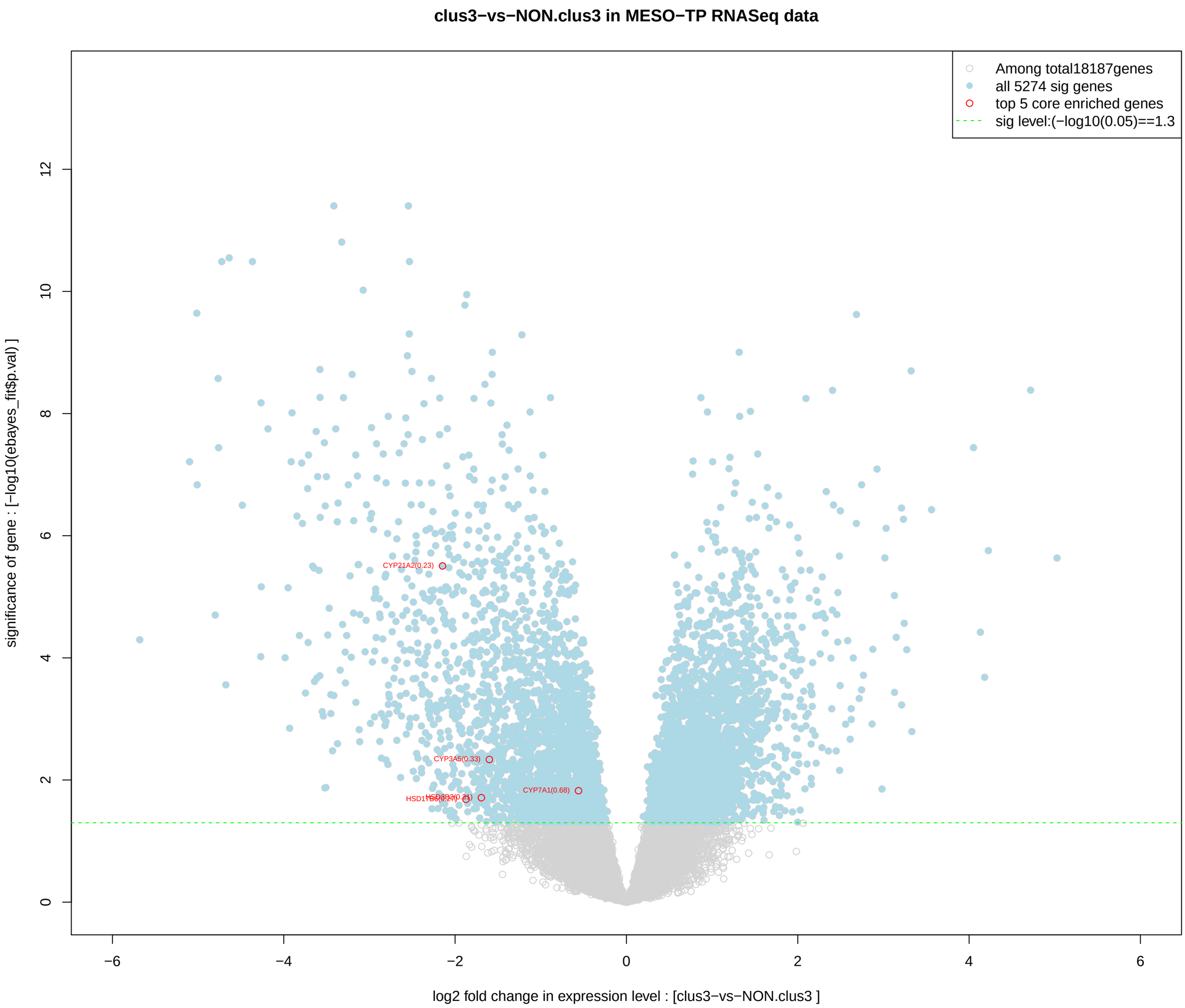
clus3−vs−NON.clus3 in MESO−TP RNASeq data
Among total18187genes
all 5274 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
CYP21A2(0.23)
CYP3A5(0.33)
CYP7A1(0.68)
HSD17B6(0.27)
HSD3B2(0.31)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus3−vs−NON.clus3]
0
2
4
6
8
10
12
−6
−4
−2
0
2
4
6