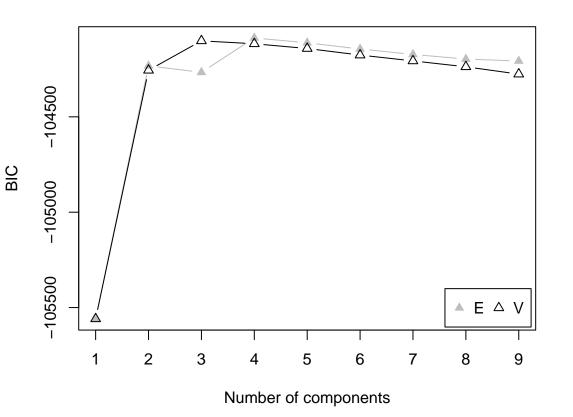
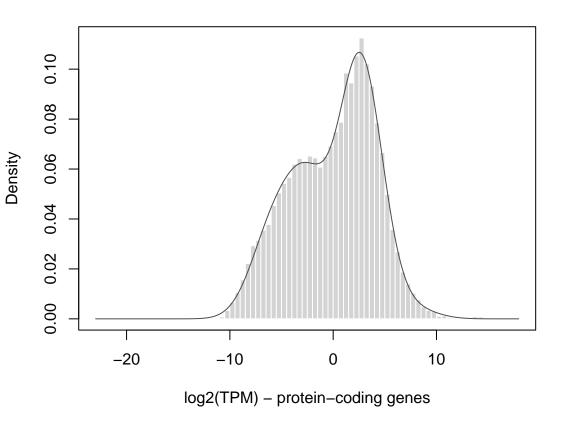
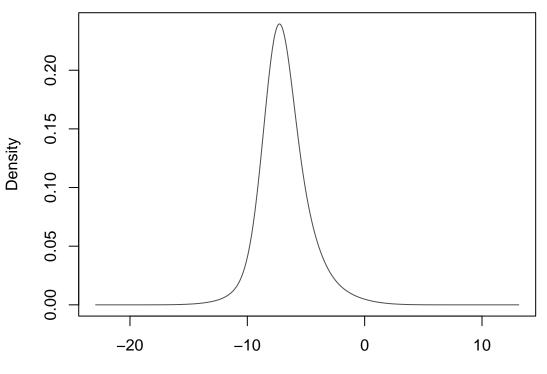


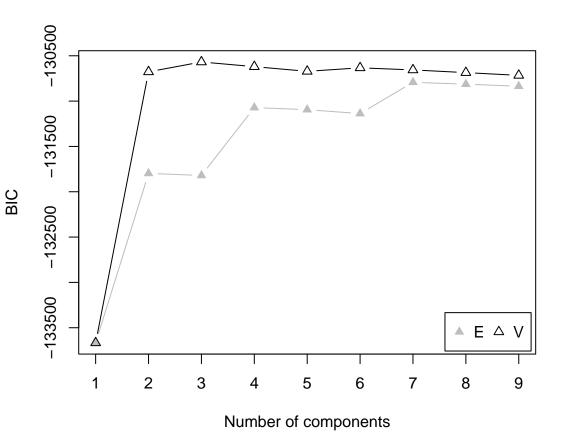
log2(summed_filtered\$tpm[summed_filtered\$biotype %in% "protein_coding"]

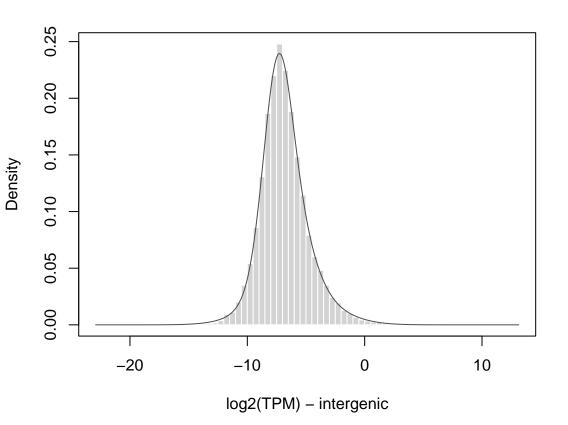




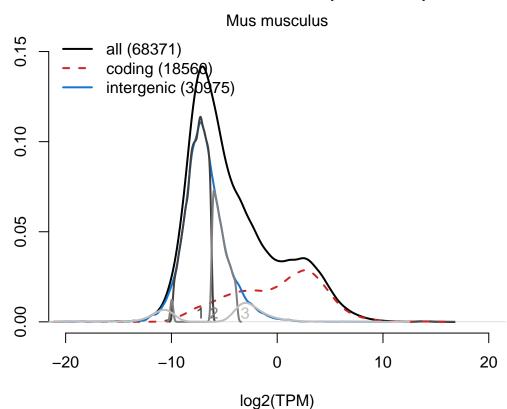


log2(summed_filtered\$tpm[summed_filtered\$type == "intergenic"])





skeletal muscle tissue (6 libraries)



Density