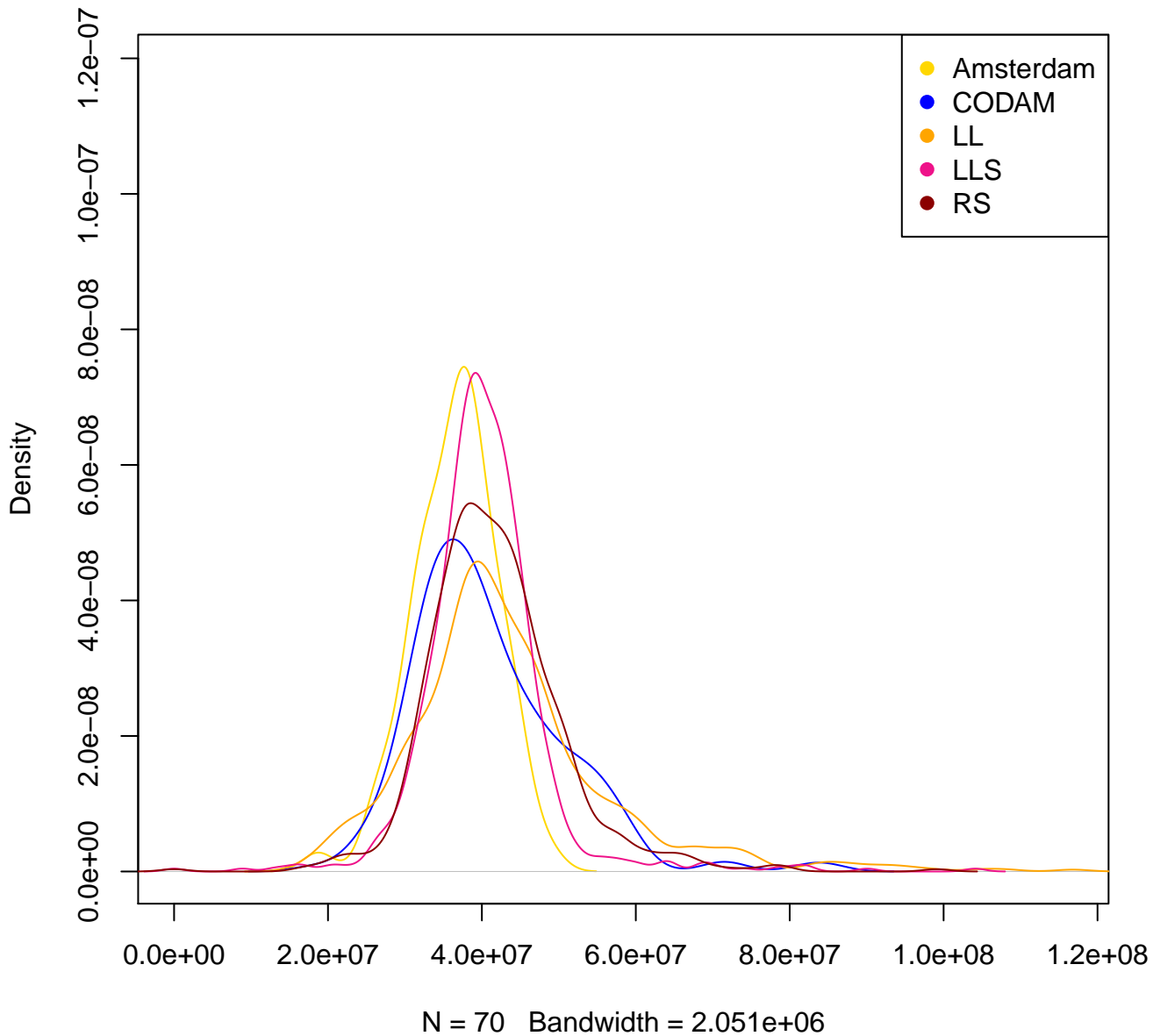
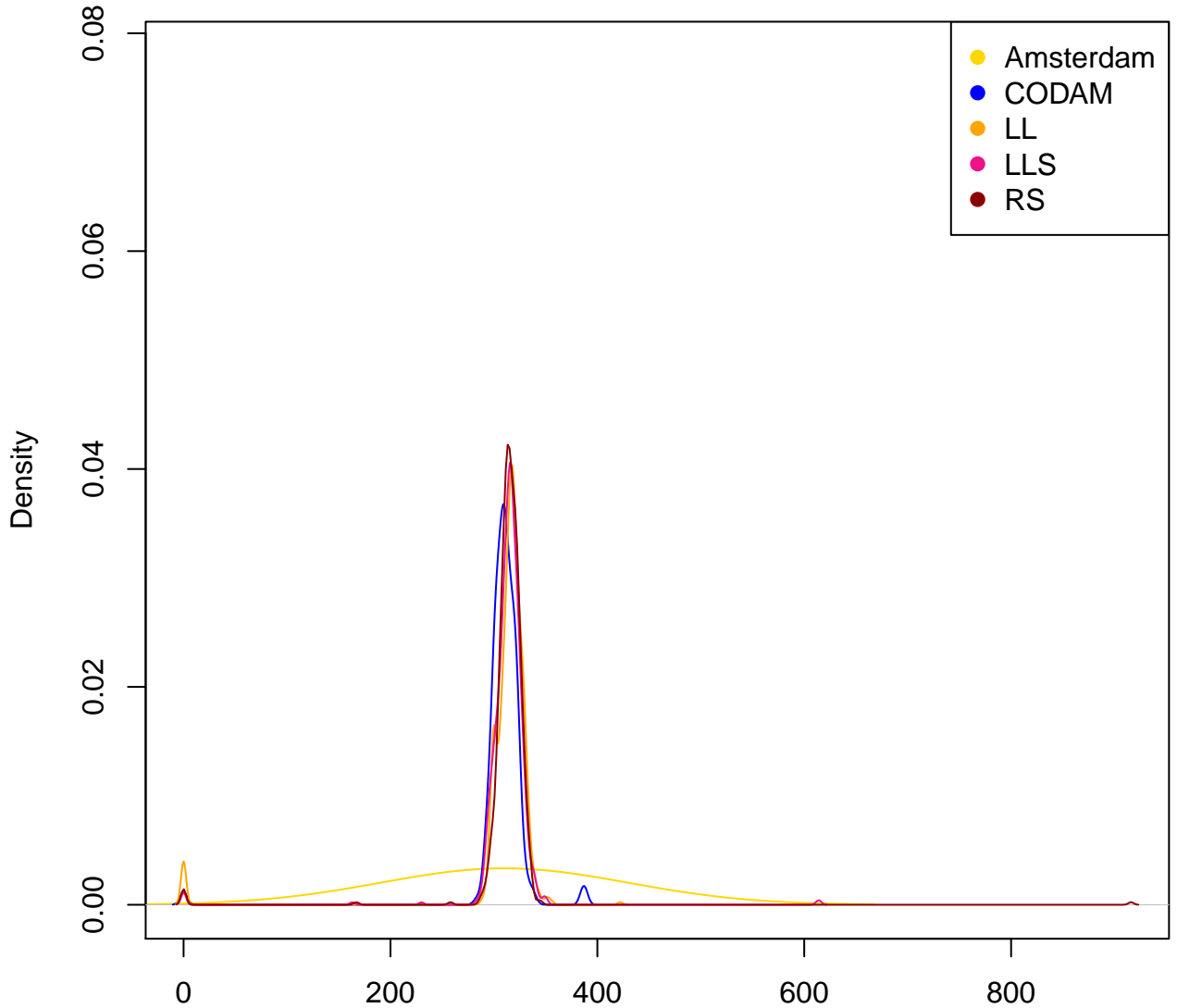


