REVIGO Gene Ontology treemap

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cytoskeleton or	ganization	actin filament–based		regulation of protein polymerization			cell redox homeostasis	energy derivation by oxidation of organic compounds	purine ribonucleosio triphosphate metabolic proc	e n e n	etyl–CoA netabolic process	coenz catal proc	bolic	oxidation proce
		process	3	microtubule	-based	process		operav	cofactor		arboxylic	res	nergy serve	ma meta
regulation of cellular component biogenesis	ion transmembra transport	Ras prot ne signa transduc	l	cytoskeletal anchoring at plasma membrane		iry or r motility	cortical cytoskeleton organization	alcohol catabolic proces	derivation by oxid catabolic ss process	L–phe	enylalanine	tyro meta	osine abolic	regul of cat
small GTPase mediated signal	cell projectior organization	assembly	filar organ	ization	ocytosis, Ilfment	muscle organ developme	protein ent localization	pyruvate metabolic process	coenzyme metabolic process	L–phe	olic process enylalanine olic process	posttranscri regulation gene expre	iptional on of fa	romatic ino acid family atabolic
transduction germarium-derived	response to inorganic substance	cytoskelet heart contraction	regulation of cellula compone organizati	ar component ar component ant movement	reproduct	of a bra	nching implantation						positiv of ubic	ve regul quitin-p
oocyte fate determination energy coupled	cellular response to reactive oxyge	imaginal disc-derived wing hair norganization	retrograde vesicle-media transport, Golgi to ER	development	cell morphogene involved in neuror differentiati	or pu	ipal process involved in	RNA splicing	mRNA metabolic	process	s translati	on	''''	ase activ de novo tein fold
proton transport, down electrochemical gradient	species maintenance of location		DNA packagir	branching morphogenes of an epithelial tube apoptotic	is regulatic embryc developr	on of onic ment organi	nediate nent celeton ization ization ization		RNA protein fold	splicin	regulation o	of cellu ty ^{me}		f negative of sequen DNA I transcrip act
macromolecular complex subunit organization	fusome organization	cell adhesion	cellular process involved i reproducti	in response t	al memb organiz o multice	orane post- zation ellular	-embryonic relopment filament-basec process	mRNA processing			weightene gemaning angebra dipas keta (antakati prosess	p ca	teasoma protein atabolic rocess	al 'de r posttran protein
hydrogen transport	spectrosome organization	response to cAMP	vesicle coating	stress cell-cell	reprod	nediated port	protein complex ogenesis		translational elo	ngation	RNA ni processing ^{bir}		ation of coxide nthetic	egative gulation of DNA m pinding 1

on-reduction process	glucose metabolic process							
malate etabolic process gulation catabolic	glu cellula carbohyo catabolic p	ar drate	netabolism carbohydrate catabolic process					
d dicarboxylic			glycogen netabolic process					
metabolic process								
gulation –protein ctivity	generation of precursor metabolites and energy							
vo'								
ative regulation quence-specific DNA binding	nicotinamide	alkaloid	locomotory	hydrogen				
ative regulation quence-specific INA binding scription factor activity de novo' translationa	nicotinamide metabolic	metabolio	behavior	hydrogen peroxide metabolism				
ative regulation quence-specific DNA binding scription factor activity de novo' translationa tein folding e cellular on amide	nicotinamide metabolic	metabolio etabolisn	behavior	peroxide				