condition_30_PrT_vs_36_PrT

DOWN: 0   UP: 0

−log10(svalue)

log2FoldChange
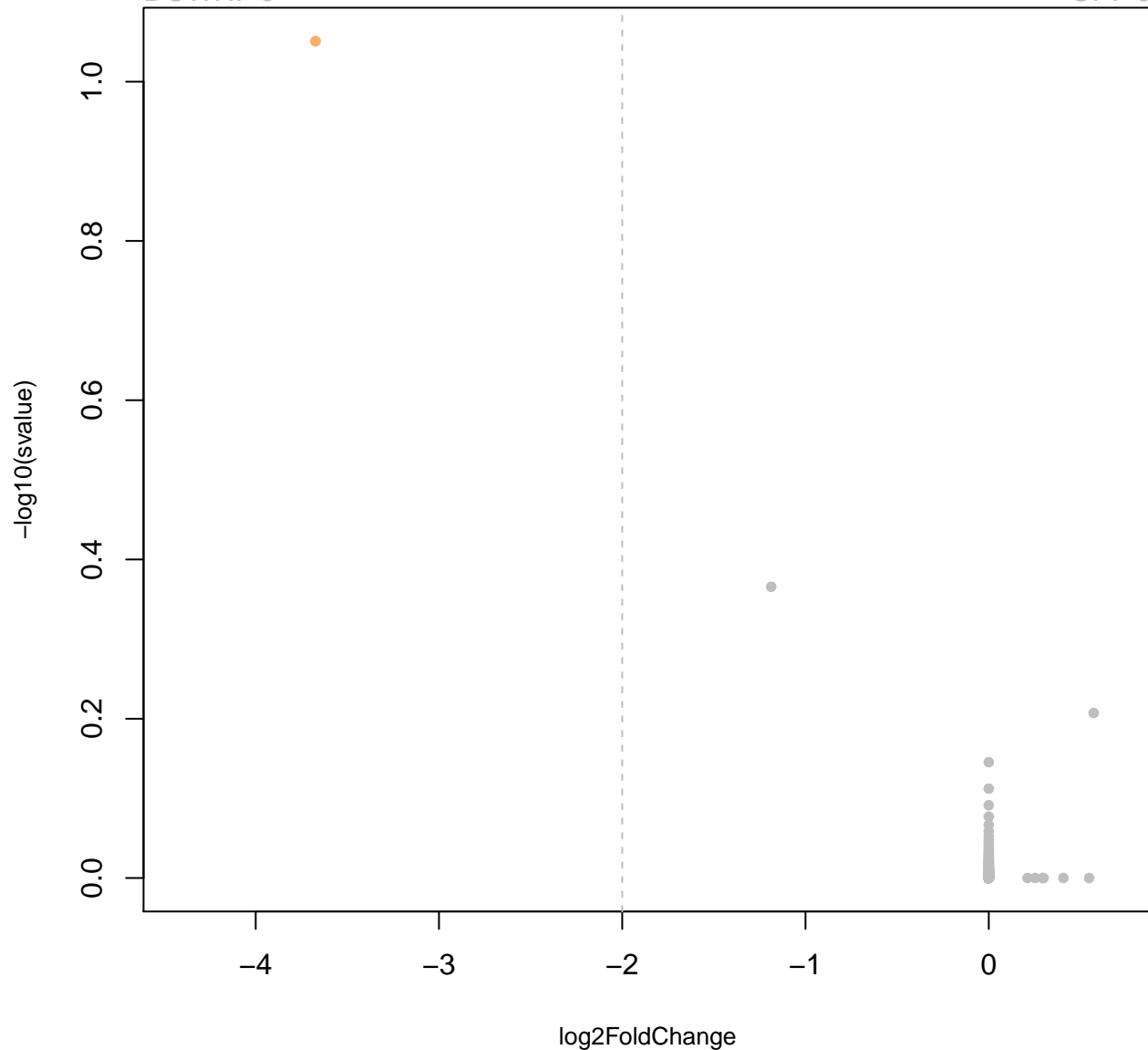
0 significant genes (L2FC > 2 & svalue < 0.005)