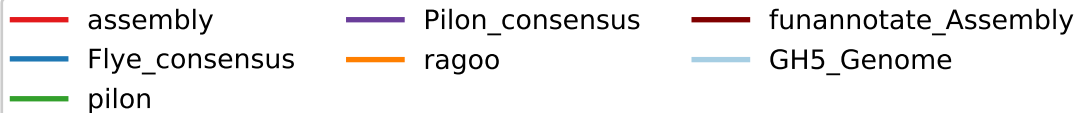
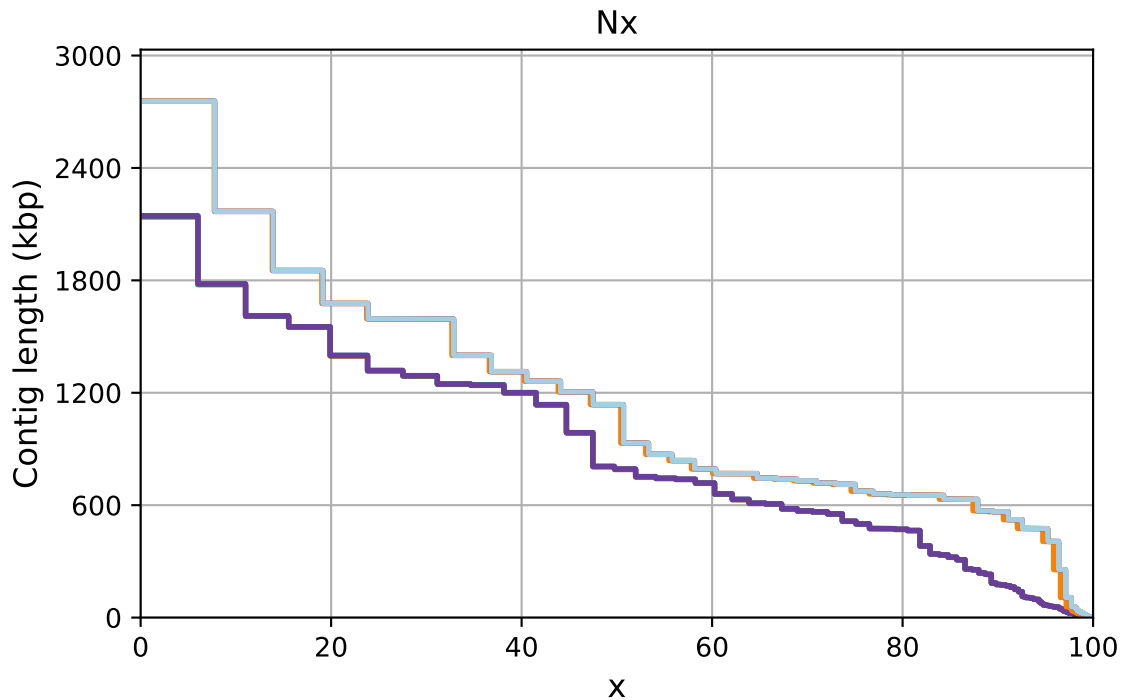


