



Supplementary Figure 3. Top networks identified using ingenuity pathway analysis of genes differentially expressed among the MA-, cocaine-, and heroin-treated. A: The top network of genes differentially expressed between the MA- and cocaine-treated groups. B: The top network of genes differentially expressed between the MA- and heroin-treated groups. C: The top network of genes differentially expressed between the cocaine- and heroin-treated groups. The intensity of the node (gene or gene products) color indicates the degree of upregulation (red) or downregulation (green). D: Node legend. MA, methamphetamine.