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

P28 Network