num_past_filter

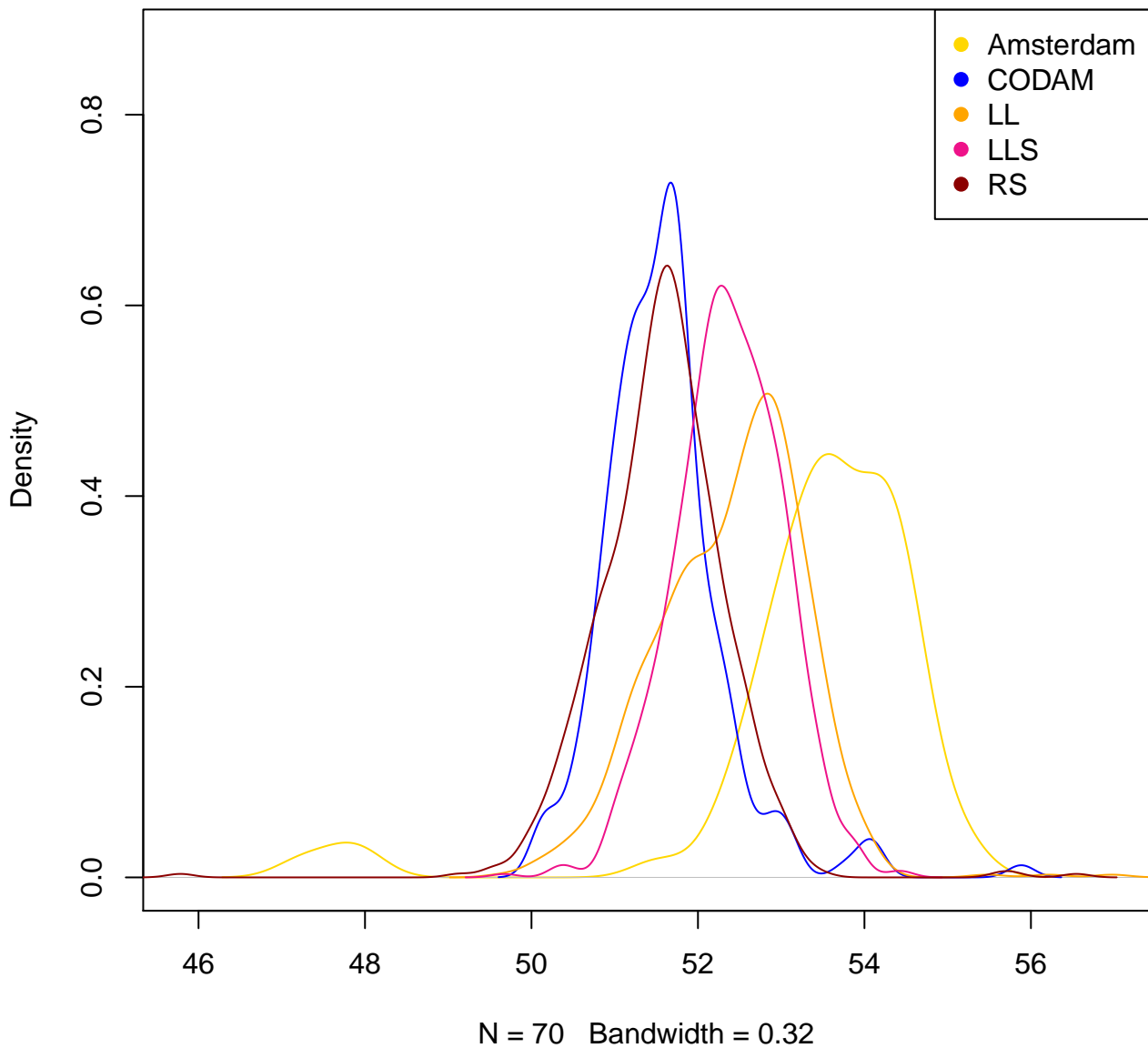


insert_size

