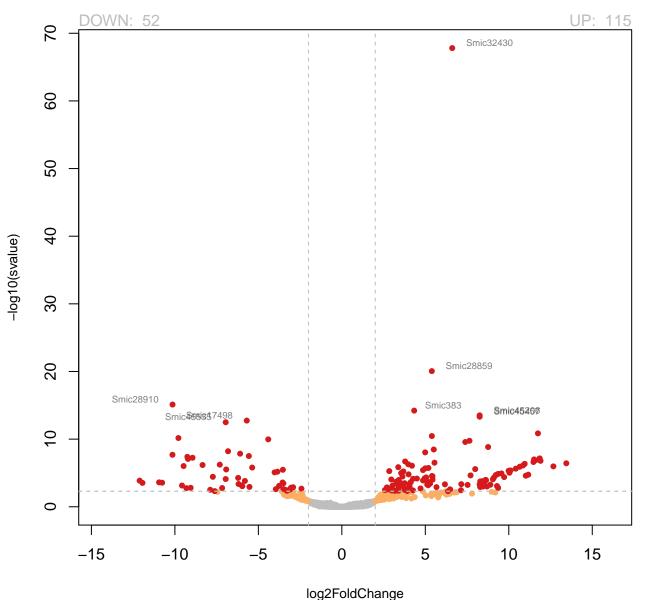
condition_36_Protected_vs_33_AF



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