Revigo TreeMap

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toxin catabolic process	defense respon	defense response to fungus		response to salicylic acid		response to wounding		positive regula of proteasor ubiquitin-depe protein catab process	nal ndent de	regulation of fense response	regulation of immune system process	glutathione metabolic process	translation	
								Calcium Ion	Diodymenotio	auxi e regulation oroce	hetic innate immune response–activating signal transduction		compound metabolic process elongation	
response to hypoxia response to water deprivation systemic acquired resistance	response to cold	response t mechanica stimulus	al response to pr		ubiquitin-de protein cat proces	tabolic	cellular esponse to salicylic acid	regulation of mitotic recombination	posubiquitin-dependent regulation of cellular regulation	regulation of salicylic acid mediated signaling pathway positive egulation of kin mediated regulation of metabolic regulation of salicylic acid mediated	elongation biosynthetic process A	bolic process intochondrial bloosynthetic process alpha-amino acid metabolic process		
		plasma membrane		esponse to calcium-mediate		esponse growth	response to comycetes	transcription, DNA-templated positive regulation	regulation of auxin	ketone metabolic process negative regulation of wax of TOR metabolic process regulation of TOR metabolic process results from transcription, DNA-templated process of positive from regulation of TOR results from transcription, DNA-templated process of positive from regulation of TOR results from transcription, DNA-templated process of the process of	signaling process pathway sitive regulation of protein exit ome endoplasmic reticulum negative negative population process negative negative population	process S-adenosylmethionin cycle protein geranylgeranylation involved in	amine metabolic process 5-phosphoribose 1-diphosphate process process tetrahydrofolylpolyglutamate metabolic process guanosine tetraphosphate	
	response to abscisic acid	2,4,6-trinitrotoluene catabolic process	cinnamic	of molecule nof fungal in	trehalose metabolism n response	response to mannitol	cell-cell	purine nucleotide	50	brane signaling calcium import of train	regulation of programmed cell death rganic acid namembrane transport regulation of maintenance maint	adial pollen-pistil ga ern interaction	biosynthetic process Acturonate osynthetic process L-pipecolic acid biosynthetic process	
	response to fungus signal transduction	camalexin biosynthetic	response to hydrogen	response to symbiotic	10 311 033	sinolate abolic nucleotide-excis	response to organic cyclic compound response to molecule of oomycetes	zinc ion	calcium ion extracellular purine nucleotide transport L-arginine import across plasma ion cart diox	n extracellular ve		ial pattern formation bi	galacturonate osynthetic process g-chain ty acid polyunsaturated polyunsaturated tynthetic fatty acid tynthetic	
		nucleotide-excision repair, DNA	response to UV-	response	ndoplasmic reticulum	nic induced systemic on resistance defense response by callose deposition		transmembpun		calcium carbon dioxide	glucosinolate transport	n n	ocess	
		duplex unwinding defense		response to carbon	methylglyoxal catabolic process to D-lactate via clactoyl-glutathione		response to nickel cation	zinc ion transport	protein targeting to membran	transport gluco	photologem loem in photologem loem loem loem loem loem loem loem lo	tosynthesis, nthesis, compound bic process that harvesting in notosystem I restauration in the state of the s	synthetic metabolic signaling	
response to chitin	response to salt stress	membrane fusion	tryptophan catabolic process to kynurenine	dioxide tri vir mitochondria-nucleus signaling pathway	cell to cell	response to zinc ion	process cellular carbohydrate	protein heterotetramerization Organizati	vesicle	ce centriole tion replication tramerization rotein cell wall		photosyn nune system process respira	tory divcosyl compound	