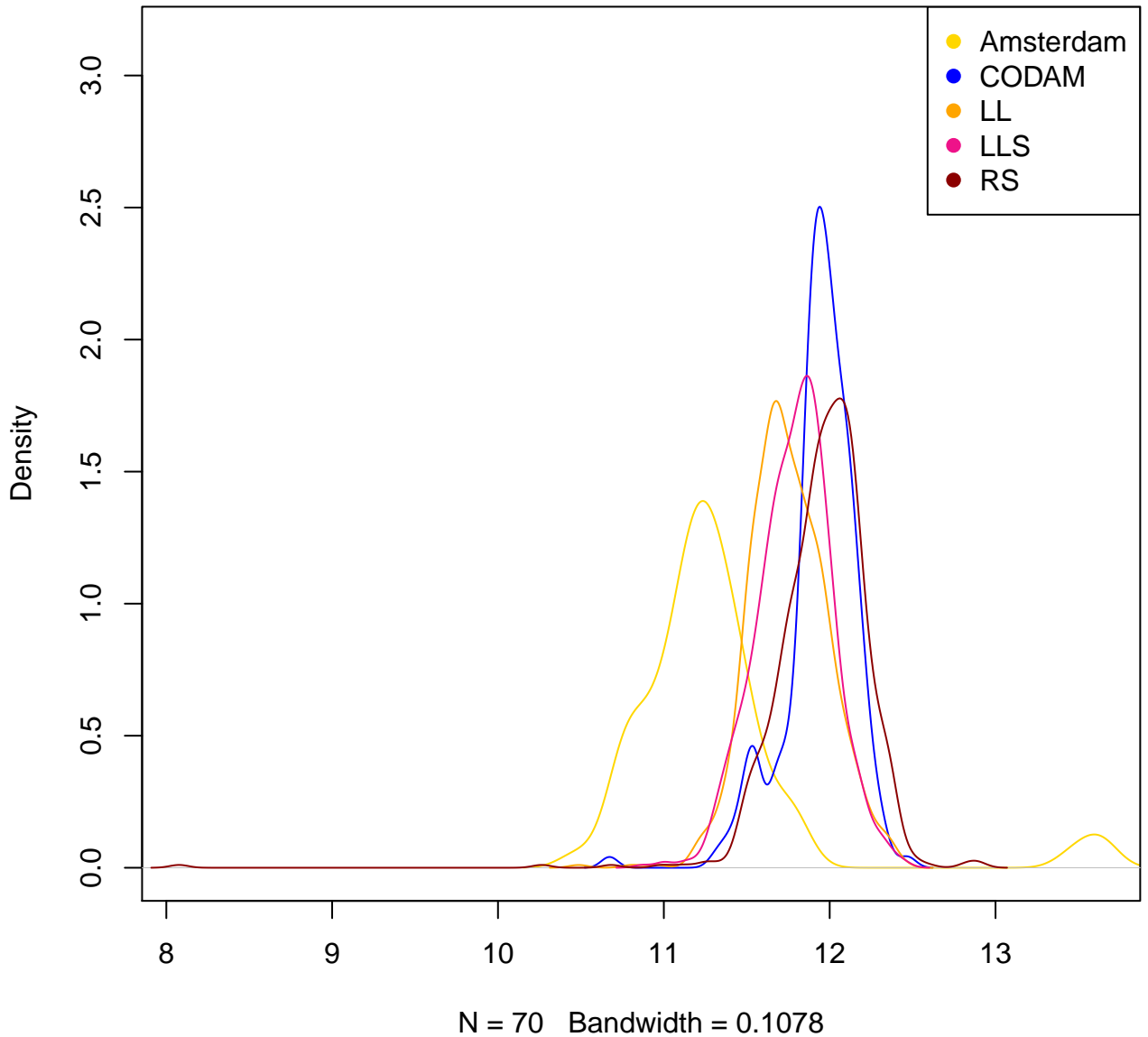


N = 70 Bandwidth = 119.3

