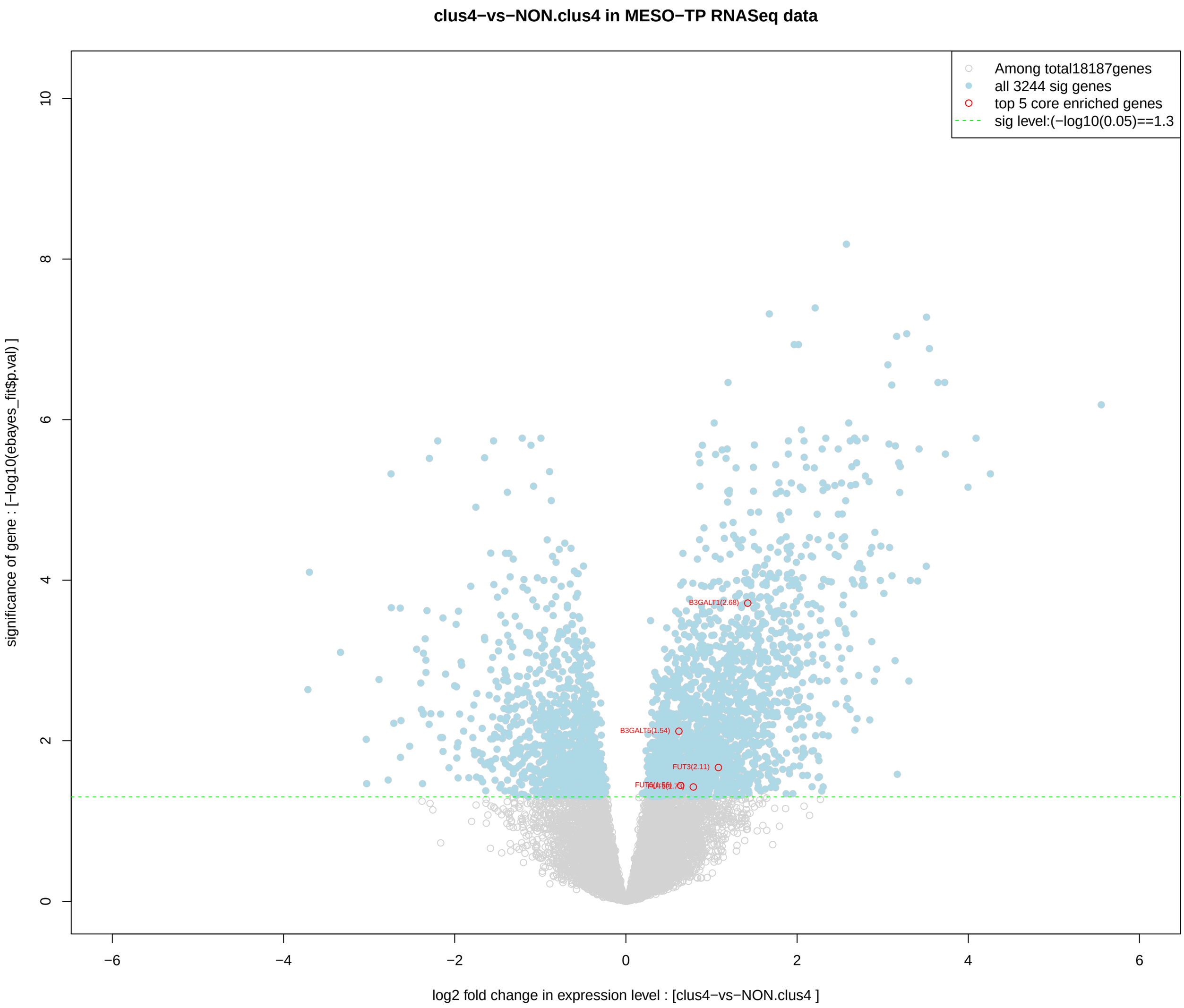
clus4−vs−NON.clus4 in MESO−TP RNASeq data
Among total18187genes
all 3244 sig genes
top 5 core enriched genes
sig level:(−log10(0.05)==1.3
B3GALT1(2.68)
B3GALT5(1.54)
FUT3(2.11)
FUT6(1.56)
FUT9(1.73)
significance of gene : [−log10(ebayes_fit$p.val)]
log2 fold change in expression level : [clus4−vs−NON.clus4]
0
2
4
6
8
10
−6
−4
−2
0
2
4
6