development	Begative regulation of average in pronephic hephron development Comparison of the comparison of transcription of nucleolar large rRNA by RNA polymerase I opening regulation of transcription from RNA polymerase II promoter in response to heat stress mRNA transcription	binding	- myosin light chain binding	
ÖĞSS	positive regulation of transcription of nucleolar large rRNA by RNA polymerase I cytokinin biosynthetic process positive regulation of transcription from RNA polymerase II promoter in response to heat stress mRNA transcription positive regulation of DNA-binding transcription factor activity dopamine biosynthetic process S-adenosylmethionine biosynthetic process phosphatidylethanolamine biosynthetic process		-sequence-specific single stranded DNA binding -sequence-specific DNA binding	
process	- hydrogen peroxide catabolic process - positive regulation of proteasomal ubiquitin–dependent protein catabolic process		RNA polymerase II cis–regulatory region sequence–specific DNA binding	
	regulation of programmed cell death		RNA polymerase II transcription regulatory region sequence-specific DNA binding	
death	e programmed cell death	ac	THE TOTAL PROPERTY OF	
proliferation	-positive regulation of cardiac muscle myoblast proliferation			
signaling	onemical cyriapus danemics.		- DNA topoisomerase binding	
ass	- polyphosphate–mediated signaling - negative regulation of lateral pseudopodium assembly - positive regulation of inclusion body assembly		guanylate cyclase activator activity	
sembly		activity	- adenylate cyclase activator activity	
meta	spermine acetylation putrescine acetylation nor-spermidine metabolic process spermidine acetylation		- acetyltransferase activator activity	
ound boli	spermidine acetylation positive regulation of mRNA polyadenylation negative regulation of double-strand break repair via nonhomologous end joining putrescine catabolic process positive regulation of apoptotic DNA fragmentation			
proce.	histone H3–K23 acetylation histone H3–K14 acetylation histone H4–K12 acetylation	carbo	N-formylglutamate deformylase activity	
: tion	- protein ubiquitination - positive regulation of protein polyubiquitination			
organization	- actin cytoskeleton organization	glyco	- hydrolase activity, hydrolyzing N–glycosyl compounds	
maturation	- synaptic vesicle maturation			
			FAD binding phosphatidylinositol phosphate binding	
process	positive regulation of cellular pH reduction over_re	epresented_pvalue		over_represented_pvalue
process	B cell activation - 0.0 - 0.0	03 02 01	glutamate 5-kinase activity -insulin-activated receptor activity -mannokinase activity -hexokinase activity -glucokinase activity	0.03 0.02 0.01
process	regulation of lipid metabolic process		- fructokinase activity	
<i>37</i>	cardiolipin acyl–chain remodeling positive regulation of positive chemotaxis to cAMP	2	RNA-3'-phosphate cyclase activity	
	chemotaxis to folate hematopoietic stem cell migration		প্রতিষ্ঠিত বিশ্ব	
	regulation of fibroblast migration sensory perception of pain		- carbonate dehydratase activity	
process	- sensory perception of chemical stimulus - cognition		- glutathione specific gamma-glutamylcyclotransferase activity	
fo			- tryptophanase activity - gamma-glutamylcyclotransferase activity	
olding	negative regulation of chaperone-mediated protein folding		- sulfide:quinone oxidoreductase activity - peroxidase activity - polyprenol reductase activity - 8—methylthiopropyl glucosinolate S—oxygenase activity	
i di con	-asexual reproduction	activity	peroxidase activity -polyprenol reductase activity -8-methylthiopropyl glucosinolate S-oxygenase activity -4-methylthiopropyl glucosinolate S-oxygenase activity -glutamate-5-semialdehyde dehydrogenase activity -peptide-methionine (R)-S-oxide reductase activity -peptide-methionine (R)-S-oxide reductase activity -proline dehydrogenase activity -proline dehydrogenase activity -oxidoreductase activity, acting on paired donors, with incorporation or reduction of molecular oxygen, another compound as one donor, and incorporation of one atom of oxygen -hydroxymethylglutaryl-CoA reductase (NADPH) activity -a-oxo-5-alpha-steroid 4-dehydrogenase activity	
	regulation of female receptivity		- oxidoreductase activity, acting on paired donors, with incorporation or reduction of molecular oxygen, another compound as one donor, and incorporation of one atom of oxygen - hydroxymethylglutaryl–CoA reductase (NADPH) activity - 3-oxo-5-alpha-steroid 4-dehydrogenase activity	
to stress	response to oxidative stress response to water deprivation response to psychosocial stress positive regulation of behavioral fear response response to hypoxia melanotic encapsulation of foreign target response to ischemia cellular response to hydrogen peroxide positive regulation of autophagy of mitochondrion in response to mitochondrial depolarization response to hypoxia	bin	- Tat protein binding	
4	positive regulation of autophagy of mitochondrion in response to mitochondrial depolarization response to hypobaric hypoxia insulin receptor signaling pathway acetylcholine receptor signaling pathway	ding	cAMP response element binding protein binding	
ansduction	insulin receptor signaling pathway acetylcholine receptor signaling pathway regulation of adenylate cyclase—activating G protein—coupled receptor signaling pathway positive regulation of TORC2 signaling interleukin—18—mediated signaling pathway negative regulation of Wnt signaling pathway Notch signaling pathway angiotensin—activated signaling pathway negative regulation of Notch signaling pathway long—chain fatty acid biosynthetic process		-1-acylglycerol-3-phosphate O-acyltransferase activity -alpha-1,6-mannosylglycoprotein 4-beta-N-acetylglucosaminyltransferase activity	
pro	I negative regulation of Notch signaling pathway long-chain fatty acid biosynthetic process gliccosplyitae biosynthetic process and the first of the signaling pathway and the signaling of Notch signaling pathway and signaling pathway and signaling pathway and signaling pathway and signaling pathway		mycocerosate synthase activity diamine N-acetyltransferase activity	
cocess	Ing-chain fatty acid hiosynthetic process Official light was the first light was the first process Official light was the first light was		UDP-alpha-D-glucose:glucosyl-glycogenin alpha-D-glucosyltransferase activity - glycogenin glucosyltransferase activity - methionine adenosyltransferase activity	
process		activ	acetylcholine-gated cation-selective channel activity fructose transmembrane transporter activity carbohydrate:proton symporter activity dehydroascorbic acid transmembrane transporter activity aromatic amino acid transmembrane transporter activity thyroid hormone transmembrane transporter activity	
	thyroid stimulating hormone secretion Eligible is it is it is it is it is a secretion into mitochondrial outer membrane		9 glucose transmembrane transporter activity 1 potassium ion antiporter activity 2 monocarboxylic acid transmembrane transporter activity	
	hyroid stimulating harmons secretion Stiff of the first of the first on into mitochondrial outer membrane Stiff of the first of th		- amino acid transmembrane transporter activity - acetylcholine receptor activity - glucose binding - neme binding - quinone binding - quinone binding	
	ne navioral response to nicetine Commonwealth		acetylcholine receptor activity fluctose binding fleme binding quinone binding ethanolaminephosphotransferase activity D5 dopamine receptor binding translation elongation factor binding leukotriene—A4 hydrolase activity TB domain binding phosphotransferase activity, for other substituted phosphate groups TORC complex binding epoxide hydrolase activity insulin binding anaphase—promoting complex binding HSp00 protein binding promoter—specific chromatin binding	
BP	De pavioral response to nightine Eligible April 1990 (1990) (199	MF	epoxide hydrolase activity - insulin binding - anaphase-promoting complex binding - Hsp90 protein binding - promoter-specific chromatin binding	
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