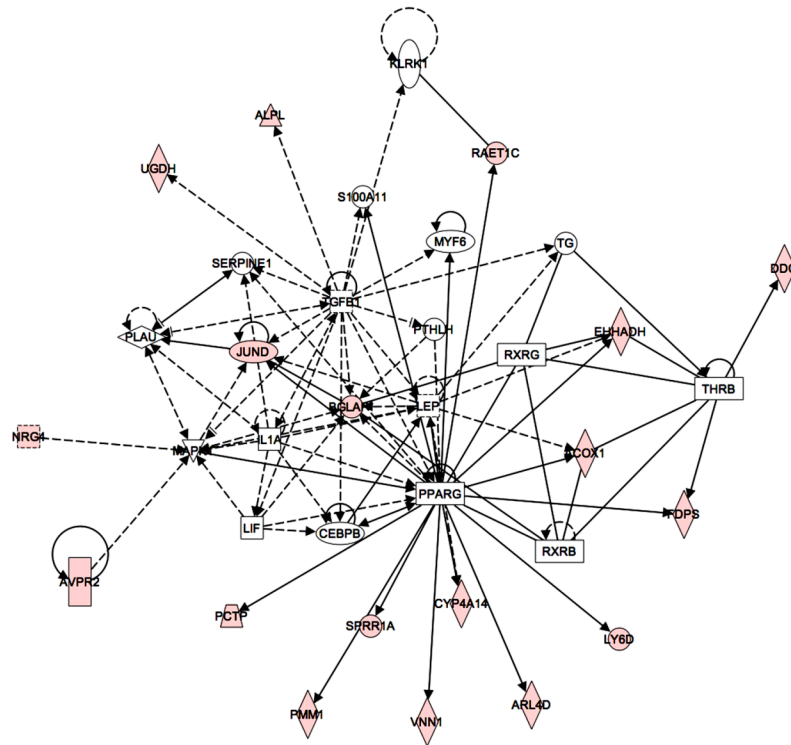


### Supplementary Figure S3. Ingenuity Pathway Analysis of P17 and P28 microarray data.

(A) Computer-assisted identified transcriptional networks denoting up-regulation of *Pparγ* and genes regulated by *Pparγ* at P17 (top) and P28 (bottom). Red shading correlates with increased gene expression, with darker shading indicating a higher fold change.

#### P17 Network



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#### P28 Network

