Genes lost in DRAW1_PELO1_Crassi BP TreeMap

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prostaglandin secretion	L-lysine transmembrane transport	angiotensin-activate signaling pathway	^d nodulati	copper ion homeostasis		single fertilization	peptidyl–cysteine S–nitrosylation	negative regulation of ribosomal protei gene transcription by RNA polymerase	involved	replication	DNA recombination	regulation of production in	nvolved in	positive regulation of interleukin–6 production		granulocyte cotory-standaring factor production
	vascular endothelial growth factor	negative regulation of mitochondrial membrane permeability	heromone–dependent signal transduction nvolved in conjugation with cellular fusion	mRNA export from nucleus	cellular response to interleukin–	3		positive regulation of translation in response	receptor recycling	ceptor regulation of tyrosine phosphorylation regu		regulation o	ion of macrophage ony-stimulating			
ascospore formation	signaling endothelin red basic	ceptor signali			regulation o post–embryor developmen	nic cell receptor	nitric oxide	to stress regulation of repositive regulation of peptidyl-tyrosine	eceptor rec regulation of receptor recycling	ycling on of protein serine/threonine kinase activity	protein K48–linked ubiquitination	facto positive regulation of		macrophage colony-stimulating regulation		amma production involved in inflammatory response
	amino acid transmembrane transport	osteoblast development	regulation of T cell receptor signaling pathway negative	regulation of mitochondr outer membrane permeabilization involved apoptotic signaling pathw	regulation	chondrocyte	biosynthetic process	phosphorylation deadenylation-dependent decapping of nuclear-transcribed	positive regulation of tyrosine phosphorylation o STAT protein negative	negative regulation of DNA-binding transcription factor activity anaphase-promoting	regulation of ergosterol biosynthetic process	interleukin–8	rei ma colon		production interleukin–1 production interleukin–1 production interleukin–1 production production	
endothelin receptor signaling pathway	dendritic transport of messenger ribonucleoprotein complex	mRNA transport	regulation of post-embryonic development sterol transport	regulation of eating behavior regulation of lymphocyte mediated	posit regulat of mem permeat regulat nematod	tition brane protein targeting to lysosome	transcription, RNA–templated	DNA integration	regulation of tubulin deacetylation histone H3-K9 modification	complex-dependent catabolic process positive regulation of mRNA splicing, via spliceosome	regulation of RNA splicing telomere maintenance	suppression by virus of host		n by t type mo	modulation by virus of host	
meiotic cell cycle		cytoly	rtolysis proteason assembly		cell	illing of s of other ganism	transport of virus in host, cell to cell	viral entry	mer ho y m	embrane with nost plasma membrane viral		MDA-5 activit	by virus of	process f host		cellular process
		lymphocy proliferati		mainto chagy mito chr	ulation of enance of tic sister romatid hesion	karyogamy		into host o	recep virio	adhesion otor-mediate n attachmen o host cell		pattern recognition receptor signaling pathway suppression by virus of host viral-induced suppression by			host cellular process	
		osome segreg	homophilic cell adhesion via ation membrane adhesion adhesion molecules ation molecules		a ativ rati	involved	establishment of integrated proviral latency	viral proce	exit of essis from host cell nucleus	viral ranscription	exit of virus from host cell nucleus by nuclear egress	cytoplasmic pattern recognition receptor signaling pathway modulation	by virus of virus of virus of host TBK1 toll-like recept signaling pathw modulation by suppression by virus of host toll-like recept signaling pathw		host STAT activity of host cell cycle modification by	
cellular bud site selection		calcium-depend cell-cell adhesi via plasma membrane ce	chromosom on segregation		ster produid utilizan autop	process utilizing autophagic mechanism	viral genome		viral budding via nost ESCRT	viral budding from Golgi membrane viral capsid	viral protein processing viral DNA genome	host immune response negative regulation of fibroblast	ost defense response syn Ilation of regular rophage macro	nses by cel	Il cycle host G ch	- 01
		regulation natural kil	regulation of proliferation natural killer cell activation response		embly regulation mainten	ance of involved in immune	integration into host DNA	viral gene , expression		viral DNA genome packaging	replication viral genome	regulation of macrophage	ation of made and a second and	crophage ation of maci	e migration rophage motaxis	leukocyte