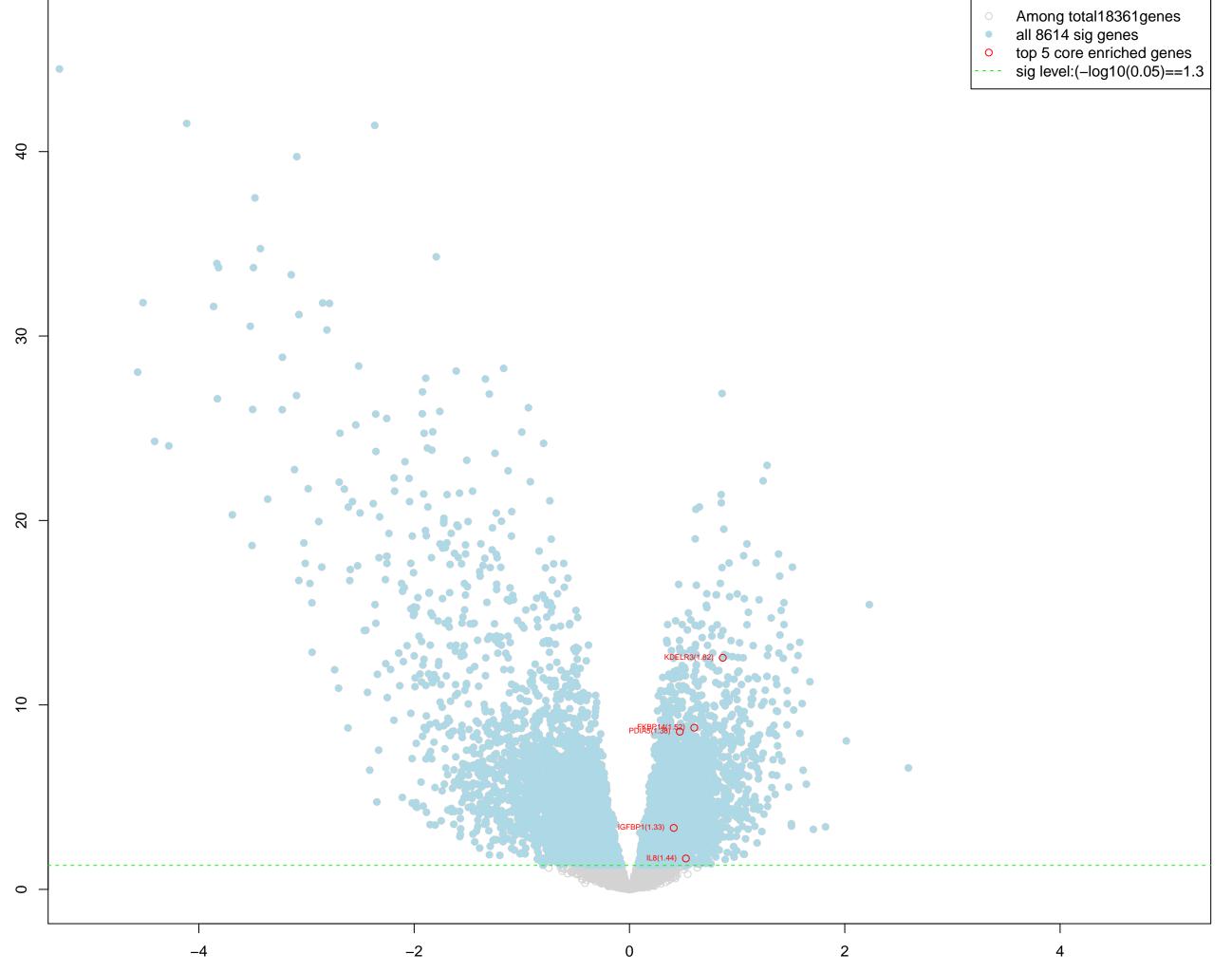
clus1-vs-NON.clus1 in HNSC-TP RNASeq data



significance of gene : [-log10(ebayes_fit\$p.val)]