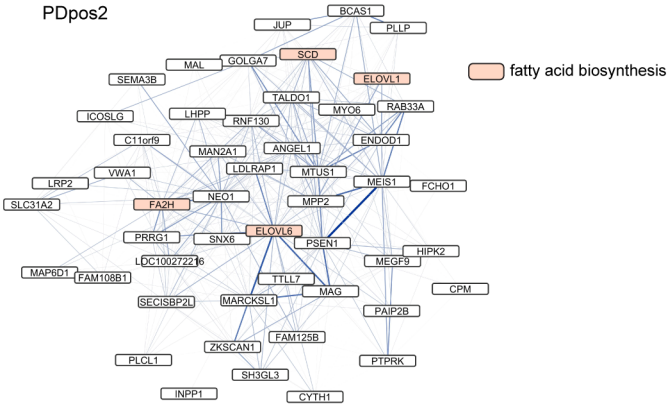

Additional data file 5. Visualization of hub genes in the PDpos2 module of the HD/PD consensus network. The 50 most connected genes (nodes) and the 500 strongest gene-gene interactions (edges) are shown. The width and the color saturation of the lines (edges) correspond to the weight of the interactions.