Revigo TreeMap

proteasome-mediated ubiquitin-dependent protein catabolic proces	tran	translation		mRNA splicing, via spliceosome	regulation of DNA–templated transcription	regulation of DNA biosynthetic process	regulation hemopoies	sis	negative regulation of apoptotic signaling pathway	negative regulation of transcription by RNA polymerase II	protein transpor	exocytosis
					regulation of translation	regulation of transcription by RNA polymerase II	regulation of circadian rhythr	re m of	regulation of protein sumoylation results of pathway		protein tr	maintenance of protein location in nucleus
cytoplasmic translation	histone a	histone acetylation		ton motive ce–driven ochondrial		positive regulation regulation of double-strand break repair via	negative of translation regulation of protein localization	negativ regulatio	gulation regulation polyamine	of mitotic	retrograde vesicle-mediated transport, Golgi to	endosomal transport
proteasome-mediated ubiquitin-dependent prote		ndent protein		e synthesis	regulation of cell cycle	homologous recombination negative regulation	to centrosome regulation of	growth positive regulation of	transport	separation	endoplasmic reticulum	
protein polyubiquitination	cellular respiration	mitochondr electron transport,		ubiquinone biosynthetic	positive regulation	of mRNA splicing, via spliceosome	process targe mitocl		g to of fibrobla	regulation of mitochondrial translation	cell cycle	rhythmic process
	roopiiation	succinate to ubiquinone		process	of transcription by RNA polymerase II	regulation of cytokinesis	regulation of regulation of DNA damage transcript initiation		regulation of proteasomal by RNA ubiquitin-dependent protein catabolic process			
protein deubiquitination	RNA splicing			tRNA processing	mitochondrial respiratory chain mito complex I assembly		autophagosome assembly		multivesicular body assembly	ribosomal small subunit assembly	DNA postrep DNA damage respo response	mitotic cytokinesis
mitochondrial translation	· · · · · · · · · · · · · · · · · · ·			on of RNA transcription		ochondrial respiratory	ribosomal si subunit bioge	mall	membrane fusion	RNA polymerase II preinitiation complex assembly	cell division	stem cell population maintenance