GO Term Enrichment − Cluster 3
actin filament organization
supramolecular fiber organization
regulation of protein modification process
locomotor rhythm
positive regulation of proteolysis
torso signaling pathway
establishment of cell polarity
regulation of filopodium assembly
segment specification
cell division
endosomal transport
response to ketone
telomere organization
response to organic cyclic compound
telomere maintenance
regulation of protein catabolic process
positive regulation of protein catabolic process
regulation of protein−containing complex assembly
innate immune response
neuroblast division
regulation of cellular localization
selective autophagy
regulation of Notch signaling pathway
negative regulation of cell motility
negative regulation of cell migration
habituation
establishment of monopolar cell polarity
establishment or maintenance of monopolar cell polarity
salivary gland cell autophagic cell death
vacuolar transport
pigment granule organization
cellular pigmentation
mRNA cis splicing, via spliceosome
protein monoubiquitination
positive regulation of organ growth
positive regulation of histone methylation
myosin filament organization
anterograde axonal transport
eye pigment granule organization
viral process
positive regulation of Ras protein signal transduction
positive regulation of small GTPase mediated signal transduction
positive regulation of immune response
positive regulation of translation
regulation of actin filament length
positive regulation of response to biotic stimulus
protein targeting to vacuole
regulation of protein polymerization
macromolecule glycosylation
protein glycosylation
regulation of protein localization
autophagic cell death
endosome organization
intercellular bridge organization
peptidyl−lysine trimethylation
actin polymerization or depolymerization
regulation of innate immune response
negative regulation of receptor signaling pathway via JAK−STAT
negative regulation of receptor signaling pathway via STAT
cytoskeleton−dependent intracellular transport
cellular response to hypoxia
regulation of stem cell differentiation
glycosylation
positive regulation of defense response
gene silencing by RNA
positive regulation of antibacterial peptide production
positive regulation of antibacterial peptide biosynthetic process
muscle cell cellular homeostasis
ovarian nurse cell to oocyte transport
regulation of proteolysis
response to nutrient levels
programmed cell death involved in cell development
chitin−based larval cuticle pattern formation
positive regulation of antimicrobial peptide biosynthetic process
salivary gland histolysis
cellular response to decreased oxygen levels
post−transcriptional gene silencing
positive regulation of protein modification process
receptor signaling pathway via STAT
cellular response to oxygen levels
receptor signaling pathway via JAK−STAT
compound eye pigmentation
germ−line cyst encapsulation
morphogenesis of follicular epithelium
positive regulation of proteasomal protein catabolic process
regulation of miRNA−mediated gene silencing
germarium−derived female germ−line cyst encapsulation
germ−band extension
epithelial cell−cell adhesion
female germ−line cyst encapsulation
establishment of spindle localization
spindle localization
cellularization
asymmetric neuroblast division
negative regulation of cellular response to growth factor stimulus
negative regulation of stem cell differentiation
positive regulation of immune system process
lymph gland development
vascular endothelial growth factor receptor signaling pathway
viral life cycle
negative regulation of cell size
response to ecdysone
response to sterol
regulation of proteasomal protein catabolic process
somatic stem cell population maintenance
germ cell migration
apical protein localization
asymmetric protein localization involved in cell fate determination
positive regulation of proteasomal ubiquitin−dependent protein catabolic process
positive regulation of axonogenesis
vesicle budding from membrane
regulation of histone methylation
regulation of ubiquitin−dependent protein catabolic process
positive regulation of ubiquitin−dependent protein catabolic process
larval somatic muscle development
regulation of histone modification
negative regulation of neuron projection development
nonassociative learning
lysosomal transport
regulation of supramolecular fiber organization
regulation of proteolysis involved in protein catabolic process
negative regulation of cell projection organization
regulation of proteasomal ubiquitin−dependent protein catabolic process
positive regulation of proteolysis involved in protein catabolic process
establishment of spindle orientation
axon extension involved in axon guidance
neuron projection extension involved in neuron projection guidance
