REVIGO Gene Ontology treemap

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translational elongation		viral gene expression		anism process	translationa terminatior		translational initiation	SRP-depende cotranslationa protein targeti to membrane	comp ng disass	onent embly	protein complex subunit organization Gold		tracellular ransport regulation	membrane organization		opioid receptor ate signaling pathway detection of response to comments.	response to oxygen-containin compound opioid of	stimulus response to oxvgen-containing stimulus
nuclear–transcribed mRNA catabolic process, nonsense–mediate	cholester biosynthe	tic multi-	organism ar process	interspecion interaction between	on comp n biosyr	ound re	positive egulation metabolic	protein localization to endoplasmic reticulum	macromo	ein single-tran	tra protein alization	alcium ion ansport synapti	i ioxin	negative	detection of hormone stimulus	response to high densit lipoprotein particle	y transcription from RNA polymerase II	response to lipoprotein particle response to ethanol
decay	heterocycle biosynthetic	regulatio of cell	n cellula n nitroga	en pred	yloid sursor isol	orenoid	cellular nitrogen	SRP-depe macromolecul complex subu organization	nit regula	tion n	inction egative ulation of	targetin organ substar	ic low-den lipoprote	growth sity negative regulation of	organic cyclic compound metabolic process	biosynthetic process	regulation homeostat process positive	lhomeostatic
RNA catabolic	process proces		on biosynth	netic meta		ocess r	metabolic process positive regulation of cAMP	single-organis cellular localiza	tion glycopr transp	otein pho	efflux	transpo actin filame organiza	nent regula	phagocytosis	cellular arorganic hydroxy compound metabolic		regulation of biological process	cholesterol lation of ostasis atic process positive regulation
process	of fat cell differentiation regulation of B cell	cell migration negative regulation of cellular	astrocyte cell	N-terminal protein	proliferation norepinephrine metabolic	process positive regulation	metabolic process regulation of cholesterol	single-organis	m prote localiza to adhe iuncti	ation me	mbrane epair	multi-orga membra organizat ative	nism reve	cell receptor signaling pathway	nitrogen compound gumetabolic express	organic hydroxy ene ession metabolic	mitochondrial depolarization	of metabolic process
nucleobase–containing compound metabolic process	multi-organism organelle organization	process pyrimidine dimer repair by nucleotide–excisior	trophoblas	regulation of activin receptor	positive regulation of	gastrulation steroid	esterification collagen catabolic	bone marrow development	ovarian follicle growth	of anatom structur orientation retinal blo	regula e on vasod	ation of lilation	tissue remodeling post-embryoni camera-type	embryo development regulation of	negative regulation of protein dephospositive reg	regulation of cyclic-nucleotide gulation of foliaterase activity	multi-organism	
biosynthetic process	negative regulation of SMAD protein complex assembly	steroid metabolic process	metabolic	mRNA metabolic process	gluconeogenesis	regulation of microtubule–based movement	process positive regulation of cellular	mammary gland epithelium development		-	wnt s	signaling ment _{ed}	eye morphogenesis organ developmen	development, heterochronic anatomical structure	phosphodiest of alkaline phosphatase		establishment	cellular component
regulation of nucleobase–containing compound metabolic process	positive regulation of amyloid precursor protein biosynthetic process	regulation of cell cycle	biosynthetic process	f positive	DNA-templated transcriptional preinitiation complex assembly	biosynthetic process	process Rho protein signal transduction	establishment of organ orientation	circulatory system process	Mulleria duct regressi	axis sp an gl devel	ecification land	axis	negative	cardiac musclo cell apoptotic cardiac musclo cell apoptotic	of cell regulation	of cell polarity	organization or biogenesis establishment or maintenance of cell polarity
compound biosynthetic process	positive regulation of RNA polymerase Il transcriptional preinitiation complex assembly	regulation of nucleobase–containing compound metabolic process	positive regulation of apoptotic process	female gamete generation	pyrimidine dimer n repair	of Rac	T cell chemotaxis	gonadal mesoderm development	brain renin–angiotensin system	myoblast commitm	fate bo	one odeling	bone trabecula formation	replicative senescence	apoptotic process	division regulation of meiosis	response to chemical	localization