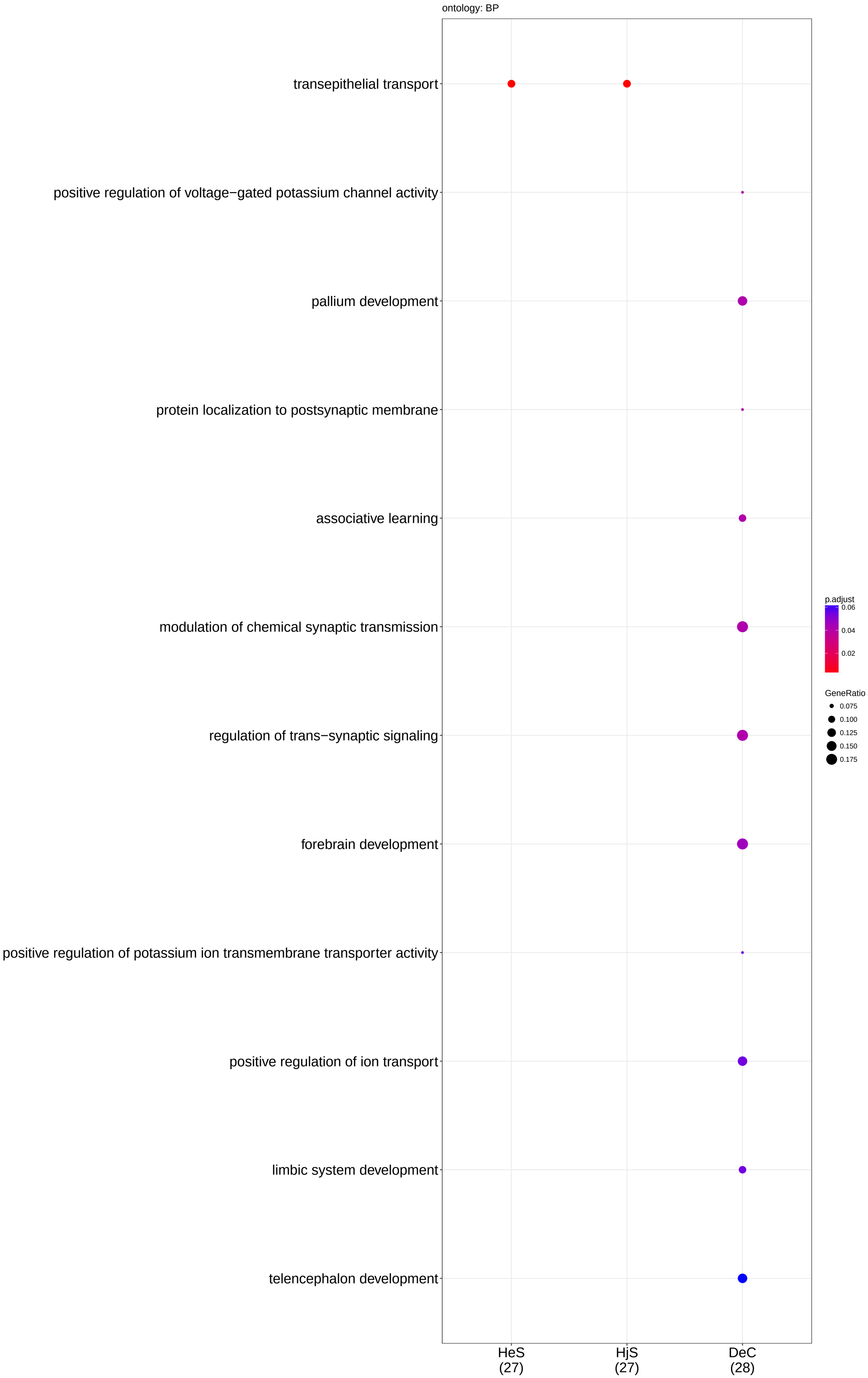
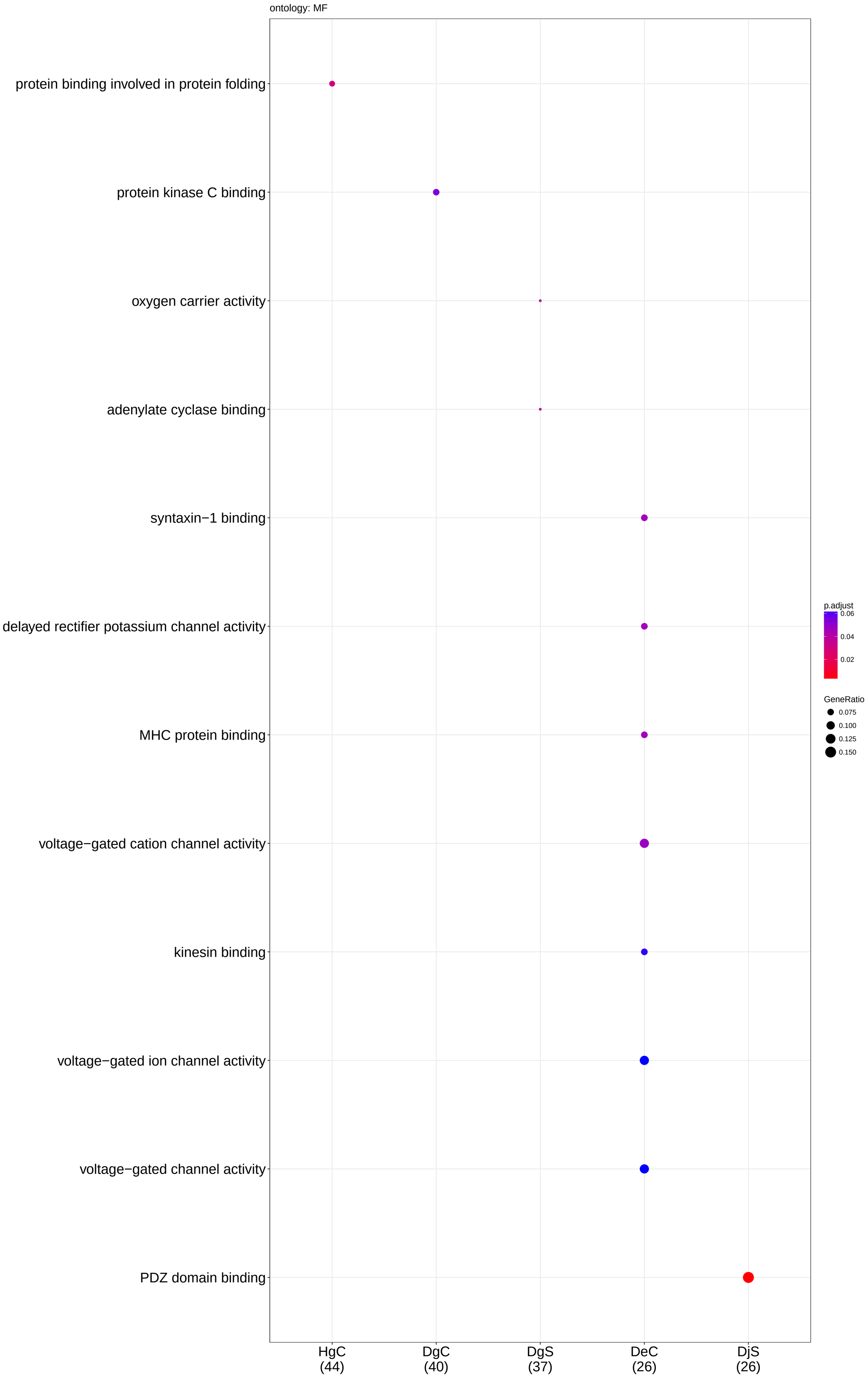
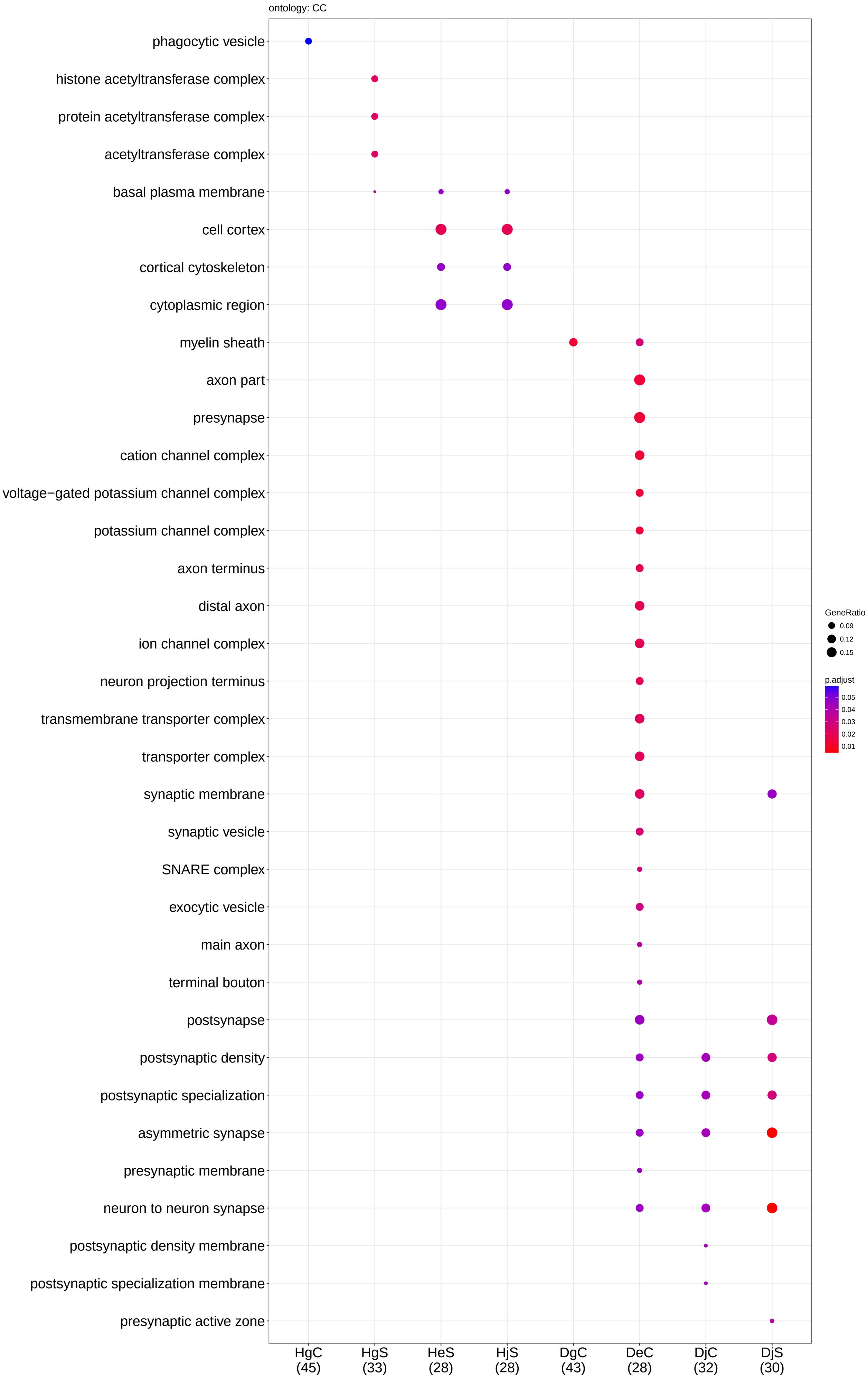
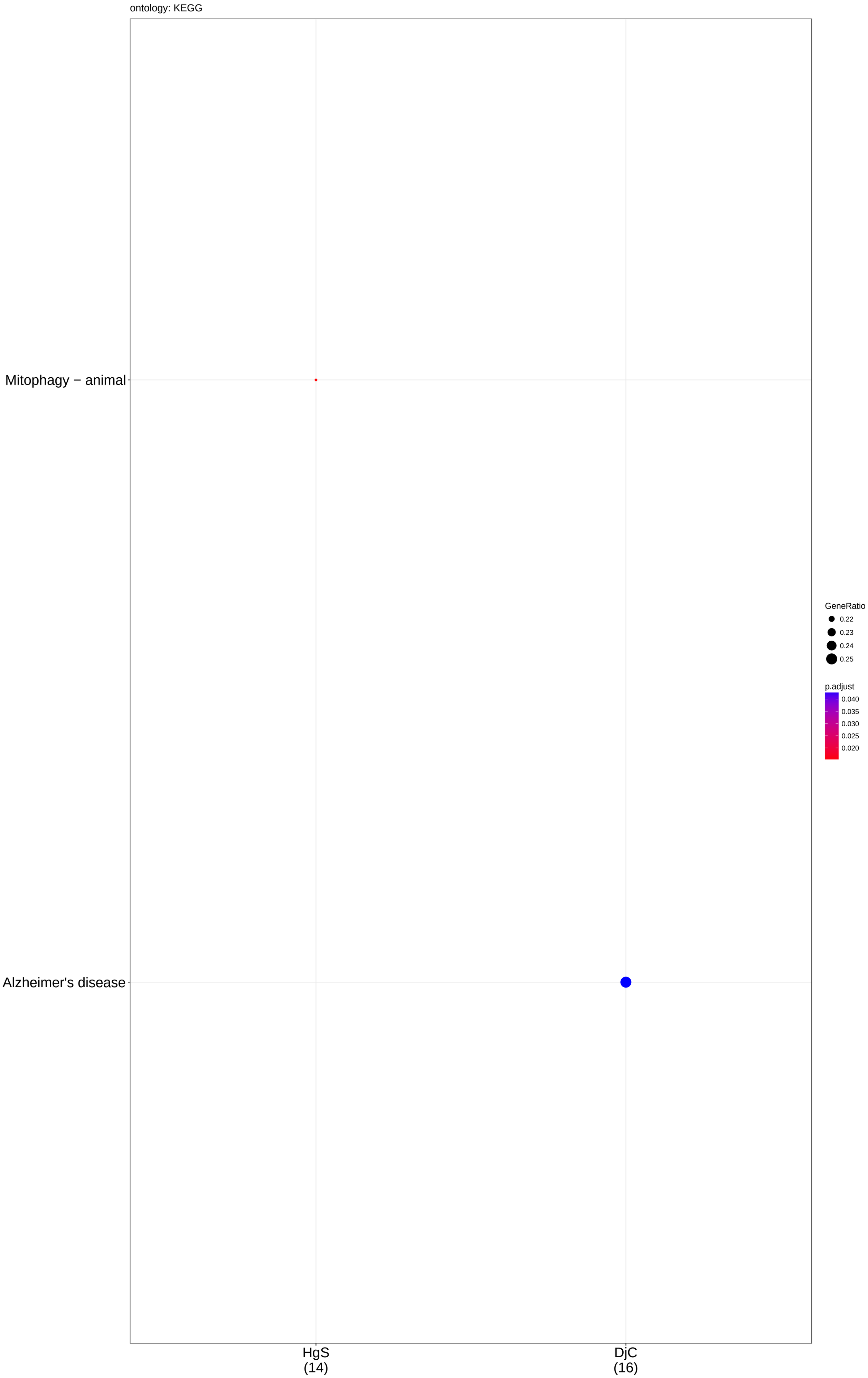

ontology: BP
transepithelial transport
positive regulation of voltage−gated potassium channel activity
pallium development
protein localization to postsynaptic membrane
associative learning
modulation of chemical synaptic transmission
regulation of trans−synaptic signaling
forebrain development
positive regulation of potassium ion transmembrane transporter activity
positive regulation of ion transport
