REVIGO Gene Ontology treemap

positive regulation of natural killer cell mediated cytotoxicity directed against tumo cell target	of cell killing	positive regulation of innate immune response	DNA excision		DN nucleotide-excision repair, DNA incision	regulation of cellular metabolic process	extrinsic apo signaling pa		gnal transductabsence of lig	ion hepatocyte and apoptotic proc	Illelliblati	e regulation of inclusion	carbohydrate metabolic process
	positive regulation of protein processing	response to hydrostatic pressure	kinin g cascade fil	negative gulation of lomerular of he growngiotensin	ation catabo	lic catabolic	apoptotic signaling pat	1.09	ulation of ep	regulation of idermal growth ERB actor receptor signaling pathway	ng biosynthetic process mitochond	respiratory	mitochondrial respiratory chain complex
response to tumor o		regulation of catabolic killer cell medi	peptidyl-glutamine methylation ated cytotoxic plasma	phosphorylation of	response to chromate inst tumor cel	tyrosine phosphorylation target 4 protein regulation of	regulation of s GTPase med signal transdu	small lut liated houction sign	einizing roomone ro	pathway pathw negative egulation of signal regulat of signa	ve cell envelope organization	negative respira respira chai interleukin–10 biosynthetic process assem	tory n inclusion body assembly
response to topologically incorrect protein	regulation of cellular catabolic process	UV protection cellular	kallikrein-kinin cascade positive	regulation of T cell mediated immunity negative regulation	cell mediated immunity cellular response to	response to biotic stimulus	regulation of communica	cell positive interleul	re regulation of kin–6–mediated aling pathway	ulation of T negativell receptor regulation of celepathway	encapsulating structure	nucleolar ragmentation compor organiza	regulation of cellular component
susceptibility to T c	regulation	response to chromate histone glutamine	regulation of fibrinolysis positive regulation of synaptic growth at neuromuscular	activity	stimulus diff	egulation of cell gonadotropin stimulus	phospholipid transport	extracellula transport	r ion transport	vir	ceptor-mediated in attachment to host cell microtubule-deper intracellular transposition of viral material towards nucleur	peroxide	cytokinetic cell cytokinetic cell
cell differentiation involved in labyrinthine layer development	cell differentiation nvolved in salivary gland development	methylation myeloid cell differentiation	compound eye corneal	primary follicle stage	smooth imm muscle sys	immune system osteoblast	anion	of reg	regulation	modulation morpholog	nsport of by virus of host material interaction of host material interaction of host material interaction of host material incompared to host materials.	biosynthesis seiemocysteine biosynthetic process	separation, rigan transition of mitotic cell cycle by negative regulation of transcription from RNA polymerase II promoter
				regulation of excitatory postsynaptic lacenta develop involved in skeletal	THORIC '	ositive gulation ovarian	ceramide 1-phosphate	localization	o	of membrane cal	oplasmic ticulum myeloid c cium ion homeosta neostasis	motabalism	
cell differentiation involved in embryonic placenta development	pancreas development	formation muscle hypertrophy	epithelial cell maturation involved in salivary gland development	regulation of skeletal	development of i	hervous development development hair follicle maturation	transport establishment of protein localization to chromatin	amino acid transport	d secretion,	regulation of of res <mark>response to metal to stimulus process.</mark>	abiotic stimulus— oolic regulation of	cellular organofluorine metabolism	ular nalogen olism killing