

Supplementary Figure 1

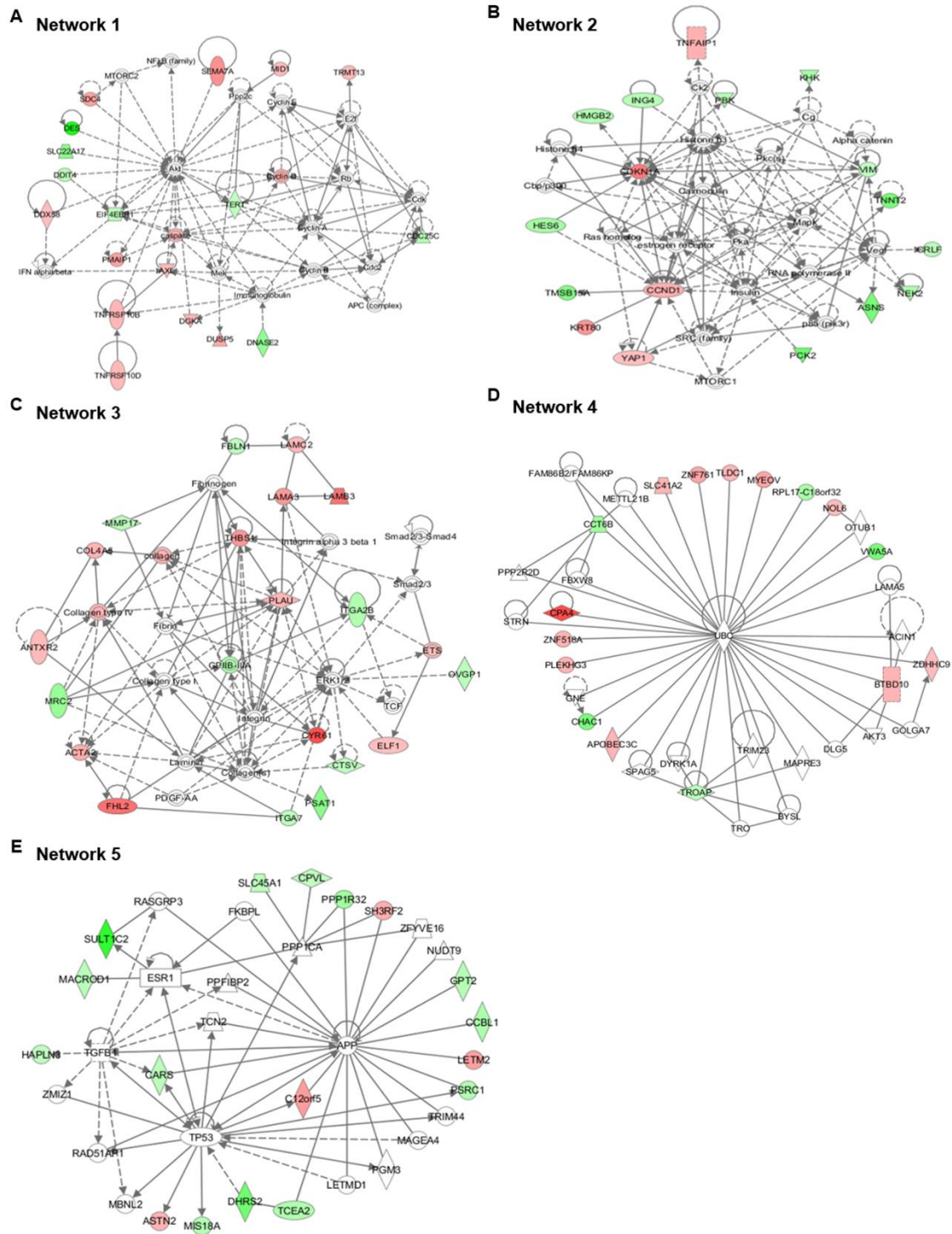


Figure S1. Functional connectivity of 159 commonly dysregulated genes in HCT119 and HT29 cells treated with RPL17siRNA for 48 h. (A-E), Five putative IPA networks with high score (>19). Up- and down-regulated genes are shown in red and green, respectively. Genes in gray are associated with the regulated genes.