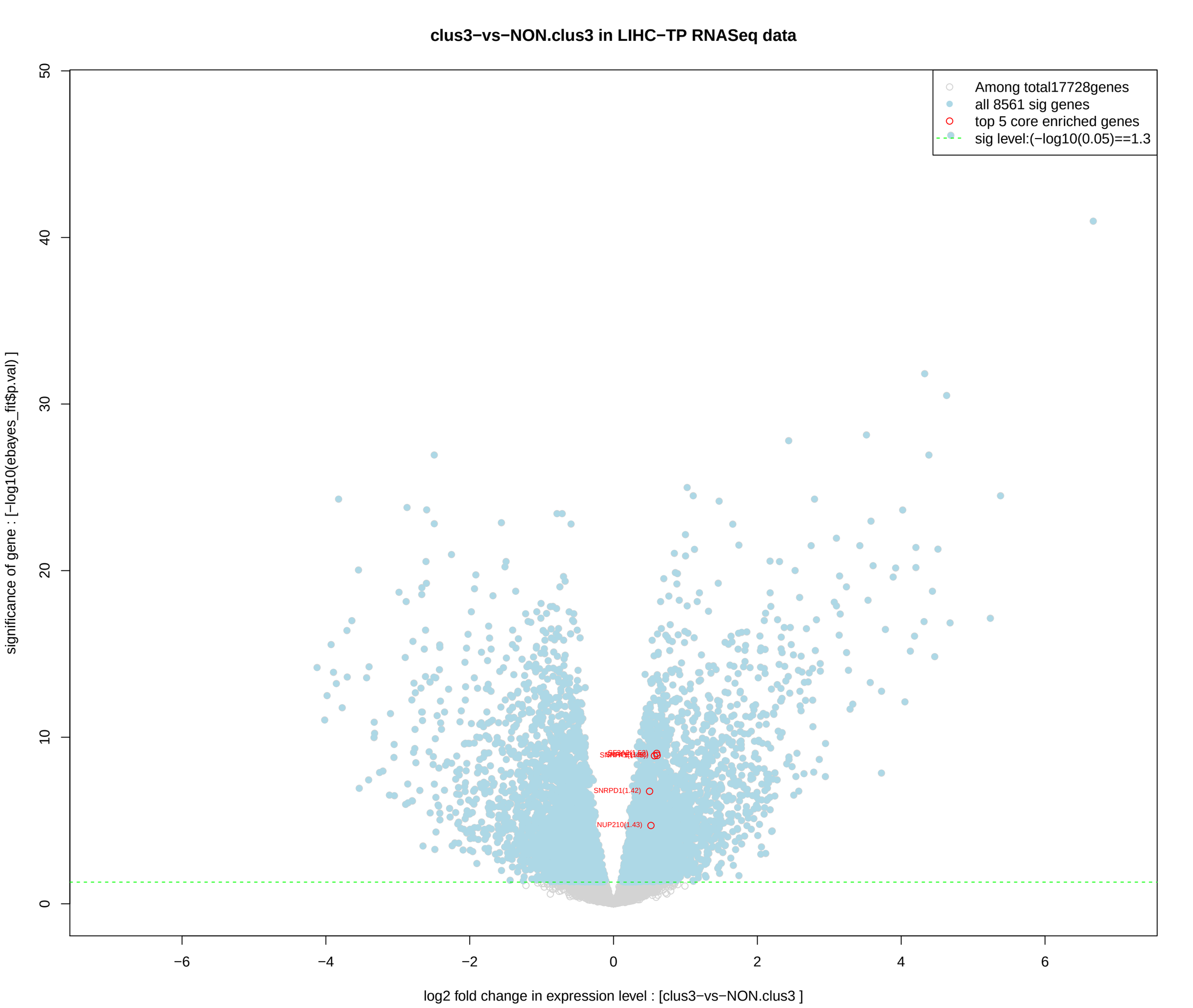

clus3−vs−NON.clus3 in LIHC−TP RNASeq data
Among total17728genes
all 8561 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
SNRPD1(1.42)
NUP210(1.43)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus3−vs−NON.clus3]
0
10
20
30
40
50
−6
−4
−2
0
2
4
6