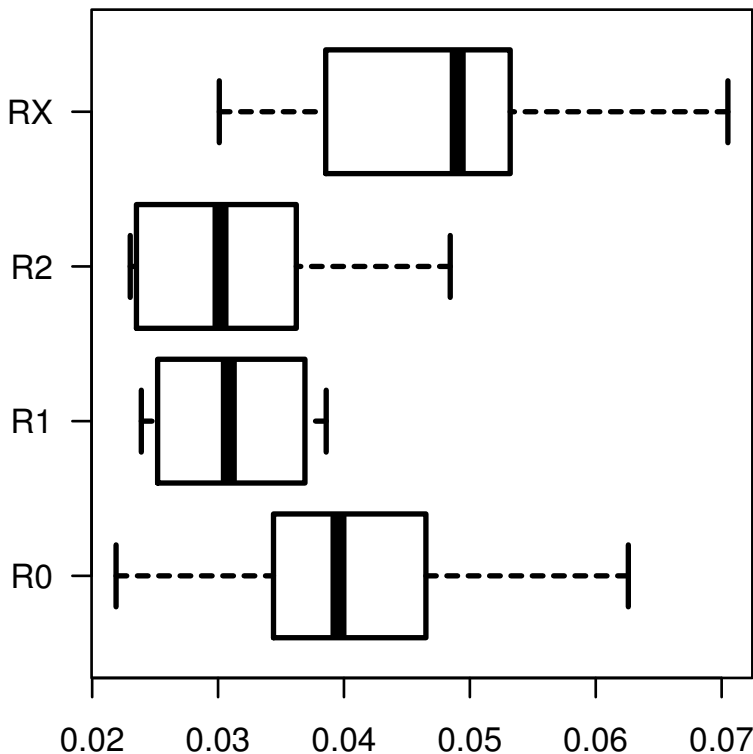


# TMEM231

RESIDUAL\_TUMOR



log<sub>2</sub>-EXPR  
Kruskal-Wallis P = 0.000109