S4 Figure. Integration of the identified dysregulated proteins into networks: Network #4 - Behavior, Cell Morphology, Cellular Function and Maintenance. Six molecules were affected and IPA score was 10. Solid lines indicate direct interaction. Dashed lines indicate indirect interactions. Red molecules were up-regulated and green molecules were down-regulated. White molecules were not user specified, but were incorporated into the network through relationships with other molecules. Of particular note were the network hubs centered on CNNG, QDPR and LAP3.