REVIGO Gene Ontology treemap

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enamel mineralization	negative regulation of cartilage development	monocyte _{ii}	nterleukin–1: production	apoptotic process involved in development metanephric	t	negative	Sertoli cell tdifferentiation	RNA proces	sing p	mRNA rocessing	organonitrogen compound metabolic process	organonitroger compound biosynthetic process	DNA-templated transcription, termination	positive regulation of vitamin D 24–hydroxylaso activity	negative regulation phospholip biosynthet process regulation	of regulation of bind	ing process	negative regulation of ligand-dependent nuclear receptor transcription coactivator activity
amelogenesis	epithelium ⁽ development	growth	cajal-Retzius cell differentiation	glomerular endothelium	regulation of mononuclear cell migration	of smooth 1	osteoclast maturation	mRNA		sulfur	guanosine–containin	positive regulation by	carbohydrate derivative	•	ulation of vit 1-monooxyger activity regulation	mitocho depolari	hydroxylas regulation guanylat	homeostasis
	mammary gland duct morphogenesis	tube ame	Heart	cell fate	sis	trabecula morphogenesis	biomineral tissue development	process	glycosaminoglyca metabolic proces transcriptio	metabolic process positive	compound metabolic process	host of viral transcription 'de novo'	metabolic process GTP	24–hydroxylaso activity	_	negative reg of transcri from RNA pol	otion Imerase during	molecule netic process
wound healing, spreading of epidermal cells s	ubstrate-dependent		corticospinal neuron axon guidance through	of an epithelial sheet	regulation striated mu cell apopto process hepaticobi	scle regulation developmen process	ntal decidualization	aminoacylation	•	translation or regulation	of axon diameter	protein folding regulation o anatomical		receptor signaling	andro	ulation of		metal ion transport
myoblast development	of tissue	regulation	nterleukin–13 biosynthetic process	embryonic skeletal joir developme	of cysteine-ty endopeptida	nent negative regulation regulation of anoiked in		ncRNA metabolic process	aminoglycar metabolic process	component	methylation or demethylation nucleobase-containing small molecule metabolic process	structure size glycosyl compound metabolic	nucleotide catabolic	t(response reg	ulation of regulation of of results of resul	lation m	valent etal ion enzyme tr	divalent inorganic cation ansport transport positive
actin filament organization	desmosom	desmos assem	ome an	atomical tructure		toskeleton _r	mitochondrion distribution	amino acid activation	termination of RNA polymerase transcription	I biosynthetic	guanosine-containing compound biosynthetic process	process chemical homeostasis	process alpha-amino acid biosynthetic process	cellular inter	esponse to feron-gamma r	esponse to lipid	RNA tra	gulation of xocytosis protein
cellular componen	regulation t of cellular	r compor	ılar po	ssembly	of synapse assembly	positive regulation of cellular component organization	protein heterotetramerization	regulation o Rac protein signal transduction	of ce	regulati	on regulation of cell communication	regulation of fibroblast growth factor receptor signaling pathwa	cascade	heterophilic cell-cell adhesion via plasma membrane heterophilic	cell cell by	skeletal m satellite activati	cell filam	actin ent-based
cell morphogenesi	protein impo	regulat actin filan	ion orga	rojection _O	regulation of synaptic plasticity		Golgi vesicle fusion to target membrane	G–protein coup glutamate receptor signaling path	prolifera	of Rac protein	signaling ng nathway nsignal tran	inflammatory response to non-antigeninsduction	negative	adhesion vi membrane ce molect cell adhesion	II adhesion ules substrate	cellula compone organizat	ent ion developmen process	cellular process
	mitochondri inner membra	from nuc	port lo	protein calization to	positive egulation of centrosome cycle	protein complex subunit organization	nuclear transport	Rac protein signal transduction	signali	beta regulation of reception recycling recycli	on paracrine otor signaline	g NK T cell proliferation	taxis	positive regula biological pro positive reg	cess Julation of	or biogen		n
ribosome biogenesis	ribonucleoprof complex biogenesis	neuro		integrin ctivation ^d	tissue evelopmentr	protein polymerization	vesicle targeting	positive regulat of response to external stimul	signa	al mediate	positive regulation chronic inflammato response to non-antigenic stimu	small GTPase	cell	biological regulation biological qu	of	biologic adhesic	vitamin metabolis proces	establishment of epithelial cell polarity