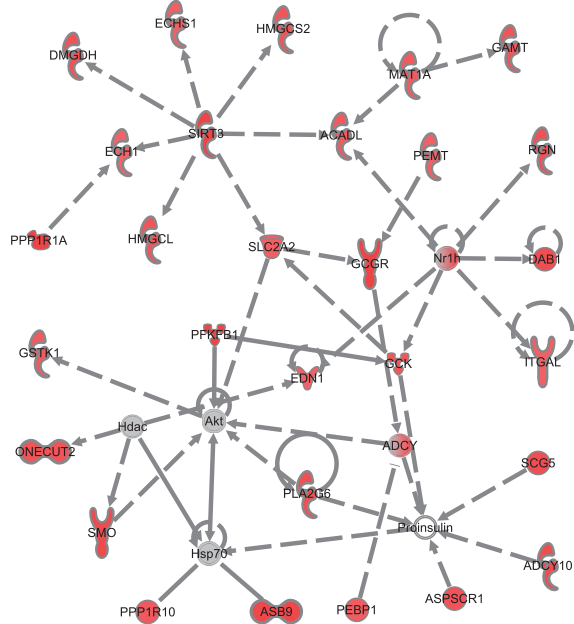
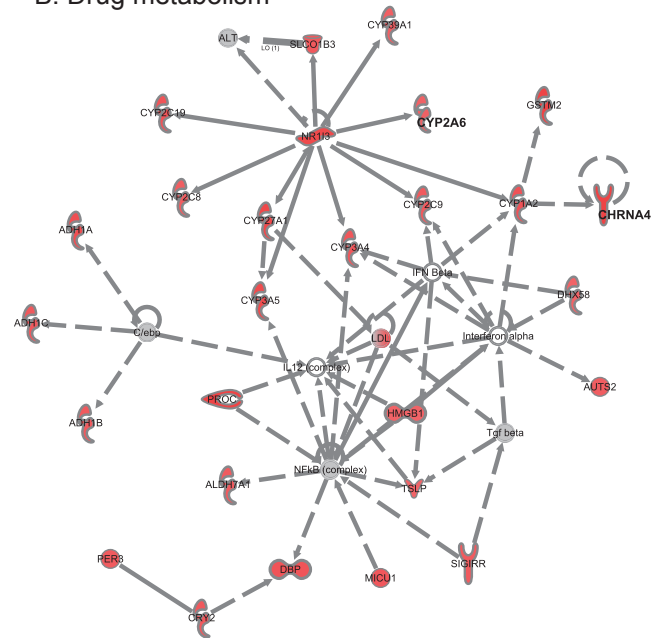


Supplementary figure5. IPA identified top3 enriched network modules in genes positively correlated to CHRNA4 in 119 human liver samples.

A: Lipid metabolism



B: Drug metabolism



C: Cell cycle

