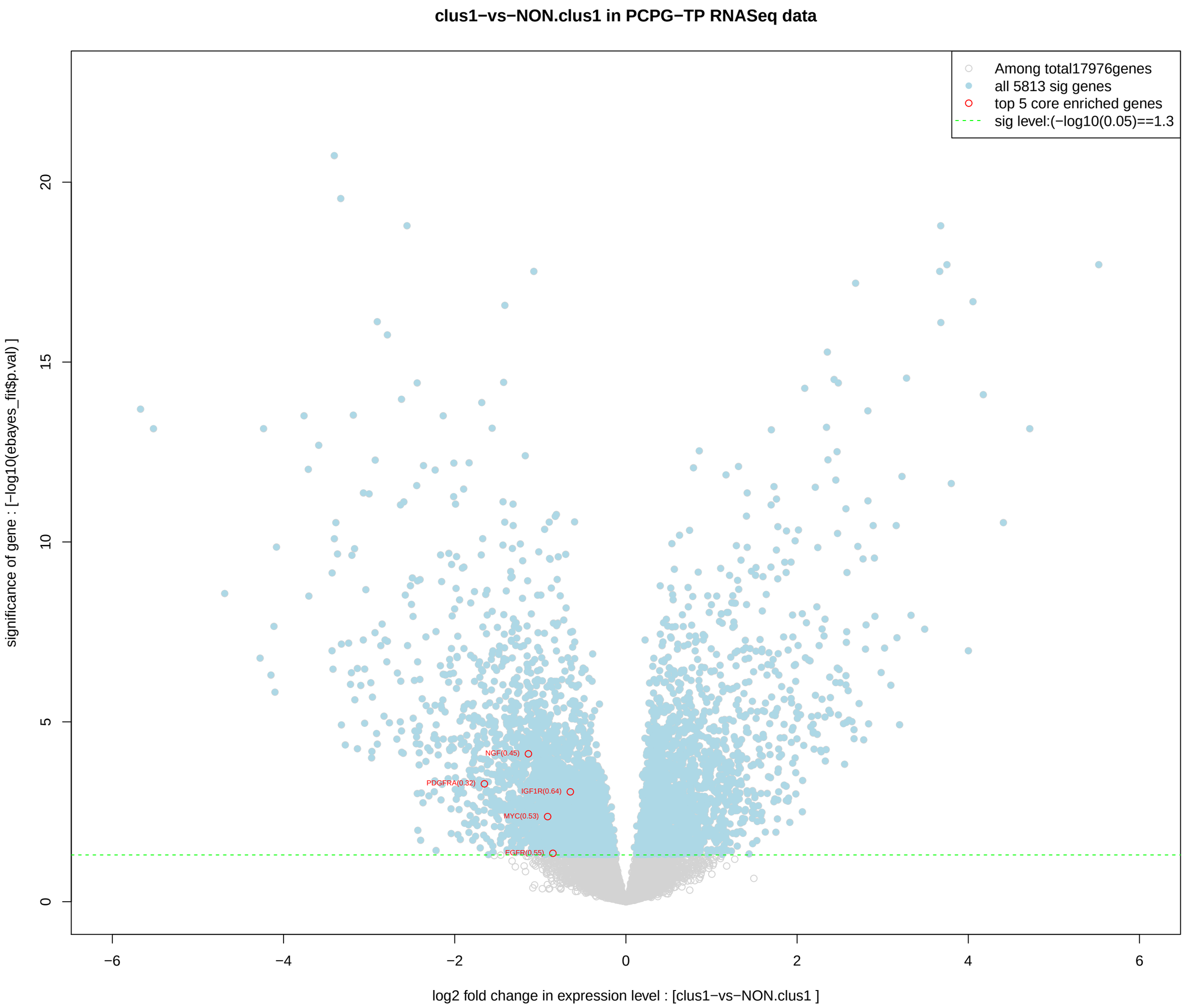
clus1−vs−NON.clus1 in PCPG−TP RNASeq data
Among total17976genes
all 5813 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
NGF(0.45)
PDGFRA(0.32)
IGF1R(0.64)
MYC(0.53)
EGFR(0.55)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus1−vs−NON.clus1]
0
5
10
15
20
−6
−4
−2
0
2
4
6