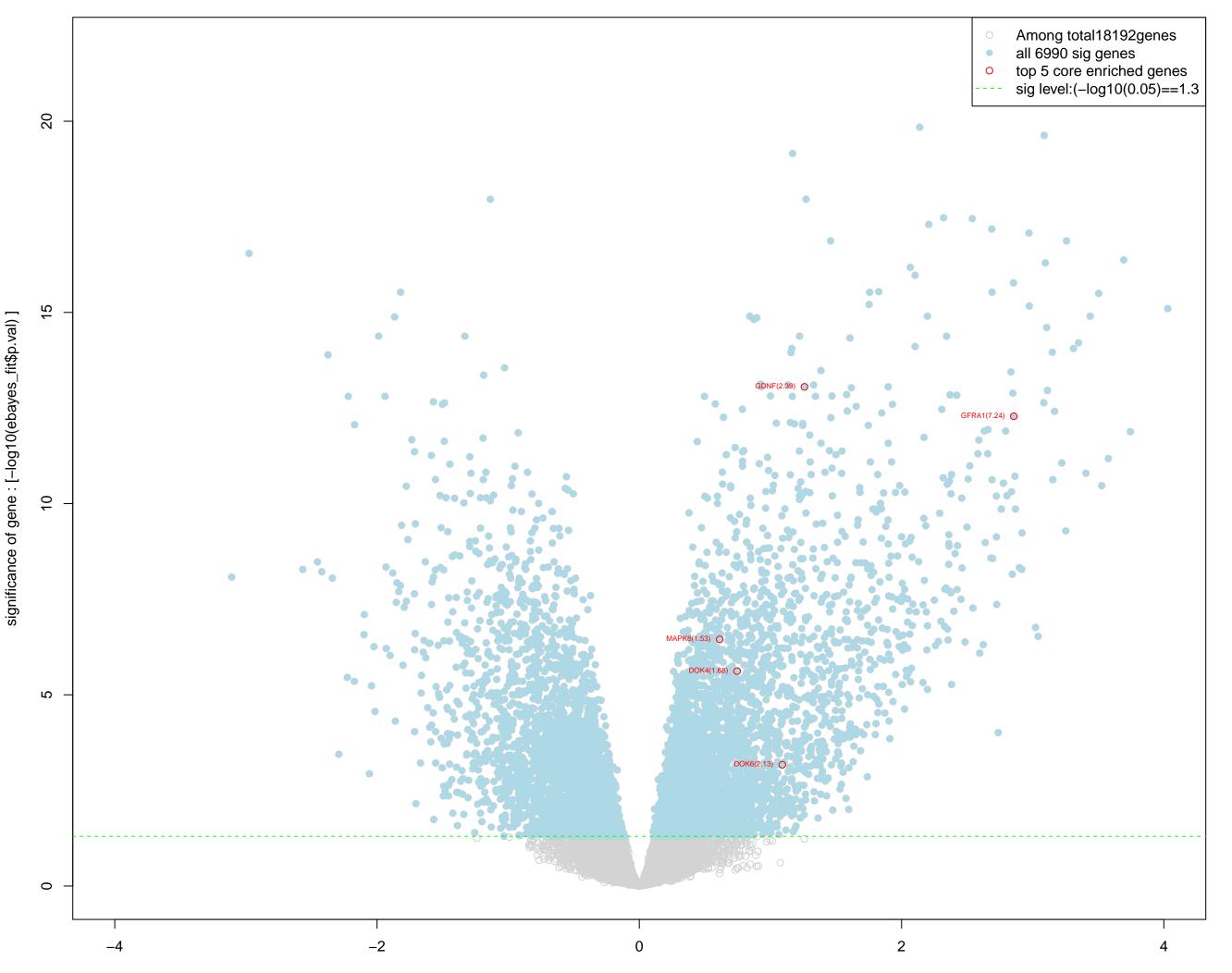
clus1-vs-NON.clus1 in GBM-TP RNASeq data



log2 fold change in expression level : [clus1-vs-NON.clus1 ]