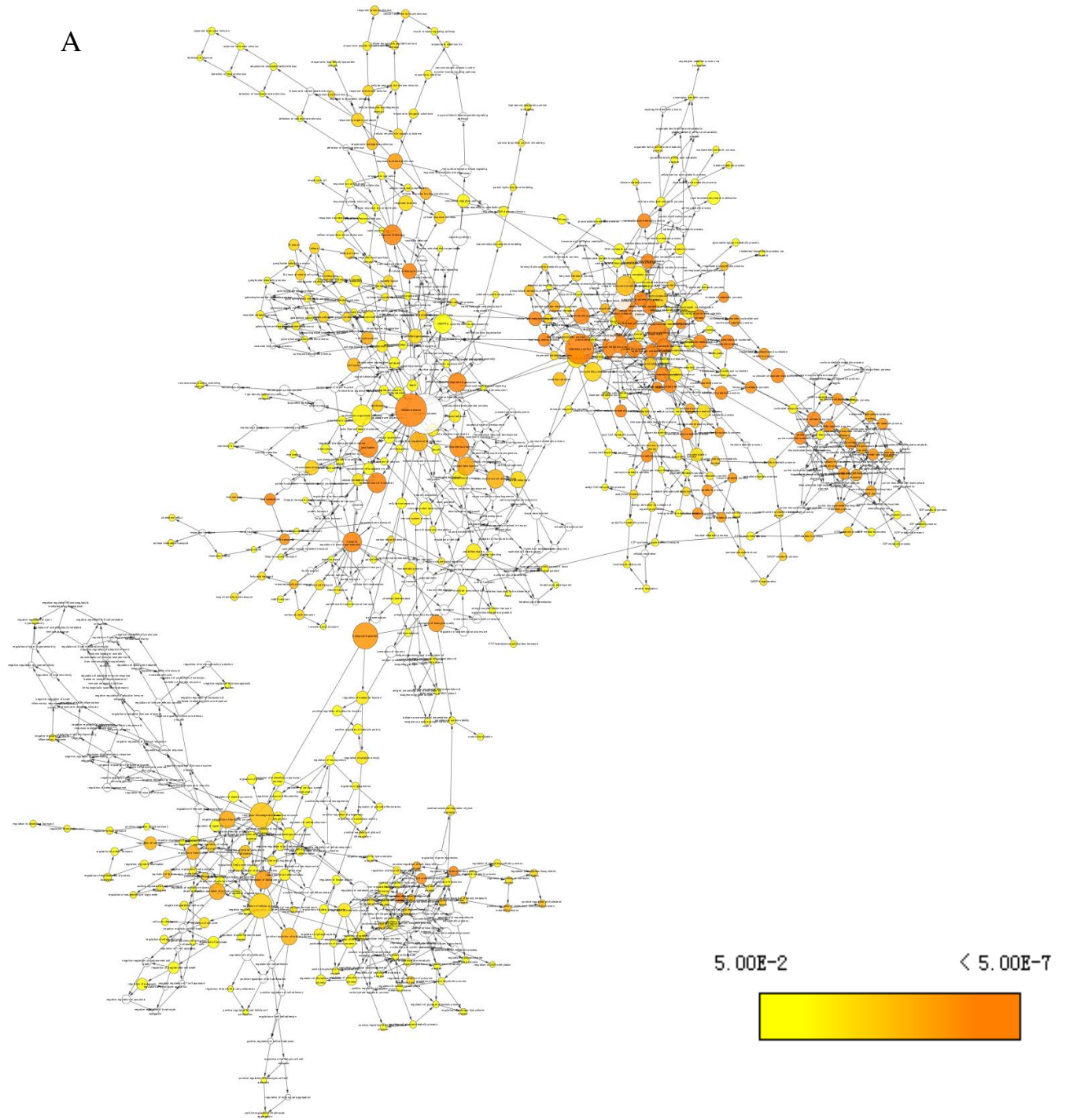


Supplementary Material

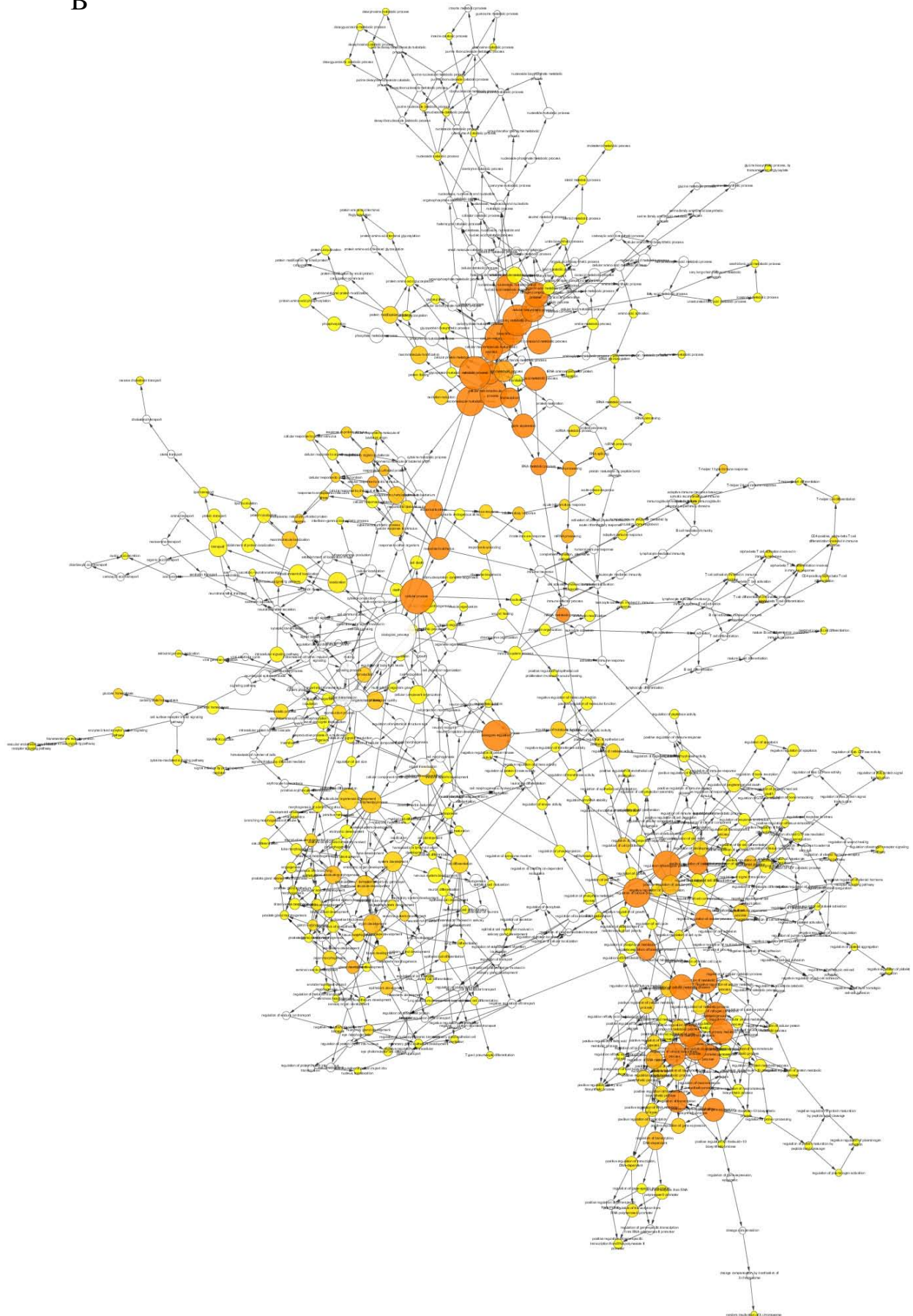
ATP Citrate Lyase and LncRNA NONMMUT010685 Play Crucial Role in Nonalcoholic Fatty Liver Disease Based on Analysis of Microarray Data