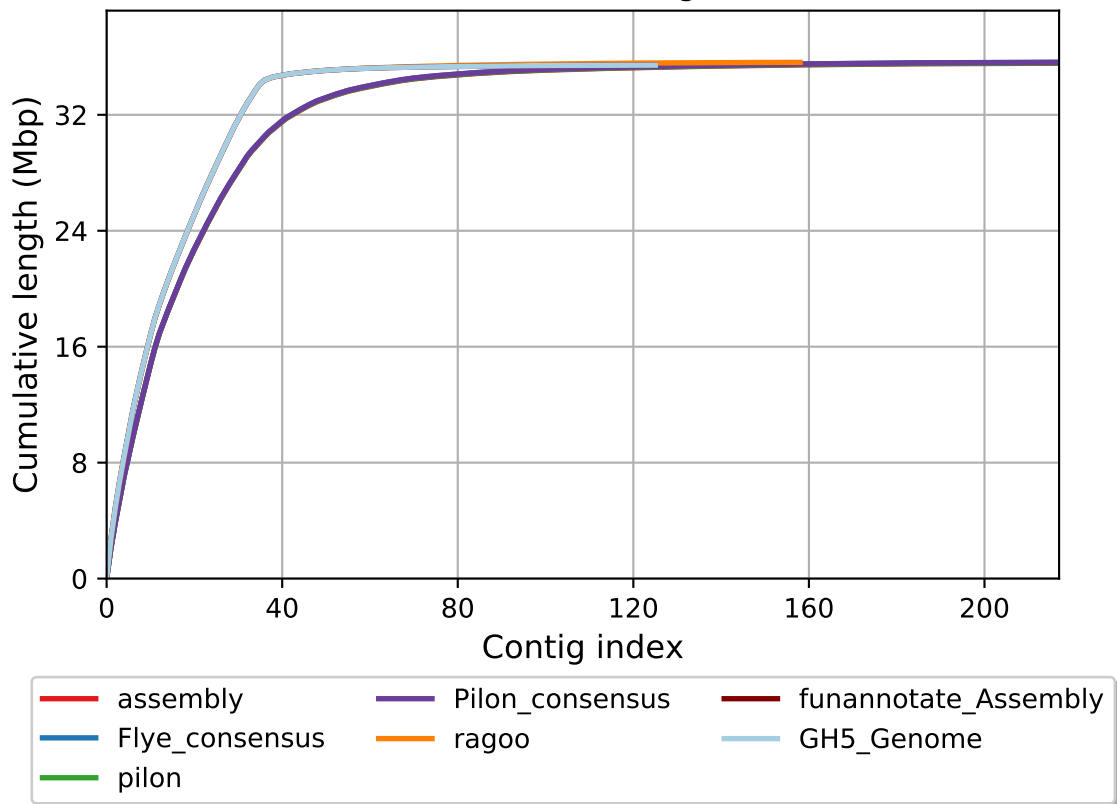
## Report

	assembly	Flye_consensus	pilon	Pilon_consensus	ragoo	funannotate_Assembly	GH5_Genome
# contigs (>= 0 bp)	217	217	217	217	158	125	125
# contigs (>= 1000 bp)	206	206	206	206	147	114	114
# contigs (>= 5000 bp)	139	139	138	138	87	73	73
# contigs (>= 10000 bp)	102	102	102	102	65	60	60
# contigs (>= 25000 bp)	76	76	77	77	49	49	49
# contigs (>= 50000 bp)	66	66	66	66	41	41	41
Total length (>= 0 bp)	35559655	35625818	35578884	35619093	35619093	35424749	35424749
Total length (>= 1000 bp)	35551722	35617828	35570977	35611150	35611150	35416806	35416806
Total length (>= 5000 bp)	35362686	35428688	35377336	35417442	35446314	35305062	35305062
Total length (>= 10000 bp)	35101182	35166874	35120692	35160732	35287947	35215027	35215027
Total length (>= 25000 bp)	34657449	34722939	34701640	34741743	35045606	35045606	35045606
Total length (>= 50000 bp)	34302326	34367274	34321642	34361609	34783812	34783812	34783812
# contigs	217	217	217	217	158	125	125
Largest contig	2139654	2145390	2140269	2143491	2755985	2755985	2755985
Total length	35559655	35625818	35578884	35619093	35619093	35424749	35424749
GC (%)	59.53	59.61	59.53	59.61	59.61	59.60	59.60
N50	790645	792898	791153	792918	1136834	1136834	1136834
N75	513527	514457	513909	514464	674535	711148	711148
L50	14	14	14	14	11	11	11
L75	27	27	27	27	23	22	22
# N's per 100 kbp	3.37	3.37	1.57	1.57	1.57	1.58	1.58

