REVIGO Gene Ontology treemap

nuclear body		micro	otubule cytoske	eleton	ton chron		tin	catalytic complex		membrane protein complex		ein	DNA packaging complex	plasma me regio		complex	postsynapse		
cytoplasmic region	transport	vesicle	lumenal si of membra		nucleo	ilus I	vacuolar nembrane	11.0 01.10.01.0 0 p. 01.01.1		oiquitin rece Sytic complex			AP-3 adaptor complex	MHC class		leading	glutama	glutamatergic synapse	
			or membra				Петилапе	protein-DNA complex	RNA effect	tor	on chanr complex	nel ATPase complex		complex	omplex			excitatory synapse	
Golgi apparatus	intrinsic component of <mark>nu</mark> endoplasmic reticulum membrane		clear body nuclear envelo	ope exocy	e exocytic vesicle		P–body	transferase complex	peptida compl	ase lex end	complex c		protein hosphatase		synapse			cell body	
	vesicle me	mbrane	nuclear periphery	fibrillar center		nucleoprotein granule	microbody					cell p	orojection toplasm				whole membrane		
mitochondrion	mitotic spindle		spliceosomal complex	endocytic membra			early endosome membrane	neuron	n n projection		cell		cell lead	ling edge enve		relope	site of polarized growth	cell division site	
perinuclear region of cytoplasm vacu		ole	intrinsic component of organelle membrane preca		lytic	heterochroma	tin organelle subcompartment	dendi			project nembra	tion nBAF complex		oring junction		membrane	cleavage furrow	midbody	