REVIGO Gene Ontology treemap

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generation of properties are		respiratory electron transport chain		oxidation-reductior process	transla double-		peptidyl-cysteine modification	single–organisr transport	n transition ion trans	metal	tablishment of protein ocalization	organic substance transport	striated muscle contraction	regulati interleu produc	on of h	neart	cardiac mu cell action potentia	scle ver on cardi	ntricular ac muscle membrane plarization
positive regulation of ubiquitin-protein	cullin	peptidyl-cyste	ine proti	ein transla	break r	repair asomal res	cellular sponse to	Golgi vesicle transport	macromolecule localization	cellular localizatio	intra-Golgi vesicle-mediat transport	l of protein	of cell	angiotensin maturation d muscle	fibroblast ;	muscle system on cess	cell membrane depola cardi	regulation of calcium ion ransmembrane transporter calciumly on potent	
cell cycle		S-nitrosylatio			tion cata	abolic cess f	peptide normone stimulus mRNA	metal ion transport	transition r water transport	métaPion t transport	intracellular cholesterol tran	lrogen establishment of vesicle localization	regulation of leukocyte mediated cytotoxicity	erception of taste	visual ^d perception	etermination of dorsal identity	membrane epolarization	nitric oxide homeostasis	telomere maintenance via recombination
positive regulation of transcription from RNA polymerase II promoter by glucose	of transcription from RNA polymera III promoter	translation	gluconed	metal proc	bolic transn	mission n	netabolic process	vesicle-mediated	divalent lo	sal protein ocalization ndoplasmic	Golgi mi	ly(A)+ IRNA port transport	RNA interference	apoptotic	egulation of re		cell redox omeostasis	cellular iron ion homeostasis	hormone catabolic process
oignainig	N-linked glycosylatior via asparagin	erespiratory	transduction involved in regulation of	transport c	sulfate catabolic chainess	tyrosine osphorylation Stat5 protein	Rac protein signal ransduction	iron ion	transport lo	reticulum calization receptor internalization involved in	positive regulation of pu	ourine regulation of clathrin-mediated endocytosis	interspecies interaction between	viral process	multi–organisı cellular process	m chemosens behavior	1	·	
post–translational protein modification	regulation of TOR signaling	cellular amide metabolic process	of cellular amino acid metabolic process	response to manganese p	of recorded resplices of respli		nuclear-transcribed mRNA catabolic process, no-go decay	transport actin filament	skeletal	canonical Wnt unaling pathway cellular	pathway protein-DN	NA regulation of cytoplasmic	organisms regulation vi	ral proces	s regulation of male		osensory nse to ticide	behavior Sponse (fungicide	multi-organism process
activation of protein kinase	TOR signaling		peptidyl-serine dephosphorylation	endodeoxyribonuclease pl	arowth factor	erpene letabolic process	wax piosynthetic process	polymerization	growth regulation of	complex assembly endoplasn	organizationic	On body assembly	response to virus by virus negative	transcription	germ cel proliferation modulation by virus	n habiti	iation I	esponse to byrethroid	
A activity peptidyl–asparagine modification	cellular ketone metabolic	second-messenger-mediated signaling	positive regulation of amyloid precursor protein biosynthetic process	protein polyubiquitination		positive regulation of phosphatase activity	regulation of rhodopsin mediated signaling pathway	nucleosome organization	lamellipodium assembly macromolecular	organizati atrial	developme	ent cell projection assembly hematopoietic	regulation of JNK cascade	of viral latency	of host process	localiz	cation cat	tabolic abolism ocess	cellular igmentation
via	process tricarboxyli acid cycle	1	positive regulation of ligand–dependent nuclear receptor transcription coactivator activity	cellular protein metabolic process multicellular organismal	lipid catabolic process 2'-deoxyribonucleoti	response to purine–containir compound	ethanol oxidation	actin filament-based movement	subunit organization aggresome	junction remodeling metanephr distal tubul	tissue growth ic mature B cell differentiation	differentiation telomere regulation of organization megakaryocyte	processing and presentation of peptide antigen via MHC class presentat		presentation exogenous ng and antige ogenous		I OT	of cell communication	wax wax metabolism process
spliceosome cellular hyperosmotic response	fatty acid beta-oxidatior using acyl-CoA dehydrogenasi	negative regulation of proteasomal protein catabolic process	pyrimidine deoxyribonucleoside triphosphate metabolic process	signaling heparan sulfate proteoglycan biosynthetic process	cellular response to caffeine	peptidyl-glutam acid	protein carboxylation	auditory receptor cell morphogenesis	chromatin assembly or disassembly	olfactory nerve developme	mesoderm p	cositive regulation of actin cytoskeleton reorganization	antige pep processing and presentation o exogenous antigen	. I processi	antigen processing ar presentatior via MHC clas	respon to activi	1-phospha metabolisi	process	regulation of development, heterochronic