## REVIGO Gene Ontology treemap

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SRP-dependent cotranslati	·	•		translational termination		NA mRNA metabo process	catabolic process regulation	multi-organism cellular process	interspecies interaction between organisms	response to ja glucagon services prostaglandin	sponse to samonic acid jasm acid timulus acid peroxide of control	drug
translational elongation	membrane organization	protein transport	translation	acromolecular complex subunit organization		g oxidation-reduction	metabolic amino acide process process pyrimidine nucleoside triphosphate metabolic catabolism, nonsens protein	metabolic process  cellular metabolic process  cellular metabolic compound metaprocess  process  e-mediated decay pyrimidine nucleoside nucleoside nucleoside process process process feemediated decay nucleoside nucleosid	de acid etic metabolic process  pyrimidine nucleoside monophosphate biosynthetic	cellular response to glucagon stimulus negative regulation of fibroblast apoptotic pregative	esponse to response to more incorrection and receptor records	
SRP-depen translational initiation	cellular dent cotranslation disassembly	protein complex macromolecule to membrane organization		cellular protein metabolic process	multi–organism metabolic process	pyrimidine-containing compound biosynthetic	respiratory peptide metabolic transport process chain regulation of glycolytic process by	process proces	ation, sical metabolic	cyclase–activating pdopamine receptor signaling pathway	Rap protein si	p protein signal asduction nse to
	11 1	actin filament dense granule organization regulation of lipid mitochondr	positive regulation of actin cytoskeleton reorganization  protein	-mediated cytosis regulation of transcription from RNA polymerase ill promoter	RNA catabolic proces	autophagy	acetylation guanosine-containing compound biosynthetic process	erpenoid organonitrogen compound metabolic	way process -density oprotein tor particle olic process l cycle	abscission of low-density lipoprotein particle clearance	ration level response extrace	rels nse to rellular
single-organism intracellular transport localization	transport by regulation of transcription from RNA polymerase II promoter substance transport chromatin	regulation of anscription from RNA olymerase II promoter hydrogen ion transmembrai transport	biogenesis ribosomal nucleoso	membrane protein-DNA complex	involved in mitotic transi	homeostasis homeos	meostasis of catalytic activity ation regulation of phosphoprotein mitochondrial		esentation of ptide antigen	catabolis	n cellul compo organiza or bioge	onent zation
	ribosomal small subunit biogenesis	ribosomal small subunit assembly  caveola assembl assembl regative regulat of transcription fac localization	y membrane budding targeting ion negative by regulation of expect for	organization  nRNA chromatin assembly or	regulation of regulation of ligase activity	on of ligase activi regulation of nega agy endopeptidase regulation	ase activity depolarization depolari	antigen process presentation of expeptide anti	xogenous igen	response to gamma radiation		neutrophil ggregation
single-organism localization	post-translational ER to Golgi protein vesicle-mediated modification transport	ER to Golgi regulation esicle-mediated apoptotic c	of regulation of modification small protein complex	orotein iffication by all protein component	signal transduction prote involved in mitotic destabili DNA damage	lmitophondriall:	enance negative regulation	and presentation process present	sing and antigen processing and reation of presentation of peptide or polysaccharide antigen antigen via MHC class I	localization	single-organism process	amino sugar etabolisr