cgmlst_ST	cgmlst_distance	cgmlst_found_loci	cgmlst_genome_match	cgmlst_matching_alleles	cgmlst_subspecies
	0.663636364	329	SRR1968150	111	enterica

fasta_filepath	genome	h1	h2	o_antigen
/home/kobina/assemblyall/S1.fasta	S1	c	1,5	1,4,12,[27]

qc_messages
INFO: Number of cgMLST330 loci found (n=329) | WARNING: Only matched 111 cgMLST330 loci. Min threshold for confident serovar prediction from cgMLST is 297.0

qc_status	serogroup	serovar	serovar_antig	serovar_cgmlst		
WARNING	B	Legon	Legon	Fanti		