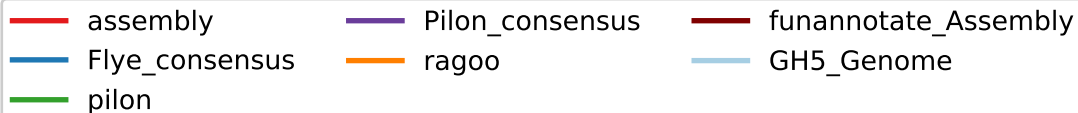
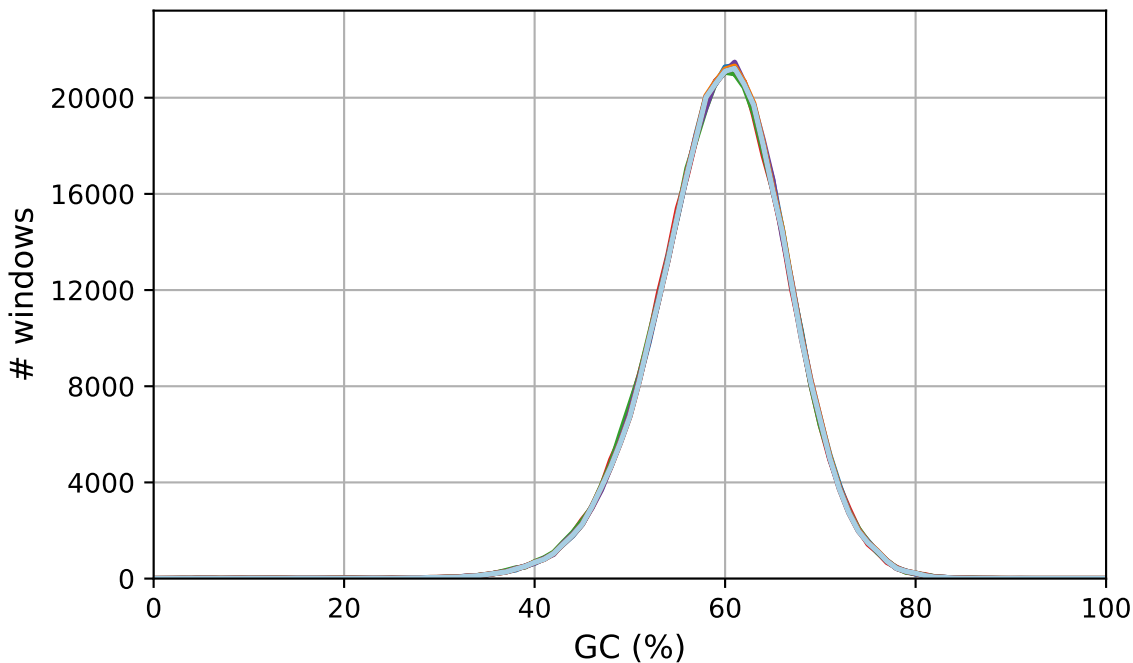
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



