

GO: Day 7 WGCNA MEbrown		
term		term
protein serine/threonine kinase activity		regulation of GTPase activity
		protein ubiquitination
		signal transduction by protein phosphorylation
		protein phosphorylation
		activation of protein kinase activity
metal ion binding		receptor recycling
		intracellular signal transduction
ubiquitin-protein transferase activity		constitutive secretory pathway
		defense response to virus
		regulation of endocytic recycling
		negative regulation of filopodium assembly
		cytoskeleton organization
		filopodium assembly
		actin cytoskeleton organization
		actin polymerization-dependent cell motility
		peptidyl-threonine phosphorylation
		regulation of anion transport
		blood circulation
		fin development
		positive regulation of inflammatory response
		positive regulation of interferon-beta production
ubiquitin protein ligase activity		autophagosome assembly
		regulation of small GTPase mediated signal transduction
		undimensional cell growth
		response to braininjury
		cell migration
		post-embryonic hemopoiesis
		megakaryocyte development
GTPase activator activity		positive regulation of I-kappaB kinase/NF-kappaB signaling
		autophagy
		positive regulation of protein catabolic process
		positive regulation of interferon-alpha production
small GTPase binding		developmental pigmentation
		peptidyl-serine phosphorylation
		transferrin transport
GDP binding		definitive hemopoiesis
		development of secondary female sexual characteristics
		retrograde transport, endosome to plasma membrane
		positive regulation of interleukin-2 biosynthetic process
		positive regulation of receptor recycling
ribonuclease activity		barbed-end actin filament capping
		positive regulation of filopodium assembly
ubiquitin protein ligase binding		regulation of transcription elongation from RNA polymerase II promoter
		macrophage migration
		primitive hemopoiesis
		negative regulation of macroautophagy
		protein K63-linked deubiquitination
transferase activity		macrophagy
		positive regulation of release of cytochrome c from mitochondria
		macrophage chemotaxis
		negative regulation of apoptotic process
		lactation
cadherin binding		endocytic recycling
		activation of JUN kinase activity
		actin filament depolymerization
		innate immune response
		positive regulation of macroautophagy
		insulin secretion involved in cellular response to glucose stimulus
phosphoprotein phosphatase activity		compound eye photoreceptor cell differentiation
		fructose 6-phosphate metabolic process
		cell morphogenesis
		interferon-gamma-mediated signaling pathway
Rac guanyl-nucleotide exchange factor activity		cellular process
		regulation of transcription, DNA-templated
		BMP signaling pathway involved in heart development
		erythrocyte differentiation
		protein autophosphorylation
		immune system process
		immune system development
		viral process
structural constituent of nuclear pore		negative regulation of cysteine-type endopeptidase activity involved in execution phase of apoptosis
		positive regulation of phosphatidylinositol 3-kinase activity
		neutrophil migration
		poly(A)+ mRNA export from nucleus
		cheating during chimeric sorocarp development
		protein export from nucleus
		response to prostaglandin E1
		positive regulation of defense response to virus by host
		platelet activation
		C-terminal protein methylation
		cytoplasmic pattern recognition receptor signaling pathway in response to virus
		regulation of lamellipodium assembly
		embryonic hemopoiesis
		negative regulation of viral genome replication
		positive regulation of cristae formation
		cellular response to brain-derived neurotrophic factor stimulus
		mitral valve morphogenesis
		regulated exocytosis
		phosphatidylinositol-3-phosphate biosynthetic process
		response to virus
		ribosomal small subunit export from nucleus
		autophagy of mitochondrion
		phagolysosome assembly involved in apoptotic cell clearance
		regulation of female receptivity
		positive regulation of epithelial to mesenchymal transition involved in endocardial cushion formation
		positive regulation of determination of dorsal identity
		positive regulation of alkaline phosphatase activity
		endocardial cushion cell fate commitment
		cardiac muscle cell fate commitment
		immune response
		in utero embryonic development
