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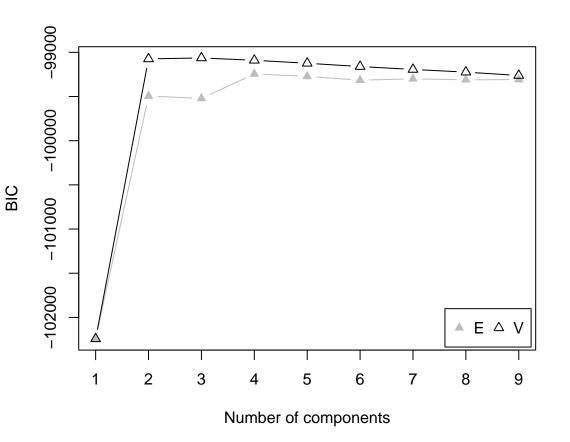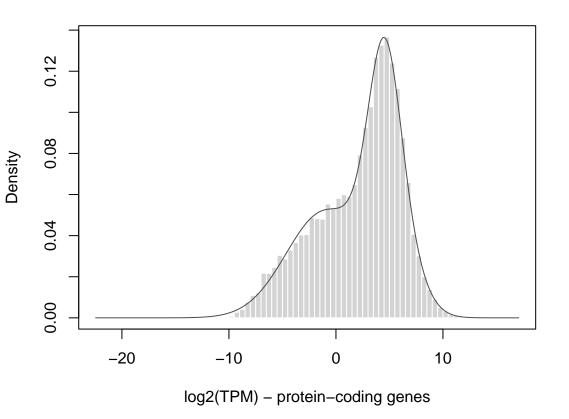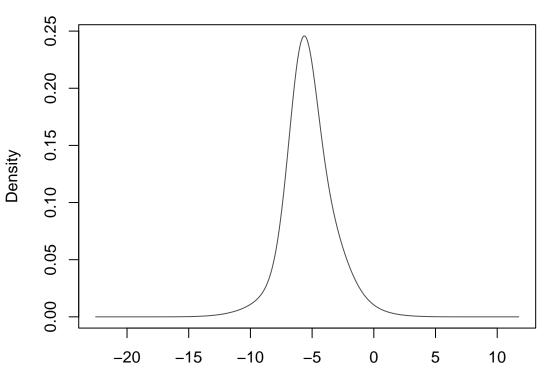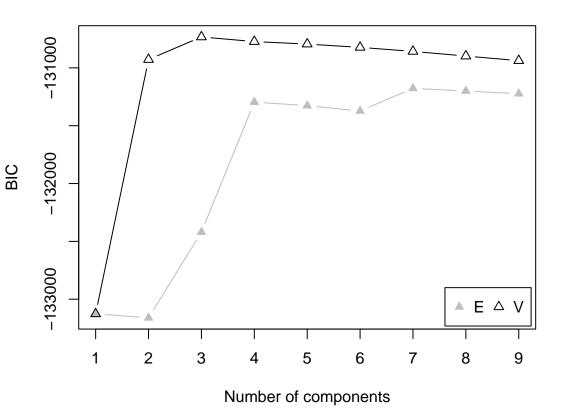
log2(summed_filtered$tpm[summed_filtered$biotype %in% "protein_coding"]

Density

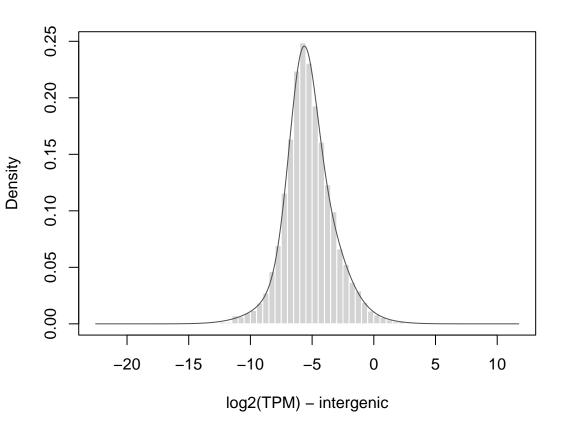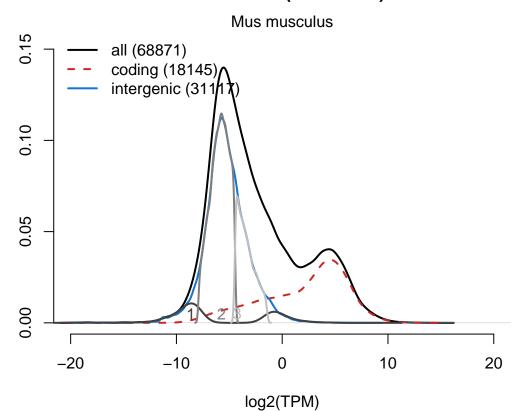log2(summed_filtered$tpm[summed_filtered$type == "intergenic"])

**neural tube (6 libraries)**

Mus musculus

all (68871)
coding (18145)
intergenic (31117)

log2(TPM)

Density