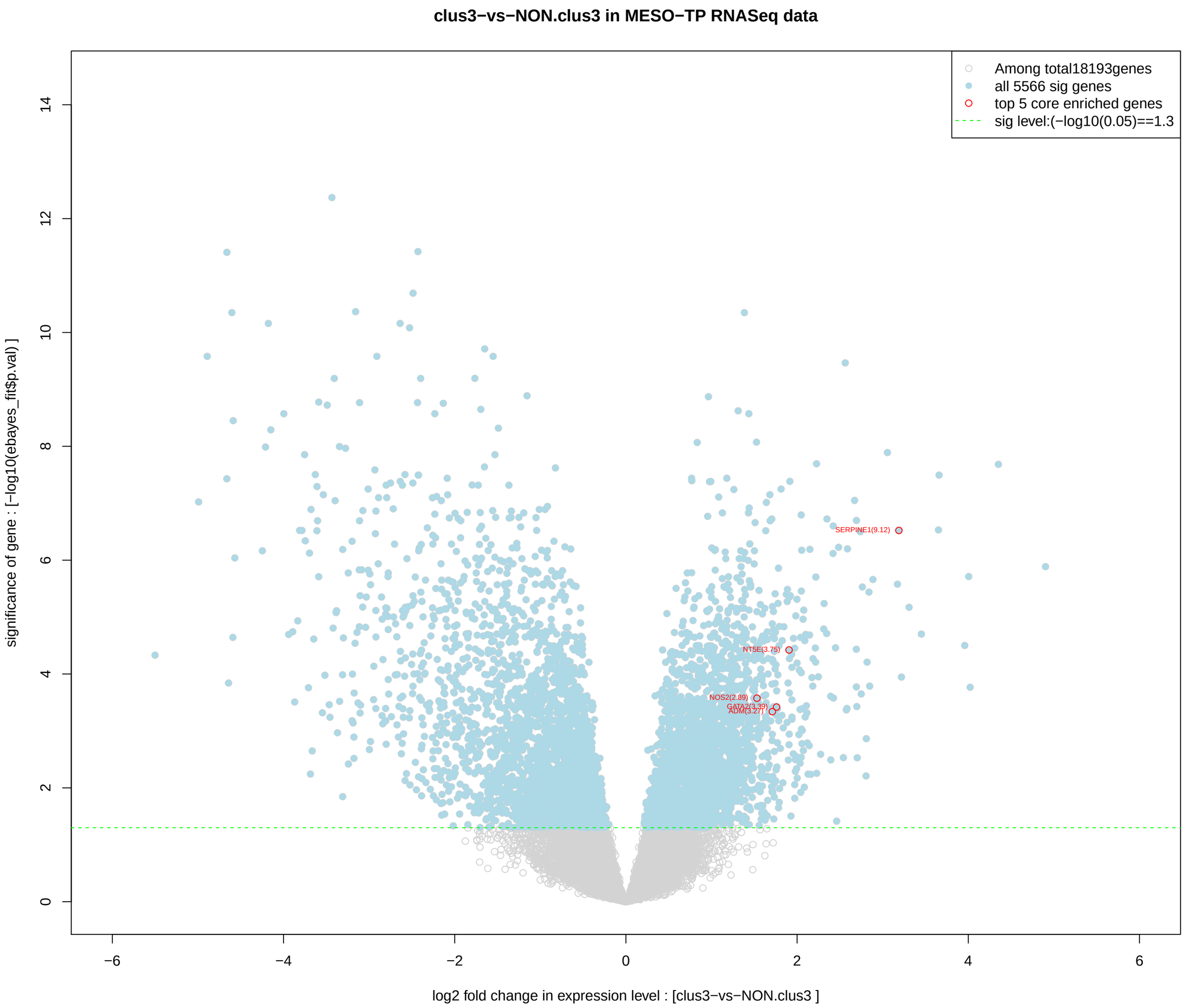
clus3−vs−NON.clus3 in MESO−TP RNASeq data
Among total18193genes
all 5566 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
SERPINE1(9.12)
NT5E(3.75)
NOS2(2.89)
GATA2(3.39)
ADM(3.27)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus3−vs−NON.clus3]
0
2
4
6
8
10
12
14
−6
−4
−2
0
2
4
6