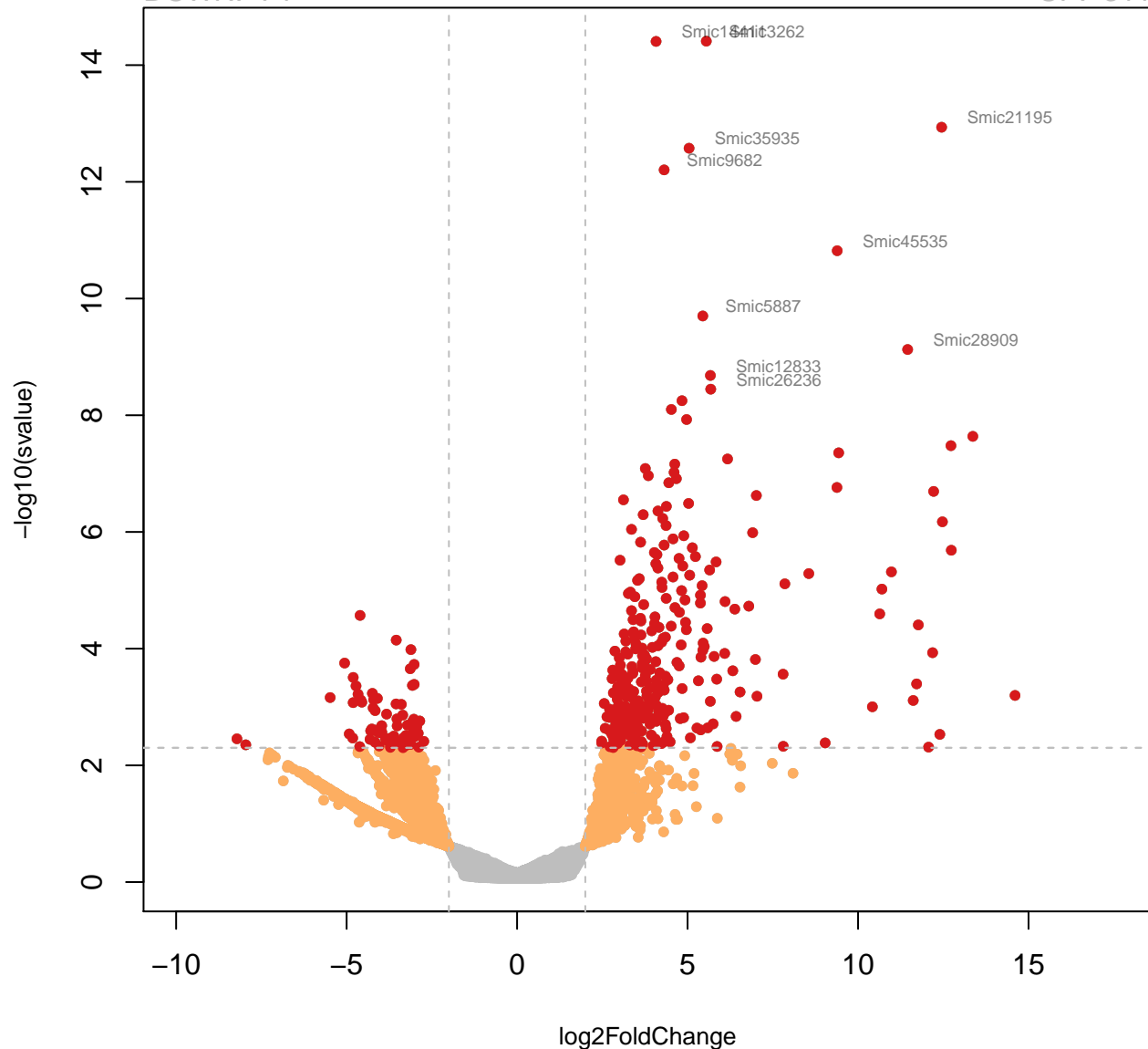


condition_33_ICN_vs_30_Protected

DOWN: 74

UP: 311



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