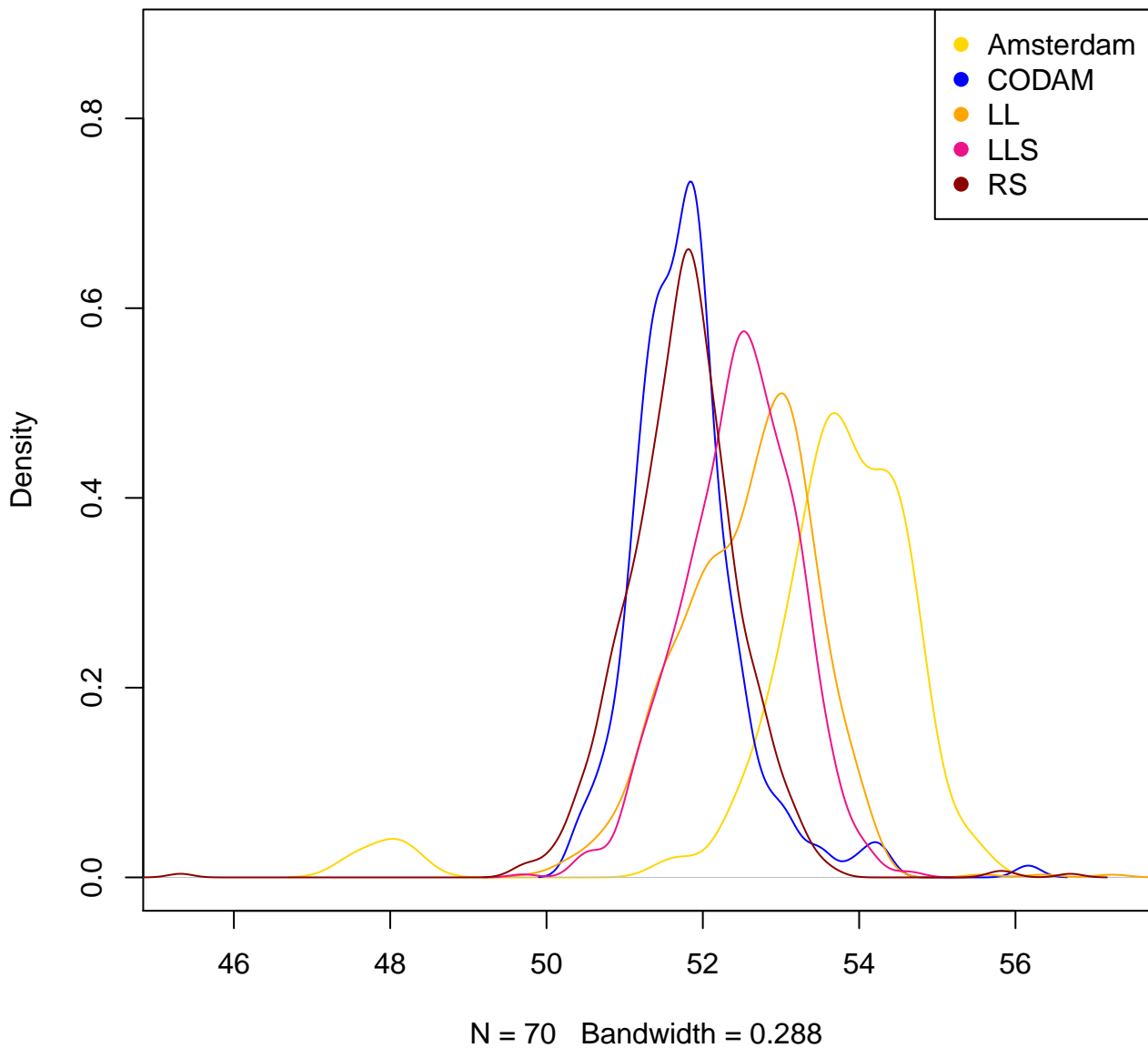
fastqc_clean_R1_clean_GC_mean



