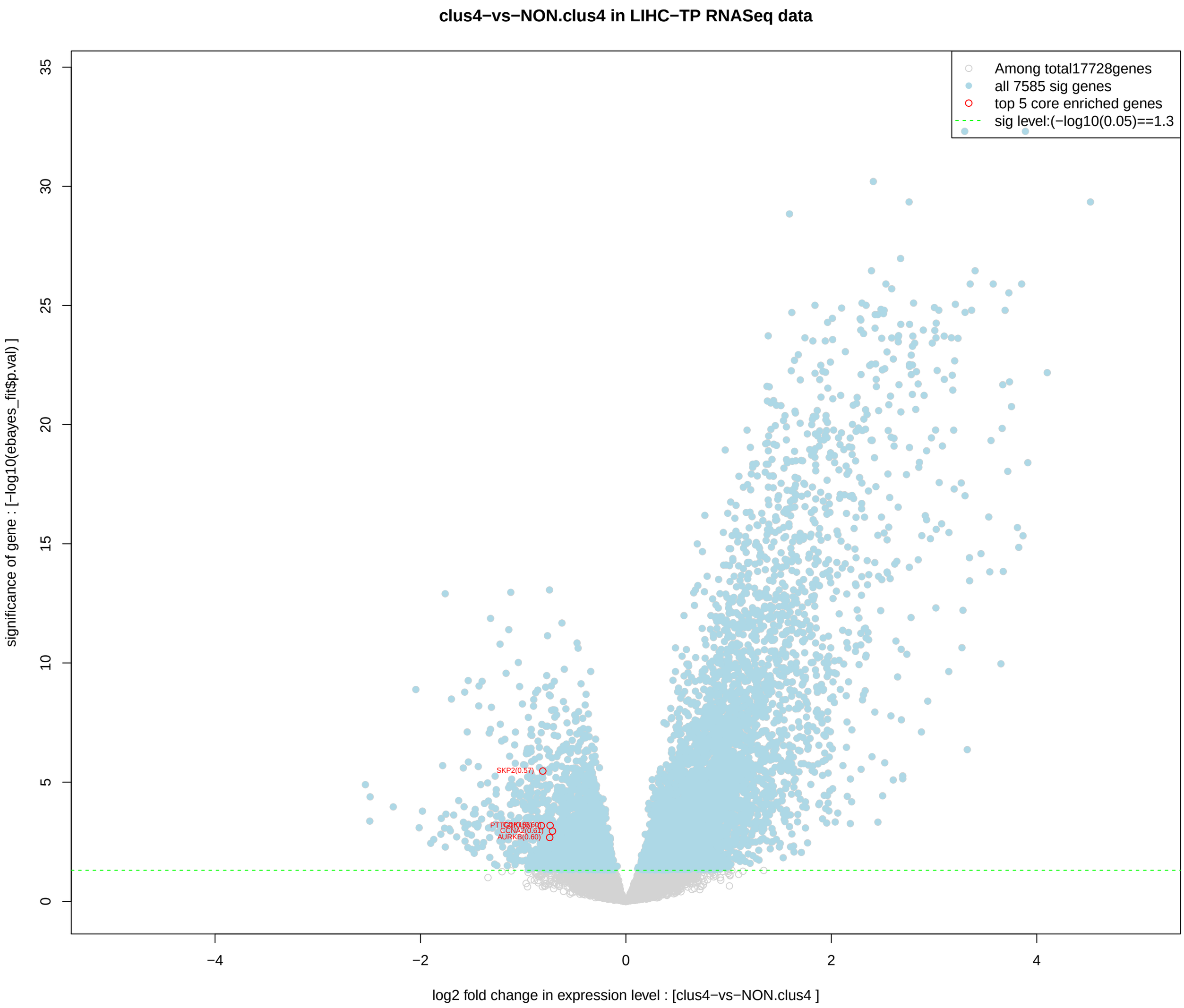
clus4−vs−NON.clus4 in LIHC−TP RNASeq data
Among total17728genes
all 7585 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
SKP2(0.57)
CCNA2(0.61)
AURKB(0.60)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus4−vs−NON.clus4]
35
30
25
20
15
10
5
0
−4
−2
0
2
4