REVIGO Gene Ontology treemap

respiratory electron transport chain			generation of precursor metabolites and energy		oxidation–reduction process		TOR signaliı	ng regulatio	on of complacting	gulation of Arp2/3 lex-mediated n nucleation	regulation of protein kinase B signaling	mature B ce differentiatio	l involved in	
							positive regulation of transcription from RNA polymerase II promoter by glucose	regulation of membrane potential positive	cellula response organonitr compou	e to intra	egulation of cellular steroid mone receptor aling pathway	negative regulation of interleukin–1 production	muscle contraction	auditory receptor cell stereocilium organization
					hydrogen ion transmembrane transport		proton-transporting ATP synthase complex assembly	regulation of lamellinodium regulation organization	signaling	muscle ce Cellular Iling homeostas	response	negatimatur regulation of interleukin–6	regulation of B cell differen membrane potential	
hydrogen transport transmembrane transport	small molecule respi metabolic process	phosphate ion	regulation of fatty acid transport ch	positive regulation of calcium ion aim	monocarbox	catabolic	proton-transporting ATP synthase complex biogenesis	two-sector ATPase complex assembly pregulation of cell-cell	response t purine–conta	ining histo H3-K	K36 iron ion lation homeostasis	myelination in peripheral	multicellular organismal movement	ovulation cycle process
		single-organism	carboxylic		mitochond	Irial	positive regulation of	adhesion mediated by integrin	response nutrient	ıtrient		nervous system regulation	regulation of ventricular cardiac muscle cell membrane depolarization	renal system process involved in regulation of systemic arterial blood pressure
	tricarboxylic acid cycle	metabolic process	metabolic process	histamine transport	transport, ubiquinol to	t, transmembrane transport	smoothened signaling pathway	cellular homeostasis	cellular response to oxygen–contain compound	ning	ISIONAIINO	of natural killer cell differentiation	ruffle organization	female gonad development
	transition metal ion transport	iron ion transmembrane transport	regulation of gonadotropin secretion	drug transport	inorgani anion transpo	azole transport	pe late viral transcription	peptidyl-asparagi modification	protein N ine glycosyla aspara	ation via	cysteine metabolic process	cullin deneddylation	peptide metabolic process	acute-phase response
ion transport	mitochondrial ATP synthesis coupled	synaptic vesicle endocytosis	reactive oxygen species metabolic process	Golgi localization	phosphatidylinositol metabolic process	negative regulation of autophagy negative regulation		arginine biosynthetic process	metat	ral transc	ellular ketone ription olic process	acid prote	mitochondrial otein processing eptidyl-histidine	pancreatic
	cellular lipid	lipid modification	carbohydrate phosphorylation	lipid oxidation	deoxyribose phosphate catabolic process	of translational initiation establishment of localization	cellular amide - metabolic process	The state of the s	NAD me	etabolic	glycosyl compound biosynthetic process	citrate	modification otein maturation protein folding	cell apoptotic process hydrogen peroxide metabolism