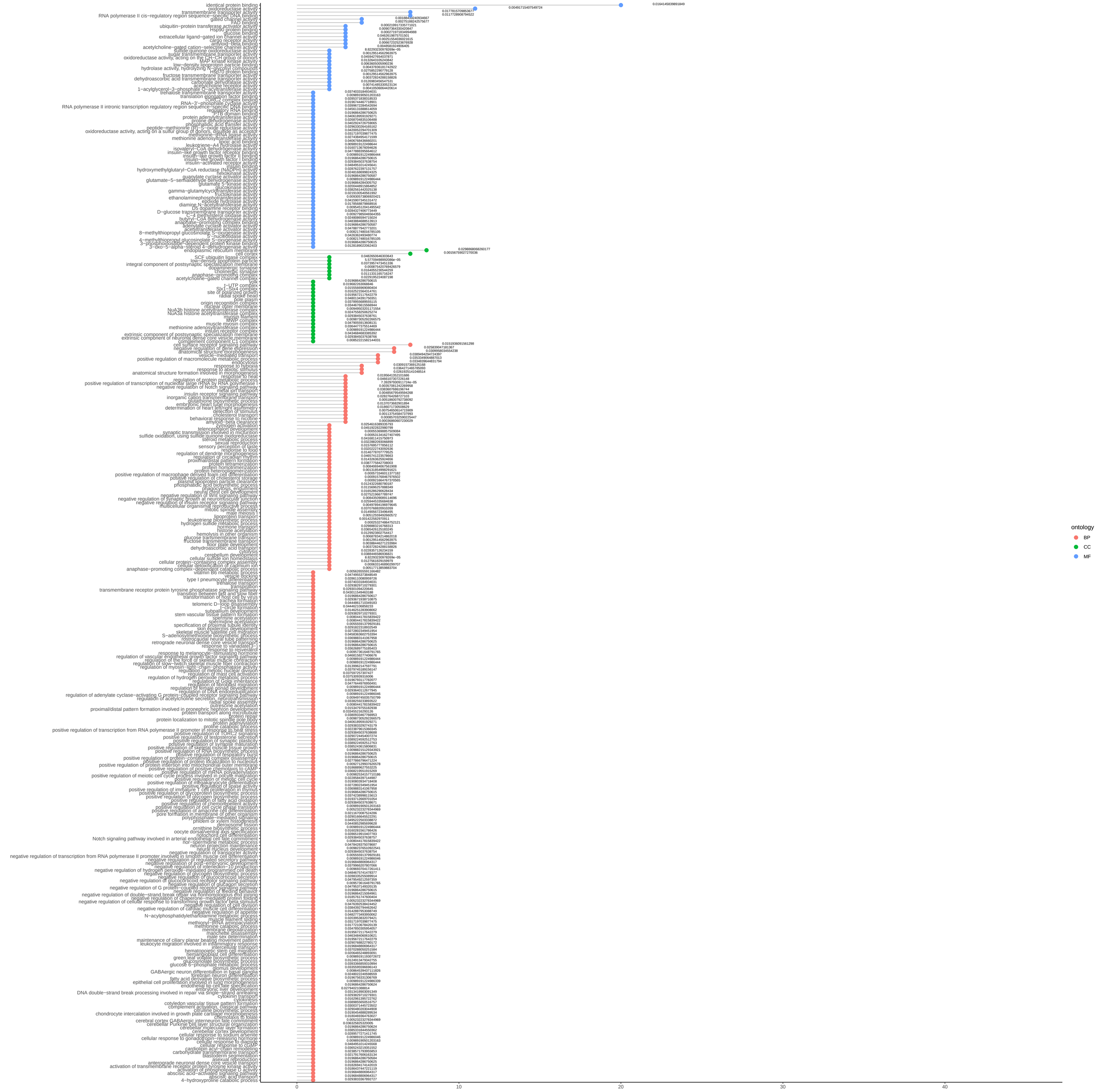
identical protein binding
oxidoreductase activity
transmembrane transporter activity
RNA polymerase II cis−regulatory region sequence−specific DNA binding
gated channel activity
FAD binding
ubiquitin−protein transferase activator activity
Hsp90 protein binding
glucose binding
extracellular ligand−gated ion channel activity
cargo receptor activity
amyloid−beta binding
acetylcholine−gated cation−selective channel activity
sulfide:quinone oxidoreductase activity
sugar transmembrane transporter activity
oxidoreductase activity, acting on the CH−CH group of donors
MAP kinase kinase activity
low−density lipoprotein particle binding
hydrolase activity, hydrolyzing N−glycosyl compounds
Hsp70 protein binding
fructose transmembrane transporter activity
dehydroascorbic acid transmembrane transporter activity
carbonate dehydratase activity
acetylcholine receptor activity
1−acylglycerol−3−phosphate O−acyltransferase activity
trehalose transmembrane transporter activity
translation elongation factor binding
TORC2 complex binding
RNA−3'−phosphate cyclase activity
RNA polymerase II intronic transcription regulatory region sequence−specific DNA binding
regulatory RNA binding
PTB domain binding
protein adenylyltransferase activity
proline dehydrogenase activity
phosphatidic acid transfer activity
peptide−methionine (R)−S−oxide reductase activity
oxidoreductase activity, acting on a sulfur group of donors, disulfide as acceptor
methionine−tRNA ligase activity
methionine adenosyltransferase activity
lipoic acid binding
leukotriene−A4 hydrolase activity
isovaleryl−CoA dehydrogenase activity
insulin−like growth factor receptor binding
insulin−like growth factor II binding
insulin−like growth factor I binding
insulin−activated receptor activity
insulin binding
hydroxymethylglutaryl−CoA reductase (NADPH) activity
hexokinase activity
guanylate cyclase activator activity
glutamate−5−semialdehyde dehydrogenase activity
glutamate 5−kinase activity
glucokinase activity
gamma−glutamylcyclotransferase activity
fructokinase activity
ethanolaminephosphotransferase activity
epoxide hydrolase activity
diamine N−acetyltransferase activity
D5 dopamine receptor binding
D−glucose transmembrane transporter activity
C−4 methylsterol oxidase activity
butyryl−CoA dehydrogenase activity
anaphase−promoting complex binding
adenylate cyclase activator activity
acetyltransferase activator activity
8−methylthiopropyl glucosinolate S−oxygenase activity
5'−nucleotidase activity
4−methylthiopropyl glucosinolate S−oxygenase activity
3−phosphoinositide−dependent protein kinase binding
3−oxo−5−alpha−steroid 4−dehydrogenase activity
endoplasmic reticulum membrane
cell cortex
SCF ubiquitin ligase complex
low−density lipoprotein particle
integral component of postsynaptic specialization membrane
dopaminergic synapse
cholinergic synapse
anaphase−promoting complex
acetylcholine−gated channel complex
Vpk
t−UTP complex
Slx1−Slx4 complex
site of polarized growth
radial spoke head
pole plasm
origin recognition complex
nuclear outer membrane
NuA3b histone acetyltransferase complex
NuA3a histone acetyltransferase complex
myosin filament
MWP complex
muscle myosin complex
methionine adenosyltransferase complex