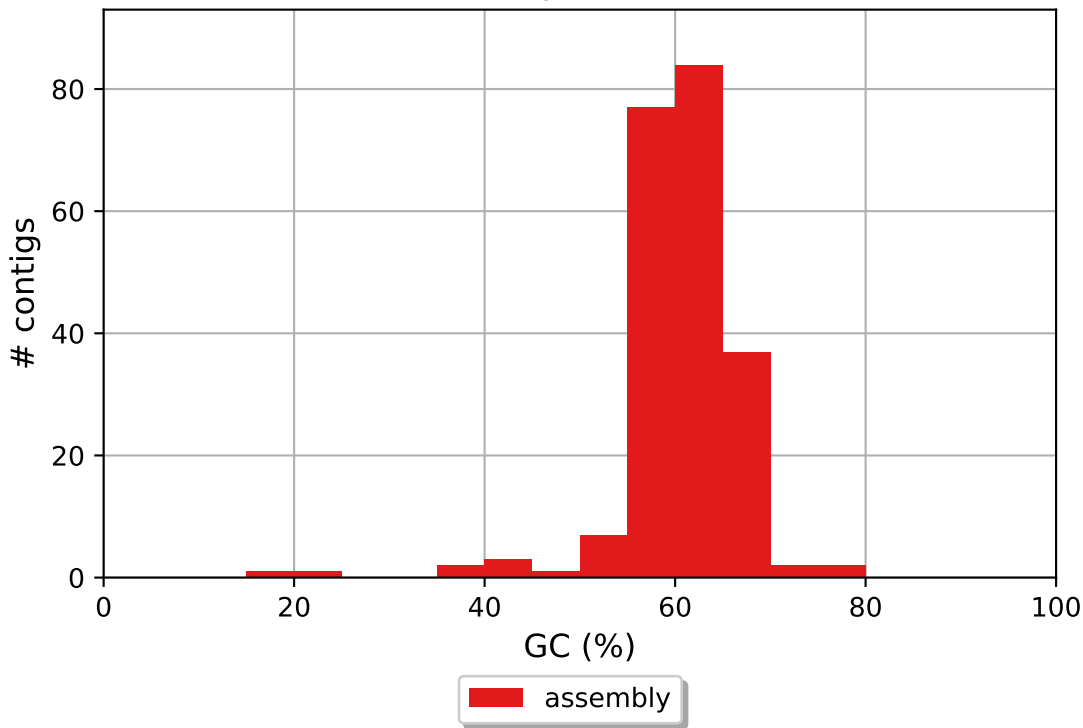
Cumulative length



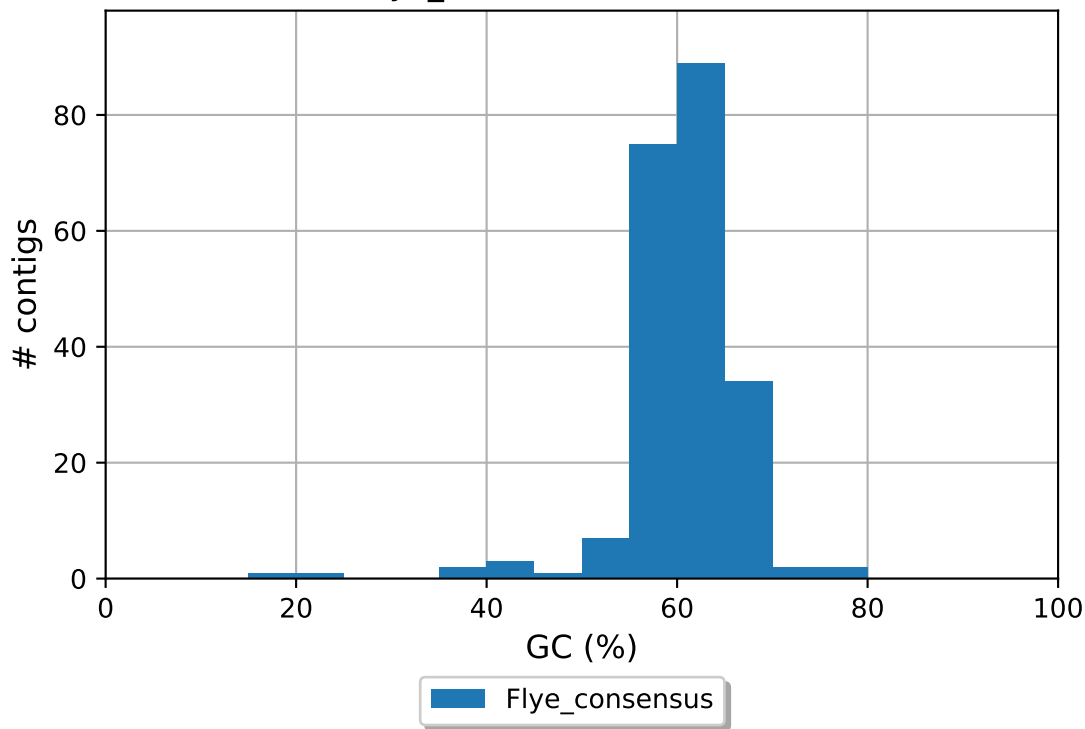
GC content



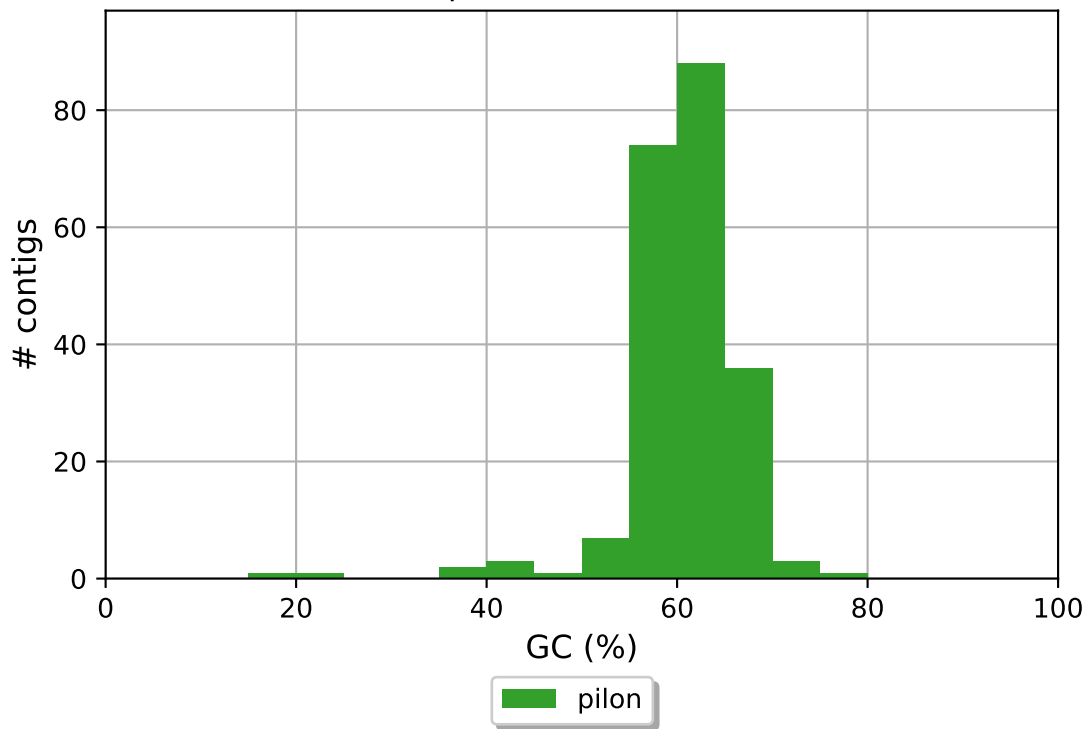
