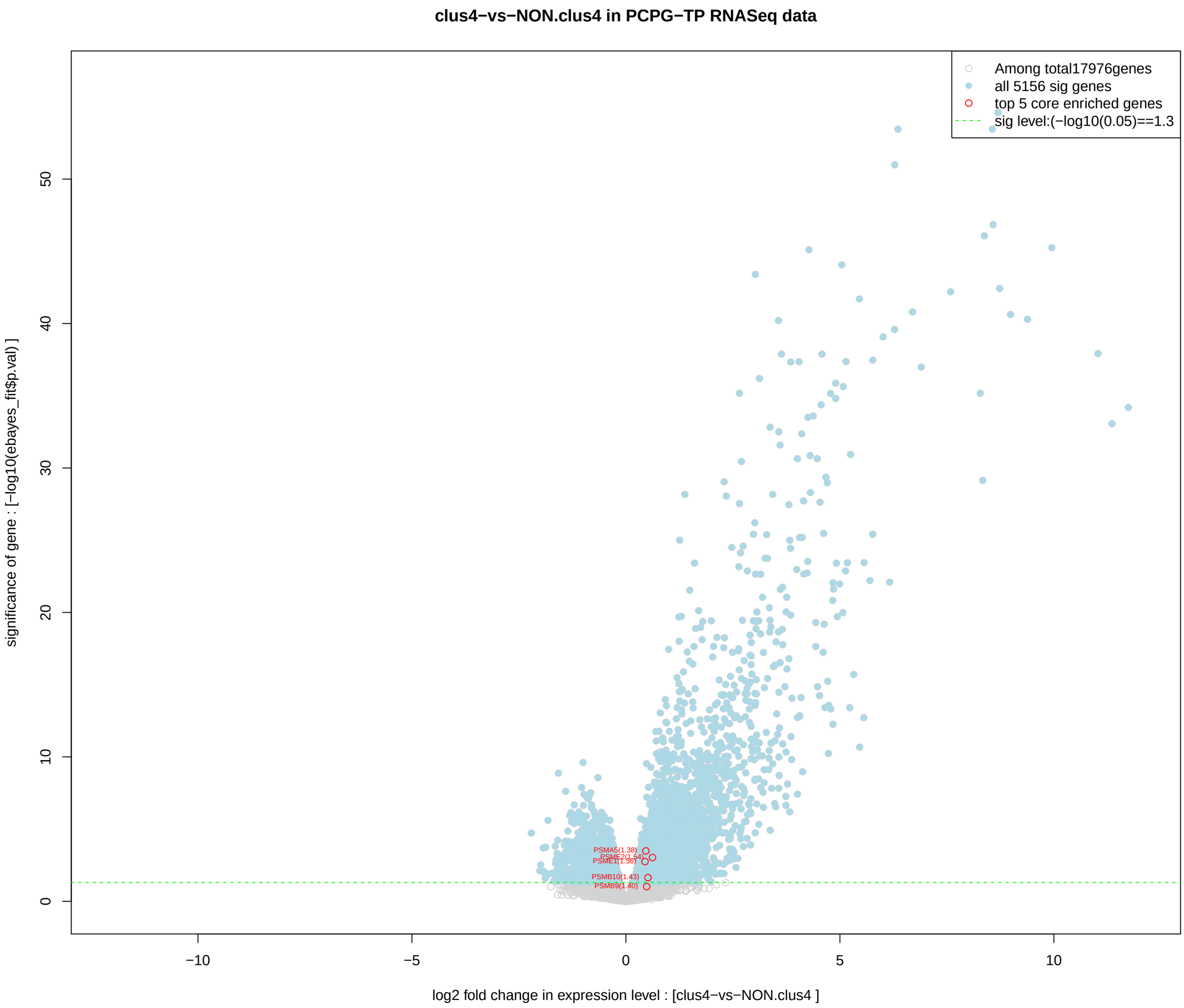
clus4−vs−NON.clus4 in PCPG−TP RNASeq data
Among total17976genes
all 5156 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
PSMA5(1.38)
PSME2(1.54)
PSME1(1.36)
PSMB10(1.43)
PSMB9(1.40)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus4−vs−NON.clus4]
0
10
20
30
40
50
−10
−5
0
5
10