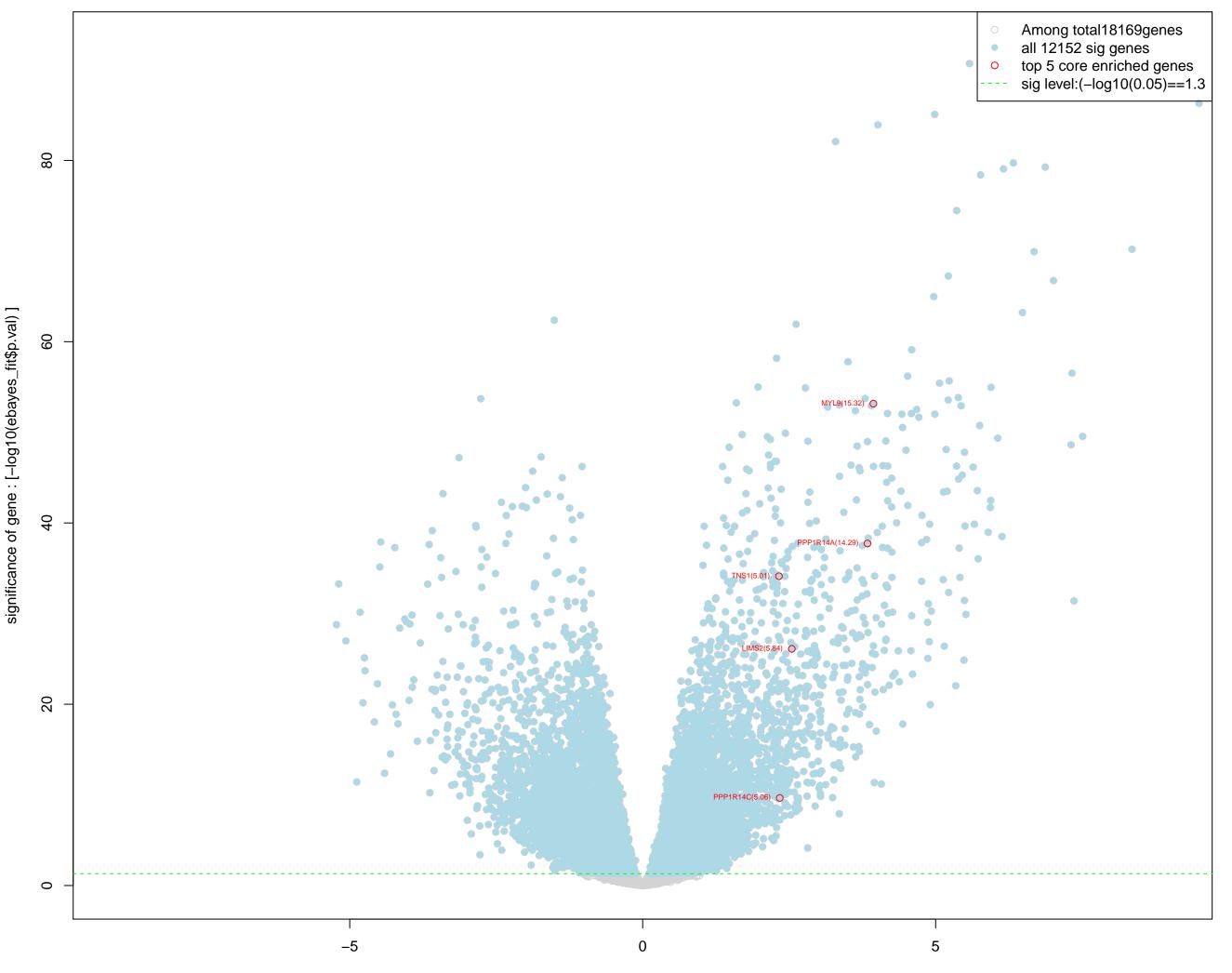
clus2-vs-NON.clus2 in SARC-TP RNASeq data



log2 fold change in expression level : [clus2-vs-NON.clus2 ]