Figure S7: GO analysis of clustered PPI network profile

iPAGE (http://tavazoielab.princeton.edu/iPAGE/) analysis of clustered protein-protein interaction data from BioGRID. Clusters include groups enriched for biological processes such as RNA processing and protein su-moylation, cellular components such as nucleoplasm and proteosomal complex, and molecular functions such as transferase activity. The columns represent protein clusters and correspond to the columns in Figure S8.