REVIGO Gene Ontology treemap

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negative regulation of gene expression		regulation of cellulus metabole proces	ar lic	regulation of metabolic process		anscription rom RNA olymerase promoter	post-embryonio organ development	B cell neg	I D CEII	D Cell Selection		-type enesis p	apoptotic process nvolved in atterning of ood vessels	establishment or maintenance of transmembrane electrochemical gradient	regulation of calcium	on transport, membrane cellul	plasma to Golgi lar blecule	regulation of protein eterodimerization activity	involved in apoptotic process by cytochrome of	fibroblast apoptotic process
positive regulation of biological	_	gene regulat of metal proces		tion bolic ess transcription from RNA polymerase promoter		aromatic compound biosynthetic	cartilage morphogenesis	abducens nerve formation	initiation o neural tube closure	call different	ation of interest in the inter	olved in nic placenta lopment	hombomere 4 levelopment	resulting in posttranscriptional regulatioestablish expression transmembra	ment or n	negative regulation naintenance ochemical gr	mRNA contact of tabilization of adient	heterodir	apoptotic	process
process						process	post–embryonic morphogenesis	semicircular canal formation	midbrain pala development develop		e n ment	anial erve nation	facial nucleus development	regulation reg	of inte	on regulation of interleukin–10 secretion exocytosis		CAAX-box regulation of mesench mesenchymal cell apoptotic process		totic nuclear receptor
heterocycle biosynthetic process	nitrog compo biosynt proce	hetic ess con bios pro	npound ynthetic ocess		etion cell arrest	of cellular process	vagina	pattern	mbryonic o ventricle development	rgan dev	elopment organ development	morphogene	regulation of Schwann sis cell proliferation	protein po	ositive Julation	ngle-organism cellular localization ngle-organism	of RNA	AAX-box apo protein pro invol	ptotic regulation of view of the properties of t	developmental programmed
intracellular signal transduction	negat regula of ce negative	tion macro	llular molecule tabolic ofsgen	metab	olecule r	positive regulation f metabolic process	pattern	cell surface receptor signaling pathway	boundary	replicative cell aging	anatomical structure	nervous system developme	endocrine pancreas nt development	response t	to res	sponse to _n	response to	respon	activity opti	okinetic ehavior
chromatin reprogramming in the zygote	negative regulation o peptidyl–serir phosphorylati	positive regulation of signal	regu n d l biolo	ılation r of	nucleic acid etabolic	nucleobase–containing compound metabolic process	specification process melanocyte	involved in heart development syncytium	limb morphogenesis		,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,	embryonic neurocranium norphogenesis		positive regula	ation	transci	F ve regulation of ription from RN erase II promo	to UV-	ponse to UV C resp abioti	ellular conse to c stimulus
negative regulation of G1/S transition of mitotic cell cycle by negative regulation of transcription from RNA polymerase II promoter	positive regulation	n metabo	ar are	ellular omatic he	eterocycle metabolic	organic substance metabolic	abducens	striated	blood	face	BMP signaling pathway involved in	endocrii systen developm	nent organ	of endopires reticulum unfo protein respo regulation	onse regul	damage pro	ved in unfolded tein response sponse to accharide	signaling pathway	in absence	response to fungus response
positive regulation of transcription on	positive regulatio	n activati	on me	rocess I GTPase ediated	process primary netabolic	process cellular nitrogen	development	muscle cell differentiation regulation	remodeling	ation of s	heart development development	genital developm proteir	nent	endoplasm reticulum unfo protein respo	olded signal by	p53 class	sponse to sucrose	op oid pat receptor c-adopam signaling	hway pholipase regulation of actin	to fungus release from viral
exit from mitosis negative regulation of	of cell communica negative regulation	ative regulation		proces		metabolic process changes to DNA methylation	positive regulation of cellular proces		ar cell growth d		complex to plas sassembly membi		ne	negative regulation of endoplasmic negative reg	lymphocyte	vitamin metabolic process	cataboli	P.	egulation	ological
biological process positive	protein kin C signalir	ase biosynthe process	etic C		of signaling egulation of		positive regulation	positive regulation mitochondrial ou membrane permeabi	n of ter	sion	organelle organization	modific		reticulum c	alcium tration	fat-solub metab		locomo	of viral	gulation
regulation of transcription, DNA-templated	repair, DNA incision, 3'-to lesion	dimer repair	by Scision		anscription	regulation of cell communication	of cellular component organization	involved in apopt signaling pathwa	otic IIPIC ay	embly m	egulation on hitochondricondricon organization	memo fusi		reticulum calcium ion homeostasis	mitochondria membrane potential		proces	cellular re to redox	sponse proce	iem