		positive regulation of G0 to G1 transition
		T cell activation
		programmed cell death
		regulation of T cell anergy
		positive regulation of vascular endothelial cell proliferation
		ubiquitin-dependent protein catabolic process
		heterochromatin assembly by small RNA
		proline transmembrane transport
		amino-acid betaine transport
		collagen and cuticulin-based cuticle development
		cellular response to salt
		positive regulation of SMAD protein signal transduction
		regulation of protein export from nucleus
		regulation of cell cycle
		cell morphogenesis involved in neuron differentiation
		endosomal transport
		mitotic cytokinesis
		cellular response to acidic pH
		BMP signaling pathway
		positive regulation of type I interferon production
		germarium-derived oocyte fate determination
		mitotic cell cycle checkpoint
		regulation of protein complex stability
		cortical actin cytoskeleton organization
		negative regulation of interleukin-1 beta secretion
		branching involved in blood vessel morphogenesis
		regulation of centrosome duplication
		ephrin receptor signaling pathway
		regulation of protein localization to cell surface
		positive regulation of peptidyl-tyrosine phosphorylation
		negative regulation of motor neuron apoptotic process
		positive regulation of transcription by RNA polymerase II
		negative regulation of cell aging
		negative regulation of DNA replication
		negative regulation of substrate adhesion-dependent cell spreading
		positive regulation of synapse assembly
		protein deubiquitination involved in ubiquitin-dependent protein catabolic process
		inhibition of cysteine-type endopeptidase activity involved in apoptotic process
		regulation of fatty acid oxidation
		positive regulation of vascular smooth muscle cell proliferation
		regulation of lateral mesodermal cell fate specification
		type I interferon signaling pathway
		positive regulation of activin receptor signaling pathway
		regulation of mitochondrial membrane permeability
		homocysteine metabolic process
		ciliary receptor clustering involved in smoothened signaling pathway
		negative regulation of ATPase activity
		Golgi to plasma membrane transport
		vascular endothelial growth factor receptor signaling pathway
		activation of protein kinase B activity
		negative regulation of cardiac muscle cell apoptotic process
		lamellipodium assembly
		execution phase of apoptosis
		protein autoubiquitination
		positive regulation by host of viral transcription
		activation of cysteine-type endopeptidase activity involved in apoptotic process
		cholesterol transport
		ubiquitin-dependent protein catabolic process via the N-end rule pathway
		negative regulation of protein kinase activity by regulation of protein phosphorylation
		apical protein localization
		response to ischemia
		negative regulation of I-kappaB kinase/NF-kappaB signaling
		positive regulation of cardiac muscle cell proliferation
		defense response
		cellular response to virus
		proline transport
		brain development
		regulation of synaptic vesicle endocytosis
		miRNA catabolic process
		I-kappaB kinase/NF-kappaB signaling
		positive regulation of hepatocyte proliferation
		negative regulation of potassium ion transmembrane transporter activity
		response to pheromone
		imaginal disc-derived wing vein specification
		T cell receptor signaling pathway
		negative regulation of G1/S transition of mitotic cell cycle
		negative regulation of signal transduction
		negative regulation of cell size
		neutrophil chemotaxis
		cellular response to arsenic-containing substance
		establishment of protein localization to membrane
		ventral midline development
		negative regulation of interleukin-6 production
		actomyosin structure organization
		growth hormone receptor signaling pathway via JAK-STAT
		pathway-restricted SMAD protein phosphorylation
		platelet-derived growth factor receptor signaling pathway
		regulation of protein phosphorylation
		positive regulation of centriole elongation
		negative regulation of extrinsic apoptotic signaling pathway via death domain receptors
		positive regulation of leukocyte adhesion to vascular endothelial cell
		maintenance of apical/basal cell polarity
		skeletal muscle tissue regeneration
		mitotic G2 DNA damage checkpoint
		ribosomal protein processing
		response to mitochondrial depolarisation
