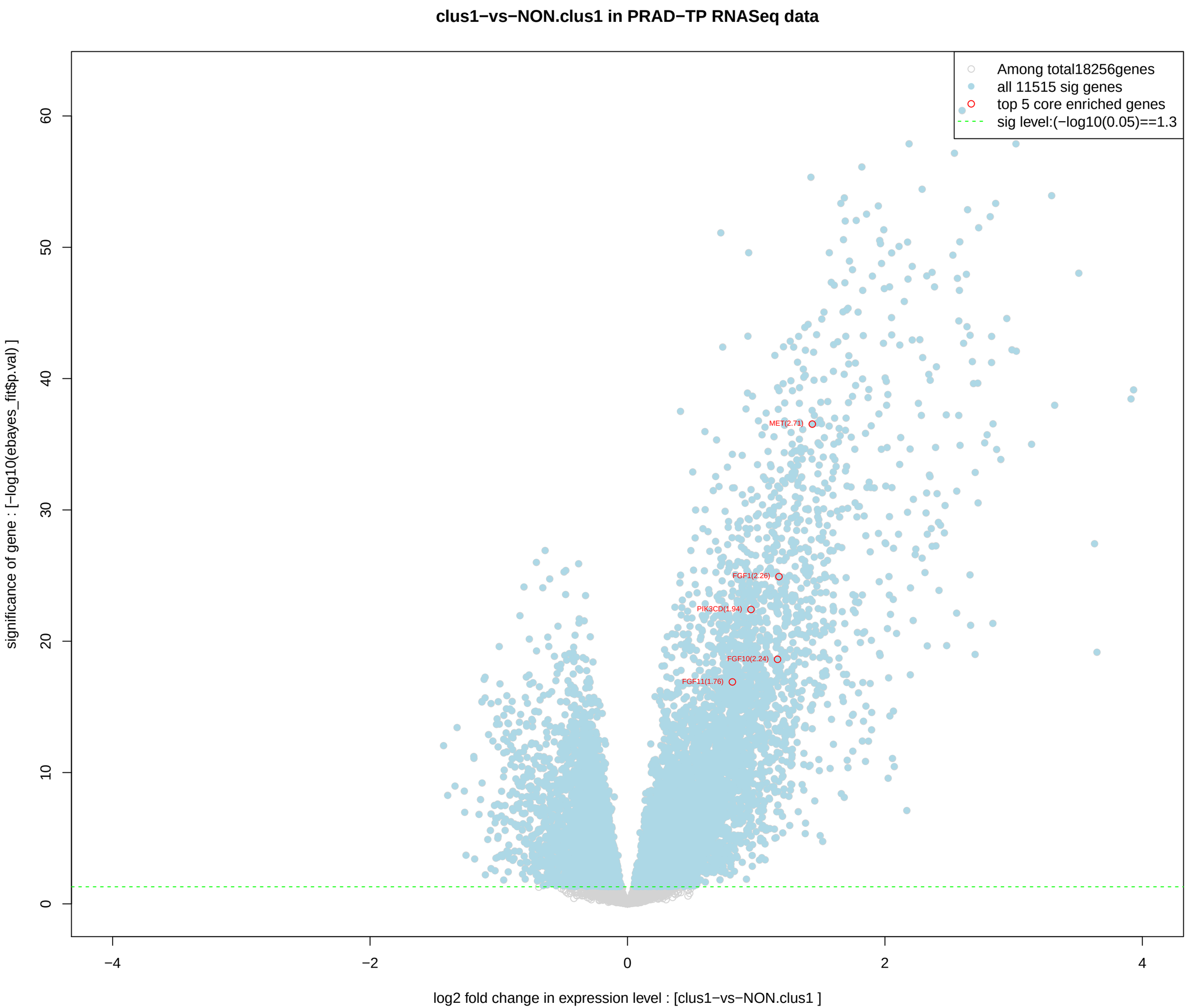
clus1−vs−NON.clus1 in PRAD−TP RNASeq data
Among total18256genes
all 11515 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
MET(2.71)
FGF1(2.26)
PIK3CD(1.94)
FGF10(2.24)
FGF11(1.76)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus1−vs−NON.clus1]
0
10
20
30
40
50
60
−4
−2
0
2
4