	anatomical structure development	accative regulation of synaptic grawth at neuromuscular junction development		cytoskeletal protein binding	myosin light chain binding	
	nthetic cess	positive regulation of transcription of nucleolar large rRNA by RNA polymerase I cytokinin biosynthetic process positive regulation of DNA-binding transcription factor activity steroid biosynthetic process dopamine biosynthetic process S-adenosylmethionine biosynthetic process phosphatidylethanolamine biosynthetic process			RNA polymerase II transcription regulatory region sequence–specific DNA binding	
	catabolic process	hydrogen peroxide catabolic process positive regulation of proteasomal ubiquitin-dependent protein catabolic process positive regulation of autophagy of mitochondrion in response to mitochondrial depolarization		DNA	sequence-specific DNA binding	
	cell adhesion	regulation of cell-cell adhesion mediated by integrin		en bi		
	cell death	regulation of programmed cell death programmed cell death		nzyme inding	DNA topoisomerase binding	
	cell population proliferation	positive regulation of cardiac muscle myoblast proliferation		en reg ac	guanylate cyclase activator activity adenylate cyclase activator activity	
	cell-ce signalir	synaptic transmission, cholinergic synaptic transmission involved in micturition neuromuscular synaptic transmission chemical synaptic transmission			acetyltransferase activator activity	
	cellula compon assemb	polyphosphate-mediated signaling negative regulation of lateral pseudopodium assembly axoneme assembly caveola assembly axonemal central apparatus assembly		hydrolase activity, acting on carbo	N-formylglutamate deformylase activity	
		positive regulation of non-motile cilium assembly histone H3-K23 acetylation histone H3-K14 acetylation histone H4-K12 acetylation		hy a a ac g		
	n develo	positive regulation of protein polyubiquitination synaptic vesicle maturation		ydrolase activity, acting on glyco	hydrolase activity, hydrolyzing N-glycosyl compounds	
	ental homeos	positive regulation of cellular pH reduction		bii	phosphatidylinositol phosphate binding FAD binding	
	tatic				iron ion binding	
	ne m	B cell activation over_represented_pvalue 0.04 0.03 0.02		kinas activ	glutamate 5-kinase activity insulin-activated receptor activity mannokinase activity hexokinase activity	ver_represented_pvalue 0.04 0.03 0.02
	lipid stabolic	regulation of lipid metabolic process lipid hydroxylation cardiolipin acyl-chain remodeling phosphatidylinositol acyl-chain remodeling			glucokinase activity fructokinase activity	- 0.01
	locomotion	positive regulation of positive chemotaxis to cAMP chemotaxis to folate hematopoietic stem cell migration regulation of fibroblast migration		ligase activity	RNA-3'-phosphate cyclase activity methionine-tRNA ligase activity	
	nervous system process	sensory perception of chemical stimulus cognition associative learning			carbonate dehydratase activity	
	protein	negative regulation of chaperone-mediated protein folding protein folding in endoplasmic reticulum			glutathione specific gamma-glutamylcyclotransferase activity tryptophanase activity gamma-glutamylcyclotransferase activity	
	protein	negative regulation of protein processing positive regulation of protein processing		oxidore	sulfide:quinone oxidoreductase activity peroxidase activity polyprenol reductase activity 8-methylthiopropyl glucosinolate S-oxygenase activity 4-methylthiopropyl glucosinolate S-oxygenase activity glutamate-5-semialdehyde dehydrogenase activity pertide-methionine (R)-S-oxide reductase activity	
	eproduction	asexual reproduction regulation of female receptivity		ductase	sulfide: quinone oxidoreductase 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 isovaleryl-CoA 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 3-oxo-5-alpha-steroid 4-dehydrogenase activity monooxygenase activity estrogen 2-hydroxylase activity estrogen 2-hydroxylase activity retinoic acid 4-hydroxylase activity	
	response to stress	response to oxidative stress response to water deprivation response to hypoxia positive regulation of behavioral fear response		transcription factor binding	Tat protein binding	
	-	response to ischemia melanotic encapsulation of foreign target insulin receptor signaling pathway acetylcholine receptor signaling pathway regulation of adenylate cyclase—activating G protein—coupled receptor signaling pathway positive regulation of TORC2 signaling adenylate cyclase—modulating G protein—coupled receptor signaling pathway negative regulation of Wnt signaling pathway interleukin—18—mediated signaling pathway Notch signaling pathway angiotensin—activated signaling pathway negative regulation of Notch signaling pathway long—chain fatty acid biosynthetic process			cAMP response element binding protein binding 1-acylglycerol-3-phosphate O-acyltransferase activity alpha-1,6-mannosylglycoprotein 4-beta-N-acetylglucosaminyltransferase activity	
process	small molecule metabolic	Notch signaling pathway angiotensin—activated signaling pathway negative regulation of Notch signaling pathway long—chain fatty acid biosynthetic process life of the signaling pathway long—chain fatty acid biosynthetic process life of the signal process life of th		transferase activity, transferring	mycocerosate synthase activity UDP-alpha-D-glucose:glucosyl-glycogenin alpha-D-glucosyltransferase activity glycogenin glucosyltransferase activity methionine adenosyltransferase activity	
		hydrogen sulfide metabolic process hydrogen sulfide metabolic process		transm trans ac	acetylcholine-gated cation-selective channel activity aromatic amino acid transmembrane transporter activity thyroid hormone transmembrane transporter activity	
		sulfide oxidation, using sulfide:quinone oxidoreductase thyroid-stimulating hormone secretion into mitochondrial outer membrane this live regulation of profession insertion into mitochondrial outer membrane this live regulation of all the chondrial membrane permeability involved in apoptotic process		brane orter ty	potassium ion antiporter activity monocarboxylic acid transmembrane transporter activity glucose:sodium symporter activity amino acid transmembrane transporter activity	
	nsport	thyroid—stimulating hormone secretion into mitochondrial outer membrane The plant of the part in the property of the property of the part		NA	acetylcholine receptor activity heme binding glucose binding duinone binding ethanolaminephosphotransferase activity D5 dopamine receptor binding leukotriene—A4 hydrolase activity alkenylglycerophosphoethanolamine hydrolase activity alkenylglycerophosphocholine hydrolase activity PTB domain binding phosphotransferase activity, for other substituted phosphate groups TORC2 complex binding epoxide hydrolase activity unfolded protein binding insulin binding	
BP	N _A	BENEVICE FOR SPIRE FOR INTERPORT IN PROJECT DE LA PRODUCTION BUTTER STATEMENT DE LA PROJECT DE LA P	MF		PTB domain binding phosphotransferase activity, for other substituted phosphate groups TORC2 complex binding epoxide hydrolase activity unfolded protein binding insulin binding	