

CS vs. HM				Set Name	Description
1d	2d	3d	8d		
Cholesterol metabolism					
				HSA00120_BILE_ACID_BIOSYNTHESIS	KEGG bile acid synthesis genes
				MONOOXYGENASE_ACTIVITY	(GO MF) integration of one oxygen atom into a compound
				CELLULAR_LIPID_METABOLIC_PROCESS	(GO BP) lipid reactions and pathways
				CARBOXYLIC_ACID_TRANSM_TRANSP	(GO MF) transfer of carboxylic acid across a membrane
				LIPID_TRANSPORT	(GO BP) transport into, out of, or between cells
				NUCLEAR_RECEPTORS	GenMAPP nuclear receptor genes
Fatty-Acid Metabolism					
				HSA00071_FATTY_ACID_METABOLISM	KEGG fatty acid metabolism pathways
				MITOCHONDRIAL_FATTY_ACID_BETA	GenMAPP fatty acid oxidation in mitochondria
				PEROXISOME	(GO CC) associated with peroxisome
				HSA03320_PPAR_SIGNALING_PATHWAY	KEGG PPAR signaling pathway
Alcohol Metabolism					
				ALCOHOL_METABOLIC_PROCESS	(GO BP) reactions and pathways involving alcohols
				HSA00071_FATTY_ACID_METABOLISM	KEGG fatty acid metabolism pathways
				HSA00980_METABOLISM_OF_XENOBIOTICS	metabolism of xenobiotics by cytochrome P450
Carbohydrate Metabolism					
				GLUCOSE_METABOLIC_PROCESS	(GO BP) pathways involving glucose
				HSA00010_GLYCOLYSIS_AND_GLUCON	KEGG glycolysis and gluconeogenesis pathways

Color legend for up-regulated <i>q</i> -values						
<i>q</i> -value	1	0.2	0.05	0.01	0.001	0.0001
Color						