RNA binding		DNA DNA binding origin binding		single-stranded DNA binding	unfolded protein binding	chaperone binding	ubiquitin binding	protein kinase binding	binding	protein-containing complex binding protein-c		o: phatidylinositol3,4,5-trisp pbssphta idylinositol4,5-bis binding binding binding phosphatidylinositol3,4,5-trisphosphat binding		chromatin binding chromatin binding		
		translation initiation IA binding activity	poly(U) RNA binding	snRNA binding	mRNA 3'–UTR binding	unfolde protein heterodimerization	d protein binding beta-catenin binding	calmodulin binding	histone	nase binding ubiquitin-specific protease binding	G-protein		os phatidylinositol—Qə lphqəsintidy lino binding bi	sitol-4-ph daphptæ tidylinositr nding binding	h <mark>dapsptatidylinositä</mark> binding nucleosomi	
mRNA binding		sequence–specific mRNA binding	mRNA 5'-UTR binding	four-way junction DNA binding	elongation factor activity	activity	syntaxin binding	cadherin binding	binding	enzyme binding	transcription coactivator	transcription	oxidase	tational	at shock atshockg	disordered domain disordered domain
		E-box response snRNA snRI		U6 snRNA binding	microtubule bindi	actin filamer ng bindin	nt tubulin binding	histone binding modification-dependent histone binding modification-dependent protein binding		atranscription activity coactivator activity transcription DNA-binding		oxidase acti phospholipid scramblase	prote	protein binding Hsp70 protein		
ubiquitin conjugating enzyme activity	peptidyl-prolyl cis-trans isomerase activity	cyclin-dependent protein serine/threonine kinase activity	tau-protein kinase activity	threonine-type endopeptidase activity flap endonuclease		micro	otubule binding alpha-tub binding	minus-end	lysine-acetylate	methylated	corepressor activity cytoskeletal	factor activity	structural constituent	protein folding	misfolded protein	phosphotyrosine residue binding
NADH	ubiquitin-ubiquitin	ribonuclease NADH dehydrogenase	DNA-directed 5'-3' RNA (guandsi polymerase activity	tRNA ne-2'-O-)-methyltrar activity	actin binding	myosir binding	l monomer	histone binding	histone	motor activity cytosko motor a		of ribosome structural molecule activity	chaperon			
dehydrogenase (ubiquinone) activity	ligase activity ubiquitin conj	endonuclease activity	ne activity protein-disulide reductase activity	phospholipase D activity	protein serine/threonine kinase activity	GTPase activator ser		nucleotide nge factor tivity protein phosphatase inhibitor activity	androgen receptor	transcription factor binding androgen	microtubule motor activity manganese	motor activity	clathrin adapto clathrini adapto activity clathrin light	activity		
nucleoside diphosphate kinase activity	tyrosine/serine/threonine phosphatase activity	ATP hydrolysis activity	endopeptidase activity	kinase activity		protein cyc	activity activator activity lin-dependent protein ATF	Pase	recepto	binding P-class nuclear	ion binding manganese id	binding	chain binding MHC cMHC class I	initiation	aynom ng.	te proteasome-activating activity
	GTPase activity	histone methyltransferase activity (H3–K4 specific)	histone acetyltransferase activity	histone demethyla activity	ase	regulator	rine/threonine activator	ubiquitin-protein transferase activator activity	recentor	protein receptor binding	magnesium ion binding	ATP binding	protein binding	g cyclospor A binding	See to the force of	2 iron, 2 sulfur cluster binding