Revigo TreeMap

gene express	sion l	nucleic acid metabolic process		omolecule etic process	small molecule catabolic process	organic acid metabolic process	cellular catabolic process	carboxylic acid catabolic process	response to oxidative stress	esponse to	response to chemical esponse to	amino acid transmembrane transport of organical organical of organical of organical orga	nbrane acid oort transport	
nucleobase–contai compound biosynth process	~	essing	metabolic ocess	rRNA metabolic process	monocarboxylic acid metabolic process amino acid catabolic process carbohydrate catabolic catabolic	catabolic process nione cellular ketone ress aldehyde catabolic process aldehyde catabolic process anide amide	alcohol catabolic process the vacuole digosaccharide process the vacuole metabolic process hydrogen sulfide catabonic metabolic process hydrogen sulfide catabonic metabolic process conosaccharide sulfide catabonic metabolic process conosaccharide sulfide catabonic process catabonic	allantoin allantoin catabolic process serine family amino acid metabolic process process serine family amino acid metabolic process splutarate asparagine catabolic catabolic allantoin catabolic process	chemical stimulus cellular response to nitrogen starvation to	l ~	levels cellular response to stimulus lar response to ethanol ess gen abiotic stimulus ative	ransport putresci transport transport establishment of localization in cell nuclear export	ne organic carboxylic	
nucleobase–containing compound metabolic process	generation cRNA precursor metabolites and energy	processing mRNA cell metabolic alde	hyde DN metal		process nicotinamide nucleotide metabolic process glutamate metabolic process non-prote amino metabolic process	autophagy of mitochondrion sinogenic acid family amino process acid metabolic n	metabolic process glucose phosphate metabolic process ydrogen sulfide netabolic process sulfide metabolic process sulfide metabolic process isocitrate metabolic process process isocitrate metabolic process isocitrate metabolic process	process tricarboxylic acid metabolic process fur metabolic process	biosynthetic process regulation of metaboli negative regulation of gene expression, epigenetic received and the proces	regulation of gene expression it is	te silve methodic insport	mall molecule tabolic process mall molecule tabolic process	cellular component organization or biogenesis	
carbohydrate metabolic process	biosynthetic process tricarboxylic	RNA acid splicing metaboric proce	metabolic process process amino acid metabolic process process process RNA splicing, via transesterification reactions processing process proces		ribonucleoprote complex biogene	ein bio	component	otein-containing mplex assembly organelle organization	regulation of small molecule metabolic process cellular homicellular homeostasis	tion of autophagy of exit of mitosis of exit of from mitosis	amide amide material metaboli	amide netabolic process contains a retrotransposition retrotransposition cycle		
macromolecule metabolic process	acid cycle translation	transport, succinate to ubiquinone DNA replication PNA recombination RNA 3'-end processir maturation of 5.8S rRNA from tricistronic rRNA transcript (SSU-rRNA, 5.8S rRNA, LSU-rRNA) 5.8S rRNA, LSU-rRNA)	RNA modification refolding RNA-templated DNA biosynthetic bios	A replication by KNA polymerase II polymeras	ribon ribosome biogen	compo	organization organization ontaining maturation of 5.8S	of gene organization expression maturation of LSU-rRNA organelle nucleus	carbohydrate homeostasis intracellula nitrogen homeostasis trehalose trehalose carbohydrate phosphorylation tracellula nitrogen homeostas	cell divisio	stimulus pyridine-containi	process metable process viral gene fructo 2,6-bispho metable proces	metabolic process establishment of cell polarity se sphate olic ss	