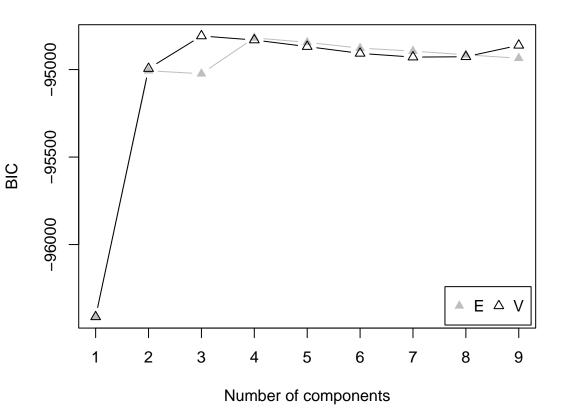
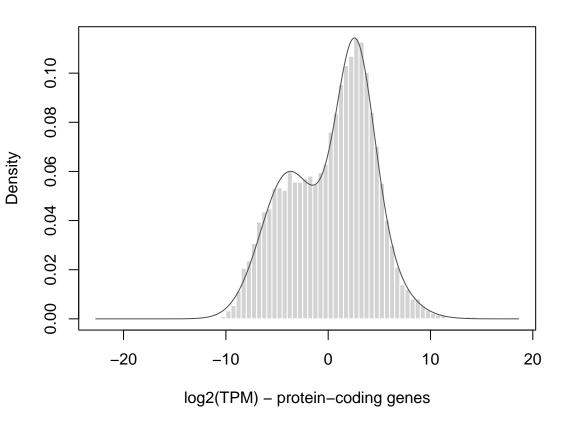
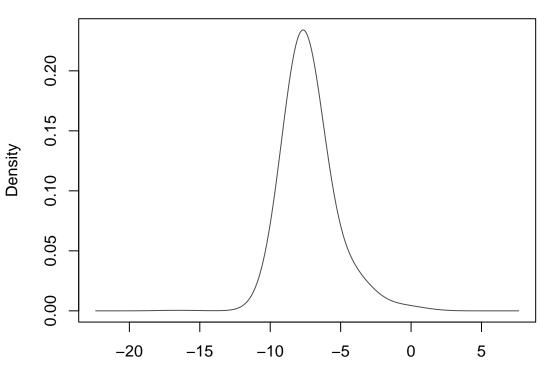


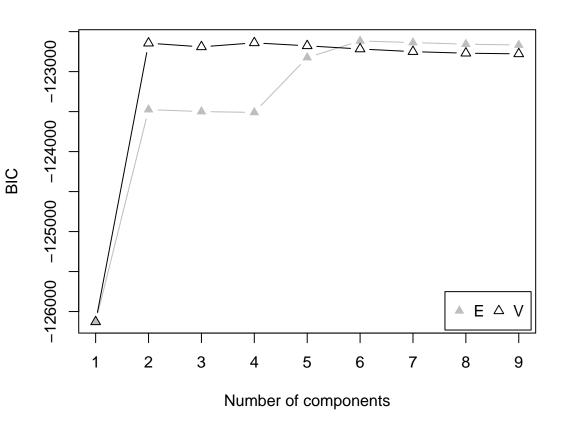
log2(summed\_filtered\$tpm[summed\_filtered\$biotype %in% "protein\_coding"]

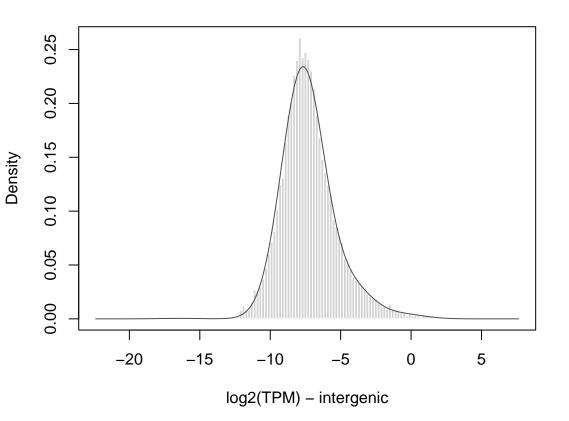






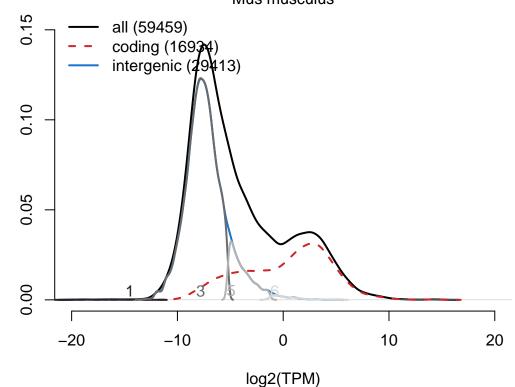
log2(summed\_filtered\$tpm[summed\_filtered\$type == "intergenic"])





## muscle tissue (5 libraries)





Density