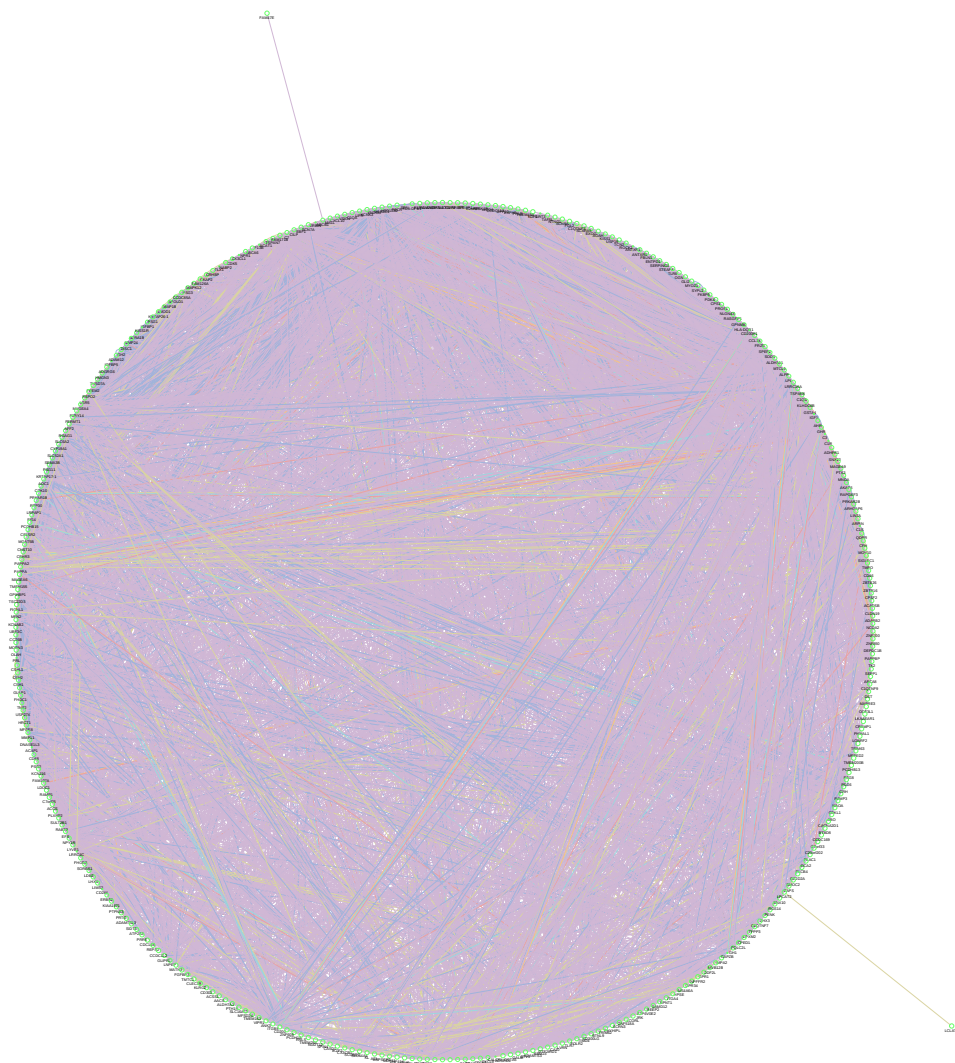


CO-EXPRESSION
CO-LOCALIZATION
PHYSICAL INTERACTIONS
GENETIC INTERACTIONS
SHARED PROTEIN DOMAINS
PATHWAY

Supplementary Figure 1: Interaction networks in differentially expressed genes from each experimental group. HPCVE were incubated during 2 and 24 h with 10^5 or 10^6 *T. cruzi* trypomastigotes. Interaction networks from differentially expressed genes ($FC \geq 2$) compared with uninfected control with GeneMANIA function prediction service plug-in in Cytoscape software. Co-expression, co-localization, physical interactions, genetic interactions shared protein domains and pathways are shown and each color represents specific interactions according to legend.



0.0 0.1 0.2 0.3 0.4 0.5 0.6 0.7 0.8 0.9 1.0

