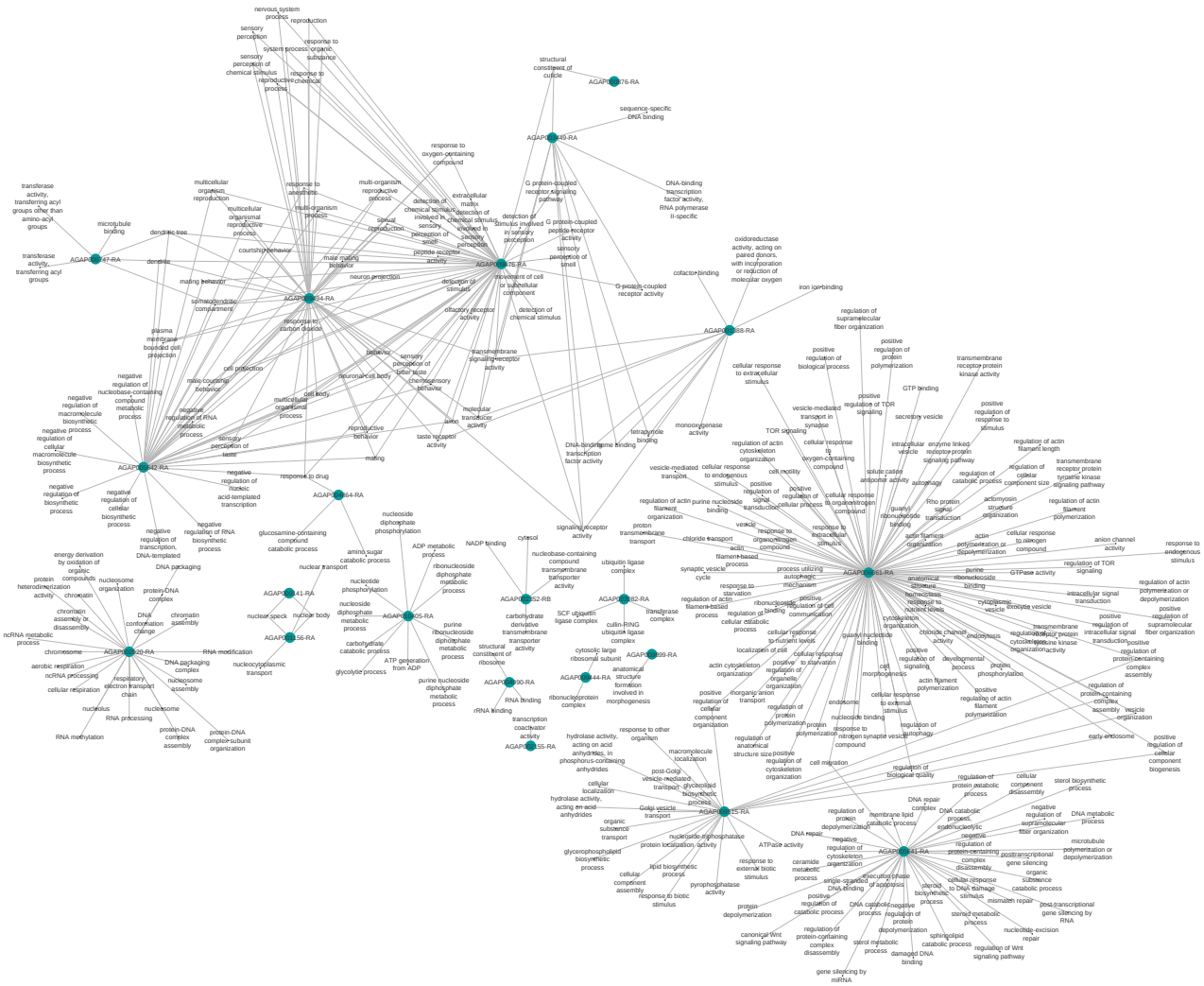
nervous system process
reproduction
sensory perception
response to stimulus
system process
organic substance
sensory perception of chemical stimulus
response to chemical
reproductive process
structural constituent of cuticle
AGAP010376-RA
sequence-specific DNA binding
AGAP010349-RA
response to oxygen-containing compound
G protein-coupled receptor signaling pathway
DNA-binding transcription factor activity, RNA polymerase II-specific
transferase activity, transferring acyl groups other than amino-acyl groups
multicellular organism reproduction
multi-organism reproductive process
response to stimulus
multicellular organismal reproductive process
multi-organism process
detection of chemical stimulus involved in sensory perception
extracellular matrix
detection of chemical stimulus
detection of chemical stimulus involved in sensory perception
detection of stimulus involved in sensory perception
microtubule binding
dendritic tree
sexual reproduction
courtship behavior
peptide receptor activity
G protein-coupled peptide receptor activity
sensory perception of smell
AGAP010347-RA
dendrite
male mating behavior
AGAP010376-RA
movement of cell or subcellular component
G protein-coupled receptor activity
cofactor binding
oxidoreductase activity, acting on paired donors, with incorporation or reduction of molecular oxygen
transferase activity, transferring acyl groups
mating behavior
neuron projection
detection of stimulus
iron ion binding
AGAP010354-RA
olfactory receptor activity
detection of chemical stimulus
regulation of supramolecular fiber organization
somatodendritic compartment
AGAP010388-RA
plasma membrane bounded cell projection
behavior
sensory perception of bitter taste
transmembrane signaling receptor activity
positive regulation of biological process
positive regulation of protein polymerization
transmembrane receptor protein kinase activity
cellular response to extracellular stimulus
cell projection
male courtship behavior
neuronal cell body
chemosensory behavior
GTP binding
positive regulation of TOR signaling
vesicle-mediated transport in synapse
secretory vesicle
positive regulation of response to stimulus
negative regulation of macromolecule biosynthetic process
negative regulation of nucleobase-containing compound metabolic process
cell body
multicellular organismal process
molecular transducer activity
TOR signaling
monooxygenase activity
tetrapyrrole binding
intracellular vesicle
enzyme linked receptor protein signaling pathway
regulation of actin filament length
negative process
negative regulation of RNA metabolic process
reproductive behavior
taste receptor activity
DNA-binding transcription factor activity
regulation of actin cytoskeleton organization
cellular response to oxygen-containing compound
solute carrier antiporter activity
regulation of catabolic process
regulation of cellular component size
transmembrane receptor protein tyrosine kinase signaling pathway
negative regulation of cellular macromolecule biosynthetic process
