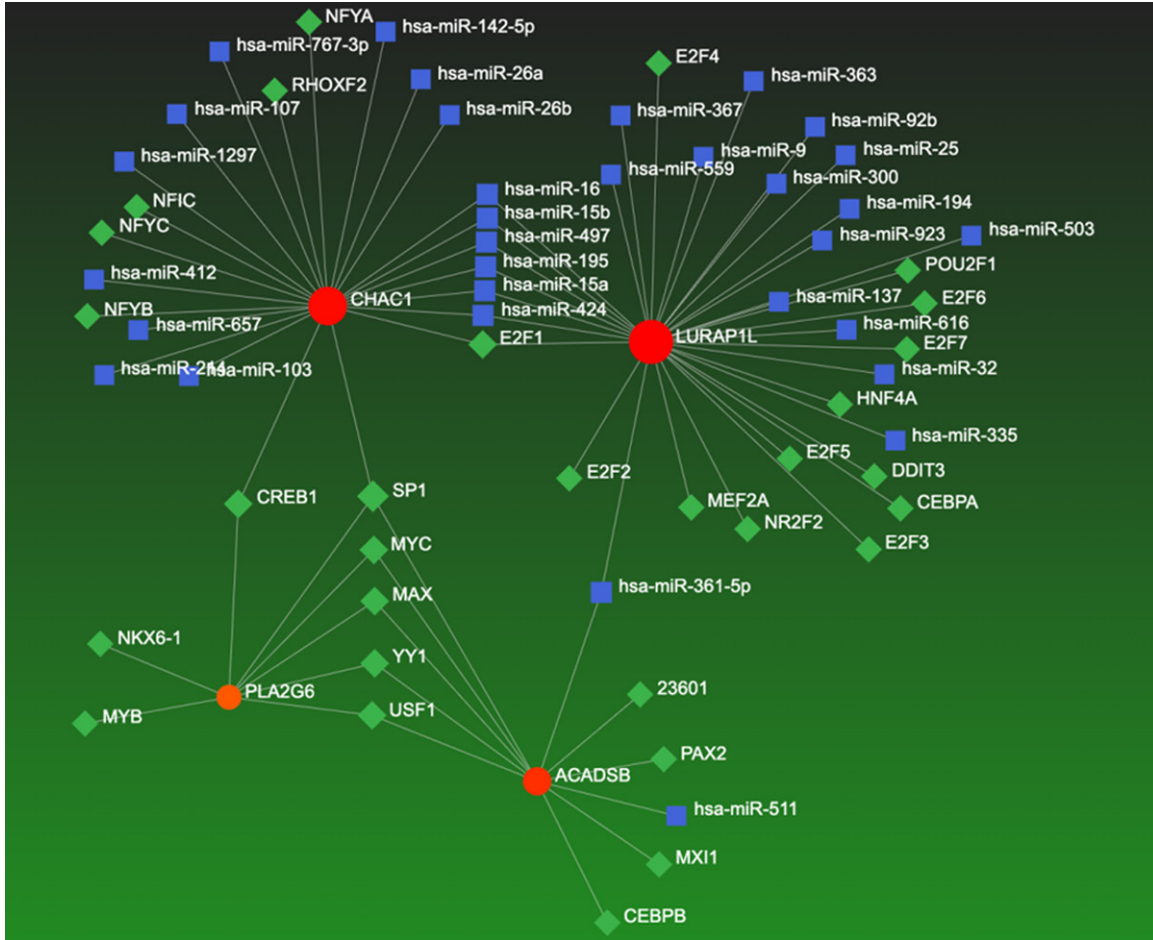


Immune ferroptosis gene in KIRC



Supplementary Figure 1. The network depicts the TF-miRNA coregulatory network. It consists of 66 nodes and 77 edges, including 29 TF-genes, 32 miRNAs, and 4 expressed genes. The nodes in red are the model expressed genes, the blue nodes represent miRNA, and the green nodes indicate TF-genes.