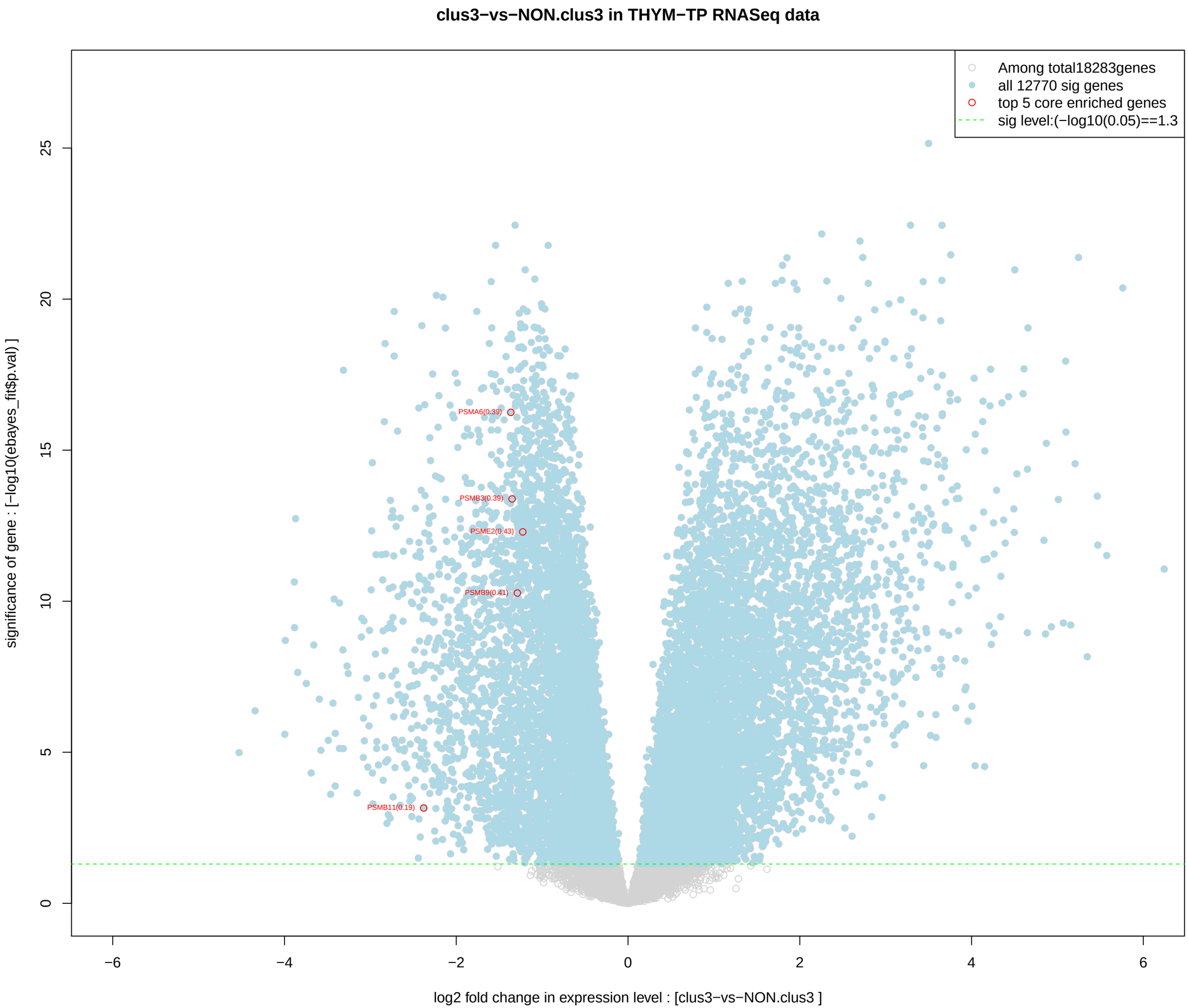
clus3−vs−NON.clus3 in THYM−TP RNASeq data
Among total18283genes
all 12770 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
PSMA6(0.39)
PSMB3(0.39)
PSME2(0.43)
PSMB9(0.41)
PSMB11(0.19)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus3−vs−NON.clus3]
0
5
10
15
20
25
−6
−4
−2
0
2
4
6