## REVIGO Gene Ontology treemap

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synapse ass	sembly	synapse organization	nervous system developmen	synaptic vesicle t clustering	multicellula organisma response to stress	of interbeta s	regulation rleukin-1 secretion ebrain ogenesis	negative regulatior of behavior	multicellular organismal reproductive behavior	negative regulation of protein kinase B signaling negative regulation of cyclin-depender	nt   Scriavioral	immune response	cellular component movement rcellular com	transmembrane transport	regulation of localization	regulation locomotio	trophoblast
platelet–derived growth factor production	regulation platelet-de growth fac production	rived bladde	r of cellu	lar lur ent develop	pment ir	cell phogenesis involved n neuron erentiation	negative regulation of amine transport	regulation of behavior	inhibition  negative regulation of cytokine-mediate	in sensory	response to cocaine sensory havior temperature	interleukin-15 of response to	transport	inorganic cation transport xenob	transport	trophob cell migra	negative chemotaxis
multi-multicellular organism process	presynapt membran organizatio	e streak	system	of primary female sexual	retina levelopment in amera–type	midgut development	mucus	circadian behavior	signaling pathwa	humoral	stimulus  regalinia regalinia of cyclic-dependent pristos secretiferatura initias scienty deviadad in CIGO translator of manda cell cycle	positive regulation of dendritic cell antigen processing and presentation	response	to norganic cat		migrati coenzyme catabolic process	glycine catabolic process
regulation of endocytosis	regulation of N-methyl-D-asparts selective glutamate receptor activity	assembly	development			inflammatory response to antigenic stimulus	mating	rhythmic behavior	reproductive	negative regulation of response to external stimul	thrombin receptor	synthase activity	cellular resp response to metal ion	ubstance ponse to zinc	cionte C		Catabolism <sup>sine</sup> Biosynthetic process
	diencephalo developme	of a branching	development or	matrix	dendrite egeneration	dendrite self-avoidance negative	detection of muramyl dipeptide negative	regulation of synaptic transmission, glutamatergic		l signaling <sup>O</sup>	regulation f membrane potential	monovalent inorganic cation homeostasis	cellular response to hydroxyisoflavon	positiv regulatio	ve D- on of c	0,00000	ydroxylysine tabolic process
regulation of vesicle–mediated transport	transporte	development	formation endocardial	lateral sprouting volved in mammary gland duct morphogenesis  positive regulation of	nicturition positive	regulation of establishment of blood-brain barrier	regulation of ribosome biogenesis regulation of	r	regulation of excitatory	egulation of h	sodium ion nomeostasis	multicellular organismal homeostasis	cell-cell a		ria e	pehavior	catecholamine metabolism
tube development	response wounding	vasculature development acrosomal	cushion cell fate commitment hepatic duct	dendritic cell cytokine production positive	regulation of type IV hypersensitivity regulation of establishmen	midgut development	chromatin disassembly	transrnis regulati	oostsynaptic   ion of synaptic	n of synaptic apoptotic n, glutamatergic cess		positive relaxation of vascular membrane membrane potential muscle		plasma membrane cell adhesion molecules		ulticellular ganismal process	microglial cell activation
positive regulation of calcium ion	positive regulation of filopodium	centrosome-templated	hepatoblast	mitochondrial DNA replication positive regulation	of blood-brain barrier regulation of nethylation-depende	transmembrane transporter activity	retinol transport	myeloid cell apoptotic	of activin receptor signaling			tissue nomeostasis membrane	cellular carbohydrate biosynthetic	dolichyl monophosph biosyntheti process	loc	omotion	biological adhesion
transport catecholamine uptake	assembly acute inflammatory response to antigenic stimulus	cerebral cortex		maturation positive regulation of nucleoside r	regulation cheurotrophi	anterior/pos patternin stress-indu mitochond	uced Irial		pathway G-protein coupled receptor signaling	enter regulation of enterm synaptic contentiation attion of thyroid alone mediated aling pathway	regulation of respiratory aseous exchange	repolarization uring SA node cell action potential gative regulation interleukin–18 production	carbohydrate metabolic	rnthesis positive regulat	ion of trar	sposition	eye blink eflex  maintenance of epithelial cell apical/basal polarity