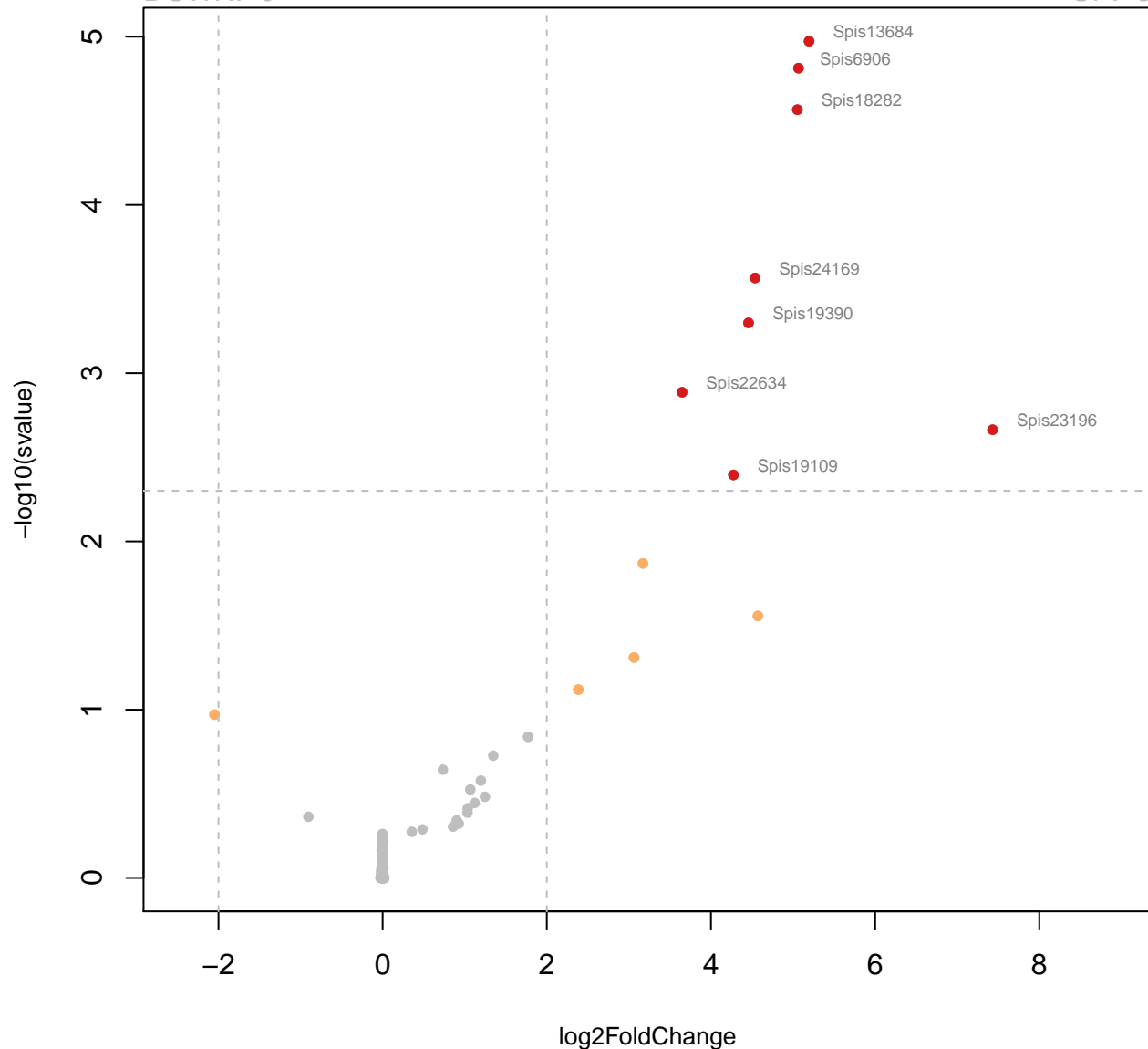


condition_36_Protected_vs_30_Protected

DOWN: 0

UP: 8



8 significant genes (L2FC > 2 & svalue < 0.005)