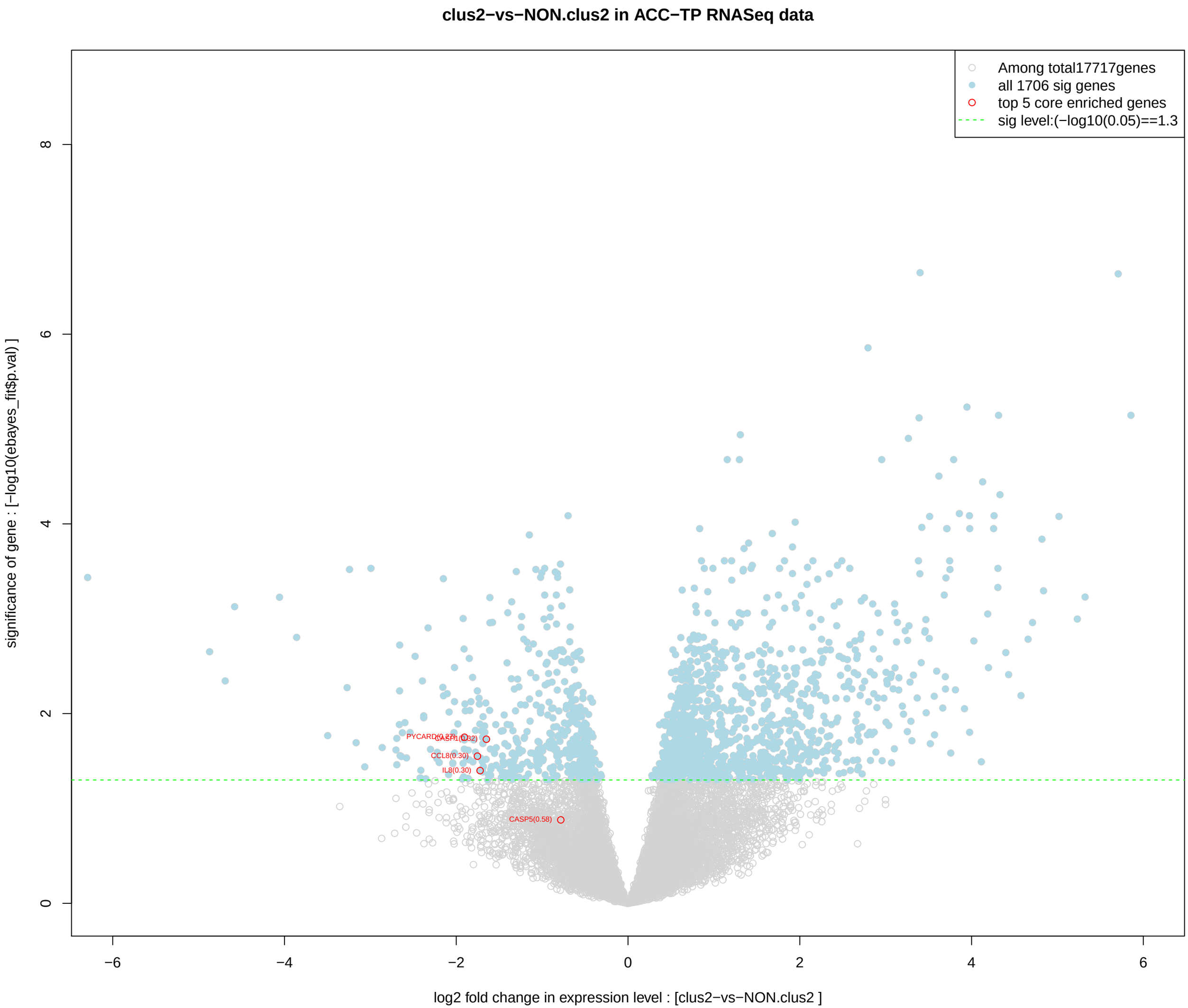
clus2−vs−NON.clus2 in ACC−TP RNASeq data
Among total17717genes
all 1706 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
PYCARD(0.27)
CASP1(0.32)
CCL8(0.30)
IL8(0.30)
CASP5(0.58)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus2−vs−NON.clus2]
0
2
4
6
8
−6
−4
−2
0
2
4
6