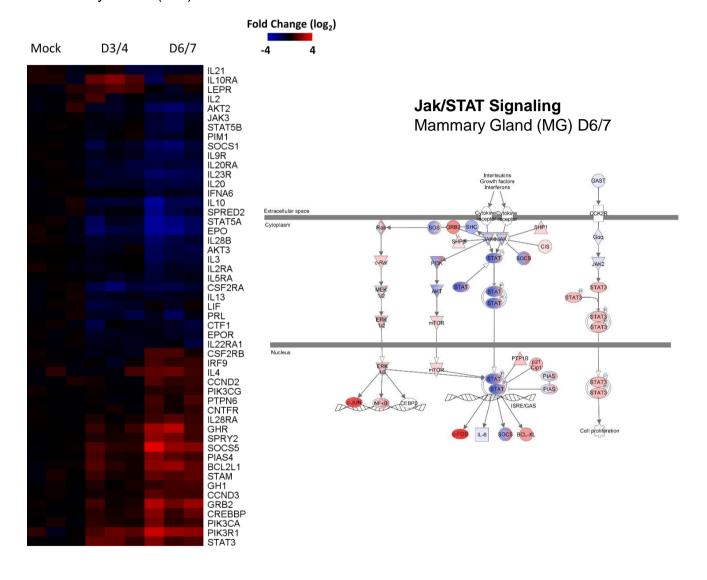
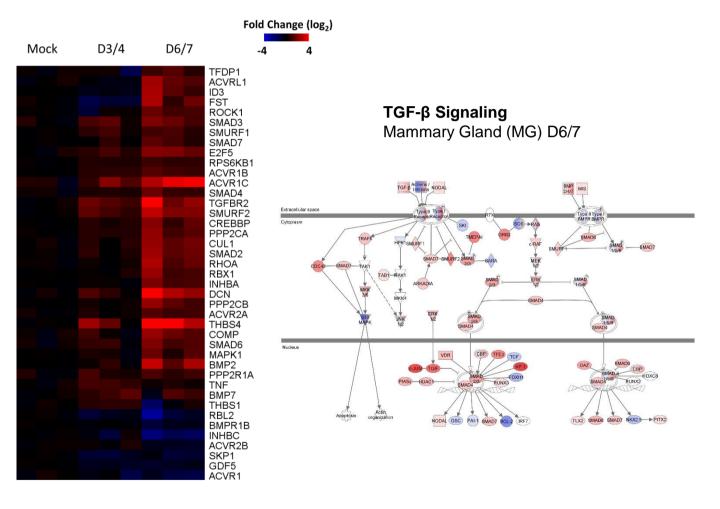
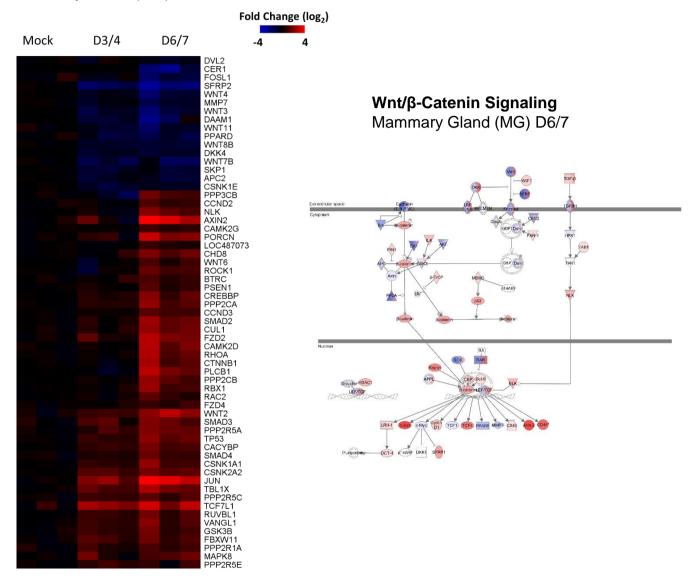
Jak-STAT Signaling Pathway; cfa04630



