	50	50	50	50	39	39	39
# contigs (>= 50000 bp)	43	43	43	43	36	36	36
Total length (>= 0 bp)	33638179	33721595	33679868	33721063	33721063	33635959	33635959
Total length (>= 1000 bp)	33630185	33713597	33671898	33713034	33713034	33628718	33628718
Total length (>= 5000 bp)	33525913	33609211	33570045	33611348	33632586	33578288	33578288
Total length (>= 10000 bp)	33329665	33412935	33373675	33415119	33552965	33541605	33541605
Total length (>= 25000 bp)	33164501	33247549	33208396	33249731	33499710	33499710	33499710
Total length (>= 50000 bp)	32910724	32993979	32954312	32996159	33406078	33406078	33406078
# contigs	132	132	132	132	92	73	73
Largest contig	2685671	2692345	2689405	2692308	2713885	2713885	2713885
Total length	33638179	33721595	33679868	33721063	33721063	33635959	33635959
GC (%)	59.50	59.47	59.51	59.48	59.48	59.48	59.48
N50	856269	858568	857384	858890	1032311	1032311	1032311
N75	637788	639273	638504	639528	672691	687670	687670
L50	11	11	11	11	11	11	11
L75	23	23	23	23	22	21	21
# N's per 100 kbp	2.68	2.37	1.70	1.70	1.70	1.70	1.70

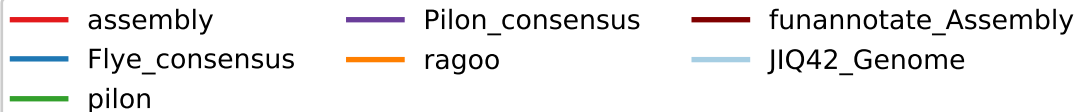
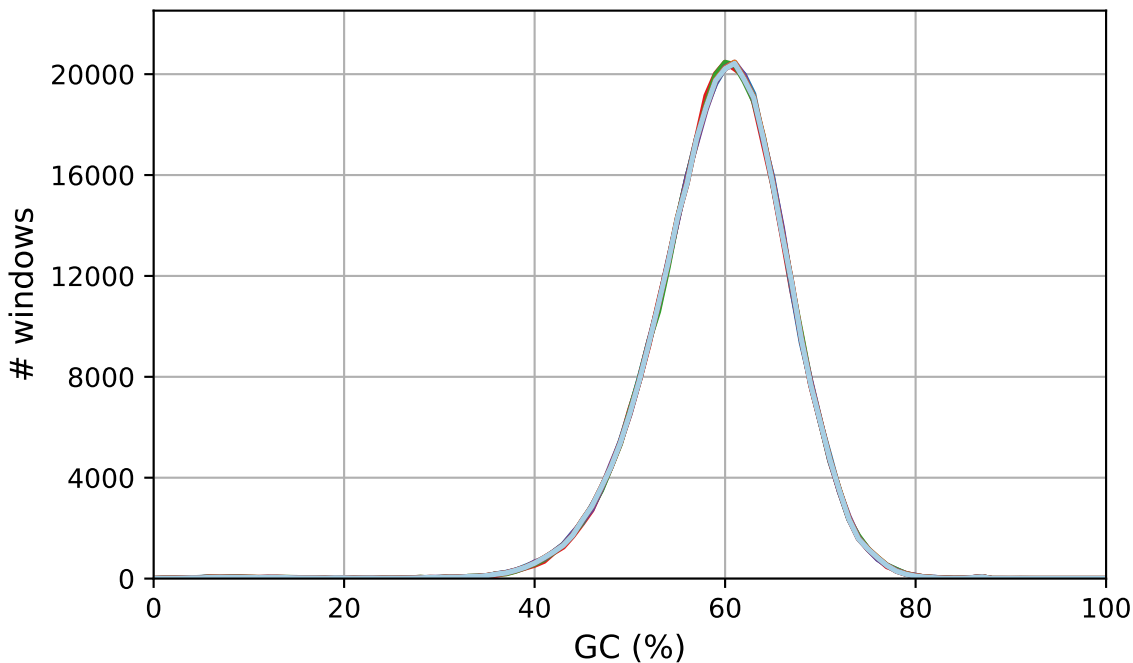
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



