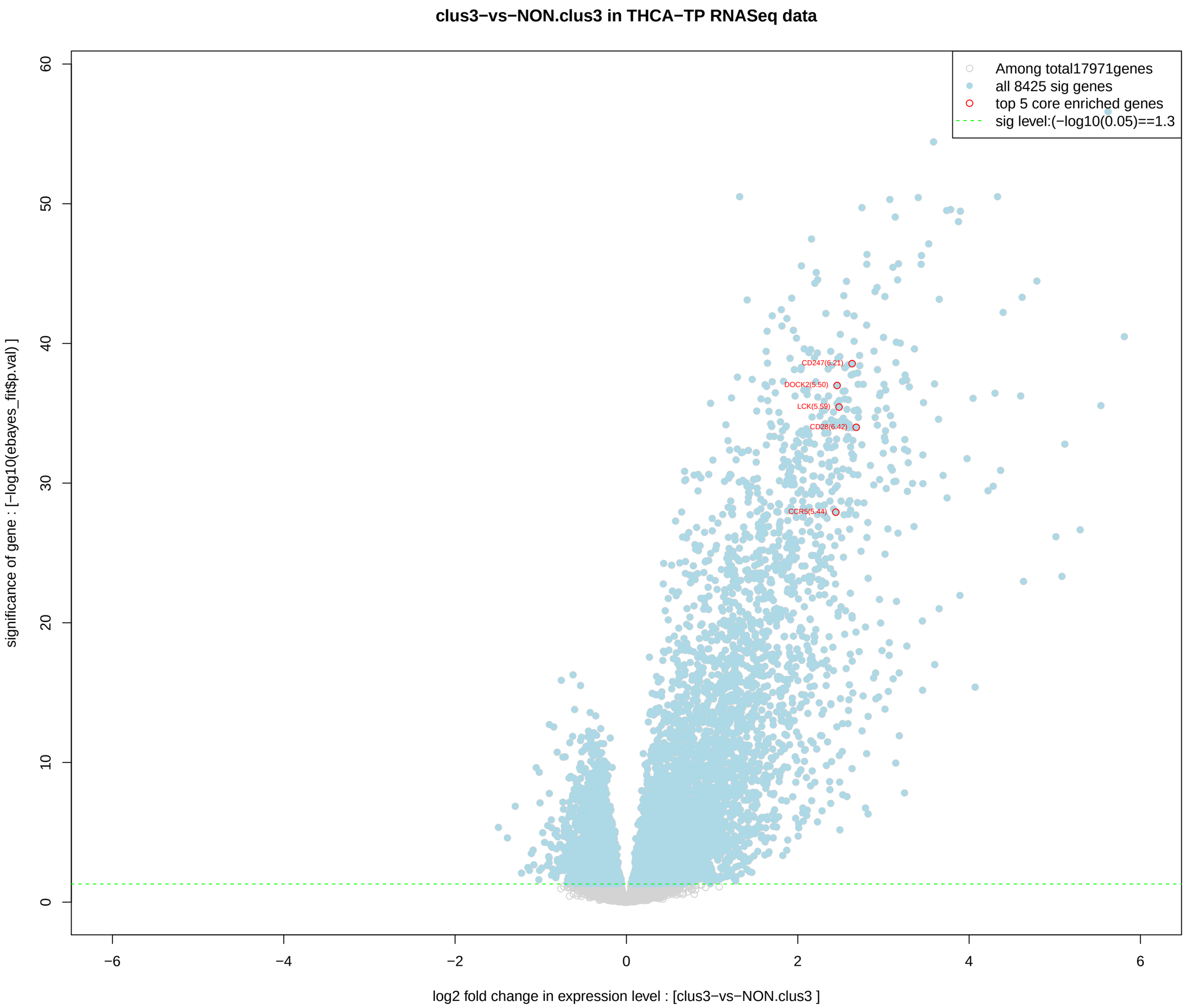
clus3−vs−NON.clus3 in THCA−TP RNASeq data
Among total17971genes
all 8425 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
CD247(6.21)
DOCK2(5.50)
LCK(5.59)
CD28(6.42)
CCR5(5.44)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus3−vs−NON.clus3]
0
10
20
30
40
50
60
−6
−4
−2
0
2
4
6