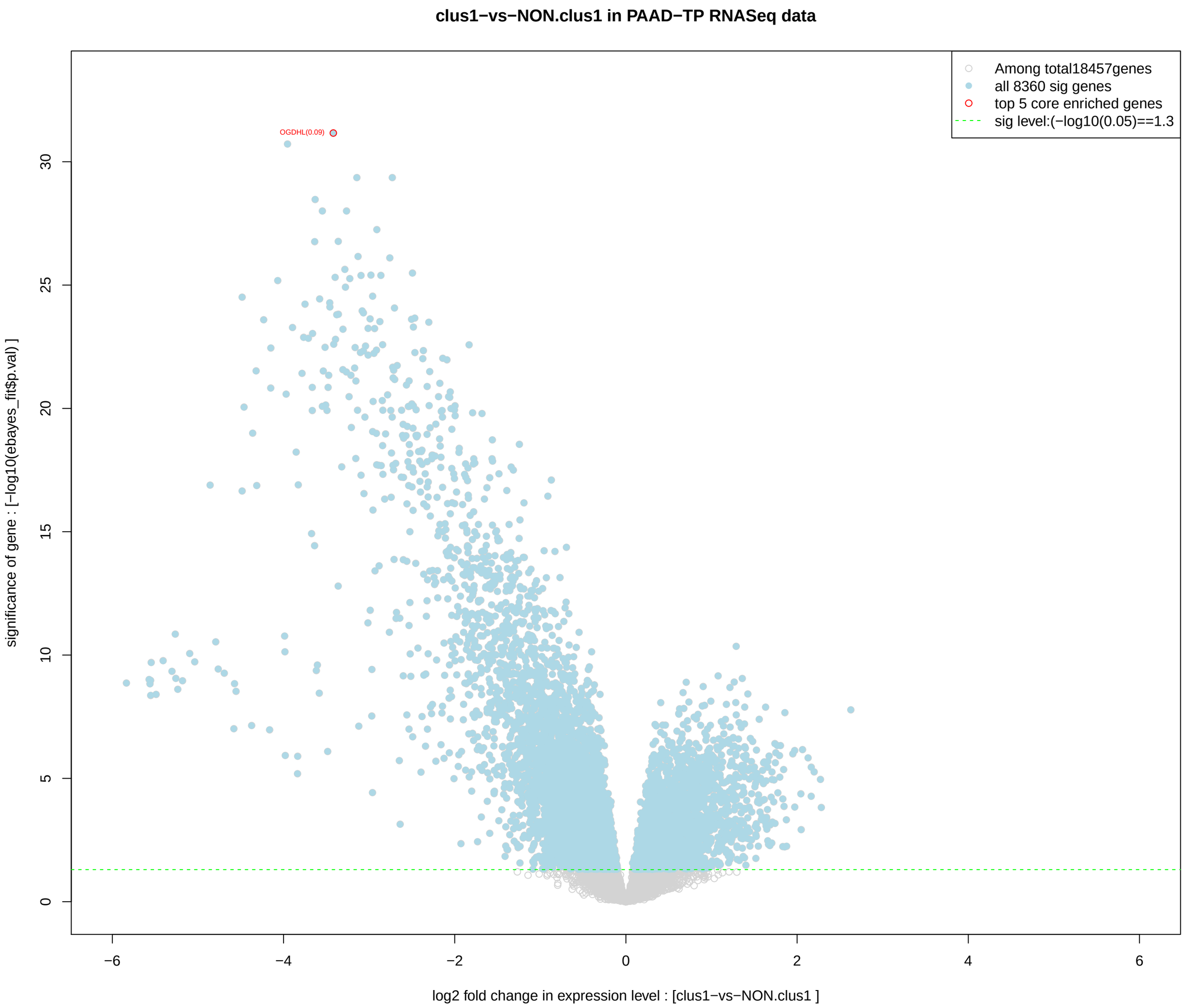
clus1−vs−NON.clus1 in PAAD−TP RNASeq data
Among total18457genes
all 8360 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
OGDHL(0.09)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus1−vs−NON.clus1]
0
5
10
15
20
25
30
−6
−4
−2
0
2
4
6