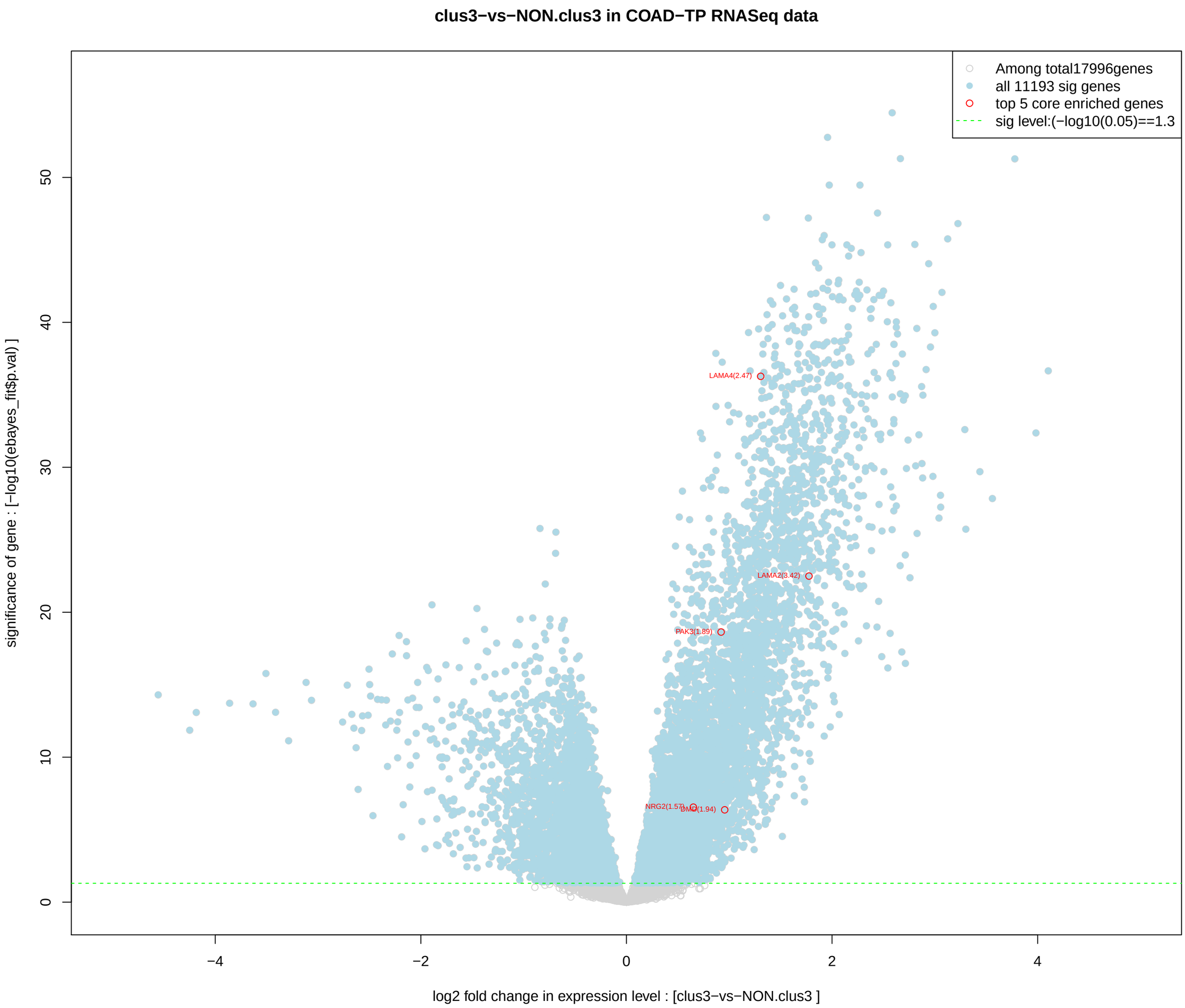
clus3−vs−NON.clus3 in COAD−TP RNASeq data
Among total17996genes
all 11193 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
LAMA4(2.47)
LAMA2(3.42)
PAK3(1.89)
NRG2(1.57)
DMD(1.94)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus3−vs−NON.clus3]
0
10
20
30
40
50
−4
−2
0
2
4