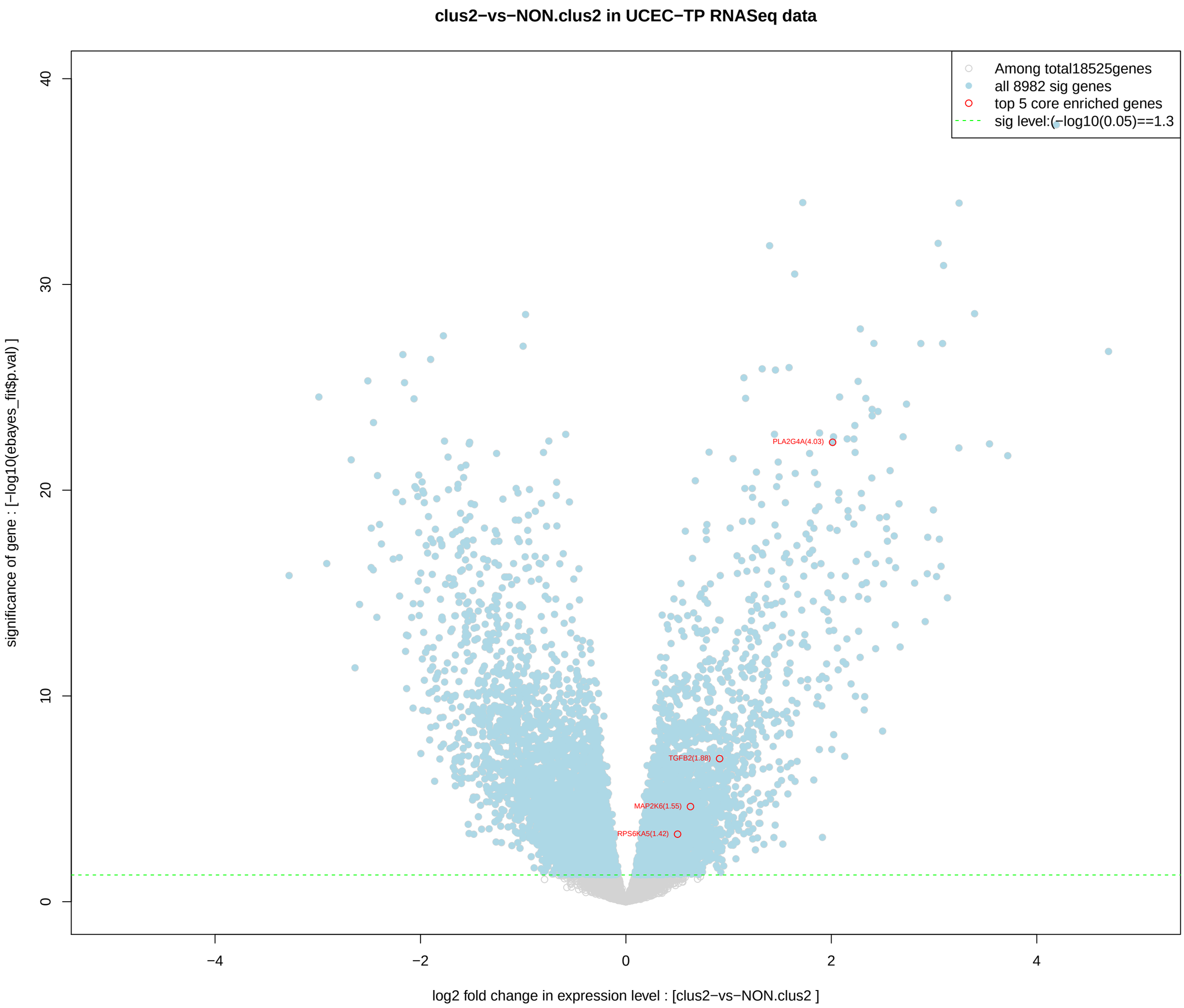
clus2−vs−NON.clus2 in UCEC−TP RNASeq data
Among total18525genes
all 8982 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
PLA2G4A(4.03)
TGFB2(1.88)
MAP2K6(1.55)
RPS6KA5(1.42)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus2−vs−NON.clus2]
0
10
20
30
40
−4
−2
0
2
4