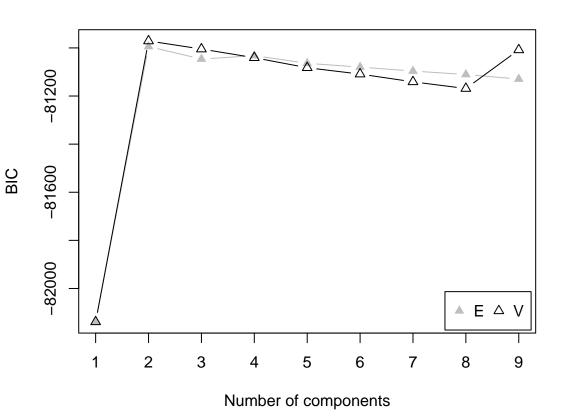
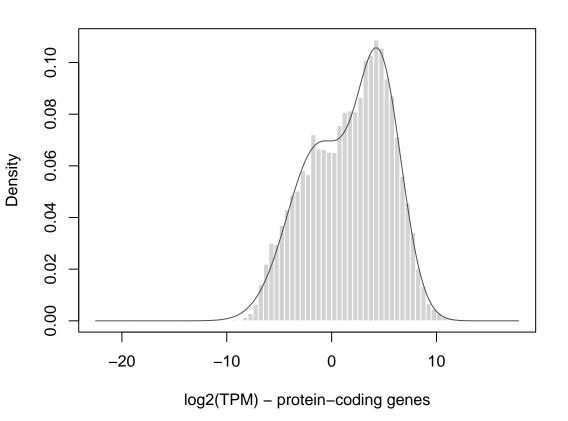
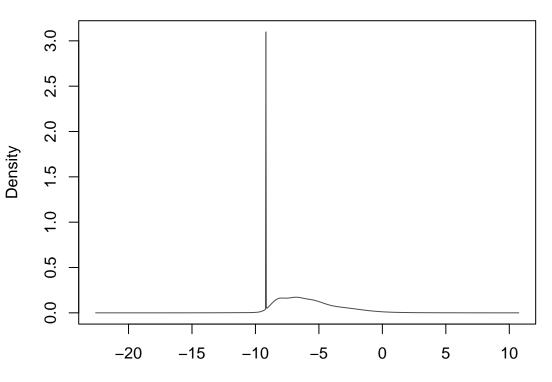


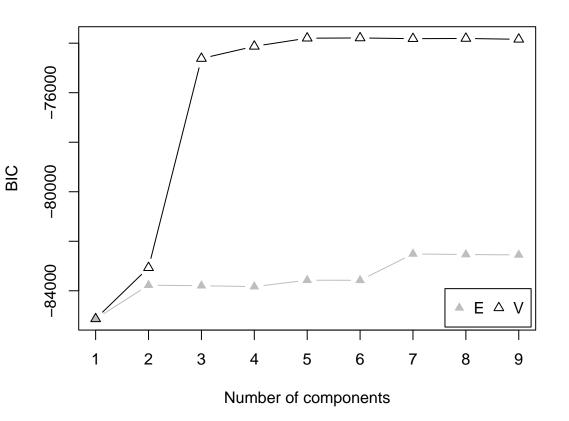
log2(summed_filtered\$tpm[summed_filtered\$biotype %in% "protein_coding"]

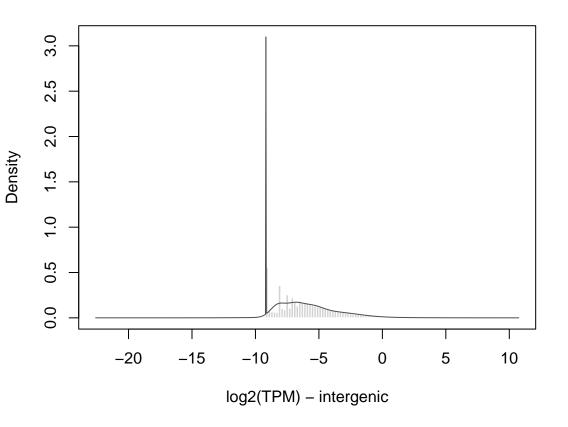






log2(summed_filtered\$tpm[summed_filtered\$type == "intergenic"])





secondary oocyte (3 libraries)

