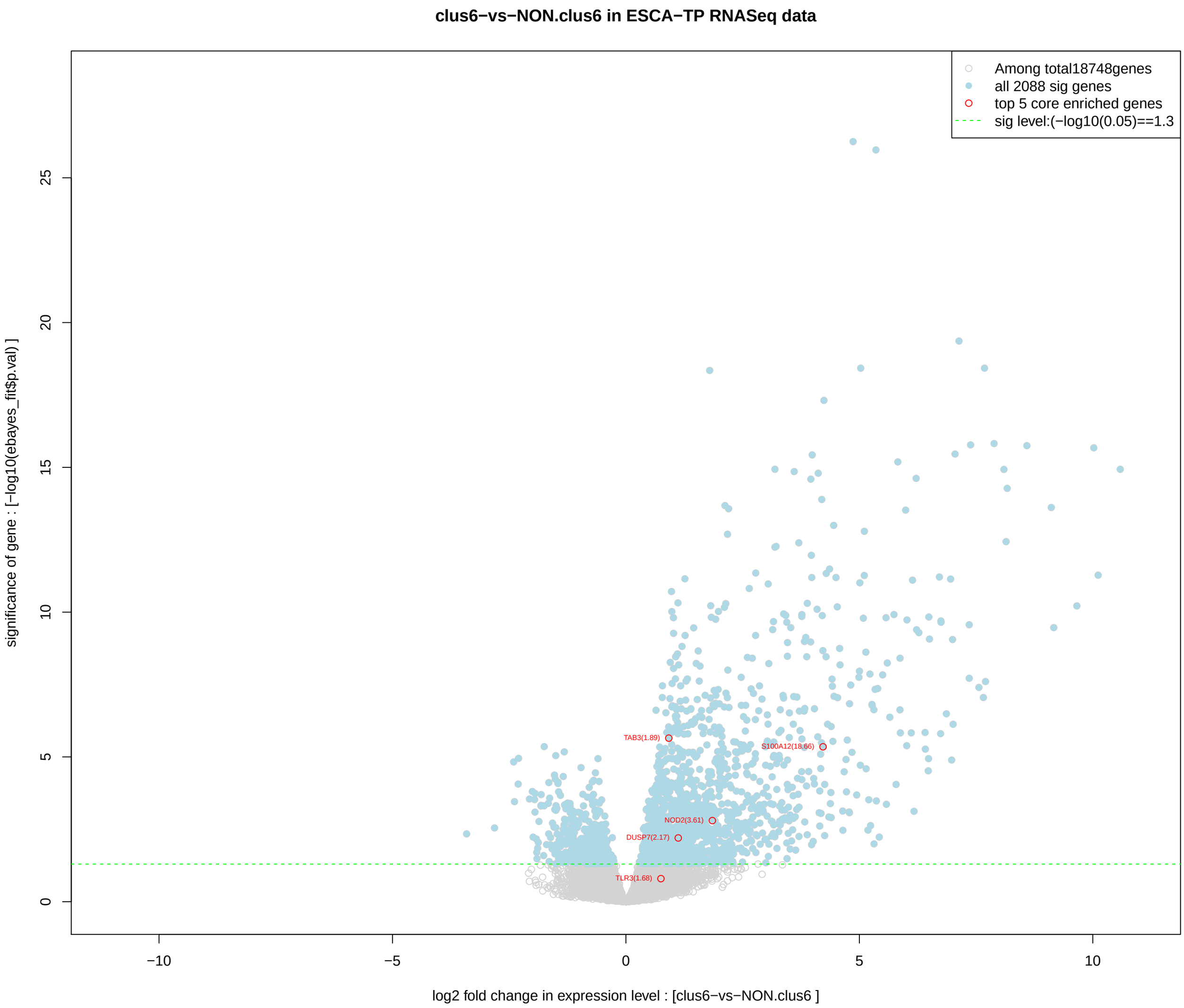
clus6−vs−NON.clus6 in ESCA−TP RNASeq data
Among total18748genes
all 2088 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
TAB3(1.89)
S100A12(18.66)
NOD2(3.61)
DUSP7(2.17)
TLR3(1.68)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus6−vs−NON.clus6]
0
5
10
15
20
25
−10
−5
0
5
10