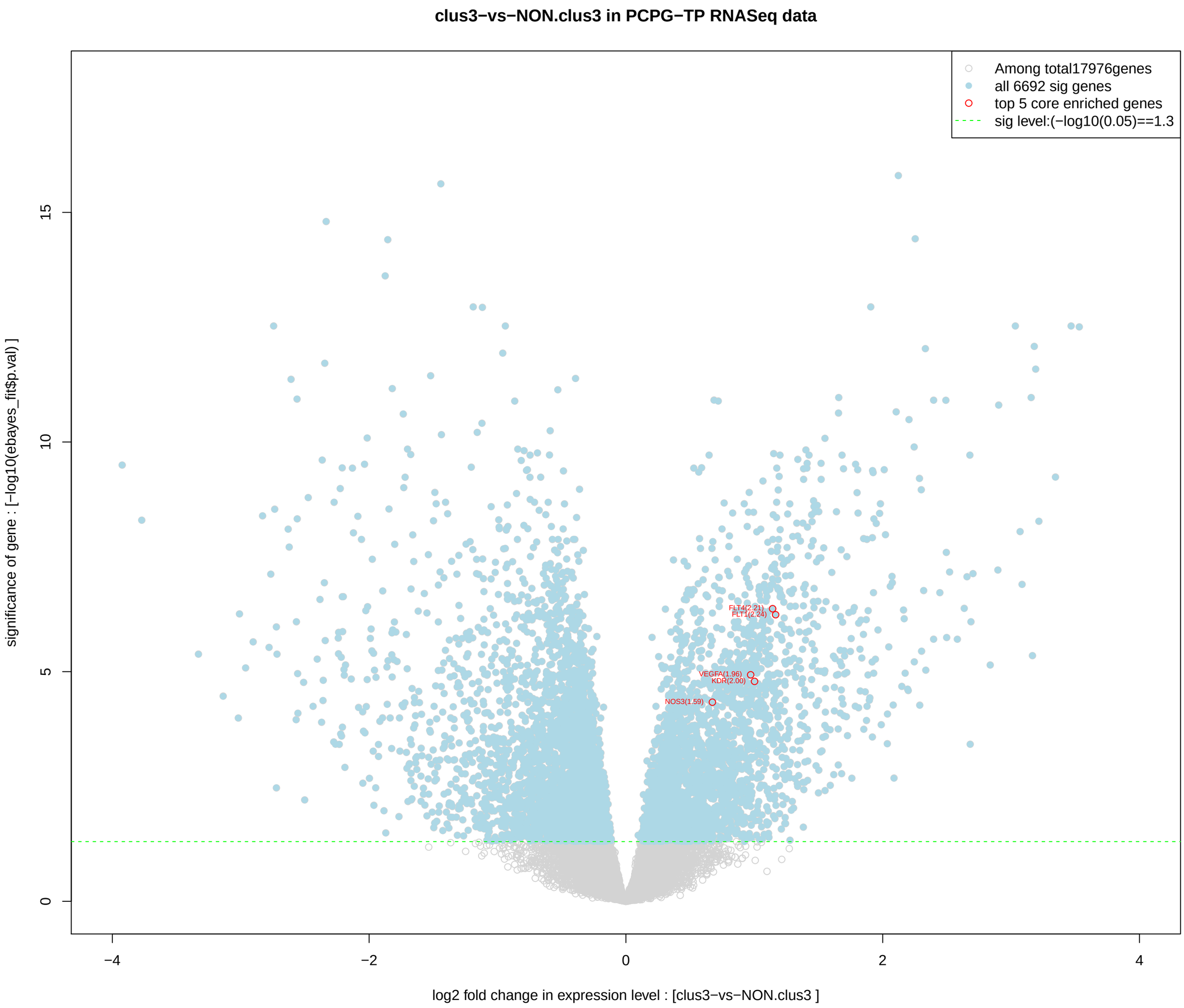
clus3−vs−NON.clus3 in PCPG−TP RNASeq data
Among total17976genes
all 6692 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
FLT4(2.21)
FLT1(2.24)
VEGFA(1.96)
KDR(2.00)
NOS3(1.59)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus3−vs−NON.clus3]
0
5
10
15
−4
−2
0
2
4