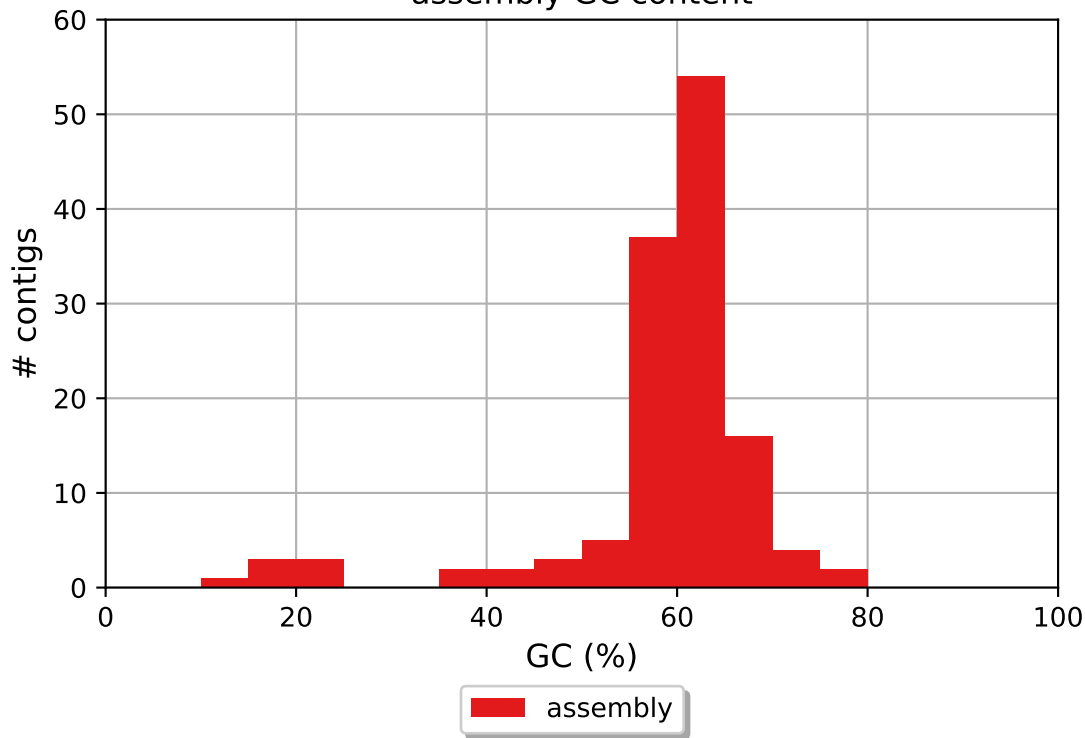
Cumulative length



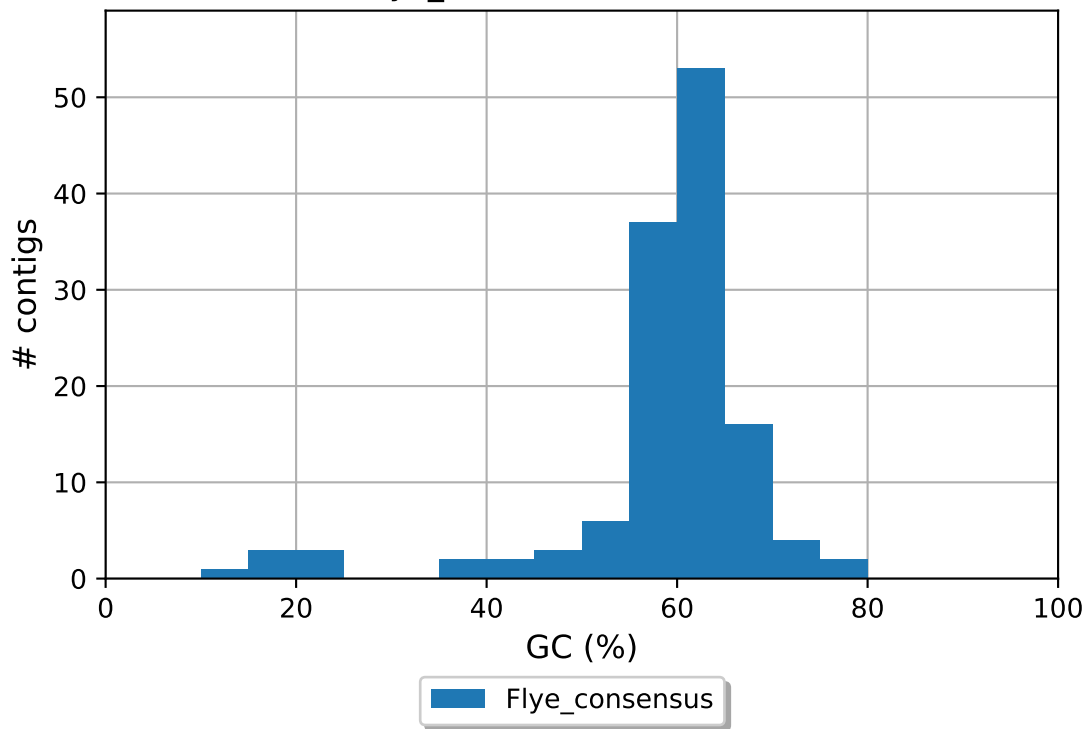
GC content



