translation		cyto	translational initiation  cytoplasmic translation		regative pulation of templated inscription regative pulation of inscription by RNA regated for the templated for the tem			endosomal transport  amino acid transport vacuolar along transport microtubule  ndent cotranslatio  geting to membra			targeting to ER  al assembly assembly rRN/ mitochondrial respiratory chain complex lassembly		
mRNA splicing, via spliceosome	mRNA processing	DNA metabolic <b>anslati</b> (	on II neg	protein N-linked glycosylation ria asparagine ative	mitochondrial translational termination positive regulation	proteir transpo vesicle	IIIItociloliulia	migration intracellular		phospholipid efflux	protein export from nucleus	subunit asse <b>ribo</b> large	emodeling osomal subunit embly
translational elongation	positive regulation of transcription by RNA polymerase II mitochondrial electron			of lation	of strand invasion regulation of lipid metabolic process	tra	cellular response to insulin stimulus	transport response to high light intensity	to starvation		localization	nı	icleosome issembly
RNA splicing	transport, NADH to ubiquinone	lipid biosynthetic process respiratory electron transport chain	DNA biosynthetic	scription bio	piRNA synthetic	cellular resporte to unfolded protein	regulati m RNA p in res		rase	II pron		of LSU-rRNA	
transcription by RNA polymerase II	opnonig, via	translation reinitiation	protein glycosylation	pho	process pspholipid synthetic process		response to glucose	response to unfolded protein			regulation cellular ami	regulation	citrate metabolic
regulation of mitotic mitotic	proc	mRNA catabolic process, nonfluclear—transcrit decay mRNA cataboli process,			peptidyl-threonine dephosphorylation			acid metabolic process process regulation of cellular amino acid metabolic process					
cycle phase transition negative regulation of	egative gulation f mitotic ell cycle	cel	le regulation		e_media cay	ated	histone H3 acetylation		İ		ATP metabolic process	process	purine nucleotide metabolic process
G2/M transition of mitotic cell cycle  regulation of hematopoietic stem cell	regulation of cell shape		of mRNA stability NIK/NF-kapp signaling	Rab pro signa transduc	l signalin	g	spindle assembly mitotic_s assen			riral cription	chromo segreo	osome	meiotic cell cycle
regulation otic aging regulation of hematopoiet stem cell differentiation			ic NIK/NF-kappal				inner dynein arm assembly		protein cell redox			regulate on of tic <sub>hydrola</sub>	ion ase
positive regulation of fat cell differentiation multicellular organism aging	development d	erythrocyte ifferentiation hymic T cell selection					folding protein f protein refolding					cell	of cytoskeleton organization