## REVIGO Gene Ontology treemap

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identical protein binding molecule bind		protein dimerizatior activity	n tub	ulin binding	ubiquitin–like protein ligase binding		transferase activity, transferring phosphorus–contain groups	ubiquitin- proteir transfera activity	n protein kinase ase activity	transferring	ion transmembrane transporter activity ion tran	zinc ion transmembra transporte activity gated chan activity smembrane	er ————————————————————————————————————	alcium ion binding	transition metal ion binding	
cytoskeletal protein binding	phosphatase binding	receptor C-	protein -terminus binding	acid	rotein binding involved in heterotypic cell-cell adhesion	GABA receptor binding	ubiquitin–like protein ligase ubiquitin– activity	protein		N-methyltransferase	transition transporter transmembrane transporter activity	passive sia transmembrane transn transporter trans	sialic acid transmembrane transporter activity  zinc ion bi		manganese ion binding	
	ident cadherin binding	heat shock ical protein bind binding	ion ling channel binding	ephrin receptor binding	almodulin binding	histone binding	transferring one-carbon	-adenosylmethionine-dependent methyltransferase activity	nexakisphosphate methyltr	itone lysine nansferase N-methyltransferase activity	cation transmembrane transporter activity	carbohydrate transme transporter activ			magnesium ion binding	
protein domain specific binding	protein	binding	haperone binding	Hsp70 protein binding	neurotrophin TRK receptor binding	RNA polymerase I core binding	groups transmembrane receptor protein tyrosine kinase activity	protein tyrosine kinase activity	activity N-acetyl	ptide transferase nucleotidytransferase activity	transcrip	tivity,	drug bind		chromatin binding	
	phosphatase binding	beta-2-microglobulin binding	53 binding	epidermal	K48 linkod	acetylcholine	hydrolase	hydrolas activity,	1	phosphatase transcriptional activator activity,				cleosome low-density lipoprotein particle binding		
receptor binding	protein heterodimerization activity	transcription as factor binding	polipoprotein binding	misfolded protein binding	phosphatidylinositol 3-kinase binding	RNA polymerase II sequence-specific DNA binding transcription factor binding	activity, acting on ester bonds	acting on a	acid hydrolas	activity	sequence-specific		ansporter activity		peptide antigen	
adenyl nucleotide binding		RNA binding  egion nucleic acid binding  large ribosor subunit rRN binding  regulatory region nucleic acid binding  telomeri		sequence	•	miRNA	ribonuclease hydrolase activity	peptidase activity, act	exopeptidase vity, acting on ester bo		enzymetregul activity		orter activ	ity	binding	
				g	DNA binding binding		ATPase activity	phosphatidylinosito bisphosphate phosphatase	poly(A)-specific ribonuclease activity	helicase activity	protein phosph activator acti	vity transc	ription ractivity	ty		
				subunit rRNA binding telomeric DNA bindin	bindir 7SK snF	ng mRNA 5'-UTR RNA binding	aminopeptidase activity	activity	phosphate phosphatase activity	trisphosphate phosphatase activity threonine-type peptidase activity	macromolec complex bind	ding			activity  binding,	