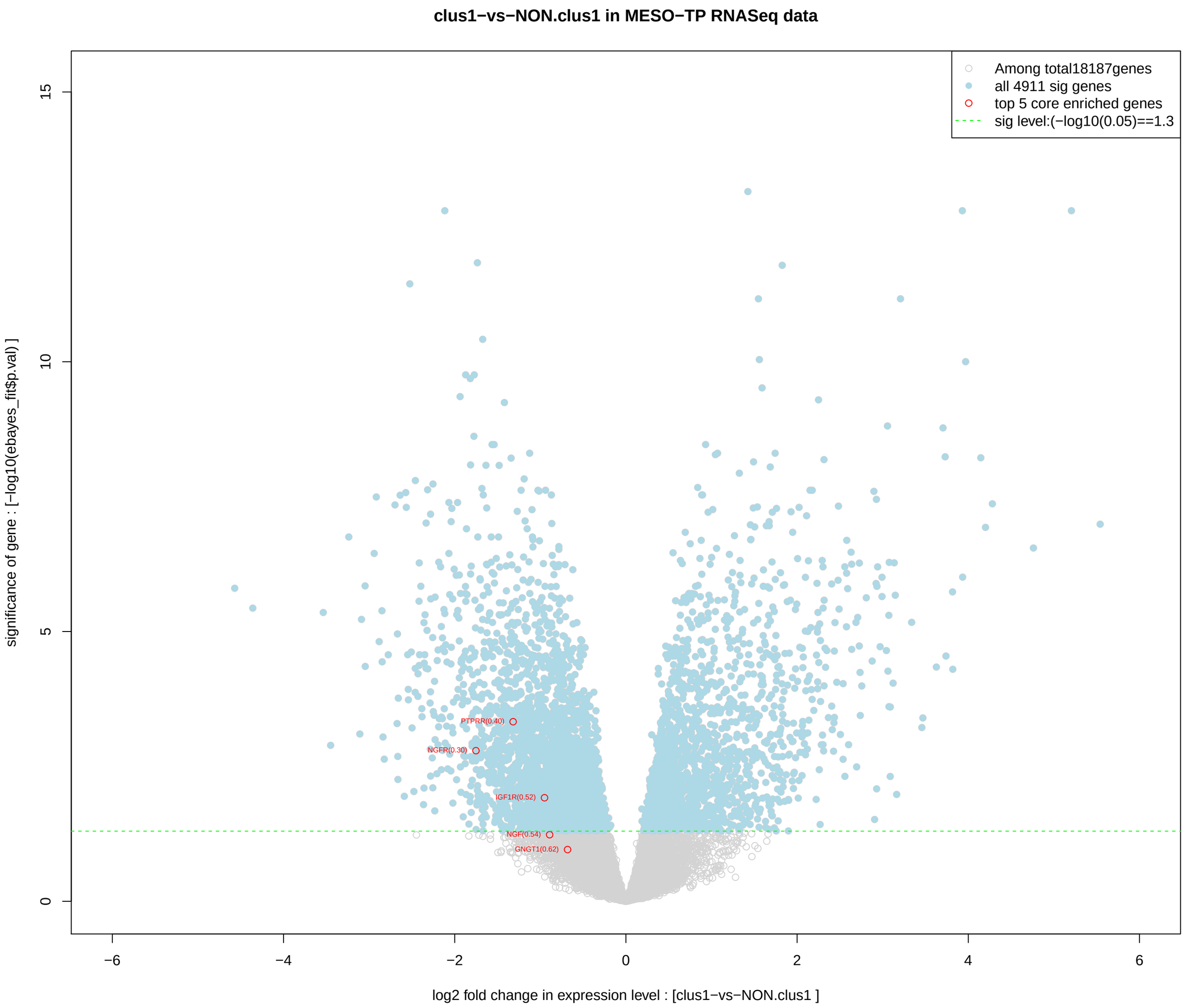
clus1−vs−NON.clus1 in MESO−TP RNASeq data
Among total18187genes
all 4911 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
PTPRR(0.40)
NGFR(0.30)
IGF1R(0.52)
NGF(0.54)
GNGT1(0.62)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus1−vs−NON.clus1]
0
5
10
15
−6
−4
−2
0
2
4
6