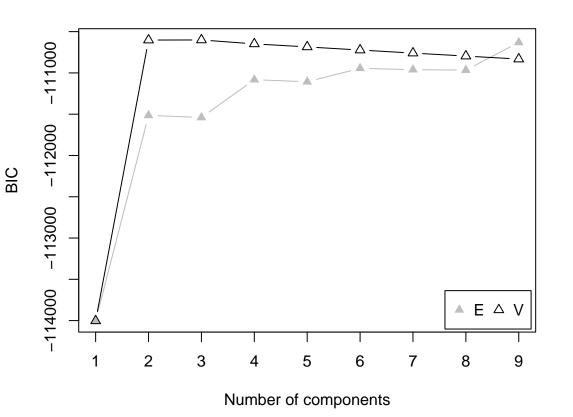
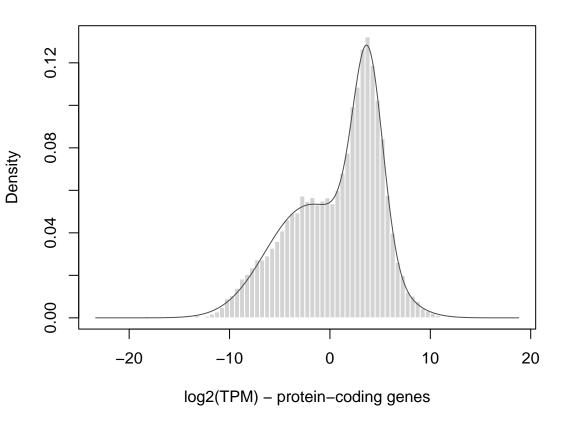
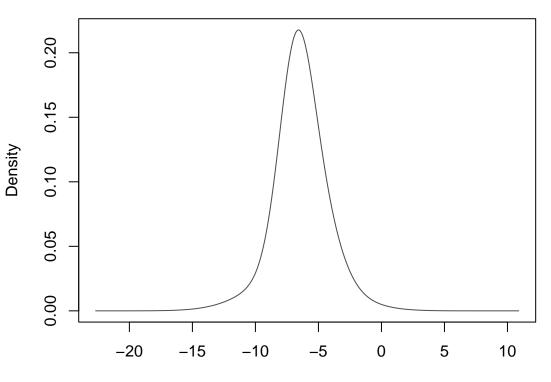


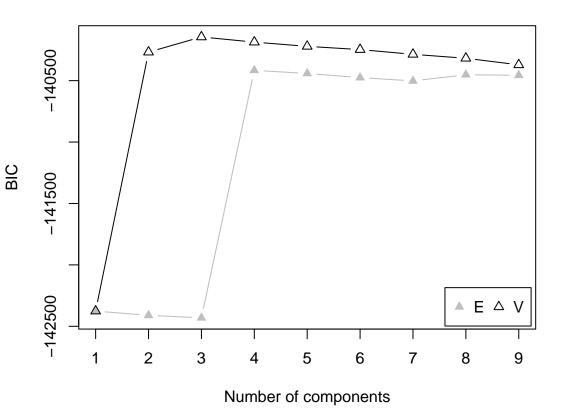
 $log2 (summed_filtered\$tpm[summed_filtered\$biotype~\%in\%~"protein_coding"]$

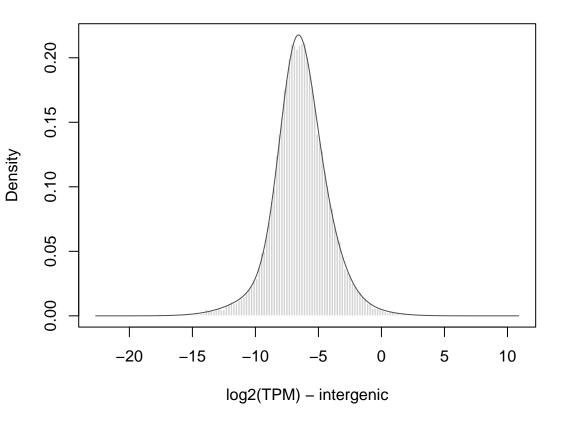






log2(summed_filtered\$tpm[summed_filtered\$type == "intergenic"])





heart (29 libraries)

