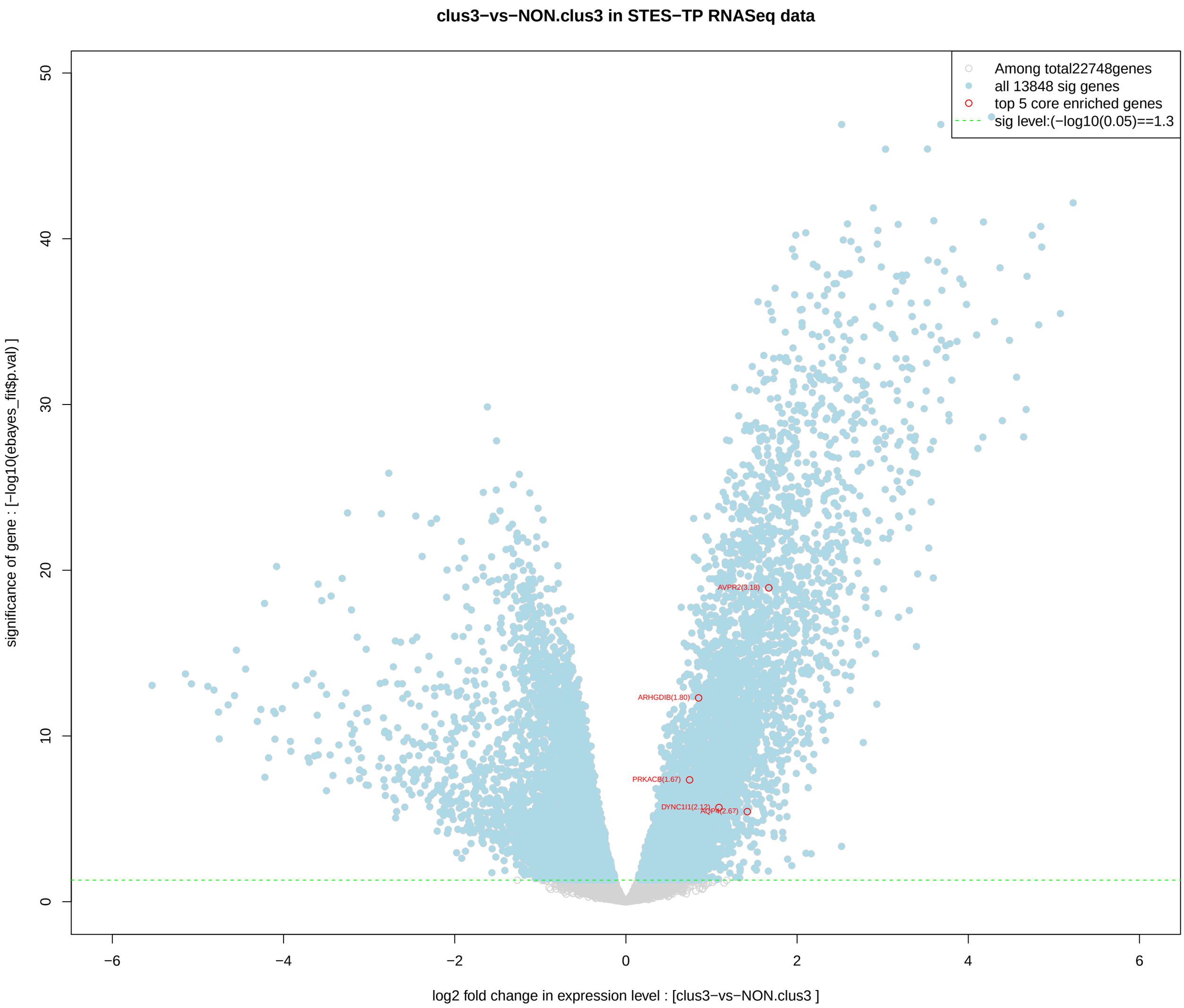
clus3−vs−NON.clus3 in STES−TP RNASeq data
Among total22748genes
all 13848 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
AVPR2(3.18)
ARHGDIB(1.80)
PRKACB(1.67)
DYNC1I1(2.12)
AQP4(2.67)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus3−vs−NON.clus3]
0
10
20
30
40
50
−6
−4
−2
0
2
4
6