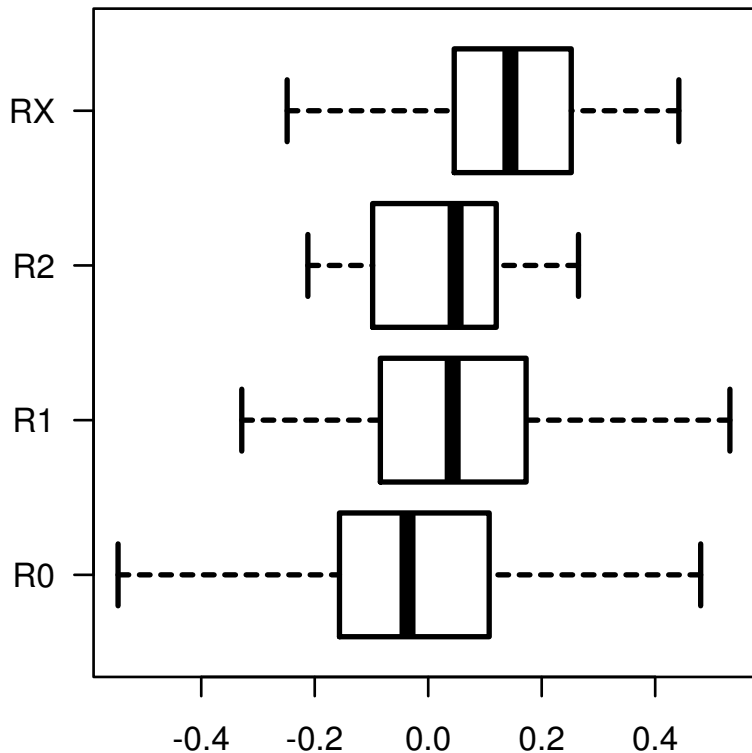


SMAD1|SMAD1

RESIDUAL_TUMOR



log2-EXPR
Kruskal-Wallis P = 0.00121