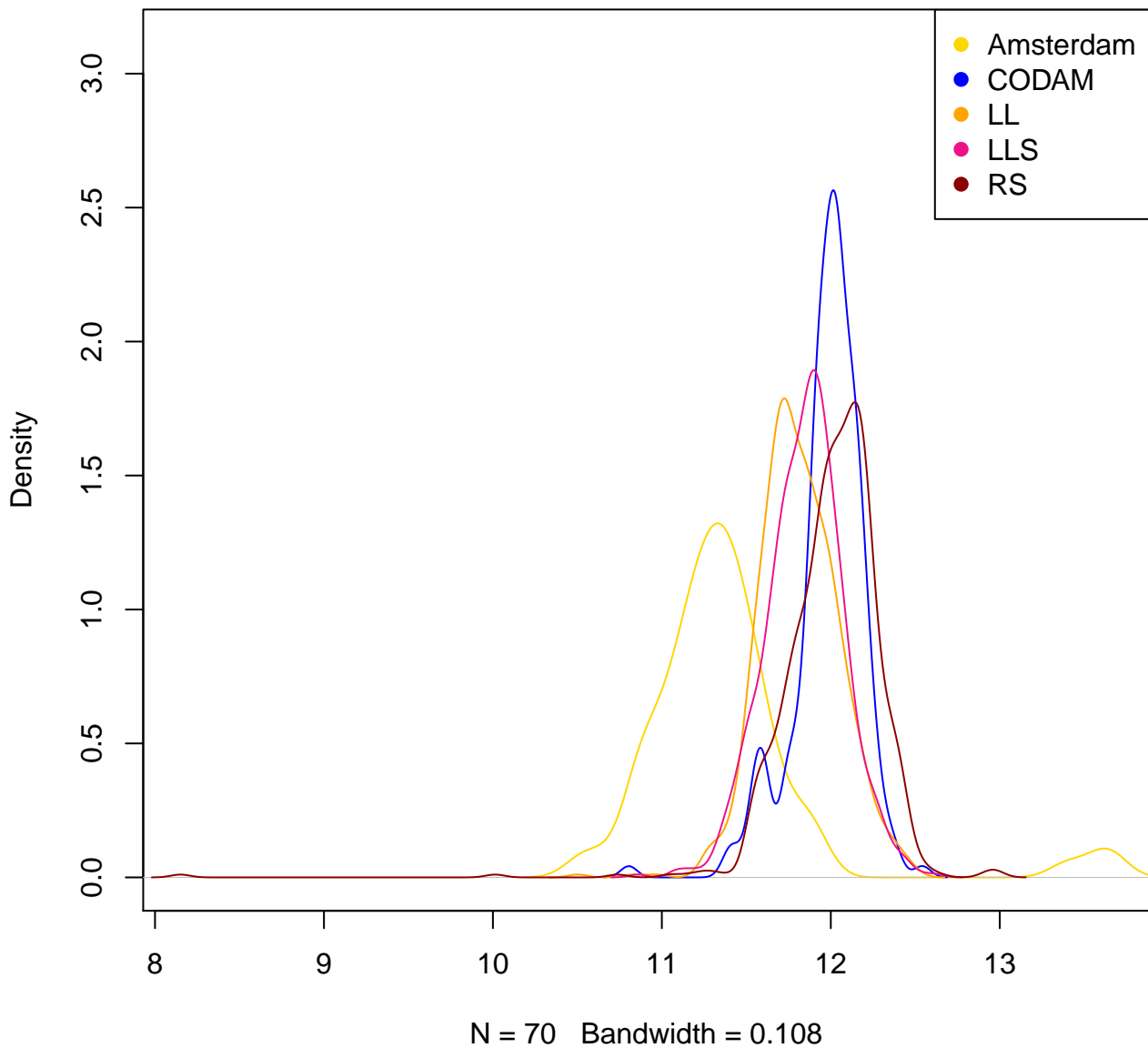
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