microtubule bindin	ng histone bii	histone binding unfolded p		aci	ctin binding	RNA binding		DNA bir	inding	chromatin binding		nucleosome binding	osphatidylinositol–3,4,5–trisp binding phosphatidylinos		nositol–4,5–bisphosphate binding risphosphate
						mRNA binding		DNA	DINA billuling				manganese ion binding		
protein heterodimerization activity								replication origin binding			cyclin-depe protein serine/three	n	transcription transcription transcription coactivatoractication activity		
	protein kinase binding microtu	actin filament binding ubule binding	transcriptio		biquitin-specific protease binding			translation initiation	sequence–specific mRNA binding	nucleoside diphosphate	diphosphate activity tau-prote kinase act	tein			transcription oractivity activity
			disordered	,	K63–linked polyubiquitin			factor activity							
heat shock protein binding chaperone binding	ubiquitin binding	tubulin binding	domain specif binding	modific	protein binding	GTPase activator activity	cyclin-dependent protein serine/threonine	ubiquitin conj	njugating	NADH d NADH dehydro		molecula	ivity ar adaptor p	complex binding	
	ubiquitin protein	misfolded	phosphotyrosi residue bindir		sine–acetylated sistone binding		kinase regulator activity	enzyme activity		((ubiquinone) a activity	activity	acti clathrin act	adaptor ribosome		
	ligase binding	protein binding	Teologic	ig		cyclin-dependent protein GTPase activator activity Serine/threofine kinase		ubiquitin conj enzyme ac							
	histone deacetylase binding	beta-catenin binding		calmodulin binding		protein phosphatase regulator activity	guanyl-nucleotide exchange factor activity	peptidyl–prolyl cis–trans isomerase activity	ubiquitin–ubiquitin ligase activity	protein tyrosine/serine/tl phosphatase a	threonine	protein fol chapero	lding cytosk one motor a	celetal co	structural constituent of ribosome