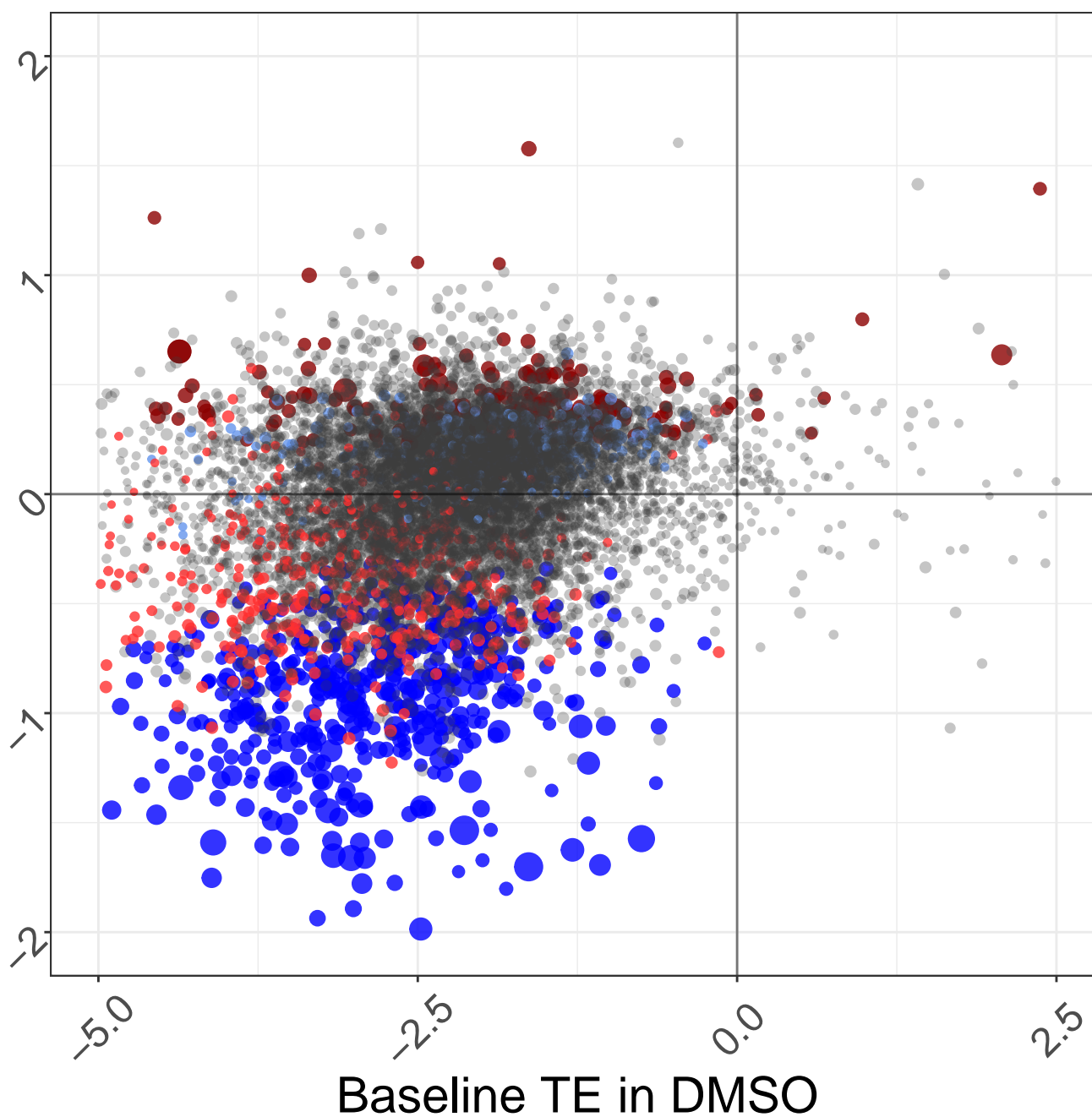


Log2 FoldChange TE
(IAA/DMSO)



$-\log_{10} \text{adj.pval}$
RiboRNA

● 10
● 20
● 30

Tx_class

● TE_up
● TE_down
● RNA_up
● RNA_down
● Not_significant