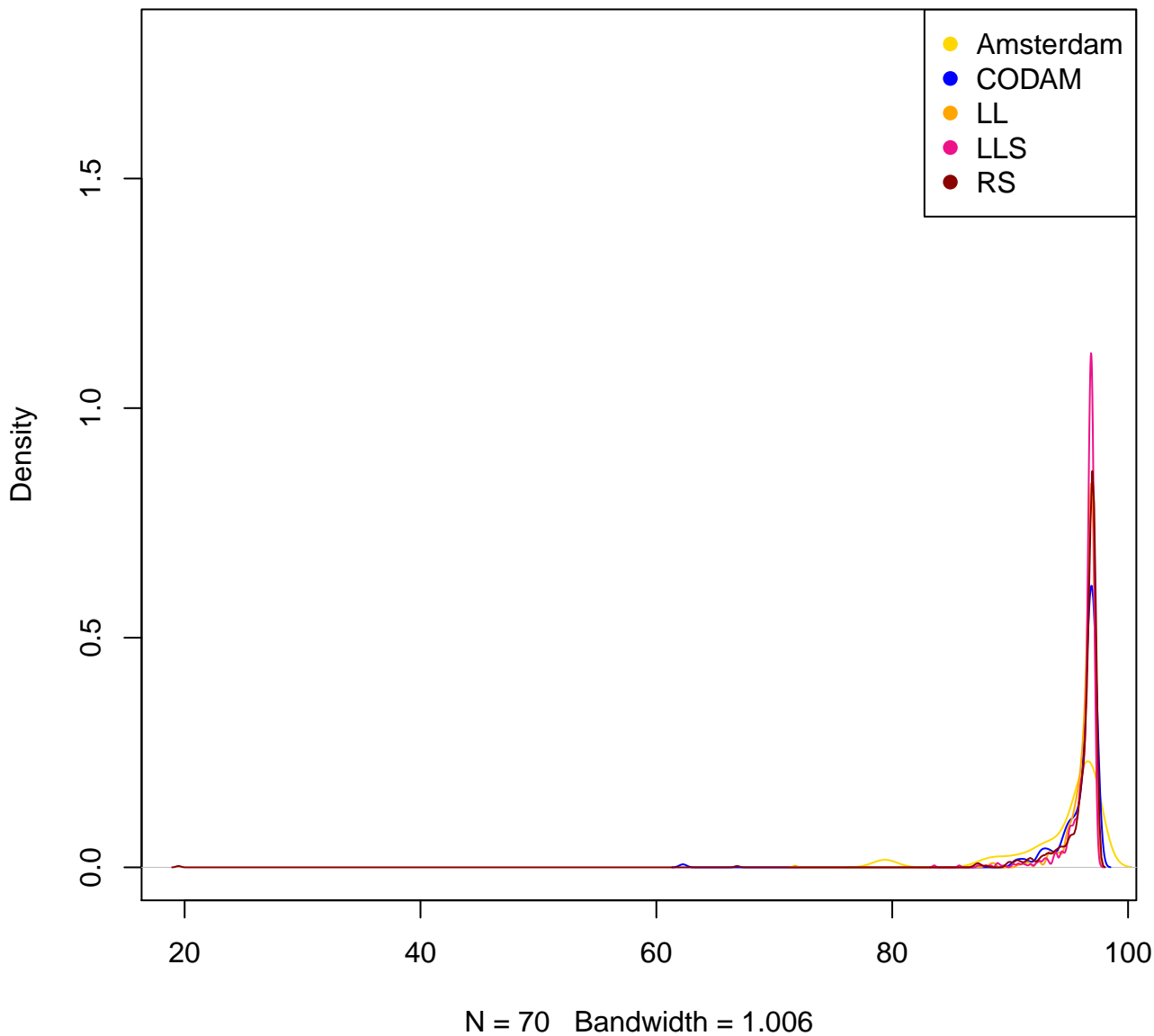
fastqc_clean_R2_clean_GC_mean



fastqc_clean_R2_clean_GC_std



bam_perc_genome_mapped



bam_perc_exon_mapped

