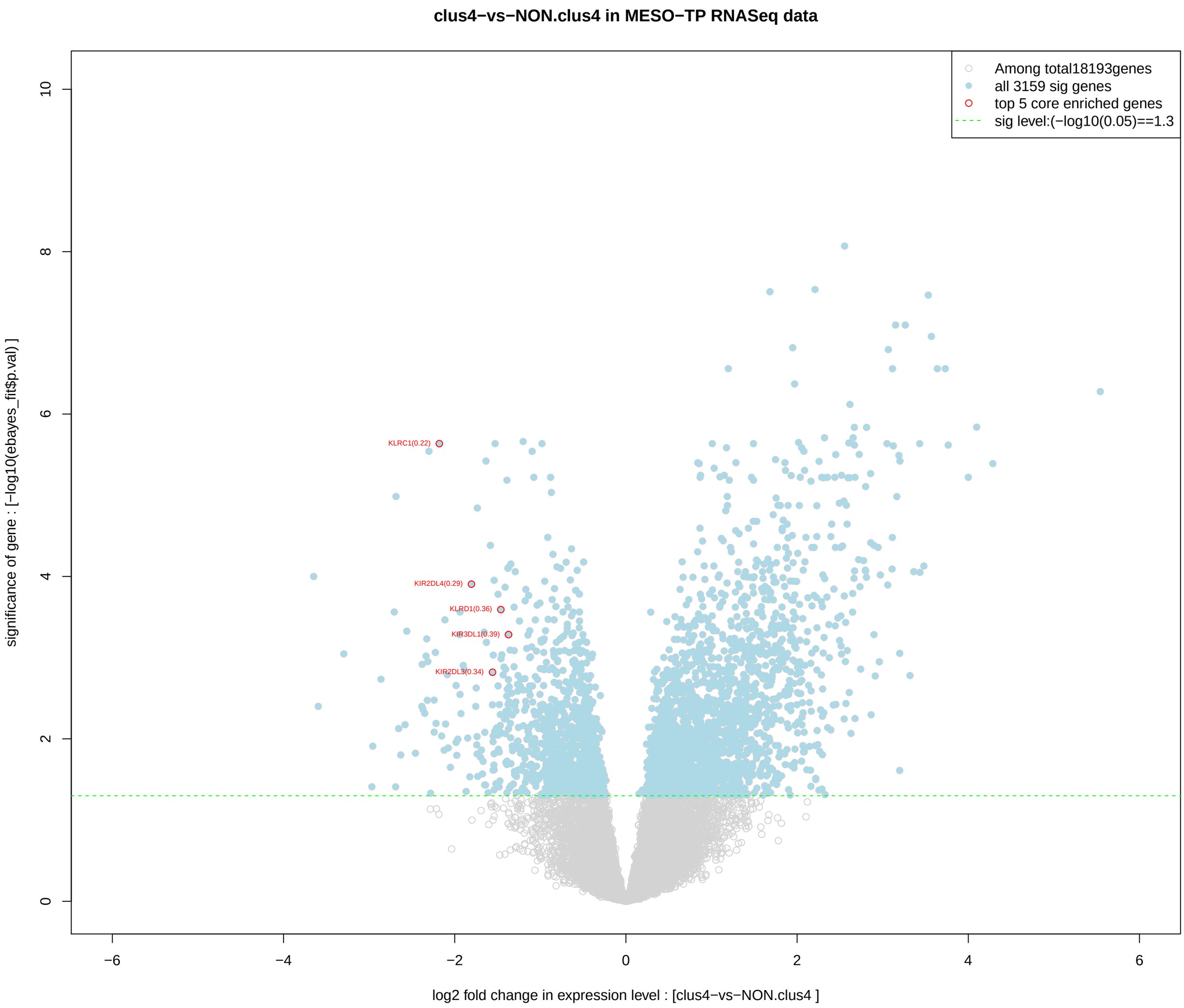
clus4−vs−NON.clus4 in MESO−TP RNASeq data
Among total18193genes
all 3159 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
KLRC1(0.22)
KIR2DL4(0.29)
KLRD1(0.36)
KIR3DL1(0.39)
KIR2DL3(0.34)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus4−vs−NON.clus4]
0
2
4
6
8
10
−6
−4
−2
0
2
4
6