assembly GC content



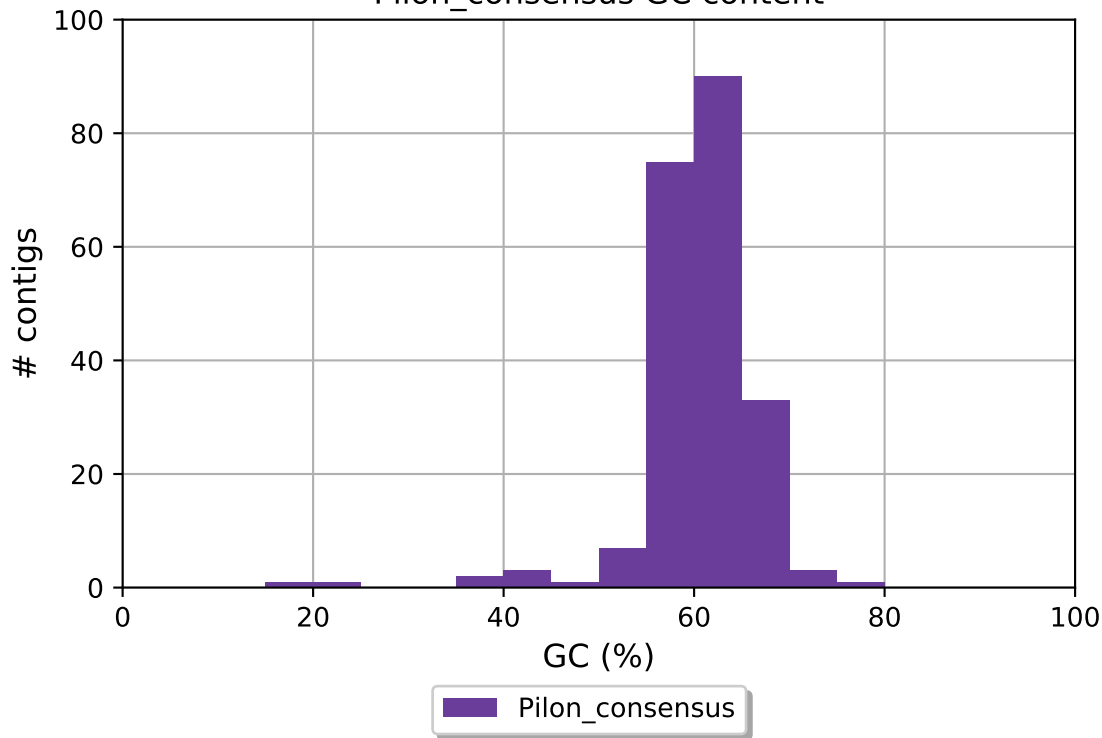
Flye\_consensus GC content



pilon GC content

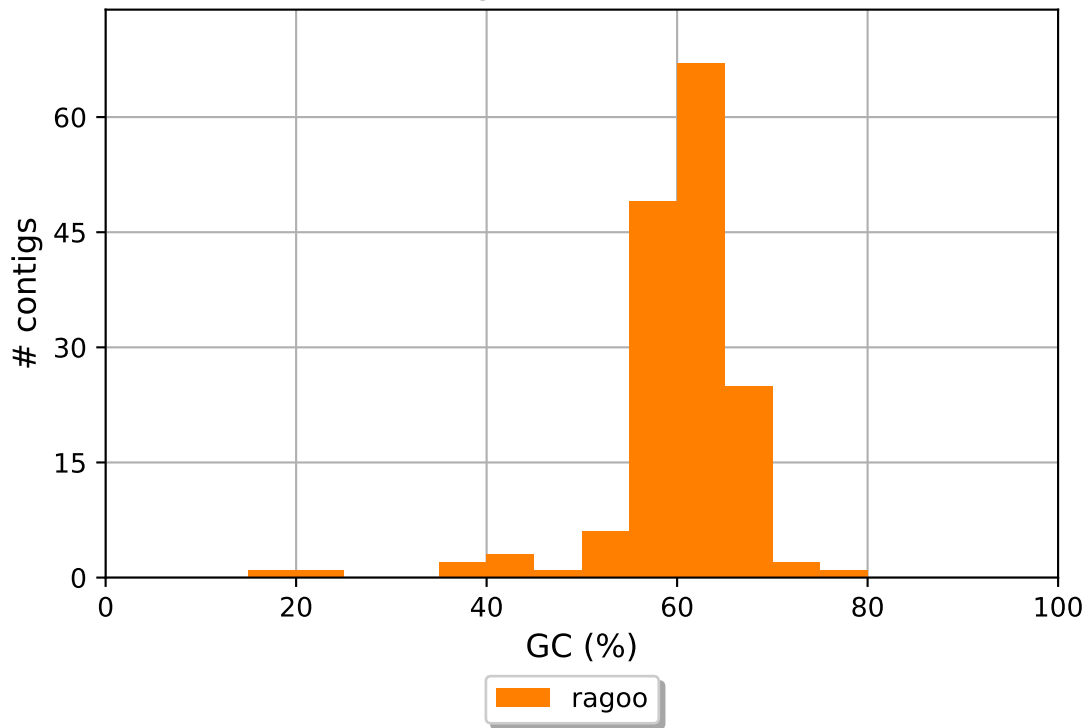


