-log10(P)	
0246 10	20
	M2674: PIGF UP.V1 UP GO:0044772: mitotic cell cycle phase transition
	GO:0043484: regulation of RNA splicing
	M121: PID MTOR 4PATHWAY M2639: GCNP SHH UP EARLY.V1 UP
	GO:0016236: macroautophagy
	GO:000082: G1/S transition of mitotic cell cycle GO:0006399: tRNA metabolic process
	R-HSA-212165: Epigenetic regulation of gene expression
	GO:0031503: protein-containing complex localization
	GO:0006281: DNA repair GO:1902115: regulation of organelle assembly
	GO:0045861: negative regulation of proteolysis M2728: RAF UP.V1 UP
	R-HSA-3700989: Transcriptional Regulation by TP53
	GO:0018205: peptidyl-lysine modification
	GO:0010638: positive regulation of organelle organization M2635: ERB2 UP.V1 DN
	GO:0060627: regulation of vesicle-mediated transport
	R-HSA-391251: Protein folding GO:0097435: supramolecular fiber organization
	GO:0032507: maintenance of protein location in cell
	GO:0034330: cell junction organization R-HSA-2173782: Binding and Uptake of Ligands by Scavenger Receptors
	R-HSA-5653656: Vesicle-mediated transport
	GO:2001233: regulation of apoptotic signaling pathway GO:0008360: regulation of cell shape
	GO:0006986: response to unfolded protein
	M2867: TBK1.DF UP R-HSA-1474290: Collagen formation
	GO:0042060: wound healing
	R-HSA-1474244: Extracellular matrix organization GO:0032801: receptor catabolic process
	R-HSA-8862803: Deregulated CDK5 in Alzheimer's disease models
	R-HSA-437239: Recycling pathway of L1 R-HSA-5626467: RHO GTPases activate IQGAPs
	GO:0002011: morphogenesis of an epithelial sheet
	M2636: ERB2 UP.V1 UP
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