		necroptotic process
		negative regulation of cysteine-type endopeptidase activity involved in apoptotic process
		response to benzene
		positive regulation of Schwann cell differentiation
		memory
		response to exogenous dsRNA
		apoptotic process
		positive regulation of protein kinase B signaling
		phosphatidylinositol phosphorylation
		regulation of lipid metabolic process
		cardiac right ventricle morphogenesis
		cytokine metabolic process
		Golgi to vacuole transport
		mitotic cell cycle
		peptidyl-glutamate ADP-deribosylation
		negative regulation of inflammatory response
		patched ligand maturation
		renal system process
		positive regulation of RIG-I signaling pathway
		activation of innate immune response
		regulation of proton-transporting ATPase activity, rotational mechanism
		regulation of necrotic cell death
		negative regulation of oxidative phosphorylation uncoupler activity
		mitochondrial outer membrane permeabilization involved in programmed cell death
		growing angiogenesis
		N-acylethanolamine metabolic process
		tolerance induction to lipopolysaccharide
		negative regulation of toll-like receptor 5 signaling pathway
		negative regulation of nucleotide-binding oligomerization domain containing 1 signaling pathway
		negative regulation of granuloma formation
		negative regulation of chronic inflammatory response
		negative regulation of CD40 signaling pathway
		negative regulation of B cell activation
		B-1 B cell homeostasis
		liver development
		regulation of type III interferon production
		regulation of steroid metabolic process
		cellular response to steroid hormone stimulus
		endonucleolytic cleavage to generate mature 5'-end of SSU-rRNA from (SSU-rRNA, 5.8S rRNA, LSU-rRNA)
		endonucleolytic cleavage in 5'-ETS of tricistronic rRNA transcript (SSU-rRNA, 5.8S rRNA, LSU-rRNA)
		severless signaling pathway
		protein localization to ciliary membrane
		melanosome transport
		synaptic growth at neuromuscular junction
		cellular response to exogenous dsRNA
		negative regulation of apoptotic process in bone marrow cell
		regulation of mitochondrial ATP synthesis coupled proton transport
		positive regulation of proton-transporting ATP synthase activity, rotational mechanism
		nucleotide-excision repair, preincision complex assembly
		positive regulation of mRNA catabolic process
		cellular response to chemokine
		negative regulation of cell motility
		de novo actin filament nucleation
		membrane invagination
		RNA processing
		Fc-epsilon receptor signaling pathway
		regulation of DNA repair
		taurine biosynthetic process
		L-cysteine catabolic process to hypotaurine
		L-cysteine catabolic process to hypotaurine
		positive regulation of hydrogen sulfide biosynthetic process
		megakaryocyte differentiation
		regulation of cell shape
		intracellular protein transport
		regulation of adaptive immune response
		positive regulation of regulatory T cell differentiation
		negative regulation of respiratory burst involved in inflammatory response
		negative regulation of epithelium regeneration
		mitochondrial ATP synthesis coupled electron transport
		organic substance transport
		regulation of protein kinase C signaling
		cellular response to peptidoglycan
		cellular response to epidermal growth factor stimulus
		regulation of histone methylation
		heterochromatin maintenance
		negative regulation of oxidative stress-induced neuron death
		positive regulation of bone mineralization
		negative regulation of gene expression
		protein de-ADP-ribosylation
		cell-cell adhesion via plasma-membrane adhesion molecules
		enucleate erythrocyte differentiation
		glutathione transmembrane transport
		nucleotide-sugar metabolic process
		negative regulation of transcription by RNA polymerase II
		low-density lipoprotein particle receptor catabolic process
		regulation of oxygen metabolic process
		nucleotide-binding domain, leucine rich repeat containing receptor signaling pathway
		negative regulation of signal transduction by p53 class mediator
		protein O-linked glycosylation
		negative regulation of macrophage activation
		meiotic cytokinesis
		meiotic chromosome movement towards spindle pole
		RIG-I signaling pathway
		positive regulation of granulocyte macrophage colony-stimulating factor production
		antiviral innate immune response