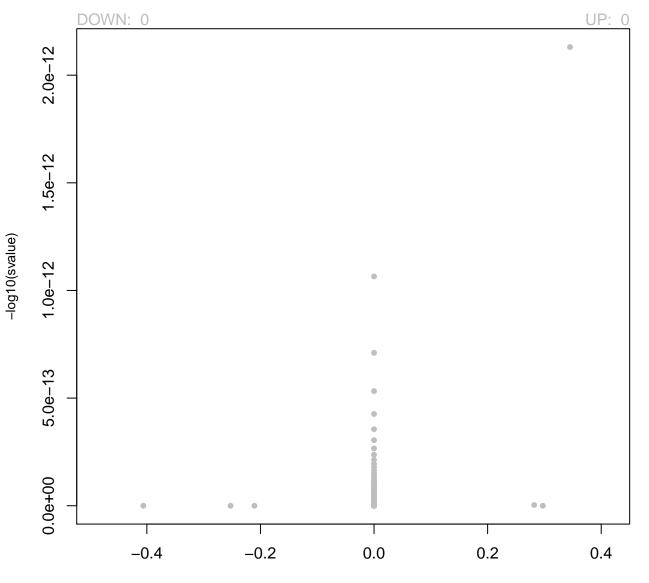
condition_36_PrT_vs_30_PrT



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