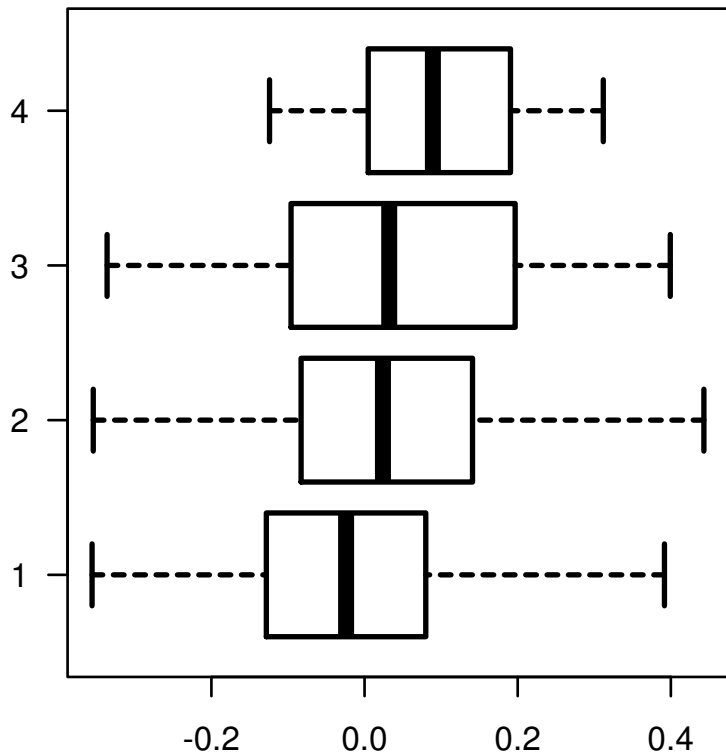


PHOLOGY\_T\_STAGE

# SMAD1|SMAD1



log2-EXPR  
Spearman.Corr= 0.179, P=0.000597