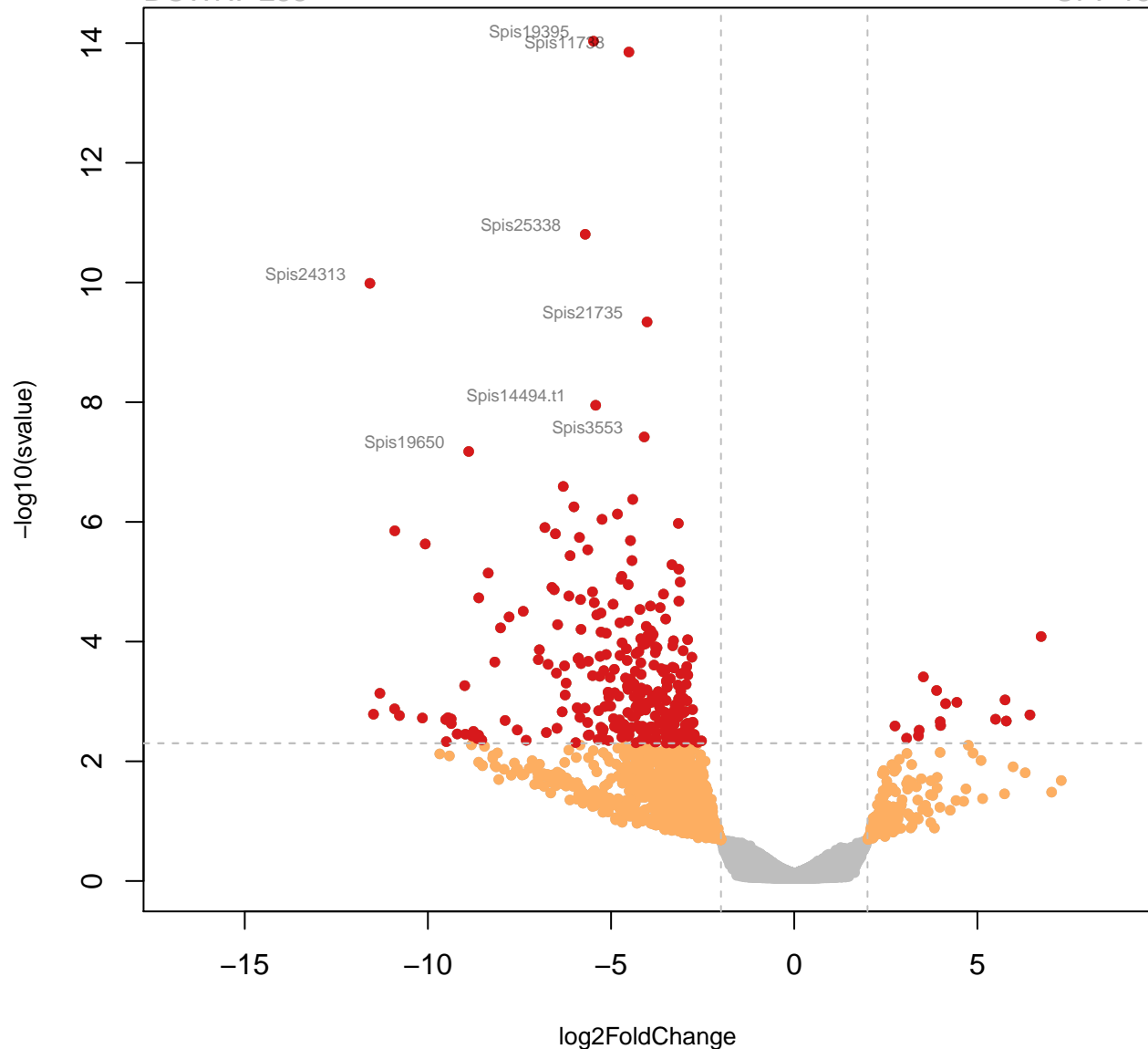


condition_30_ICN_vs_30_Protected

DOWN: 288

UP: 15



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