REVIGO Gene Ontology treemap

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cellular component organization	multicellular organism development	regulation of multicellular organismal process	cellular developmental process	cell differentiation	cellular component assembly	cellular response t stimulus	to stress	response to organic substance	regulation of response to stimulus response	transport	localiza	tion macr	romolecule calization	regulation of cell communication	signaling	signal transduction cell surface	regulation of biological process	of regulation of cellular process
organelle organization	developmental process regulation of multicellular	neurogenesis	complex subunit organization regula anato	mical ture ribonucieopro	organ	cellular response to organic substance	defense response to cytoki	ne response to interferon-gamm response response to organic	external stimulus response	localization in cell		alization lo	of calization import sis into	positive po	intracellular signal transduction odulation costille regulation to the cost of	receptor signaling pathway	bibiologi	negative lation of lation of processical
regulation of cellular component organization	organismal devchromo membrane organization	nervous system development	Anization—guiauoii immune system effe	iune regulation of cytoki	ne projection organization		response to type I interferon signaling pathway	cytokine compound response to response to topologically	to lipid eron onse response to drug	protein to	tablishment of protein ocalization organelle protein regu	retion regular of transpellular	ation vesicle-mediated transport in synapse	regulation of new intracellular regulation of regularization of regularization regularization regularization regulation of regulation of regulation of regulation of regulation of regulation of new interesting regulation regulation regulation of new interesting regulation regula	of respondent to stree egative gulation esponse stimulus	negative regulation of cell	process negative regulation of cellular proces	function Interaction
anatomical structure development development	cytoskeleton organization positive regulation of multicellular organismal process	fiber organization diffi	positive egulation of cell erentiation cell morphoge morphoge involved differential differential	anatomical structure formation str	activation of immune response central nervous system development	to chemical	response to biotic stimulus response to unfolded	cellular response to oxygen-containing compound response to nitrogen inor	response onse to ganic to	substance transport regulation	membrane nitrogen ompound ransport cellular	ocytosis localization of co	wlation cell migration	regulation of cell communication	egative positive regulation of signal transduction qualing	modulation regulation of 1-kappaß of synaptic transmission regulation of 1-kappaß chasseNF-kappaß signaling	regulation of biological quality	function process negative regulation of macromolecule metabolic process
organionina ogoni	cellular macromolecular complex assembly organic	cellular	cellular ortin cytoskele organiza	cellular	oductive ribonucleoprotein complex assembly	response to chemical stimulus	response to endogenous stimulus response to DN. damage stimulu	regulation of A cytokine-mediated s signaling pathway	antibiotic ulation sponse tokine nulus antibiotic response to steroid hormone	of primary metabolic process regulation of macromolecule	metabolic process regulation of protein metabolic	ic met s pro regulation of cellular protein	positive regulation	biologic	al_process	cellu	ılar process	biological regulation
metabolic bio	process positive hete egulation molecular	ynthetic com	process Janic cellul rclic nitrog pound compo abolic catabo	ar aromatic compoun catabolic	J MKNA	metabolic p	n regulation	rocess	etabolic rocess cellular aromatic	regulation of nitrogen compound	regulation of	cellular protein	regulation of proteolysis	cellular com organizati biogene	ion or Vira	al process	localization Si	gnaling cell communication
nucleobase-confucteot compound metabolic process	organic cyclic composition base contains synthetic concess	ase-containing of of containing c	pomolecule com pound ca process	proces	lic transcription, DNA-templated	compou metabol proces	lic process	c compound metabolic abolismess biosynthetic	compound metabolic process catabolic	metabolic process positive regulation of cellula	process positive regulation of cellular	positive regulation of gene	protein ubiquitination regulation of organelle organization		betw dev	interaction peen organisms relopmental process	multicellular organismal process	cell cell proliferation
metabolic process —	obbase-containing cound biosynthetic process reg	ulation atabolic me	etabolic ocess pro- lar amide etabolic ocess trans	ellular regulation of cataly activity ation of cappaB cription activity activity activity	tic hydrolase / activity June 10 DNA Indicate metabolic	macromole metabolic pr cellula	process	process	phosphorus metabolic process	process	positive gulation of ap nucleobase-containin compound metabolic	positive pototic pr of cell death	positive rocession of apoptotic process	metabo respons		mmune system process	cell cycle	postatic of cell or subcellular component component
nitrogen co compound bio	ompound reg osynthetic nucleob	of C	ulation pro atalytic cata	otein rRN/	regulation of sequence–specific DNA binding transcription	macromole metabolic pr		macromolecule modification	phosphate-containing	positive regula of macromoled metabolic prod	regulation of cellular	apoptotic signaling pathway	positive regulation of cell proliferation	stimul	lus mul	ti–organism process	cycle filame	ctin locomotion nt-based prowth