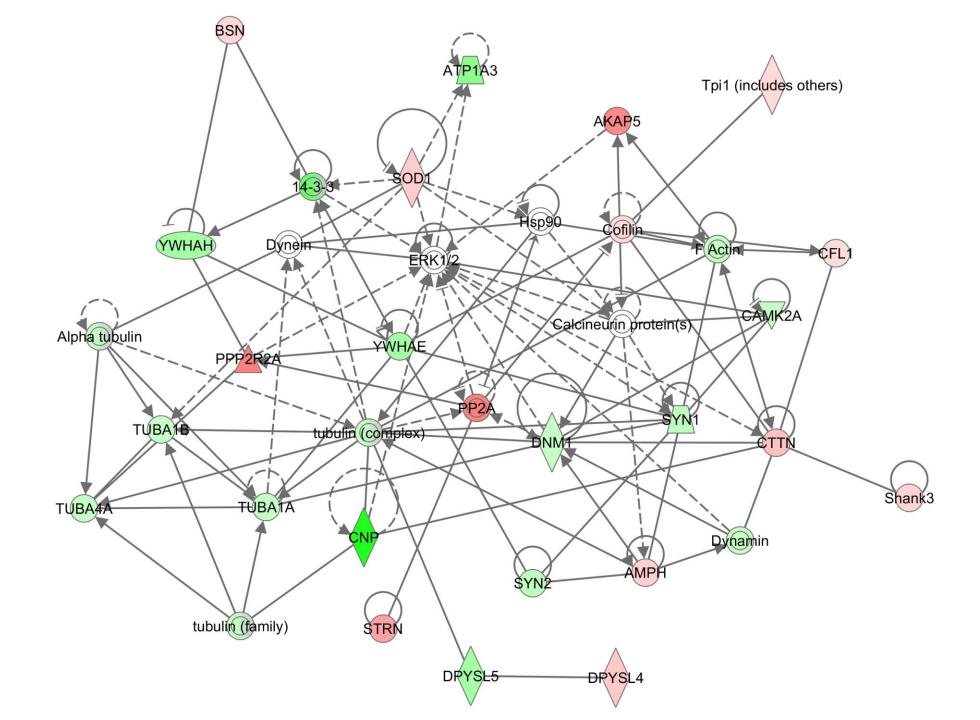
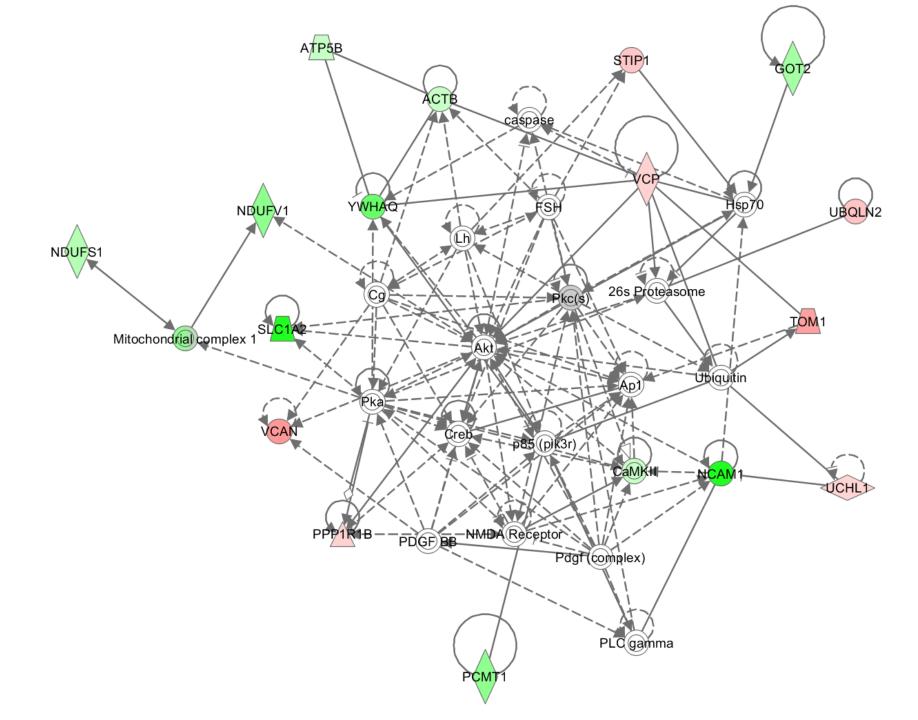
Supplementary Figure S3. Networks of differentially expressed proteins involved. (A) Cell-to-cell signaling and interaction, nervous system development and function and cellular assembly and organization, IPA score = 55; (B) Cell morphology, cellular assembly and organization and cellular development, IPA score = 31; (C) Cellular compromise, cell morphology, cellular assembly and organization, IPA score = 21. Proteins shaded in green indicate down-regulation and red means up-regulation. The intensity of the shade corresponds to the degree of up (*red*) or down (*green*) regulation. Proteins in white are those identified through the Ingenuity Pathways Knowledge Base. The shapes denote the molecular class of the protein. A solid line indicates a direct molecular interaction, and a dashed line indicates an indirect molecular interaction. The cut-off score of network identification was 20.

Network (A)



Network (B)



Network (C) MKK1/4 Focal adhesion kinase PI3K (complex) ARHGEF25 Histone n3 PPM1E S(TMN4 ALDOA NEKB (complex) PEBP1 PAK1IP1 Calmodulin MYLK3 **Ж**НО)Н Cathrin MDH2 Tpm3