REVIGO Gene Ontology treemap

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oxidation-re proce		organic acid metabolic process	small molecu catabolic process	ıle regula lipid ca prod	tabolic	monosacchario metabolic process	transr	naptic mission, matergic	pyruvate transpor	vesicle-me transpo	diated of rt exocyto:	pyruvat	of horm	none inc	novalent organic cation neostasis	of renal sodium	of mast cel	regulation of inflammatory response to antigenic stimulus inflammatory
		regulation of excitatory postsynaptic	pentose biosynthetic process	regulation of lipid metabolic	carbohy catab	Olic transmer	mbrane me	ofactor etabolic	hydrogen transport negative	regulation of cellular component mopyruvat	positive regulation of receptor internalization e transport organic	transmembr transpor	anatom structu regu	ical regu	ell size	mucus secretion ne levels	_	response to antigenic stimulus regulation of l activation
generation of precursor metabolites and energy		membrane potential mitochondrial	tricarboxylic	glutamate receptor signaling	lipio	d potassiu	um ion keto	cellular	regulation of amine	transport	acid transr	negative regulation	oort regulat of membra	and biosy ane pro	androgen biosynthetic process multic		signaling pathway lo	positive regulation of humoral immune
carboxylic metabolic p		ATP synthesis coupled proton transport	rboxylic acid n pentose metabolic	pathway netabolism nitochondrial	acid		glucan	G-protein coupled receptor	monocarboxylic acid transport	plasma membrane long-chain fatty acid transport	transport neurotransmitter transport	ion transp monoami transpo	hormo ne metabo	one production destate	otein bilization	organismal omeostasis glucose omeostasis	negative regulation of transposition	response positive regulation of response to external stimulus
synaptic transmission		lipid metabolic process	process	transport carbohydrate p	compound		oumarin	signaling pathway receptor guanylyl cyclase	cellular resp to cadmiun			num regul	ation bel	cadian havior	rhythm behavi	ior metab	phate organophosphate biosynthetic process	ntecholamine metabolic process bile acid metabolic process
		small molecule metabolic	ribose	positive	biosynthetic process cellular carbohydrate	process pregulation of short-term	gulation of tinoic acid receptor	signaling pathway	response cellular nicotine		onse xenob cadmium io vation proce	n ^{lic}	_regulation	ticellular n of beha anismai oductive	hehavi avior fear respor	oral glycos	chosphate ctabolism single-organism colic biosynthetic	metabolic metabolic metabolic
respiratory electron transport chain	cell-cell signaling	negative regulation		lipid catabolic process energy	biosynthetic process	synaptic plasticity po	oathway lysaccharide metabolic process	interconversion de single-organism	response alkaloid	,	to to resp	regular of belicosteroid mulus	ation bel	havior havioral efense sponse	reprodu	rior calciu	um-dependent	regulation of
		of synaptic transmission	biosynthetic process	metabolic process	transport	movement 1	anion transport	catabolic	synapse	diencepha	substai	ntia cata	bolic cata	IIIGIIL	egulatior locomotic	n of we	via plasma embrane cell	regulation of monooxygenase activity oxidoreductase
excretion	retion startle dendritic cell response cytokine production		ith mouth forming second platelet-deriv growth factor production	or vessel morphogenesis	positive regulation of interleukin–1 beta secretior	retina	developme	vagina development nt	assembly sy	developm napse orga	qevelopr	nent p cofa	ctor catabo	re	egulation ocomoti	ion m	regulation of myeloid cell ototic process apoptotic	regulation of biological quality
myelin maintenance	central nervous system myelin maintenance	vesicle duri	regulation or platelet—derive growth facts production on potential	of urinary ved bladder	regulation of respiratory gaseous exchang by neurological system process	development ir camera-type ey bone marro developme	regulation of angiogenesis		synapse organizatio	n mitochon DNA replic mitochon fission	drial morphoge nervo	us m NADI	oolic process P metabolic process	biosynthetic process trop	phoblast	hoblast cell	process	