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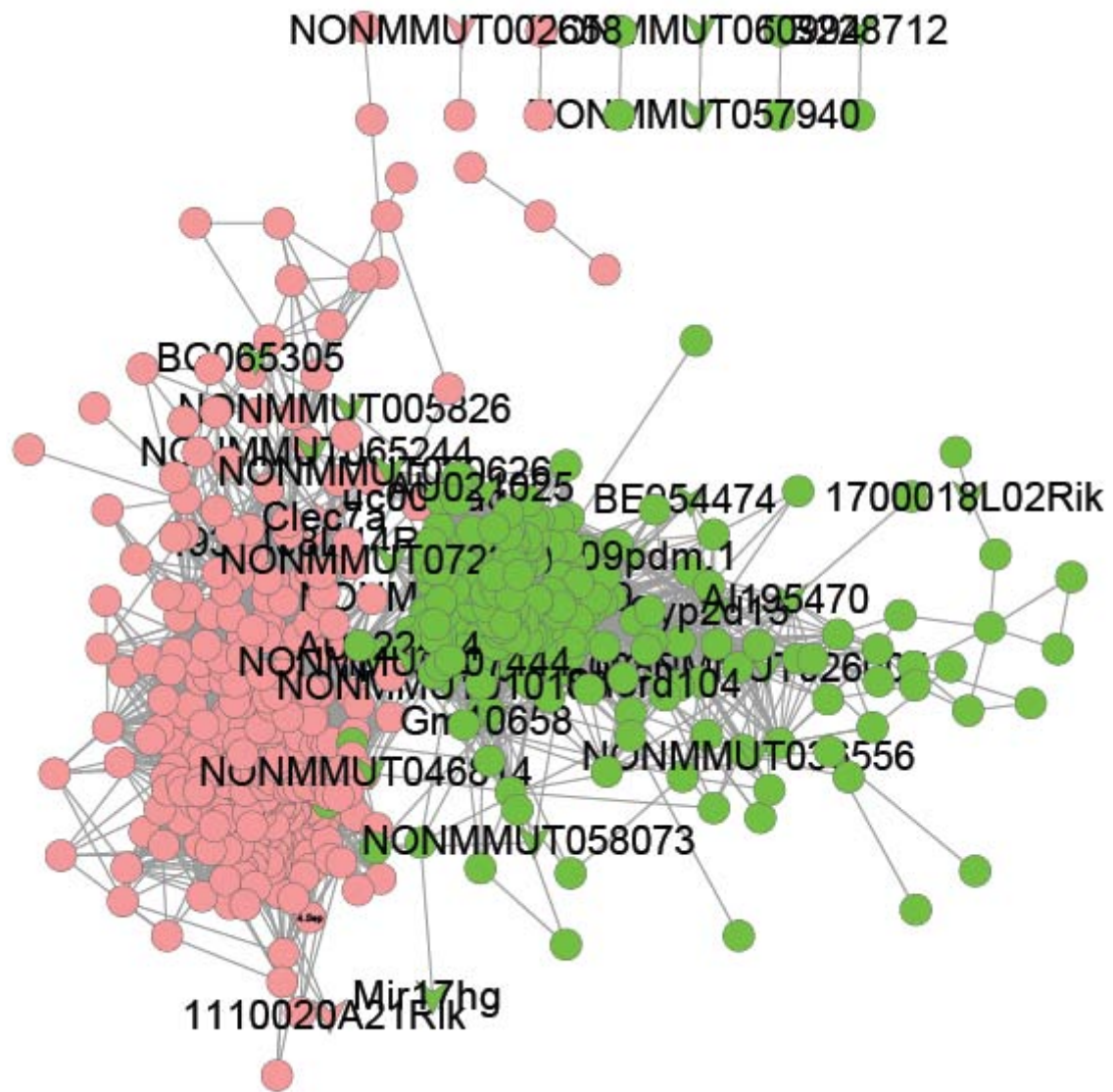


