REVIGO Gene Ontology treemap

cell adhesion molecule binding	identical protein binding	tubulin binding	phosphatase binding	protein phosphatase binding	transferase activity, transferring	N–acyltransferase activity	transferase activity, transferring acyl groups	hydrolase activity, acting on ester bonds		ribonuclea activity	IATPase activity
					phosphorus–containing groups						
protein domain specific binding	cell adhesic ubiquitin–like protein ligase binding	protein heterodimerization activity on molecule binding	receptor binding	histone binding	transferase activi peptide N–acetyltransferase activity	transmembrane receptor protein tyrosine kinase	protein kinase	hydrolase activity.		ity, acting of activity	exonuclease activity, nester bonds rribo- or deoxyribonucleic acids and producing 5'-phosphomonoesters
		chaperone binding	RNA polymerase core enzyme binding	beta-2-microglobulin binding	ubiquitin–like protein transferase activity		rane receptor			phosphatase activity	phosphatidylinositol bisphosphate phosphatase activity
cadherin binding	cytoskeletal protein binding	protein dimerization activity	heat shock protein binding	protein C–terminus binding	chromatin binding	g	macromolect		drug bind	ding	enzyme regulator activity
RNA binding regulatory reg		adenyl nucleotide bind		ılatory region ic acid binding	nucleosome bindir	binding					NADH
		on nucleic acid binding		ecific miRNA	zinc ion ion	transition metal ion	transcriptio	on	peptide antigo	en binding	dehydrogenase activity
		ribonucleotide bindin	sequence–sp DNA bindi		transmembrane transmembrane transmembrane transporter transporter activity	rane I sporter activity er transporter	cofactor acti	vity	transporter	activity	manganese ion binding