Pilon\_consensus GC content

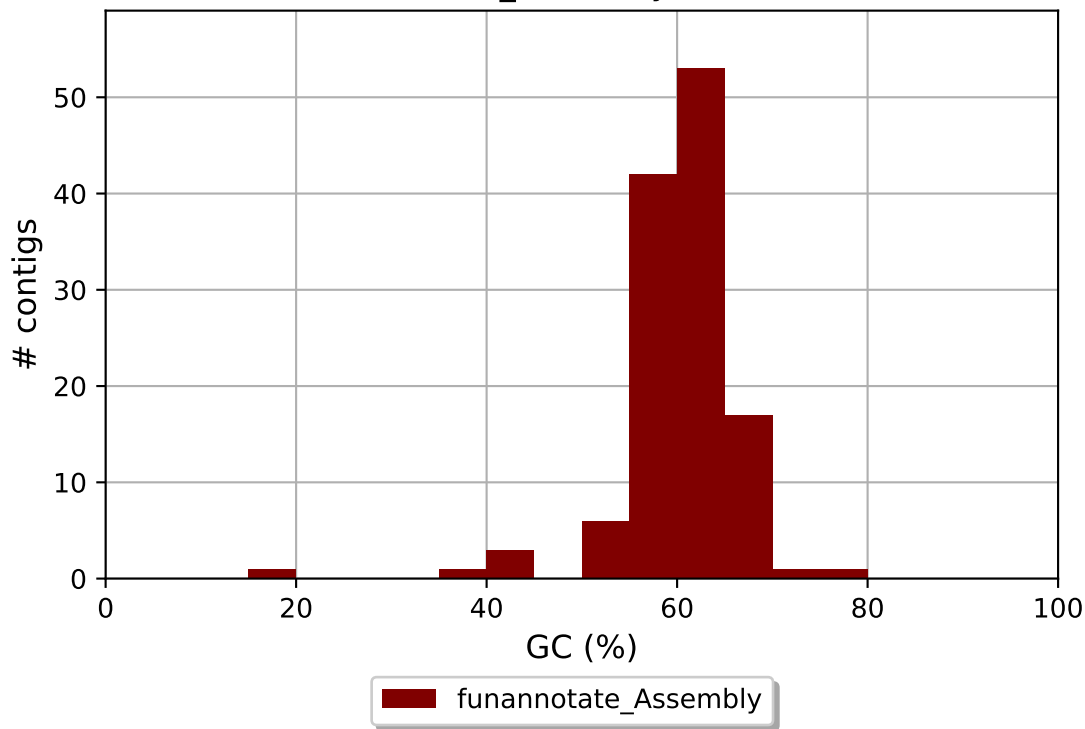




ragoo GC content



funannotate\_Assembly GC content



GH5\_Genome GC content

