REVIGO Gene Ontology treemap

patterning of blood vessels	regulation of interferon–gamma production	nucleus accumbens development	anatomical structure formation involved in morphogenesis regulation of	of MDA-5 signaling pathway	development	of skeletal muscle cell	posterior mesonephric tubule development	embryonic axis specification	urate metabo process	glycine decarboxylatio via glycine cleavage system	inosine catabolic process	mitochondria RNA modification	NAD biosynthesis via nicotinamide riboside salvage pathway	mitochondria transport	ll transmembrar transport	ne ion transpoi	protein rt targeting to mitochondrion
camera-type eye morphogenesis	production	olfactory pit development	embryonic development adrenal	regulation of kidney development endocrine	positive regulation of developmental process	sensory organ development	immune system development myoblast	morphogenesis of a branching structure positive	nicotinamide riboside catabolic process	small molecule metabolic process	organic acid metabolic process	D-ribose catabolic process	heme oxidation	secretion by tissue	amino acid transmembrane t transport	creatine ransmembrane transport	taurine transport
post-embryonic camera-type eye	negative regulation of metanephric glomerular mesangial cell proliferation	visceral serous pericardium development	cortex formation patterning of branchiomeric skeletal	system development blood vess	system	voltage-gated sodium channels secretion of	fate commitment apoptotic process	regulation of myoblast fusion cardiac muscle	pyridine	methionyl-tRN/ aminoacy <mark>urat</mark>	pyridine–containing compound		flavin-containing compound metabolic process	body fluid secretion	positive regulation of mitochondria translation	I transport	protein localization to mitochondrion
leukocyte differentiation	negative regulation of Notch signaling pathway involved in somitogenesis	positive regulation of multicellular organismal	muscle development coronary artery morphogenesis	secretion negative regulation of type 2 immune	homeostasis	lysosomal enzymes regulation of developmental	involved in patterning of blood vessels G1 to G0	cell fate commitmen positive regulation	process threonylcarbamoyladenosine metabolic process	urate biosynthetic process	fructose 6-phosphate metabolic process	process	positive regulation of cholesterol biosynthetic process	transferrin transport	proton transport	positive regulation	regulation of phospholipid transport
post-embryonic organ development	Notch signaling pathway involved in	process macrophage differentiation	G1 to G0 transition involved in cell	response primitive erythrocyte	cardiac muscle fiber	growth embryonic pattern specification	negative regulation of interleukin–12 production		mitochondrial RNA	oxidation-reduction process	nucleobase-containing small molecule catabolic process	retinoic acid biosynthetic process tRNA wobble	UDP-N-acetylglucosamine biosynthetic process organonitroger compound	trivalent inorganic cation transport positive regulation	hydrogen transport establishment or maintenance of	riboflavin transport endosome to	organic substance transport nitrogen
positive regulation of cell–substrate	positive regulation of epidermal growth factor receptor	basophil	VEGF-activated neuropilin signaling	positive	development of primary	cellular respiration	B cell negative selection	B cell selection	metabolic process	metabolic process	metabolic process	uridine modification	metabolic process	of endocytic recycling ositive gulation	transmembrane electrochemical gradient single-org metab	janism prec	compound transport ation of ursor
adhesion regulation of protein kinase	retinal ganglion cell axon	neuropilin signaling pathway	positive regulation of reproductive	positive regulation of trophoblast	positive regulation of	peptidyl-tyrosine autophosphorylation	trophoblas cell migration	positive regulation of adaptive immune response based on somatic recombination of immune receptors built from immunoglobulin superturally domains	mitochond organizati positive	of histo	one peptidyl-se lation phosphoryla	rine proce	cell apopto	of cell process or neuron necro	ptotic proce	ar carbol metab	energy hydrate polism abolic process
C signaling regulation of	guidance protein kinase	positive regulation	liecebroi broteii	phospholipase C-activating	from RNA polymerase II promoter in response to hypoxia	positive regulation of cell morphogene involved in differentiation	heterotypi cell–cell	of peptidyl-lysine modification to peptidyl-hypusine	activity	regulation on drion organical process	ATP synthase anization lex biogenesis	cellula	ntation p	rocess path	organic sul	ostance zym	ogen vation
lymphocyte activation positive	C signaling regulation of generation of		tyrosine kinase signaling pathway positive	receptor signaling pathway regulation	regulation of cell adhesion	mast cell migration	regulatio choleste storag	n of Prol positive regulation of	mitochondrial proton-transportir ATP synthase complex assembl	two-sector ATP	ribosom assembl	e homeost	asis permea	cular neuronal a neuronal a neuronal a neuronal a	cell activ	respon mycot	
regulation of leukocyte migration	precursor metabolites and energy	T cell extravasatio	regulation of cellular respiration	of cell	osteoclast proliferation	regulation of	positive reg of endopla reticulum ur protein res	ulation cell proliferation nfolded	negative regulati of phosphate metabolic proces	elastic lib	dense gran	mitochon membra	drial mitocho	rane regulatio	cell by prolifera	righting	reflex