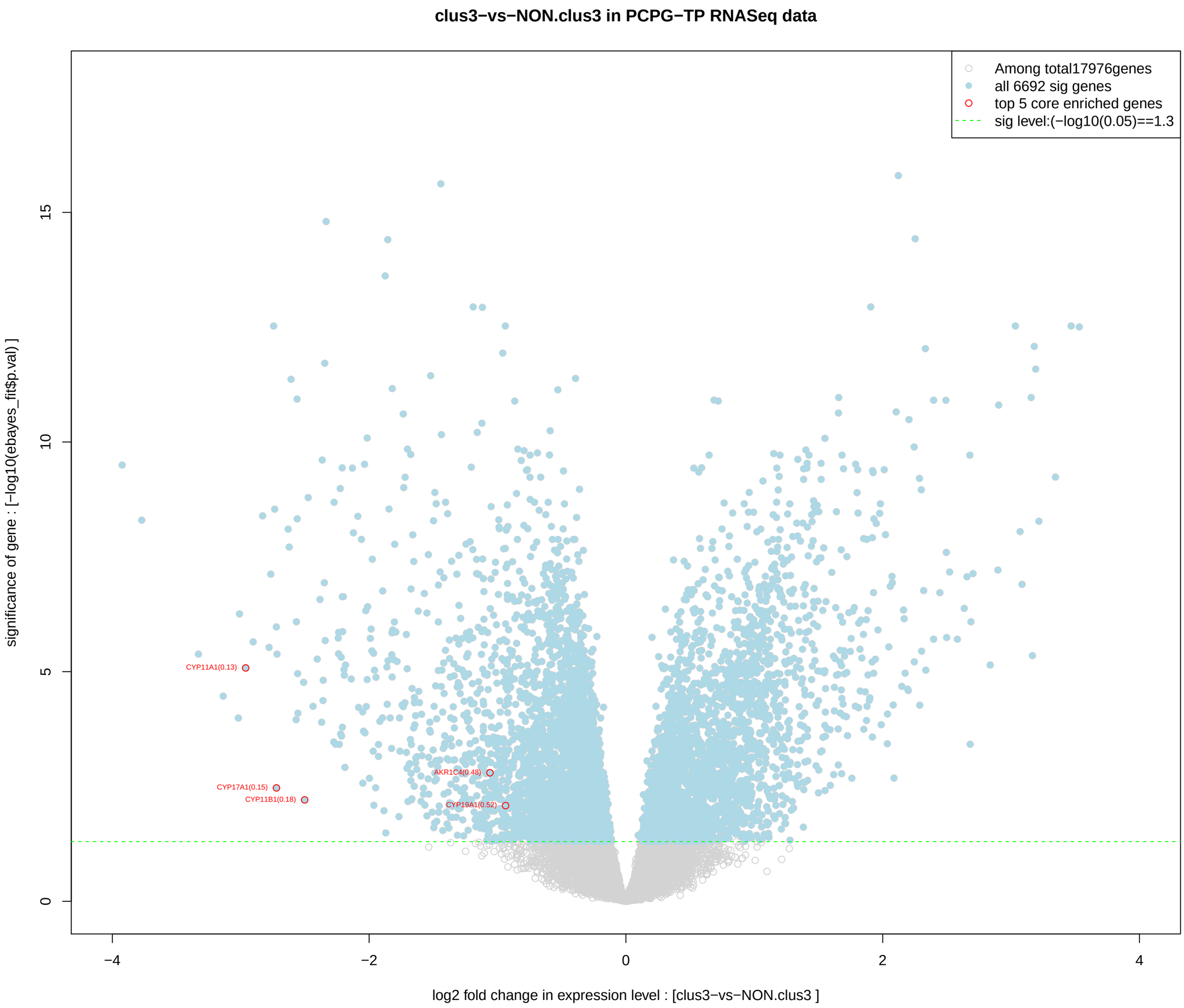
clus3−vs−NON.clus3 in PCPG−TP RNASeq data
Among total17976genes
all 6692 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
CYP11A1(0.13)
CYP17A1(0.15)
CYP11B1(0.18)
AKR1C4(0.48)
CYP19A1(0.52)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus3−vs−NON.clus3]
0
5
10
15
−4
−2
0
2
4