condition_33_Protected_vs_36_Protected

DOWN: 0

UP: 0

−log10(svalue)

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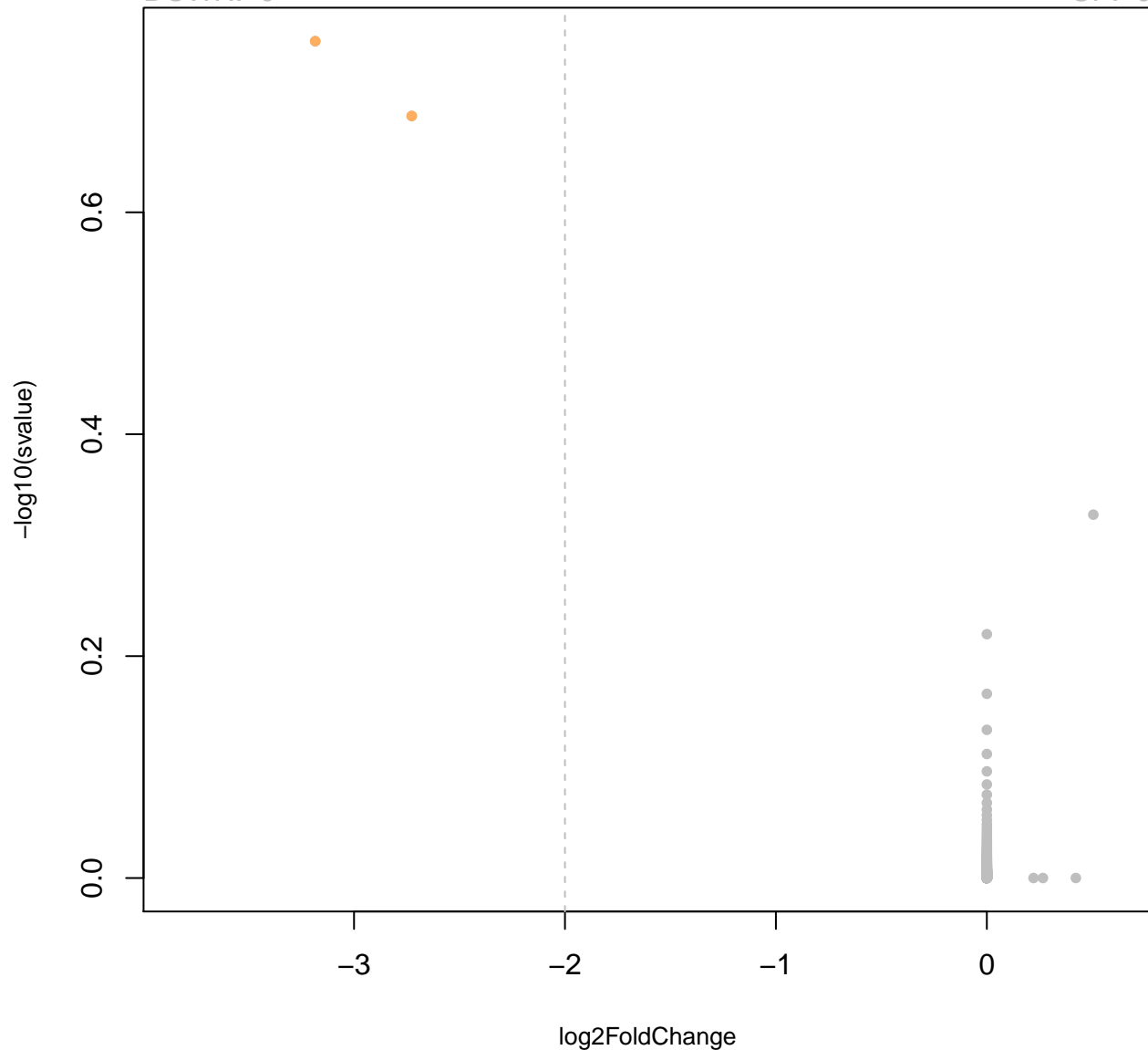
0 significant genes (L2FC > 2 & svalue < 0.005)