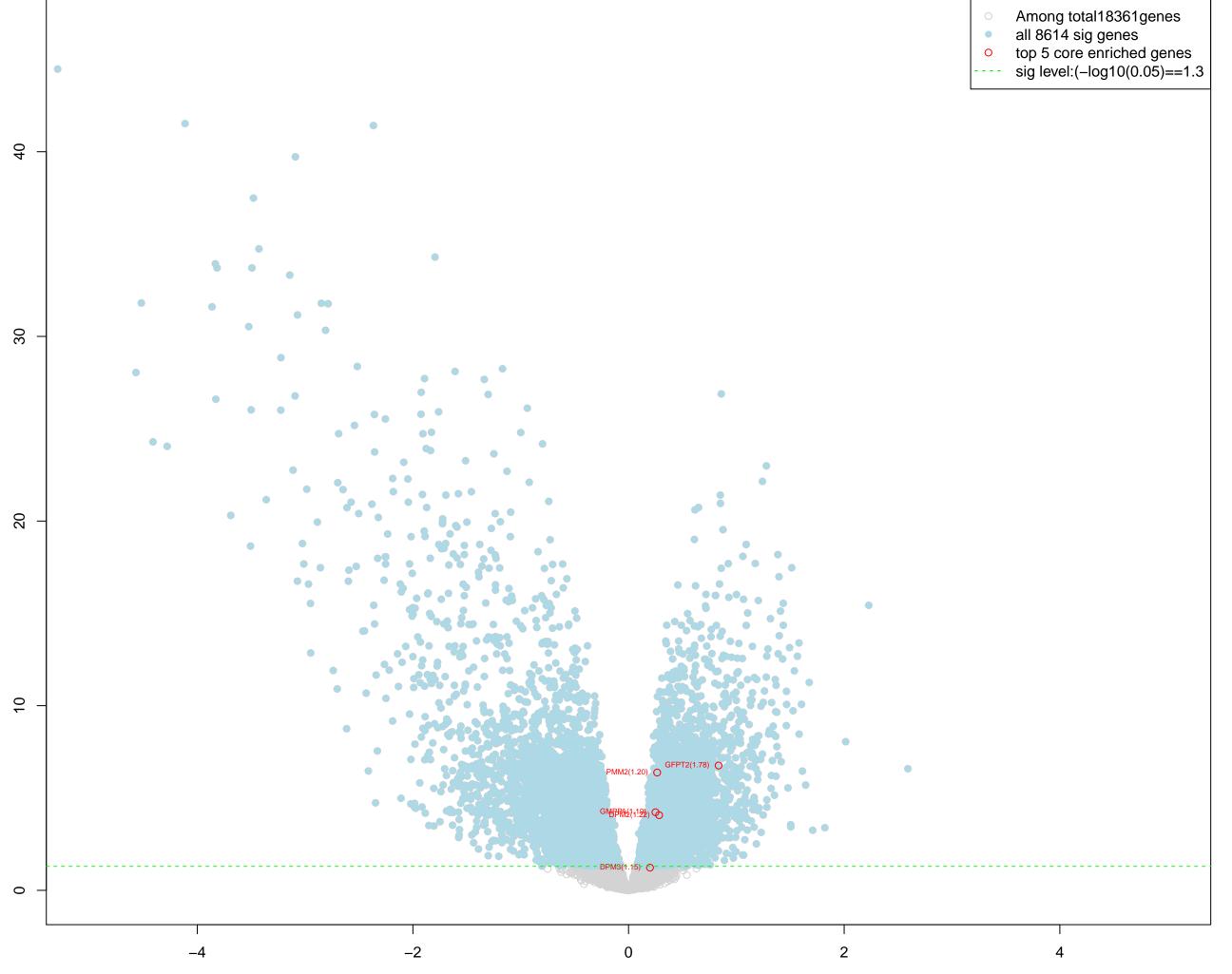
clus1-vs-NON.clus1 in HNSC-TP RNASeq data



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