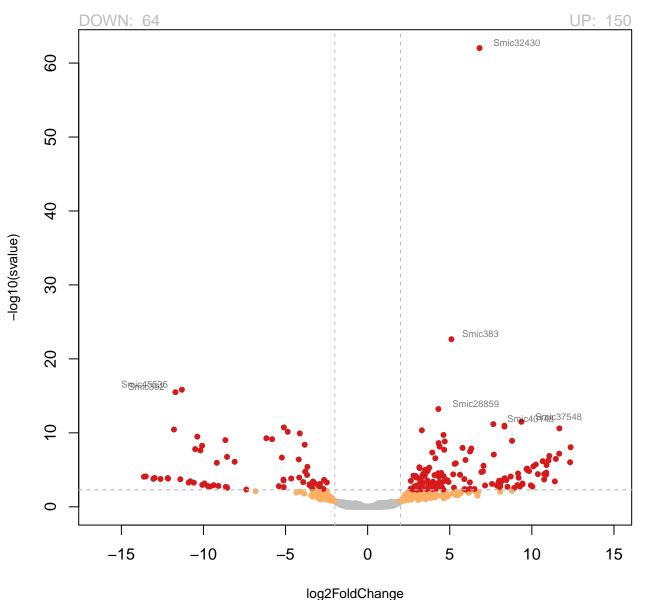
condition_30_Protected_vs_30_AF



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