establishment of organelle localization
mRNA modification
positive regulation of protein metabolic process
organelle localization
positive regulation of cellular amide metabolic process
vesicle organization
endosome to lysosome transport
regulation of cell growth
SNARE complex assembly
negative regulation of locomotion
regulation of post−transcriptional gene silencing by RNA
regulation of gene silencing by RNA
histone ubiquitination
negative regulation of axonogenesis
histone H3−K4 methylation
negative regulation of cell growth
regulation of hydrolase activity
regulation of ERBB signaling pathway
regulation of epidermal growth factor receptor signaling pathway
positive regulation of catabolic process
positive regulation of cellular catabolic process
regulation of catabolic process
regulation of cellular catabolic process
regulation of axon extension
regulation of extent of cell growth
axon extension
neuron projection extension
long−term memory
regulation of axonogenesis
negative regulation of cell differentiation
negative regulation of growth
negative regulation of nervous system development
cell growth
negative regulation of response to stimulus
regulation of Wnt signaling pathway
regulation of canonical Wnt signaling pathway
regulation of cellular component size
regulation of cell size
negative regulation of developmental growth
regulation of GTPase activity
organ growth
regulation of organ growth
positive regulation of molecular function
positive regulation of GTPase activity
negative regulation of organ growth
ERBB signaling pathway
epidermal growth factor receptor signaling pathway
negative regulation of ERBB signaling pathway
negative regulation of epidermal growth factor receptor signaling pathway
peptidyl−tyrosine modification
peptidyl−tyrosine phosphorylation
positive regulation of hydrolase activity
positive regulation of catalytic activity
regulation of catalytic activity
regulation of molecular function
dorsal/ventral pattern formation
maintenance of cell number
stem cell population maintenance
negative regulation of canonical Wnt signaling pathway
negative regulation of Wnt signaling pathway
negative regulation of cellular component organization
dorsal/ventral pattern formation, imaginal disc
gastrulation
unidimensional cell growth
ovarian follicle cell migration
wing disc dorsal/ventral pattern formation
regulation of neural precursor cell proliferation
positive regulation of cell development
gastrulation involving germ band extension
gastrulation with mouth forming first
positive regulation of neurogenesis
gland morphogenesis
salivary gland morphogenesis
cellular response to growth factor stimulus
response to growth factor
head involution
regulation of chemotaxis
border follicle cell migration
positive regulation of cell projection organization
regulation of Ras protein signal transduction
regulation of neuroblast proliferation
peptidyl−serine phosphorylation
positive regulation of programmed cell death
regulation of actin polymerization or depolymerization
positive regulation of response to external stimulus
regulation of stem cell proliferation
Notch signaling pathway
programmed cell death
cell death
regulation of cellular response to growth factor stimulus
regulation of actin filament organization
regulation of actin cytoskeleton organization
stem cell proliferation
negative regulation of neuromuscular junction development
negative regulation of synaptic assembly at neuromuscular junction
negative regulation of cell junction assembly
negative regulation of synapse assembly
negative regulation of synapse organization
positive regulation of cell differentiation
regulation of actin filament−based process
imaginal disc−derived wing vein morphogenesis
homeostasis of number of cells
intestinal stem cell homeostasis
negative regulation of cell adhesion
regulation of cytoskeleton organization
positive regulation of multicellular organismal process
regulation of cell junction assembly
regulation of synaptic assembly at neuromuscular junction
adherens junction organization
regulation of synapse assembly
regulation of neuromuscular junction development
regulation of synapse organization
larval development
regulation of small GTPase mediated signal transduction
regulation of neurogenesis
negative regulation of cell communication
negative regulation of signaling
negative regulation of signal transduction
regulation of cell development
Ras protein signal transduction
adherens junction assembly
small GTPase mediated signal transduction
Cluster_5
Cluster_27
Cluster_28
Cluster_20
Cluster_29
Cluster_7
Cluster_2
Cluster_8
Cluster_25
Cluster_24
Cluster_13
Cluster_17
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Cluster_22
Cluster_23
Cluster_14
Cluster_16
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