## REVIGO Gene Ontology treemap

KLVIGO Gerie Ontology treemap																
epidermis development		skin development		single–multicellular organism process		negative regulation of smooth muscle cell migration	interleukin–1–mediat signaling pathway	ed localiza of ce	ation regulation os diffe	rentiation	defense response to Gram-negative bacterium	intrinsic apoptotic signaling pathway in response to osmotic stress by p53 class mediator	negative regulation of vascular wound healing negative	cell-matrix adhesion regulation of		regulation of smooth muscle cell-matrix adhesion cell-cell adhengative negative nega
						negative regulation of	olfactory bulb axon guidance	of leukotriene production involved in inflammatory response	nucleosi	nucleoside transport monocyte		I TO	of cellular	heterotypic cell-cell adhesion	regulation of monocyte aggregation	regulation of cell adhesion mediated by integrin chorio–alla
organ development tissue		volonment	epithelium developmei	) Va	lomerulus asculature evelopment	fibrinolysis	keratinocyte proliferation	antigenic	of glutama	hormone	nyroid response to carbon etion monoxide	mycophenolic acid negative regulation of collagen biosynthetic	response	granule	yaluranon regula cable hyalu assembly ca	response response
		velopment	ossification	1	lens velopment in nera-type eye	of smooth muscle cell proliferation	cytoskeleton-dependent intracellular transport	vestibular reflex	interleukin–8 fa	-8 factor-medial	organelle transpo				regulation of	cerespons
renal system outer me vasculature development	g duct myo	ompact	nerve	nelial cell	organ morphogenesis	intrinsic apoptotic signaling pathway by p53 class mediator			positive regulation o	oth muscle		negative regulation of		g	cable assemap junction assembly	hyperoxia regulation resp
regulation of cell developmental differentia	muscle	cell chronolo	· ·	cloacal septation	female genitalia morphogenesis	muscle cell	cellular response to isoquinoline	organelle transport along	protein localization to adherens junction	cell-cell signaling	keratinocyt migration	homeostasis	positive regulation of fibroblast apoptotic process	organization	otubule anchoring act microtubule rganizing center	to osmotic stress str
skeletal odontogene system	l IIIcuui	la morphoge	I of planar I	establishmen of tissue polarity	urinary bladder development	cell migration	l my conhonolia	by p53 class	protein localization to cell	response regulation o multicellular	f wound healin	secretion	cytokine-mediated signaling pathway	multicellu organismal p		cellula opmental ocess
ectoderm and cochle mesoderm	renal ou ea medul nent developr	la developr	n of a polarized ment epithelium	central nervous system vasculogenesis	lobar bronchus development T cell receptor	muscle cell proliferation	cellular response to sorbitol	mediator acute-phase response	junction regulation of cell projection size	organismal developmen cellular response heparin	heart	follicle-stimulating hormone stimulus  direct  n ossification	ossification  negative regulation of lipopolysaccharide-mediated signaling pathway			single-orga cellular pro
interaction establishment or maintenance of polarity of	-1 7	syste developr	ment differentiation apoptotic	of mesoderm development trachea	· ·	negative regulation of endopeptidase	negative regulation of elastin biosynthetic	nentide	hyaluronan metabolic		extracellular olysaccharide metabolic process	peptidyl-glutamine modification transcription	negative regulation of metabolic process	regulation o multicellula organismal		epidermal interme filament-proces
entifyonic epithelium negative regulation cardia	differentia	develop	ment involved in patterning of blood vessels collecting	development	establishment of blood-brain barrier vound healing,	activity negative	regulation of	regulation	•	extracellular polysaccharide	carnitine ase activity process elastin	from RNA polymerase II promoter negative	positive regulation of MAPK cascade DNA damage	biological	locomotion	
of mesoderm muscle development developn	cell development	Lodontoge	nesis duct development	process	spreading of epidermal cells	3	Janus kinase	of gene expression		biosynthetic process	metabolic process	I of molecular I	induced protein phosphorylation	adhesion		response establishme maintenand cell polar