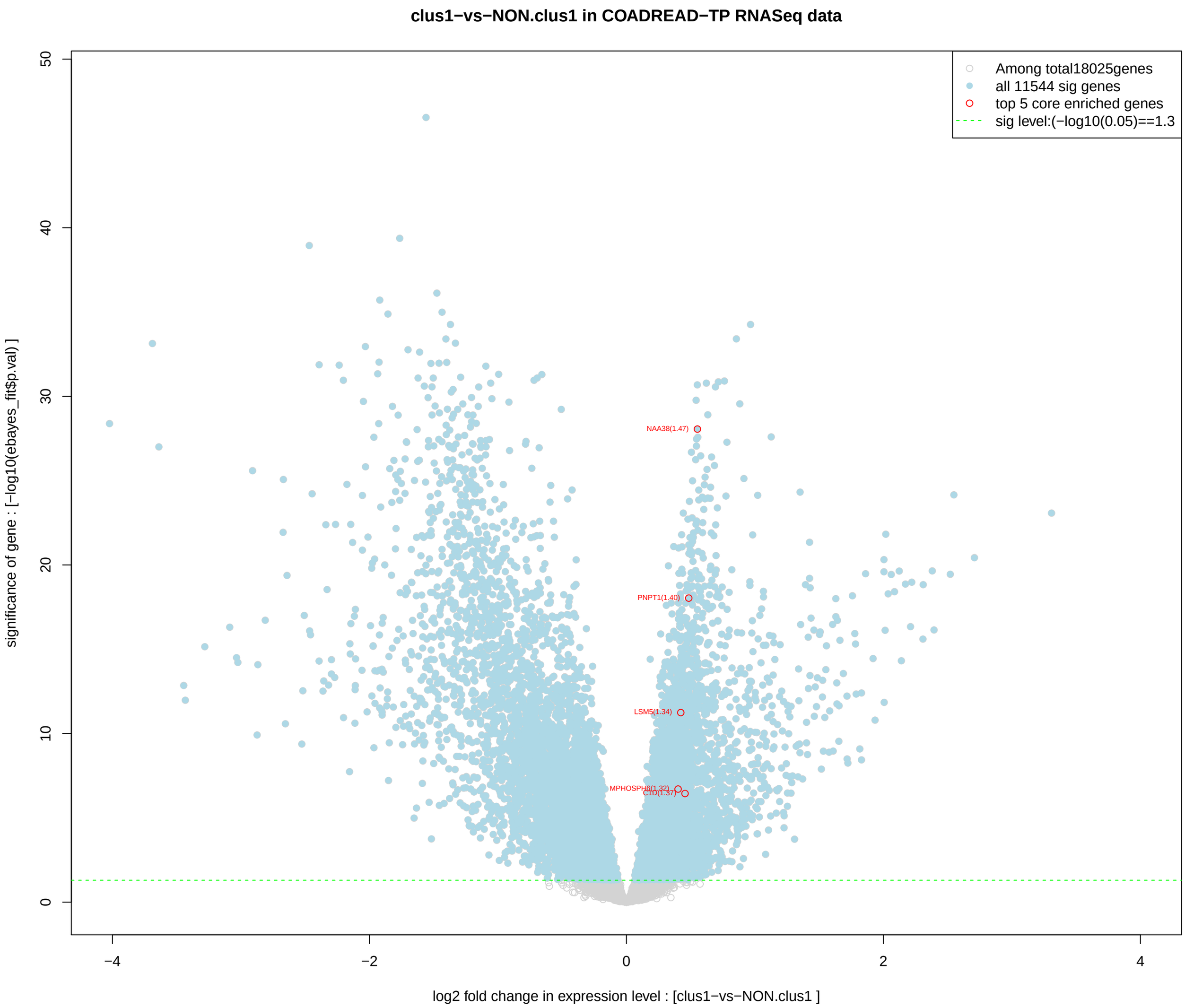
clus1−vs−NON.clus1 in COADREAD−TP RNASeq data
Among total18025genes
all 11544 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
NAA38(1.47)
PNPT1(1.40)
LSM5(1.34)
MPHOSPH6(1.32)
C1D(1.37)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus1−vs−NON.clus1]
0
10
20
30
40
50
−4
−2
0
2
4