

Figure S1. Overview of the RNA-Seq analysis. (Top) Overview of the PCA analysis. (Middle) Overview of the MDS analysis. (Bottom) The volcano plot shows the differential expression analysis of the comparison lyco-s vs. buf-s

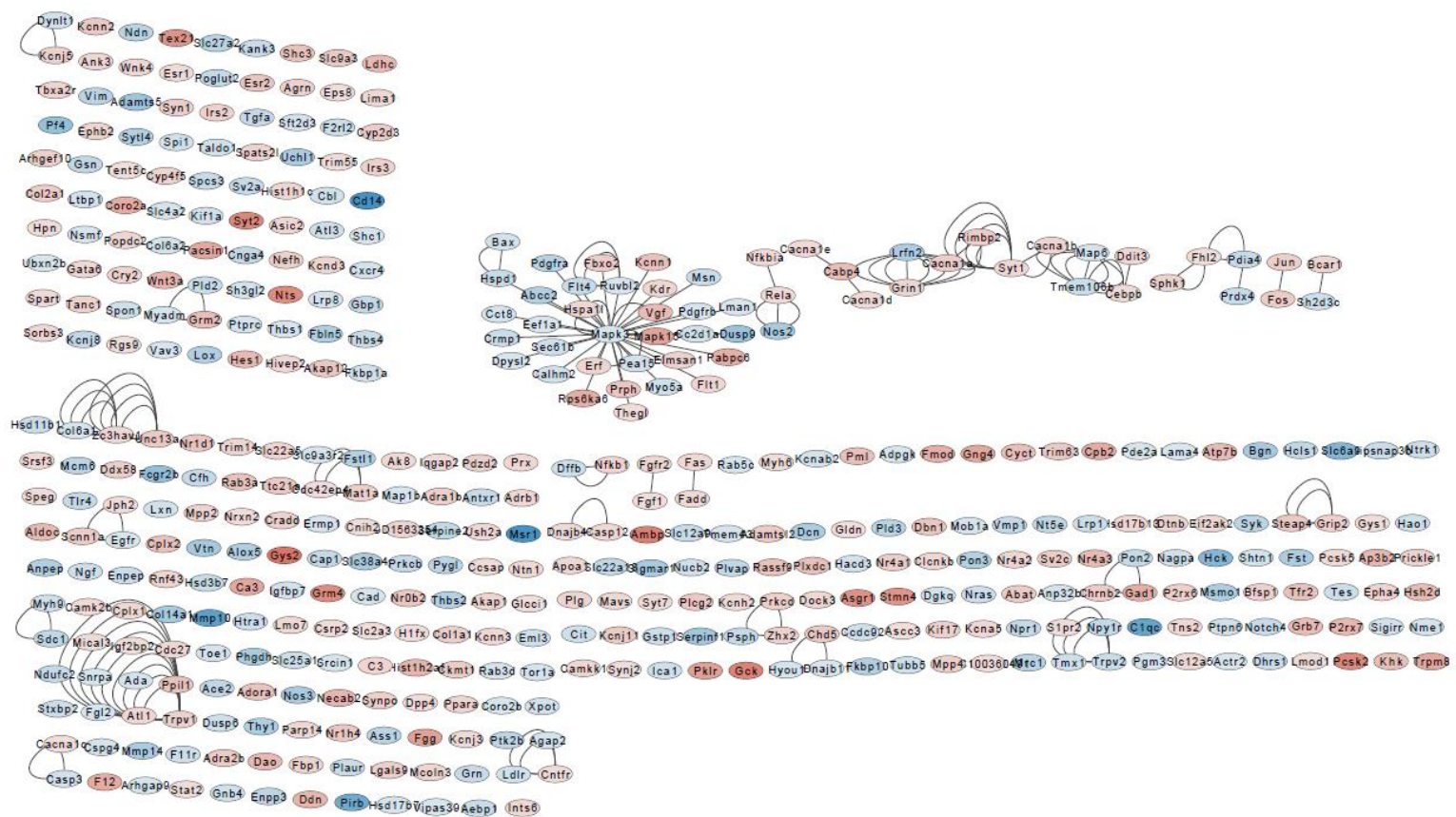


Figure S2. Mapping of the DEGs to the rat interactome results in a molecular interactome of 423 nodes and 113 edges (each node represents a DEG, edges interactions; color scaling shows the deregulation of genes (blue: downregulated, red: upregulated))