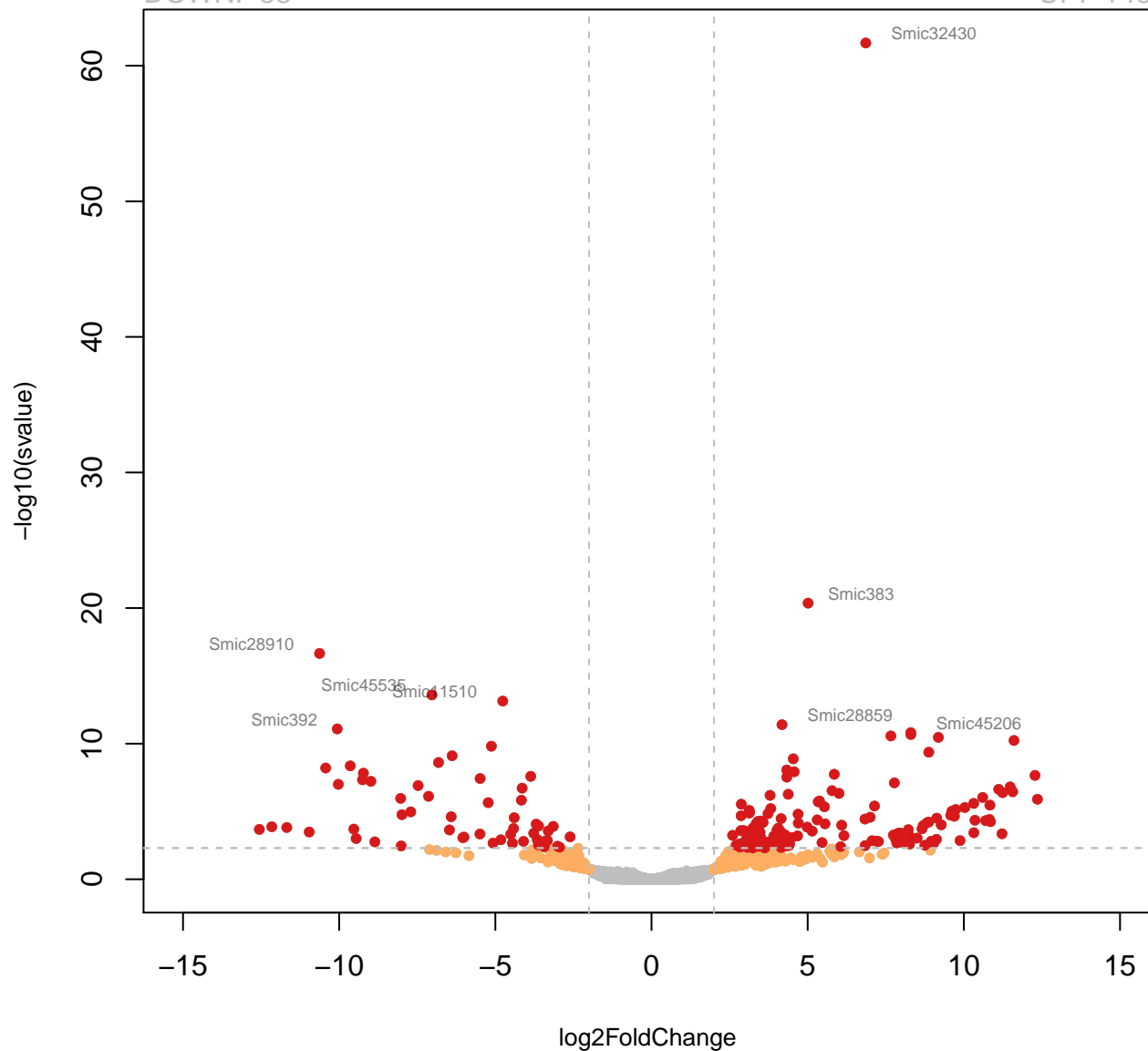


condition_36_Protected_vs_30_AF

DOWN: 58

UP: 146



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