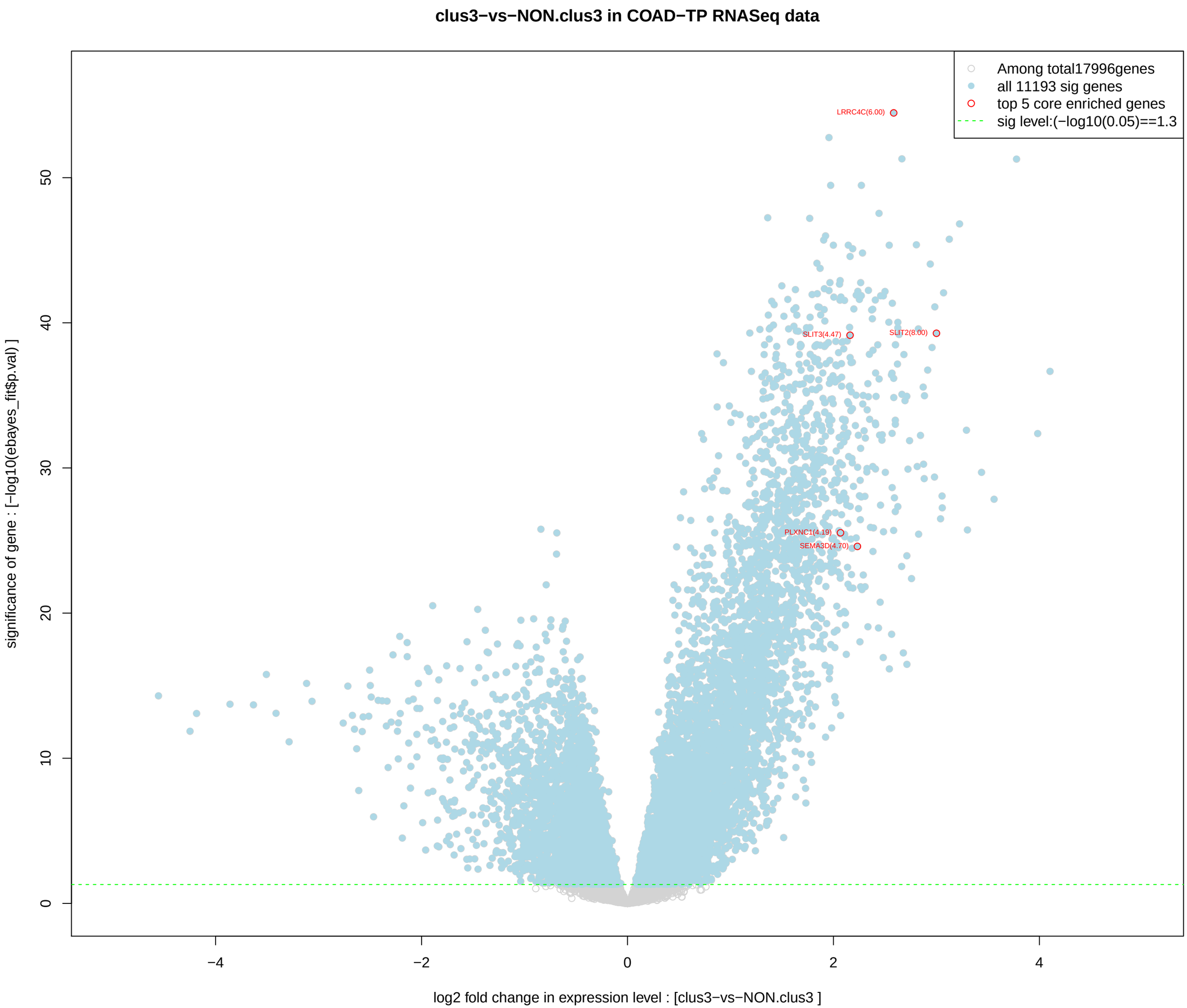
clus3−vs−NON.clus3 in COAD−TP RNASeq data
Among total17996genes
all 11193 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
LRRC4C(6.00)
SLIT3(4.47)
SLIT2(8.00)
PLXNC1(4.19)
SEMA3D(4.70)
significance of gene : [−log10(ebayes_fit$p.val)]
0
10
20
30
40
50
−4
−2
0
2
4
log2 fold change in expression level : [clus3−vs−NON.clus3]