REVIGO Gene Ontology treemap

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post–anal tail morphogenesis	axial mesoderm formation	primitive hemopoiesis embryonic	macrophage differentiation cell migration	soma	atic		activation of JUN kinase activity	l regulati	l of	protein phosphorylation	regulation of multicellular organismal process	cellular response to interleukin–4	response to estrogen	response to	l response l
negative regulation of fat cell	cardiac muscle cell differentiation	organ development	sprouting	rophoblast giant cell of immunog gene seg	structura I ^{lobulin} organizati	eosinophil	regulation of transferase	transcription	regulation of intracellular	0.00	signal ransduction by hosphorylation	hepatocyte growth factor receptor signaling	platelet-derived growth factor receptor signaling	stress-activated protein kinase signaling	signal transduction
differentiation myeloid cell	negative F regulation of cellular process	post ana tail r hemopoiesis	pituitary norphogenes development _{ol}	ganization	chord regulation of cells ketone metabolic py negative regulation transcription from polymerase II pron	rocess cell fate	activity positive regulation of	elongation positive regulation of (regulat	of	ositive gulation regulation regulation	negative regulation of oxidative phosphorylation	pathway SREBP signaling cellu	pathway positive regulation of RNA lar response	regulation of response	antibiotic n ⊣4 ansport
differentiation	cell differentiation	axial mesoderm development	development di	secretory transcell from	nRNA scription m RNA merase comoter migra		biological process	regulation of cellular process	negative regulation of Rho-dependent	in endoplasmic	regulation of ton–transporting TPase activity,	response to organic cyclic	process basophil	positive	response to chromate
immune system development	regulation of developmental process	coronary vasculature development	in wound	ifferentiation involved in eletal muscle egeneration Wnt-signalii inv	ein neuron Ommitment de	cle structure velopment	phosphorylation	keratinocyte proliferation	kinase activity positive regulation of peptidyl-tyrosine autophosphorylation S	regulation of tho-dependent protein regulation cregulation of the position regulation crime/threonine crime/th	on of regulation of keratinocyte	response to sterol	cellular	to tellurium	cell surface receptor naling pathway egulation of
mitochondriai	nitochondrion respondent	ponse to orga	organism anelle ization	l of cellula	n desmosom organizatio	transcription ne from nmitochondrial promoter	regulation of response to stimulus	regulation of phosphorus metabolic process	regulation	regulation of immune system process skinase activity factor at	promeration	depletion ER-nucleus signaling pathway		/EGF-activated neuropilin	lular response to stress esponse to alcohol
regulation of	cell cycle in	lucose II proi	ription creating n from transmemb ymerase transpo noter	mitochondria translation	f polysaccharide	component	cholest biosynthetic	erol	small molecule biosynthetic process		regulation of B cell activation	immune	cell activat	cell ion vation	growth
organization	modulation of programmed cell death in other	arvation cargo I of mitochondri COPII- ves	al membrane coated expor	regulation permeability t localization	body fluid secretion	protein insertion into membrane			lipid biosynthetic process	regulation of type 2 immuno	T-helper	of establishment	peptidyl–lysine hydroxylation to b–peptidyl–lysine	cellular response to	cell proliferation
		transition respo	ion demethyla	e tissue	nucleocytoplasmic transport	response to vitamin	isoprenoid biosynthetic process	sterol bios diphosphate biosynthetic process	U6 snRNA uridine 3'-end metabol processing process	response	response positive regulation	polarity ve	hydroxylation to 5-hydroxy-L-lysin pepticiyi-iysin hydroxylation		ithelial
negative regulation of mitochondrion organization	process	neostasis number actin fi of cells burn asse	actile histoned histo	B DNA-templated transcription,	nuclear endo transport recy	regulation of focal	negative regulation of fatty acid biosynthetic process	dimethylallyl diphosphate metabolic process	vitamin D catabolic process	regulation regul of growth	developm ation of gro positive regulate of anace	regulation of synaptic growth at neuromuscular junction	death	adhesio regulation beta-amyl	to low light intensity stimulus