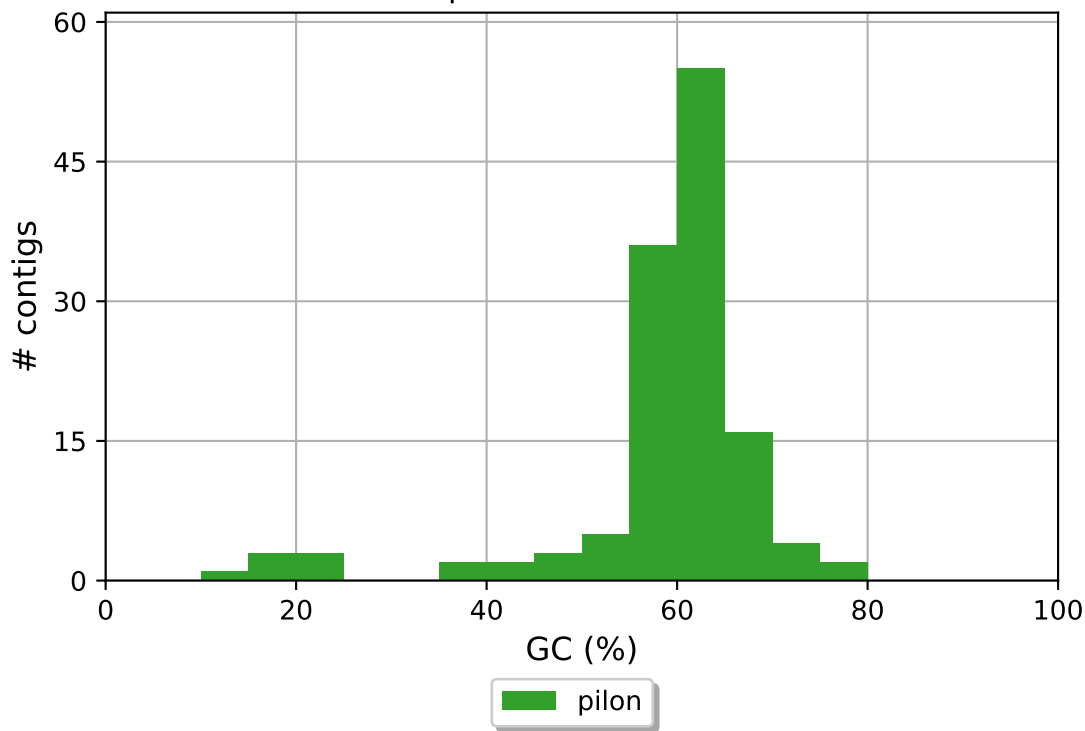
assembly GC content



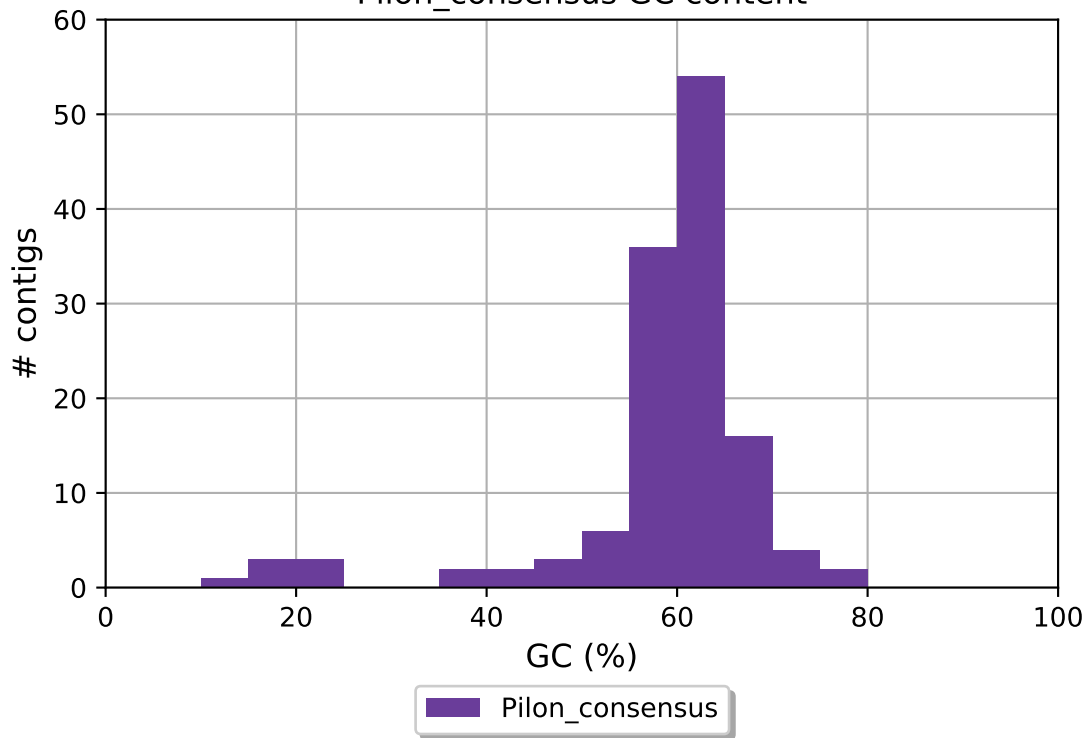
