REVIGO Gene Ontology treemap

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kidney morphogen		tion of cell entiation	urogenital system developmen	regulatio dendritic : t morphoge	spine olig	l oligodendrocyte	semaphorin–plexin signaling pathway	mitochondrion s	response to DNA image by p53 class	negative regulation of MAPK cascade	lipoprotein transport	regulation of histone H3–K9 methylation	regulation of alkaline phosphatase activity	dephosphorylation	synapse organization	
regulation o			e-multicellular nism process	artery norphogene	sis re:	havioral sponse cocaine	negative regulation of protein export from nucleus	cholesterol efflux	meiotic mismatch repair	membrane raft polarization	fatty acid		malonyl-CoA	deoxycytidine metabolic process orylation positive regulation of	low-density lipoprotein sparticle synapse or remodeling	cortical microtubule
spinal cord motor neuron differentiation	proximal/dista pattern formation	neuron migration	regulation of cardion muscle of differential	ac ciliu cell asser	ım nbly ⁱ	velopmental growth nvolved in rphogenesis	mismatch repair	carbon dioxide	positive regulation of plexin sign peptidyl-lysine acetylation	aling pathwa	ay /asopressii secretion	apoptotic	biosynthetic process positive regulation of ARF			dendritic spine organization
photoreceptor cell	xenobiotic	adult feedir	acillular raana	hormo regulation of the fo	onal tion re	negative gulation of oranching rphogenesis	Wnt signaling pathway	fructose transport	proline transmembrand	cyclic-nucleotide-mediated signaling	DNA catabolic process	macromolecular complex remodeling	GTPase activity mating	regulation of fusion of sperm to egg plasma membrane	response	osmosensory
differentiation	negative regulation of	kidney morphogene positive regulation		s contract positive regulation	contraction positive regulation of trail		regulation of Wnt signaling pathway negative	hydroxyproline transport	response to triacyl	regulation of actin	regulation of protein sign	nal on localization	mā	tingnatid nucleus elongation	to light signaling stifesponse to light stimulus	
development	cardiac muscle cell myoblast differentiation	ell myoblast differentiation		H biosynth	netic involv	rmerase II promoter red in smooth muscle sell differentiation	regulation of inflammatory response	inactivation of MAPKK activity	hacterial	cytoskeleton	single organ		copulation	spermatid differentiation	respons	onse to pH
xenobiotic metabolic process	negative regulation of muscle hyperplasia	regulation of kidney size	response to methyl methanesulfor	SKIII	enesis e	gulation of extent of ell growth	fatty acid beta-oxidation using acyl-CoA oxidase	lipid cataboli process		formaldehy biosynthe	tic catabolic	lipoprotein particle	macrophage activation	locomotic	on I	ction of mulus
negative regulation of cell projection organization	negative regulation of neuron maturation	response to	Cell late	morphoger		cilium anization	biosynthetic	lipoprotein ty acid beta-o catabolic process	xidation usin	positive pos	oxidase dine	ne retinoid	regulation of multicellular	homophilic cell adhesion via plasma membrane adhesion molecules	amino-acid betaine metabolism	to
regulation of developmental process	negative regulation of T cell anergy	I filloxetine		response to cAMP	telencephalor cell migration		receptor catabolic process	unsaturated fa acid biosynthe process	· I IIPIG	aromatic a acid far metabo	amino mily olic is	oprenoid polic process	multicellular	superoxide		signaling