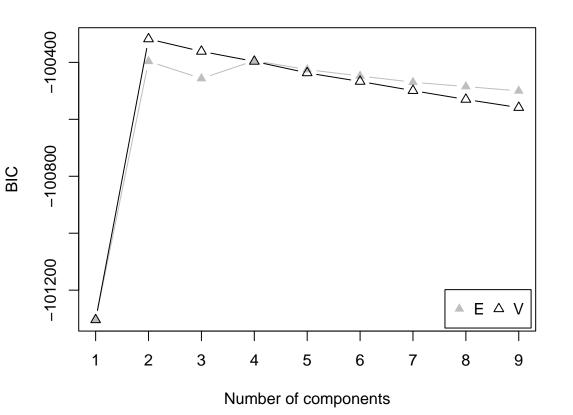
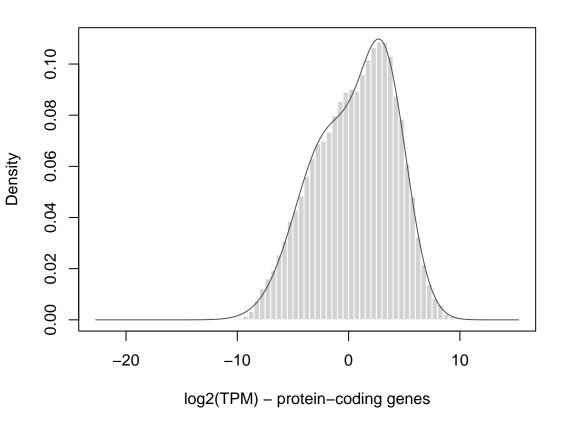
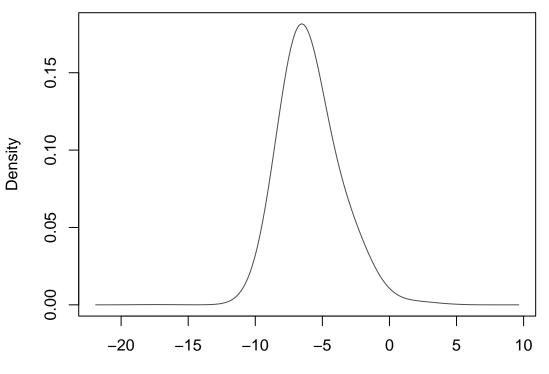


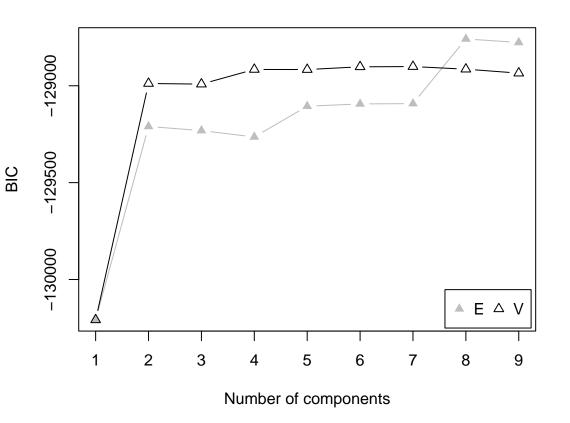
log2(summed_filtered\$tpm[summed_filtered\$biotype %in% "protein_coding"]

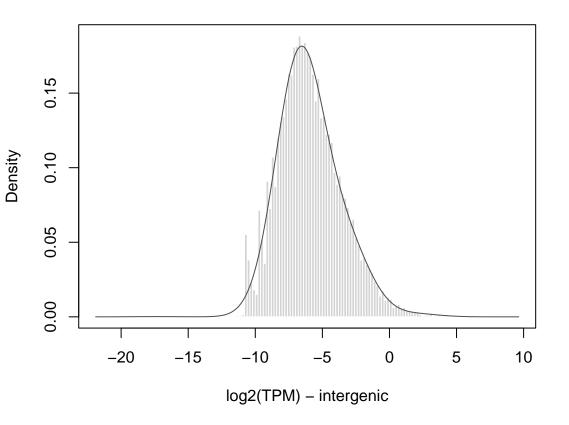






log2(summed_filtered\$tpm[summed_filtered\$type == "intergenic"])





spermatocyte (1 libraries)

Mus musculus