sensory perception of taste
mating
AGAP010342-RA
response to drug
vesicle-mediated transport
cellular response to endogenous stimulus
cell motility
positive regulation of signal transduction
positive regulation of cellular process
cellular response to organonitrogen compound
autophagy
actomyosin structure organization
regulation of actin filament polymerization
negative regulation of biosynthetic process
negative regulation of cellular biosynthetic process
negative regulation of toxic acid-templated transcription
AGAP010364-RA
regulation of actin filament organization
purine nucleoside binding
Rho protein signal transduction
guanyl ribonucleotide binding
nucleoside diphosphate phosphorylation
proton transmembrane transport
signaling receptor activity
vesicle
response to organonitrogen compound
response to extracellular stimulus
actin filament organization
actin polymerization or depolymerization
cellular response to nitrogen compound
anion channel activity
response to endogenous stimulus
negative regulation of transcription, DNA-templated
negative regulation of RNA biosynthetic process
glucosamine-containing compound catabolic process
cytosol
NADP binding
chloride transport
actin filament-based process
synaptic vesicle cycle
process utilizing autophagic mechanism
AGAP010361-RA
regulation of TOR signaling
GTPase activity
DNA packaging
amino sugar catabolic process
ADP metabolic process
nucleobase-containing compound transmembrane transporter activity
ubiquitin ligase complex
response to starvation
regulation of actin filament-based process
ribonucleoside binding
positive regulation of cell communication
analogical structure homeostasis
purine ribonucleoside binding
intracellular signal transduction
regulation of actin polymerization or depolymerization
energy derivation by oxidation of organic compounds
nucleosome organization
nuclear transport
nucleotide phosphorylation
AGAP010341-RA
AGAP010352-RB
AGAP010362-RA
response to nutrient levels
regulation of cellular catabolic process
cytoplasmic vesicle
endocytic vesicle
protein heterodimerization activity
chromatin
protein-DNA complex
nucleoside diphosphate metabolic process
carbohydrate derivative transmembrane transporter activity
transferase complex
cytokinesis organization
positive regulation of intracellular signal transduction
regulation of supramolecular fiber organization
chromatin assembly or disassembly
DNA conformation change
chromatin assembly
nuclear speck
nuclear body
AGAP010305-RA
purine ribonucleoside diphosphate metabolic process
SCF ubiquitin ligase complex
cullin-RING ubiquitin ligase complex
localization of cell
cellular response to nutrient levels
chloride channel activity
endocytosis
transmembrane receptor protein tyrosine kinase activity
ncRNA metabolic process
AGAP010356-RA
carbohydrate catabolic process
structural constituent of ribosome
cytosolic large ribosomal subunit
AGAP010309-RA
positive regulation of organelle organization
cellular response to starvation
