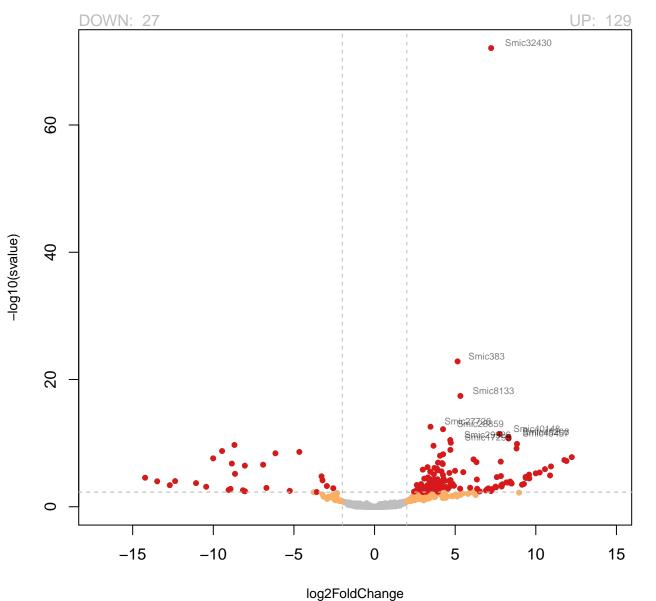
condition_30_ExT_vs_30_AF



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