	g	-chronological cell aging			l ¬≞.o.∣	- DNA-binding transcription factor activity, RNA polymerase II-specific	
	anatomical structure development	regulation of cell morphogenesis serotonergic neuron axon guidance dopaminergic neuron axon guidance planar cell polarity pathway involved in axon guidance regulation of dendrite extension negative regulation of dauer entry post—embryonic animal organ morphogenesis axonal fasciculation nervous system development			ling		
	biosynthetic process	lipoxin biosynthetic process			ion binding	- zinc ion binding	
	catabolic process	-positive regulation of autophagy				- phosphatidylinositol–3,5–bisphosphate binding - endonuclease activity	
	cell	-homophilic cell adhesion via plasma membrane adhesion molecules			clease	- type II site–specific deoxyribonuclease activity - RNA–DNA hybrid ribonuclease activity	
	cellular nitrogen compound metaboli	BNA integration involved in BNA mediated transposition  FY 15 - 16 15 16 16 16 16 16 16 16 16 16 16 16 16 16			nucleot	ribonuclease activity - RNA-directed DNA polymerase activity	
	mod pr	- negative regulation of cytoplasmic transfation - negative regulation of protein kinase activity by protein phosphorylation			idyltransferase activity	-DNA-directed DNA polymerase activity	
	cytoskeleton organization	-cortical microtubule organization	over_represented_pvalue  0.04 0.03 0.02 0.01		oxidoreductase pep activity ac	very-long-chain-(S)-2-hydroxy-acid oxidase activity medium-chain-(S)-2-hydroxy-acid oxidase activity long-chain-(S)-2-hydroxy-long-chain-acid oxidase activity	
	growth	-positive regulation of invasive growth in response to glucose limitation				glycolate oxidase activity  L-lactate dehydrogenase activity	represented_pvalue
	homeostatic process	- cellular cation homeostasis				aspartic-type endopeptidase activity  - 0.04 0.03 0.02 0.01	.04 .03 .02
	nmur yster	-mast cell activation				phosphatidylinositol-3,4,5-trisphosphate 5-phosphatase activity	
	locc	negative regulation of macrophage cytokine production			Ф	inositol-1,3,4,5-tetrakisphosphate 5-phosphatase activity	
		granulocyte chemotaxis			Se	inositol-1,4,5-trisphosphate 5-phosphatase activity	
	onse	positive regulation of cellular response to amino acid starvation cellular stress response to acidic pH ATF6-mediated unfolded protein response negative regulation of ATF6-mediated unfolded protein response			RNA	inositol-polyphosphate 5-phosphatase activity  translation elongation factor activity  selenocysteine insertion sequence binding	
	signal transductic	-interleukin–18–mediated signaling pathway				- single-stranded RNA binding	
	small molecule n metabolic process	ketone catabolic process -ketone body catabolic process -cellular ketone body metabolic process -keratan sulfate catabolic process -lactate oxidation -lipoxygenase pathway			structural molecule activity	-structural constituent of virion	
	symbiotic process	viral genome integration into host DNA establishment of integrated proviral latency viral entry into host cell suppression by virus of host gene expression virion assembly				NAADP-sensitive calcium-release channel activity	
	ranspor	fluid transport borate transport regulation of protein localization by the Cvt pathway			ansmembrane transporter activity	- active borate transmembrane transporter activity - bicarbonate transmembrane transporter activity - voltage–gated sodium channel activity - kainate selective glutamate receptor activity	
	transposition	-transposition, RNA-mediated				ligand-gated sodium channel activity  nucleic acid binding peptidase activity	
BP	n NA	post-transcriptional gene silencing by RNA larval feeding behavior. oxidative photosynthetic carbon pathway innate immune response-activating signal transduction cellular response to histidine amine metabolic process pharyngeal pumping cellular response to benomyl detection of virus		MF	NA	- DNA binding - RNA binding - 3-oxoacid CoA-transferase activity - cAMP receptor activity - 3-hydroxyisobutyryl-CoA hydrolase activity - pectin acetylesterase activity	