Flye\_consensus GC content



pilon GC content

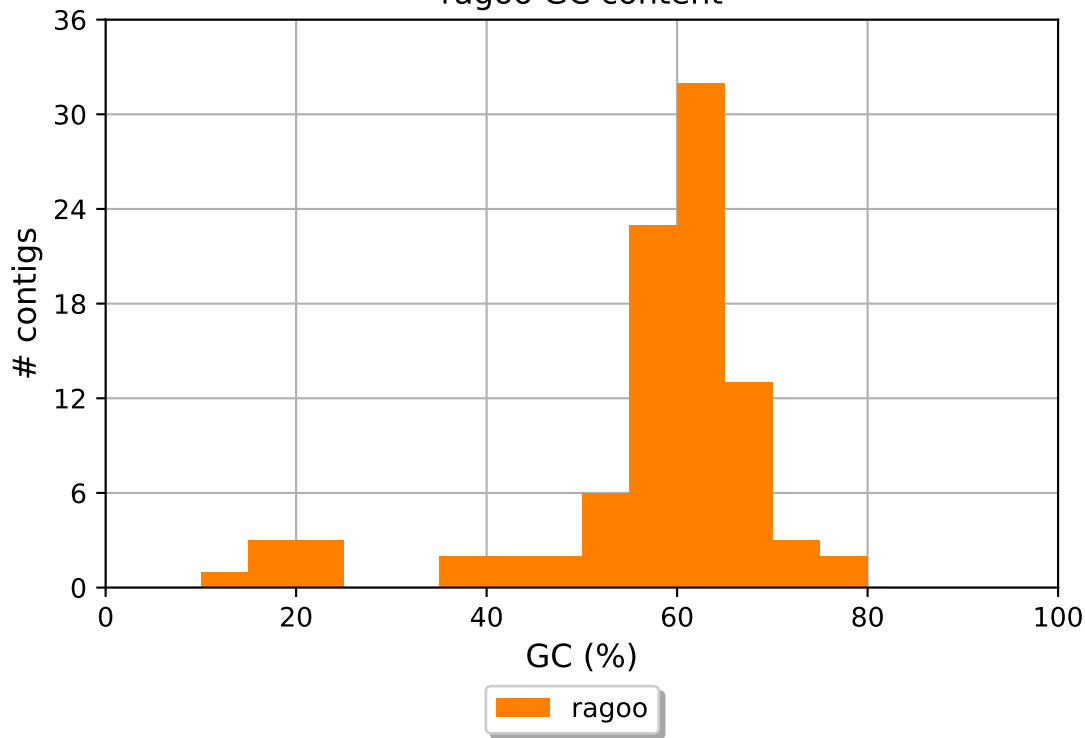


Pilon\_consensus GC content

