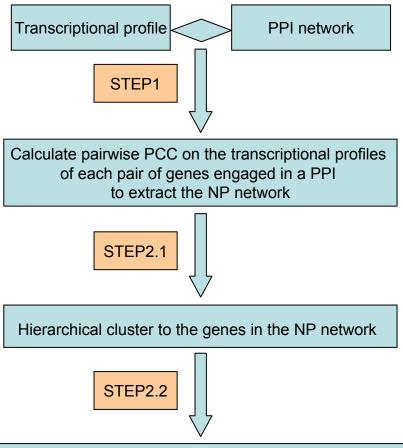
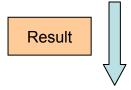
Supplementary Figure 1



Manually dissect the largest anti-correlated clusters or automatically scan from the top of the hierarchical tree for clusters that have <1% intra-cluster anti-correlated interactions



Modules within which gene expression profiles are similar, and correlated interactions are maximally enclosed and in between which anti-correlated interactions are optimally distributed