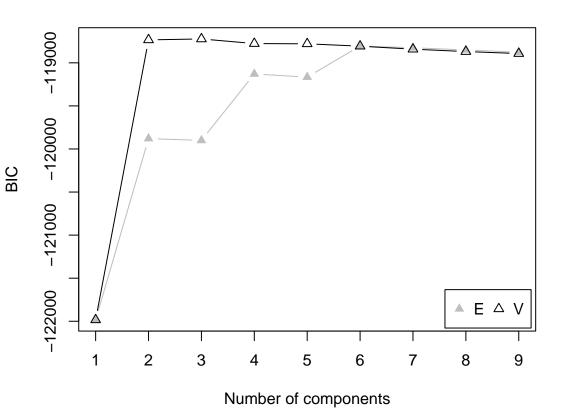
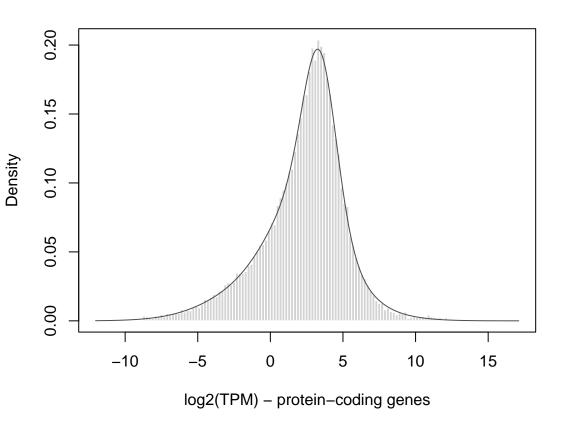
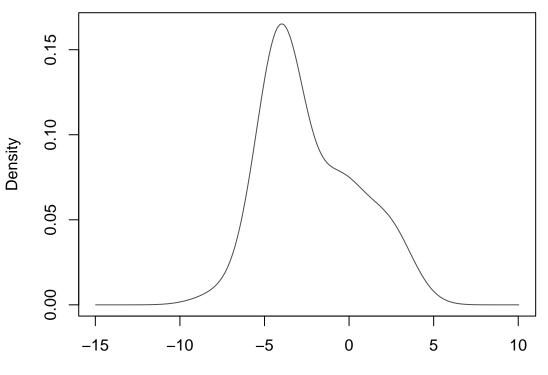


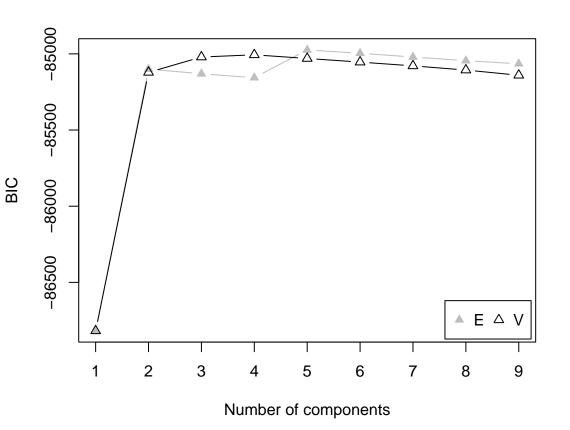
 $log2 (summed_filtered\$tpm[summed_filtered\$biotype \% in\% "protein_coding"]$

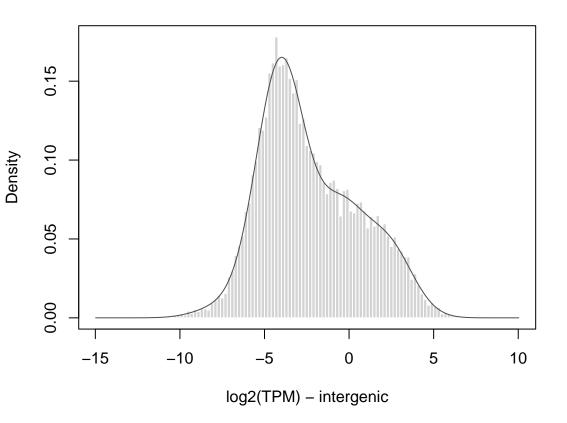






log2(summed_filtered\$tpm[summed_filtered\$type == "intergenic"])





Esox lucius (24 libraries)

