phosphatidylinositol dephosphorylation	protein peptidyl–prolyl isomerization	regulation of macroautophagy	protein K63–linked deubiquitination	positive regulation of cell growth	maturation of SSU–rRNA from tricistronic rRNA transcript (SSU–rR 5.8S rRNA, LSU–rR	NA, actin org	cytoskeleton Janization	lysosome localiza	ion lysosome tra	nsport estat	establishment of planar polarity	f neural tube closure
positive regulation of DNA replication	protein deubiquitination	RNA splicing	piRNA processing	chromatin remodelingos	regulation of sitivelregulation of	mitochondrial ell growth chain complex I assembly	rRNA processing	regulation of	Golgi to plasm membrane prot			
								regulation of synaptic v€ lysoso n exocytosis	•		establishment of _{negativ} nar polarity	
						_			positive regulat of endocytosi	on of	gulation ⁻ neuron ojection	neuron development
negative regulation of NF–kappaB transcription factor activity	_	positive dephosphorylation nuclear-transcribed mRNA poly(A) tail shortening	regulation of gene expression	endosome organization	mitochondrion organization	ribosomal large subunit assembly		positive regulation		dev	elopment	
	phosphatidylinositol of pre–miRNA processing							of protein import into nucleus	endoplasmic reticulum to Go vesicle-media	lgi	axonogenesis	
				cellular response to glucose starvation	negative regulation of non–canonical NF–kappaB signal transduction	epidermal growth factor receptor signaling pathway			transport	nsport		
positive regulation of transcription by RNA polymerase I	mitochondrial translation	positive regulation of protein ubiquitination	regulation of translation					2–oxoglutarate metabolic process		cell–cell adhesion		
				cellula	positive regulation r response to glucose starvation signaling pathway					negative regulation of		
GPI anchor biosynthetic process		negative regulation of translational initiation		response to lipopolysaccharide	defense response to virus	signaling	patnway			mitotic cytokinesis		fibroblast proliferation
	positive regulation of nuclear–transcribed mRNA catabolic process deadenylation–dependen decay					transformi factor beta	gulation of ng growth a receptor pathway	blood vessel ren	nodeling n			