TGF-β Signaling Pathway; cfa04350



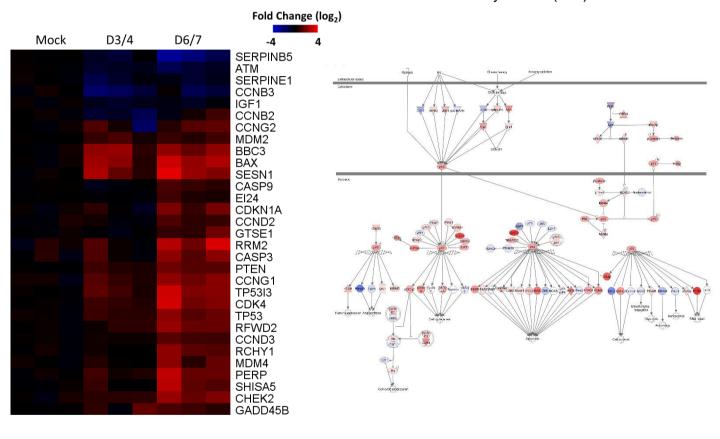
Wnt Signaling Pathway; cfa04310



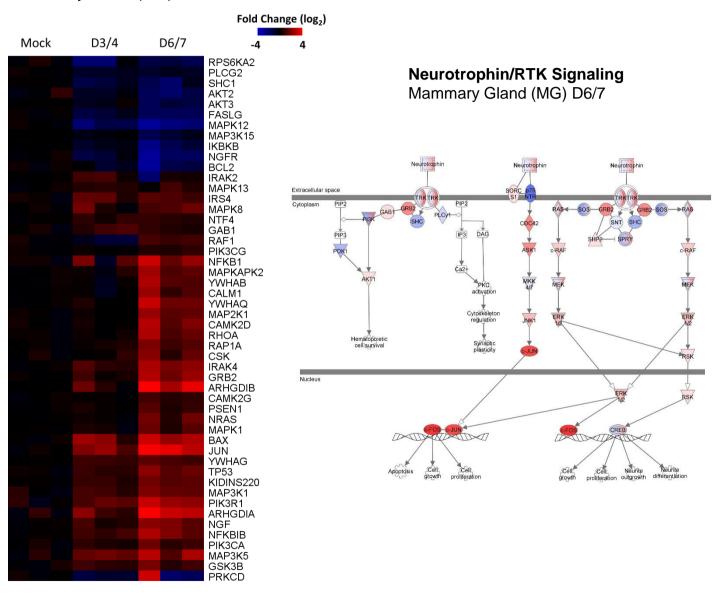
p53 Signaling Pathway; cfa04115

Mammary Gland (MG)

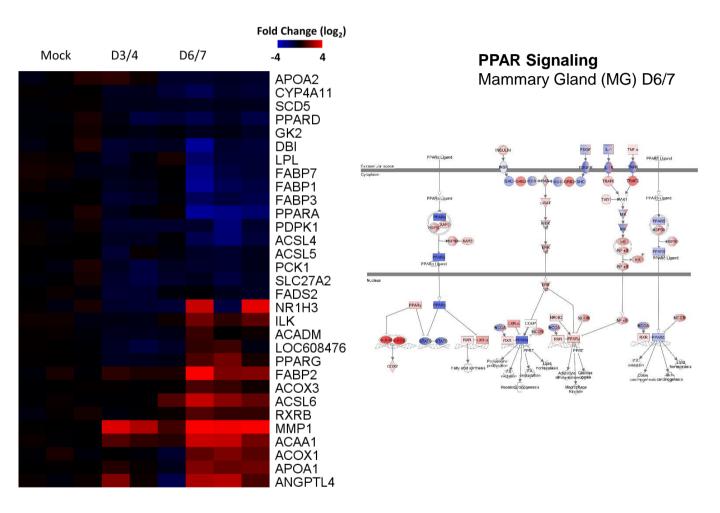
p53 SignalingMammary Gland (MG) D6/7



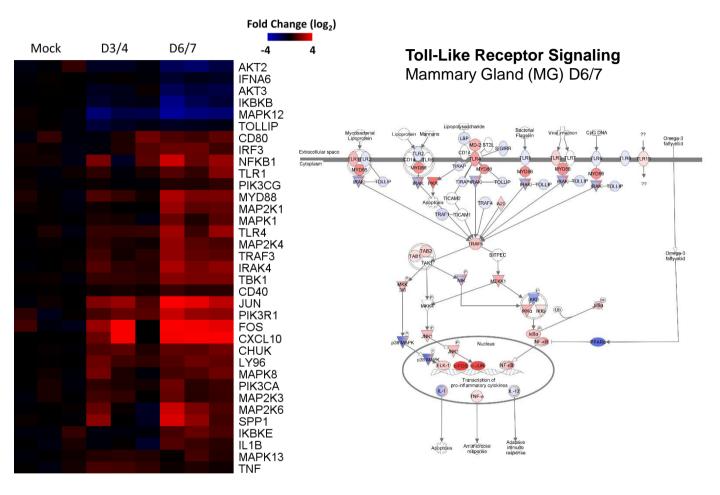
Neurotrophin Signaling Pathway; cfa04722