limbic system development
telencephalon development
HeS (27)
HjS (27)
DeC (28)
p.adjust
0.06
0.04
0.02
GeneRatio
0.075
0.100
0.125
0.150
0.175


ontology: MF
protein binding involved in protein folding
protein kinase C binding
oxygen carrier activity
adenylate cyclase binding
syntaxin−1 binding
delayed rectifier potassium channel activity
MHC protein binding
voltage−gated cation channel activity
kinesin binding
voltage−gated ion channel activity
voltage−gated channel activity
PDZ domain binding
HgC (44)
DgC (40)
DgS (37)
DeC (26)
DjS (26)
p.adjust
0.06
0.04
0.02
GeneRatio
0.075
0.100
0.125
0.150


ontology: CC
phagocytic vesicle
histone acetyltransferase complex
protein acetyltransferase complex
acetyltransferase complex
basal plasma membrane
cell cortex
cortical cytoskeleton
cytoplasmic region
myelin sheath
axon part
presynapse
cation channel complex
voltage−gated potassium channel complex
potassium channel complex
axon terminus
distal axon
ion channel complex
neuron projection terminus
transmembrane transporter complex
transporter complex
synaptic membrane
synaptic vesicle
SNARE complex
exocytic vesicle
main axon
terminal bouton
postsynapse
postsynaptic density
postsynaptic specialization
asymmetric synapse
presynaptic membrane
neuron to neuron synapse
postsynaptic density membrane
postsynaptic specialization membrane
presynaptic active zone
HgC (45)
HgS (33)
HeS (28)
HjS (28)
DgC (43)
DeC (28)
DjC (32)
DjS (30)
GeneRatio
0.09
0.12
0.15
p.adjust
0.05
0.04
0.03
0.02
0.01


ontology: KEGG
Mitophagy − animal
Alzheimer's disease
HgS
(14)
DjC
(16)
GeneRatio
0.22
0.23
0.24
0.25
p.adjust
0.040
0.035
0.030
0.025
0.020