positive regulation of protein-containing complex assembly
chromosome
AGAP010320-RA
RNA modification
glycolytic process
ATP generation from ADP
AGAP010344-RA
anatomical structure formation involved in morphogenesis
actin cytoskeleton organization
developmental process
protein phosphorylation
aerobic respiration
DNA packaging complex
nucleocytoplasmic transport
AGAP010330-RA
ribonucleoprotein complex
inorganic anion transport
actin filament polymerization
positive regulation of actin filament polymerization
ncRNA processing
respiratory electron transport chain
nucleosome assembly
rRNA binding
RNA binding
transcription coactivator activity
positive regulation of cellular component organization
regulation of protein polymerization
endosome
cellular response to external stimulus
nucleoside binding
early endosome
vesicle organization
cellular respiration
nucleolus
nucleosome
RNA processing
AGAP010355-RA
hydrolase activity, acting on acid anhydrides, in phosphorus-containing anhydrides
response to other organism
macromolecule localization
protein polymerization
response to nitrogen-containing compound
regulation of anatomical structure size
positive regulation of cytoskeleton organization
cell migration
positive regulation of cellular component biogenesis
RNA methylation
protein-DNA complex assembly
protein-DNA complex subunit organization
post-Golgi vesicle-mediated transport
glycerolipid biosynthetic process
regulation of biological quality
cellular localization
hydrolase activity, acting on acid anhydrides
Golgi vesicle transport
AGAP010315-RA
DNA repair complex
regulation of protein catabolic process
cellular component disassembly
sterol biosynthetic process
membrane lipid catabolic process
DNA catabolic process, exodeoxyribonucleolytic
negative regulation of supramolecular fiber organization
DNA metabolic process
organic substance transport
nucleoside-triphosphatase activity
protein localization
regulation of protein depolymerization
DNA repair
ATPase activity
negative regulation of cytoskeleton organization
AGAP010341-RA
negative regulation of protein-containing complex disassembly
microtubule polymerization or depolymerization
glycerophospholipid biosynthetic process
lipid biosynthetic process
response to external biotic stimulus
ceramide metabolic process
execution phase of apoptosis
postranscriptional gene silencing
organic substance catabolic process
cellular component assembly
response to biotic stimulus
pyrophosphatase activity
single-stranded DNA binding
cellular response to DNA damage stimulus
sterol metabolic process
positive regulation of catabolic process
DNA catabolic process
negative regulation of protein depolymerization
mismatch repair
post-transcriptional gene silencing by RNA
protein depolymerization
sterol metabolic process
nucleotide-excision repair
canonical Wnt signaling pathway
regulation of protein-containing complex disassembly
sphingolipid catabolic process
damaged DNA binding
regulation of Wnt signaling pathway
gene silencing by miRNA