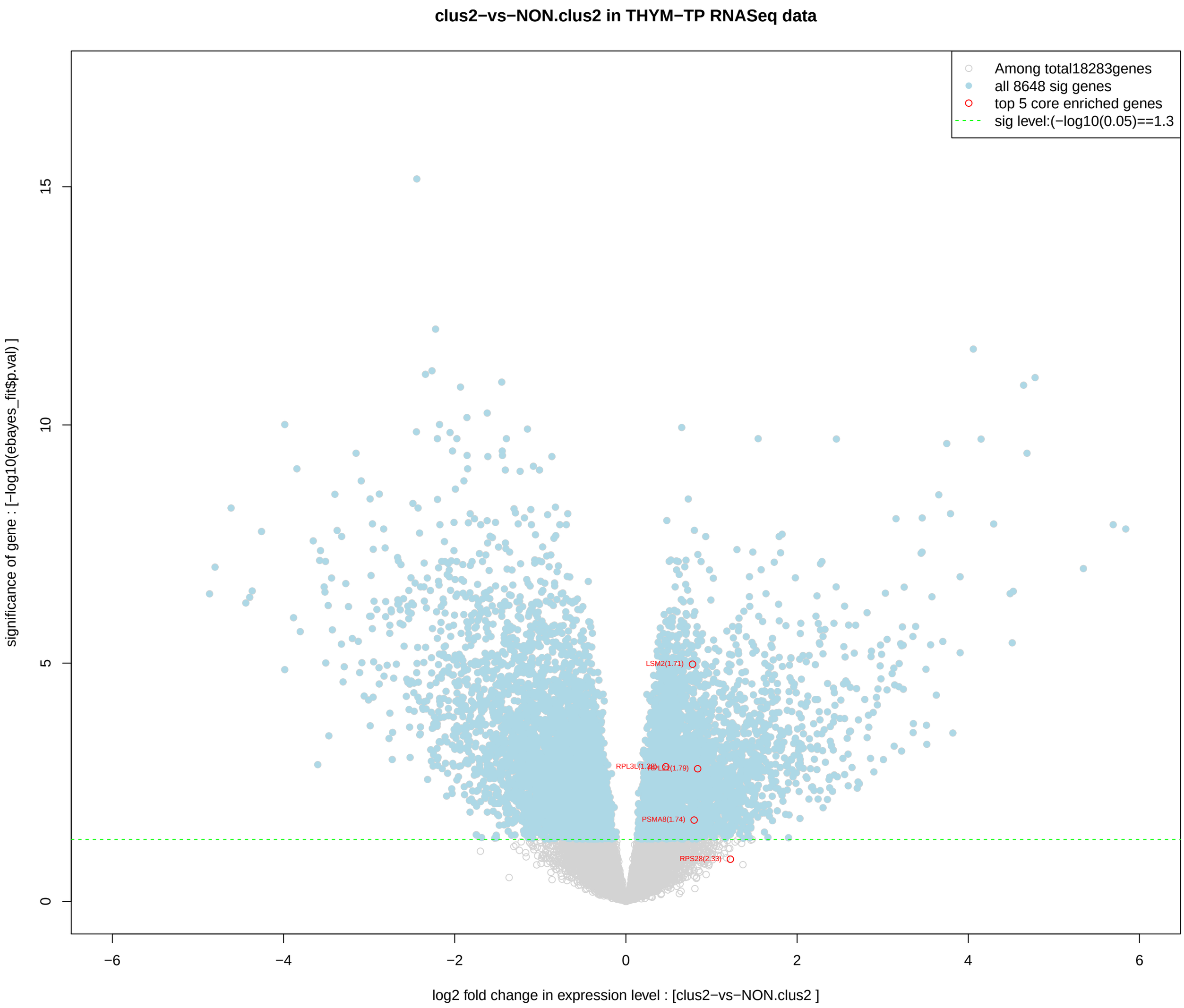
clus2−vs−NON.clus2 in THYM−TP RNASeq data
Among total18283genes
all 8648 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
LSM2(1.71)
RPL3L(1.39)
RPL22(1.79)
PSMA8(1.74)
RPS28(2.33)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus2−vs−NON.clus2]
0
5
10
15
−6
−4
−2
0
2
4
6