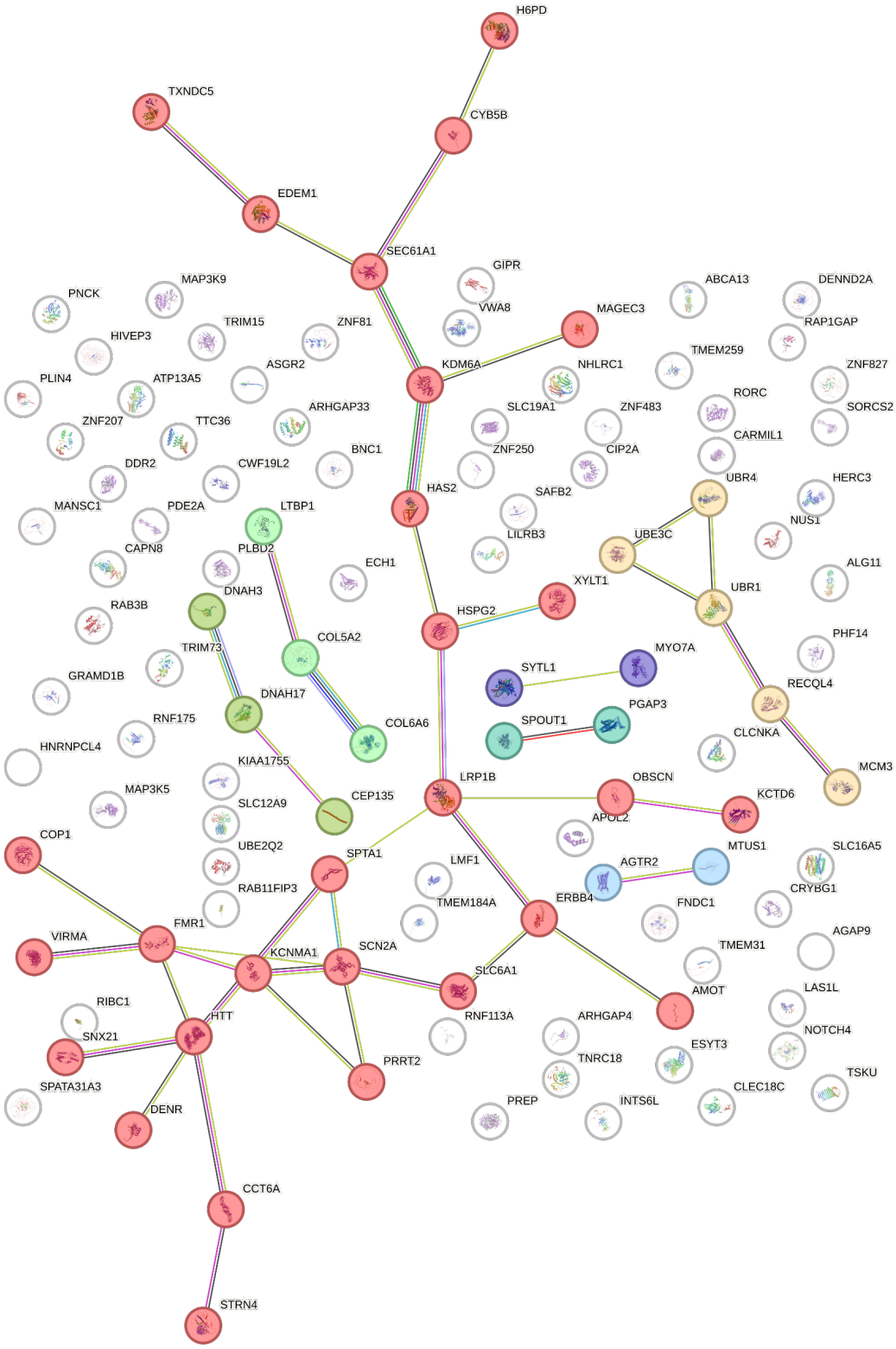

Suppl. Figure S1: STRING interaction enrichment analysis of the candidate genes identified two dominant enrichment networks represented by red (synaptic membranes) and yellow nodes (ubiquitin pathway).

Suppl. Fig. S1