## REVIGO Gene Ontology treemap

_	NEVIGO Gene Ontology treemap													
regulation of acid-sensing ion channel activity		ng CD40 signalii pathway	~	toll–like receptor 9 signaling pathway		esion i	ipoprotein ocalization	phosphatidylcholine acyl–chain remodeling	1 20 20 20 20 20	metah	enylalanine olic process	ribosome–associa ubiquitin–depende protein catabolic process	nt rescue	of stalled osome
	lipoprotein transport	sodium ion export from cell	positive regulation of macrophage activation	positiv regulatio protein targ to membr	of cytokine signaling	g pathway e	negative regulation of ndothelial cell optotic process	pentose–phosphate shunt	chondroitin su biosynthetic pr	ılfate 6- ocess r	glucose phosphate netabolic process	membrane protei ectodomain proteolysis	prot	ane protein eolysis
	establishment or maintenance of transmembrane electrochemical gradient	JAK-STAT cascade regulatior	I–kappaB kinase/NF–kappaB signaling of acid–sensing	10.7 400 4	ctivity nlasma	sium ion t across membrane	interaction with symbiont	glycosaminoglycan biosynthetic process <sup>p</sup>	bhosphatidylcholine mu metabolic process		ing triglyceride metabolic process			nt
	positive regulation by host of viral process	STAT cascade	regulation of ATPase activity	modification of morphology or physiology of other organism	phospholipid transport	ion transpor	response to cytokine	glycerolipid metabolic process	glucose metabolic process	proteoglycar metabolic process	aminoglycar metabolic process	process	oroteolysis	cellular macromolecule biosynthetic process
	cellular potassium ion homeostasis	chemical homeostasis	Sodium ion	ransition meta ion transport	cellular response to lipopolysaccharide	positive regulation of I–kappaB kinase/NF–kappa signaling	nuclear export		neutral lipid metabolic process		nolecule c process	organonitrogen compound biosynthetic process	macromolecule biosynthetic process	
	monocyte differentiation	iron ion transport	cholesterol homeostasis	metal ion transport	lipid homeostasis	cellular response to biotic stimulu	ester transport	organonitrogen compound roganonitrogen compound metabolism cellular amide metabolic process	glomerular viscera epithelial cell differentiation	al sulfur com metabo	ipound <mark>carbo</mark> lism meta	biosy bhydrate abolism	rotein nthesis n rhythm	lipoprotein lipoprotein metabolism process