Table S4: GO enrichment of core proteins relative to all interactome proteins. Analysis was performed using the DAVID functional annotation chart with GO term Level of 5. Only terms with a corrected p-value < 10^{-5} with at least 50 genes in the query set are reported. DAVID identified 3,899 genes out of the 4,989 genes in the core interactome. The analysis was performed against a background list of 7,909 genes identified by DAVID out of 11,225 genes in the global interactome.

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