Revigo TreeMap

RNA binding <b>mRNA bindin</b> g	mRNA binding	tRNA binding	histone H3K14 acetyltransferase activity histone H3K14 a	acetyltransferase		JUN kinase kinase activity ivity	kinase activity ubiquitir		histone binding	TBP-class protein binding
		nucleic acid binding	histone H4K8 acetyltransferase activity	kinase activity		acetyltransferase activity	cadherin binding		RNA polymerase II complex binding	methylated histone binding
	telomerase RNA binding	rRNA binding	transcription coregulator		transcription activity vator activity	ATP hydrolysis activity bydro		hydrolase activity, hydrolyzing O_alycosyl tivity compounds	y, cing	
protein binding					33.111,	cysteine-type deubiquitinase activ		nase activity		
			protein phosphatase <b>protein phosphatase</b> regulator activity	enzyme regulator activity regulator activity		NADH <b>NADH dehy</b> activity	drogenase	ctron <b>activity</b> tivity	molecular adaptor activity	4 iron, 4 sulfur cluster binding
				ornithine decarboxylase inhibitor activity		S-adenosyl-L-methionine binding		onine	chromatin binding	