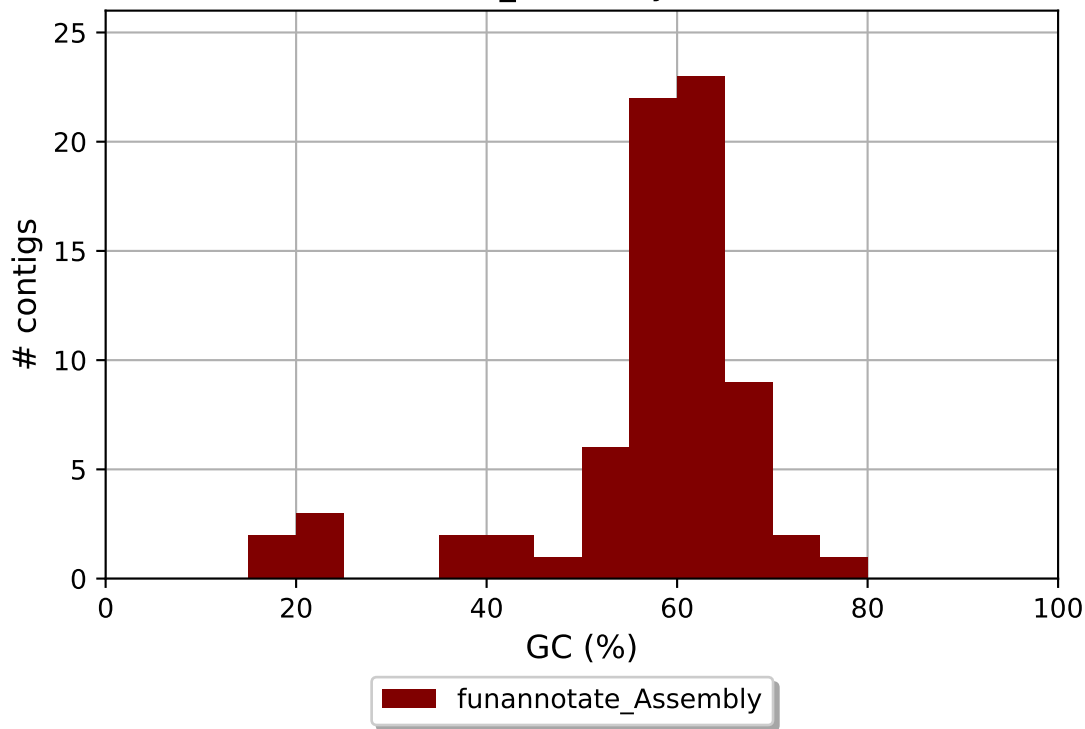




ragoo GC content



funannotate\_Assembly GC content



JIQ42\_Genome GC content