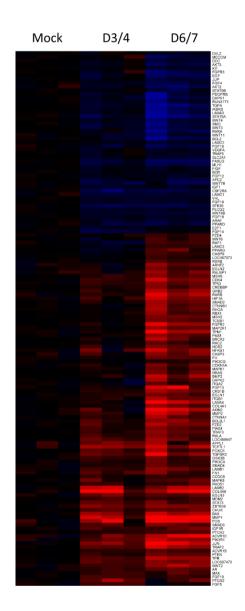
PPAR Signaling Pathway; cfa03320



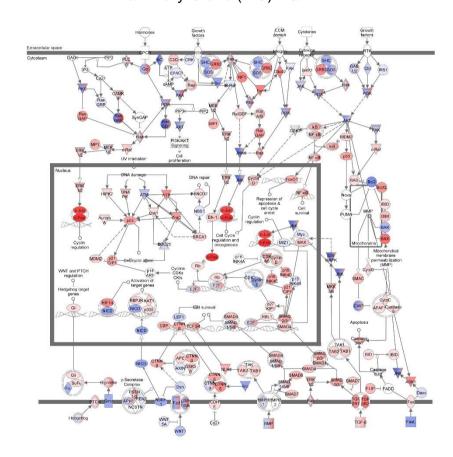
Toll-Like Receptor Signaling Pathway; cfa04620



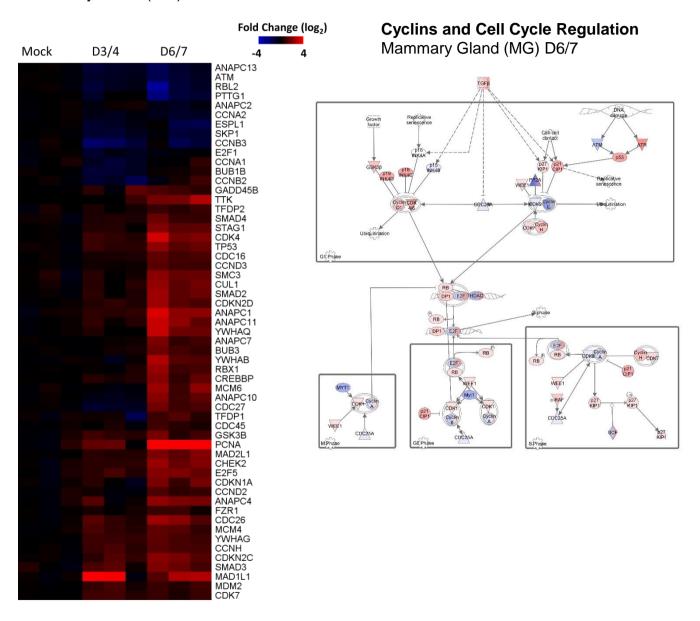
Pathways in Cancer; cfa05200



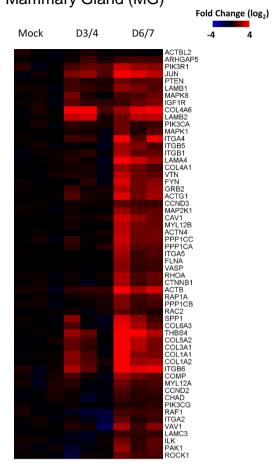




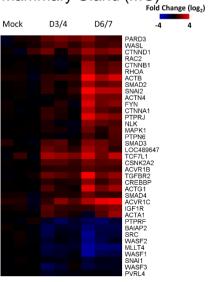
Cell Cycle; cfa04110 Mammary Gland (MG)



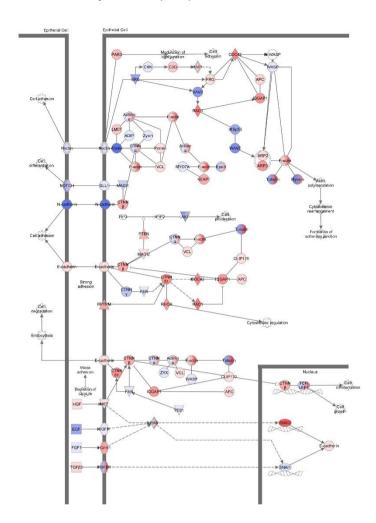
Focal Adhesion; cfa04510 Mammary Gland (MG)



Adherens Junction; cfa04520 Mammary Gland (MG)



Epithelial Adherens Junction Signaling Mammary Gland (MG) D6/7



Network Shapes	
	Cytokine
	Growth Factor
	Chemical / Drug/ Toxicant
\Diamond	Enzyme
	G-protein Coupled Receptor
	Ion Channel
∇	Kinase
	Ligand-dependent Nuclear Receptor
\Diamond	Peptidase
Δ	Phosphatase
\bigcirc	Transcription Regulator
\bigcirc	Translation Regulator
0	Transmembrane Receptor
	Transporter
	microRNA
0	Complex / Group
0	Other

*For dataset files that contain only identifiers (i.e. no expression values), the color gray identifies the Focus Genes from that dataset. Red User input molecule that is upregulated (ie. has a positive (+) expression value) and whose expression value meets the user defined cutoff. User input molecule that is downregulated (ie. has a negative (-) expression value) Green and whose expression value meets the user defined cutoff. Gray User input molecule. Neither up nor down-regulated or does not meet the userdefined cutoff. Molecule that is not user specified, but incorporated into the network through White relationships with other molecules. [Blue] For canonical pathways, molecules that are members of the network being examined are outlined in blue.