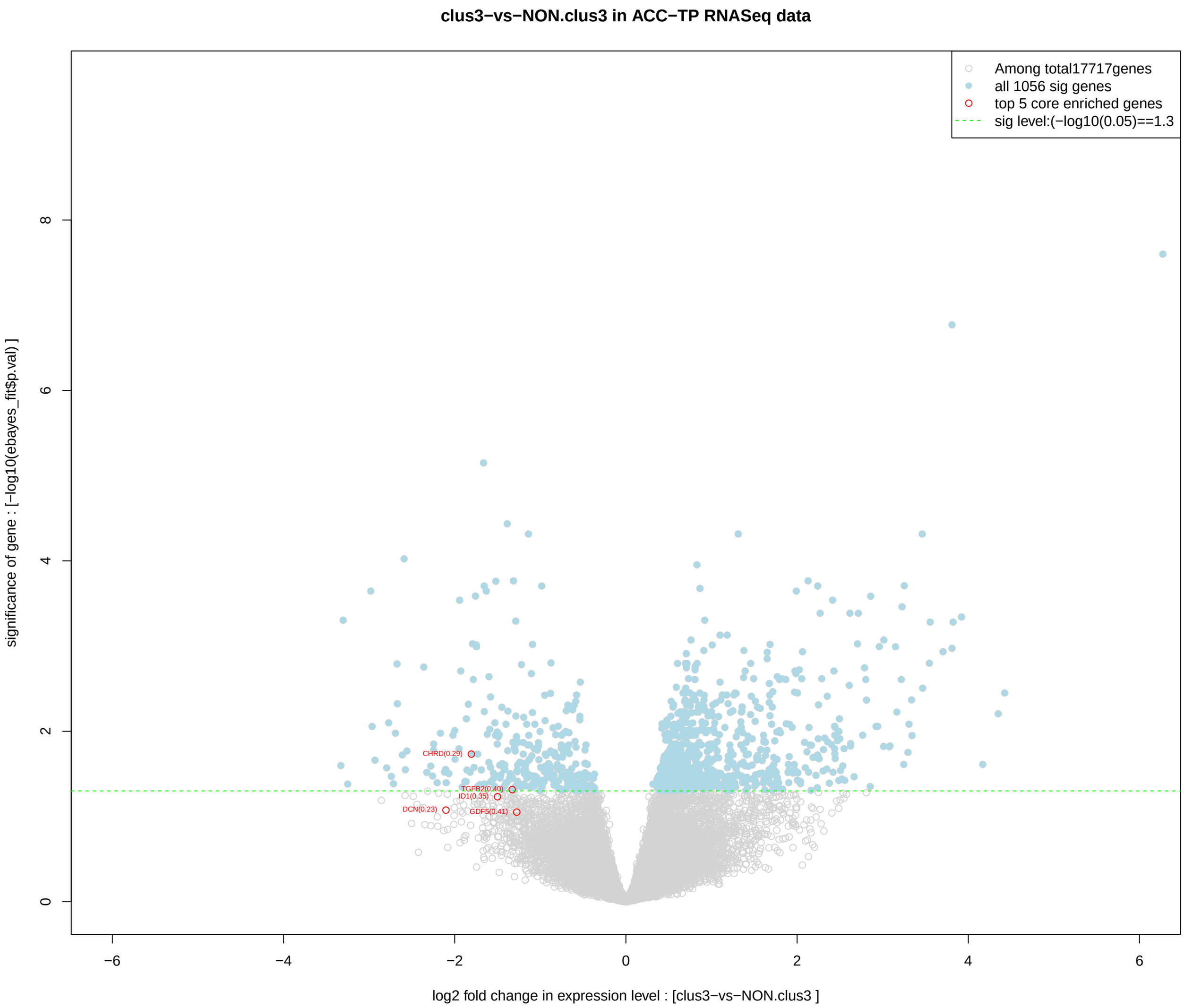
clus3−vs−NON.clus3 in ACC−TP RNASeq data
Among total17717genes
all 1056 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
CHRD(0.29)
TGFB2(0.40)
ID1(0.35)
DCN(0.23)
GDF5(0.41)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus3−vs−NON.clus3]
0
2
4
6
8
−6
−4
−2
0
2
4
6