B



Down regulated genes GO-net

A



B

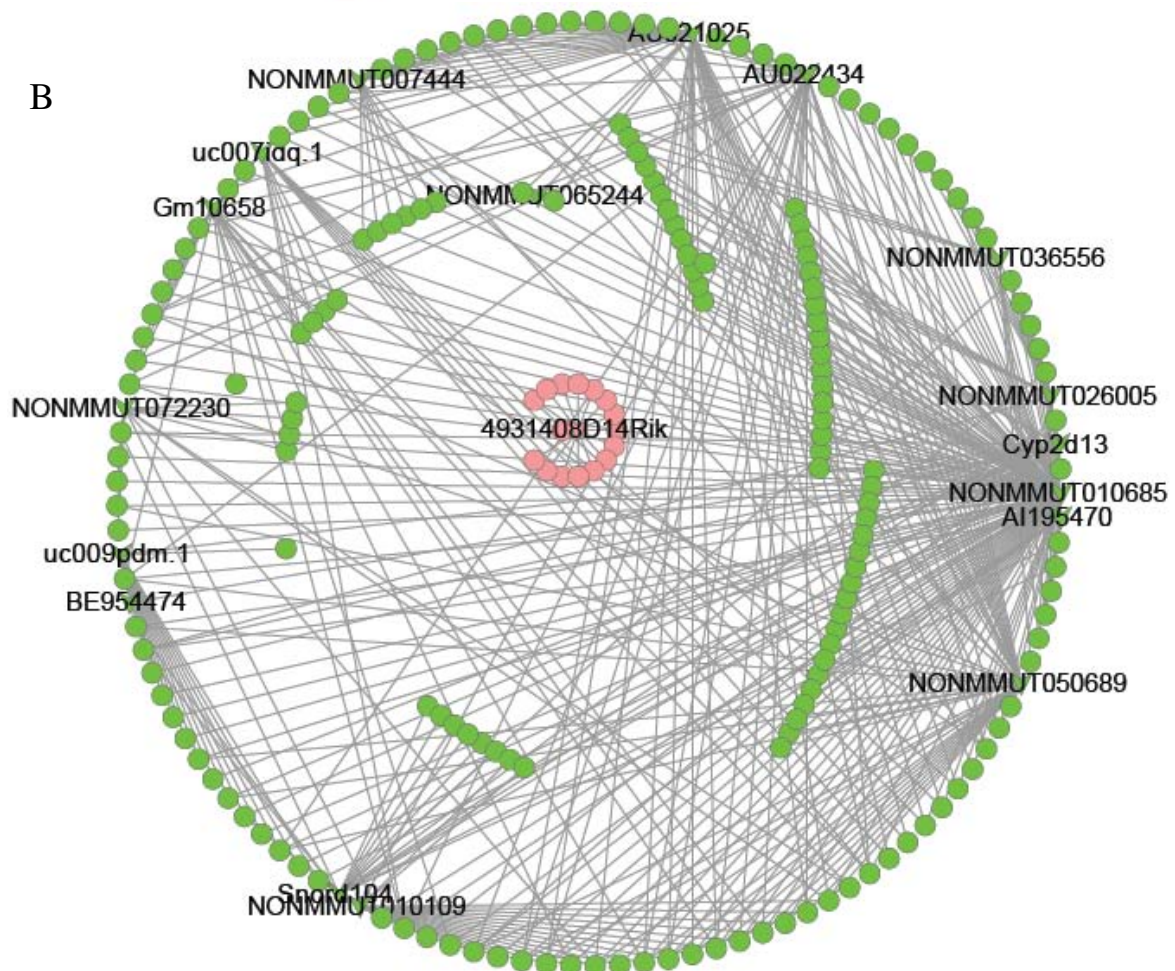


Fig. S3 Ma et al

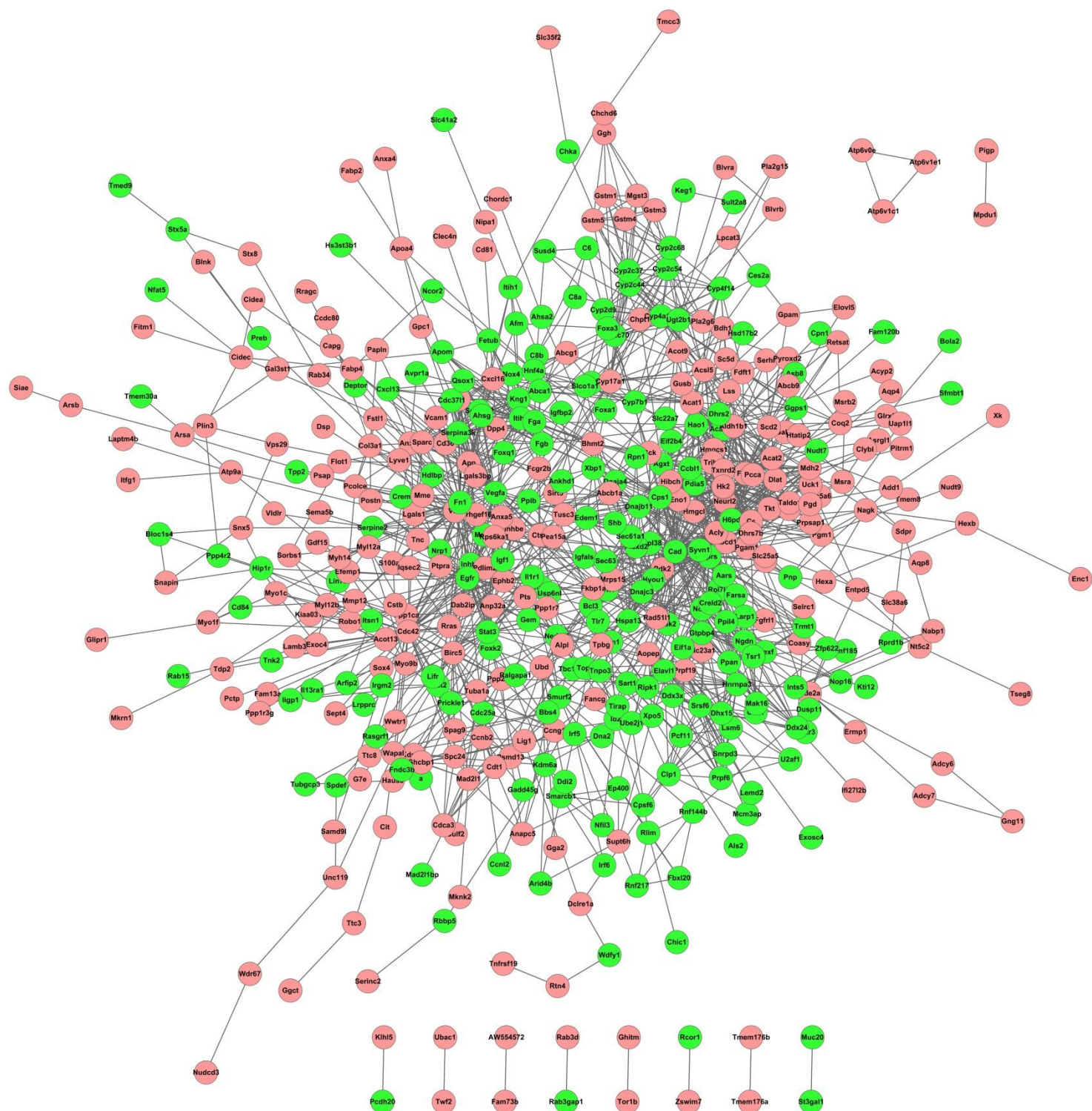


Fig. S1. Go-net analysis of differentially expressed mRNAs.

- A. Up-regulated genes GO-net.
- B. Down-regulated genes GO-net.

Fig. S2. Co-expression between differentially expressed lncRNAs and mRNAs.

- A. Co-expression network with statistically differentially expressed lncRNA and mRNA
- B. Subnetwork of lncRNA node degree>10

Fig. S3. Significantly up and down-regulated genes from the protein-protein interaction network. (Up-regulated genes were labeled in red, down-regulated genes were labeled in green.)