insulin receptor complex
extrinsic component of postsynaptic specialization membrane
extrinsic component of neuronal dense core vesicle membrane
complement component C1 complex
cell surface receptor signaling pathway
negative regulation of gene expression
anatomical structure morphogenesis
vesicle−mediated transport
positive regulation of macromolecule metabolic process
endocytosis
response to hypoxia
response to abiotic stimulus
anatomical structure formation involved in morphogenesis
response to heat
regulation of protein metabolic process
positive regulation of transcription of nucleolar large rRNA by RNA polymerase I
negative regulation of Notch signaling pathway
metal ion transport
insulin receptor signaling pathway
inorganic cation transmembrane transport
glutathione biosynthetic process
embryonic heart tube morphogenesis
determination of heart left/right asymmetry
detection of stimulus
cholesterol transport
behavioral response to nicotine
amyloid−beta clearance
zymogen activation
telencephalon development
synaptic transmission involved in micturition
sulfide oxidation, using sulfide:quinone oxidoreductase
steroid metabolic process
sexual reproduction
sensory perception of taste
response to food
regulation of dendrite morphogenesis
regulation of circadian rhythm
proximal/distal pattern formation
protein tetramerization
protein homotrimerization
protein heteroligomerization
positive regulation of macrophage derived foam cell differentiation
positive regulation of cholesterol storage
plasma lipoprotein particle clearance
phosphatidic acid biosynthetic process
phagocytosis, engulfment
neural crest cell development
negative regulation of Wnt signaling pathway
negative regulation of synaptic growth at neuromuscular junction
negative regulation of insulin receptor signaling pathway
multicellular organismal reproductive process
mitotic spindle assembly
male meiosis I
lipoprotein transport
leukotriene biosynthetic process
hydrogen sulfide metabolic process
hormone transport
histone acetylation
hemolysis in other organism
glucose transmembrane transport
fructose transmembrane transport
floor plate development
dehydroascorbic acid transport
cytolysis
cerebellum development
cellular sulfide ion homeostasis
cellular protein−containing complex assembly
cellular detoxification of cadmium ion
anaphase−promoting complex−dependent catabolic process
vitamin B6 metabolic process
vesicle docking
type I pneumocyte differentiation
trehalose transport
transpiration
transmembrane receptor protein tyrosine phosphatase signaling pathway
transition between fast and slow fiber
transformation of host cell by virus
trachea formation
telomeric D−loop disassembly
t−circle formation
subpallium development
stem vascular tissue pattern formation
spermine acetylation
spermidine acetylation
specification of proximal tubule identity
skin epidermis development
skeletal muscle satellite cell migration
S−adenosylmethionine biosynthetic process
rostrocaudal neural tube patterning
retrograde neuronal dense core vesicle transport
response to vanadate(5−)
response to resveratrol
response to melanocyte−stimulating hormone
regulation of vascular endothelial growth factor signaling pathway
regulation of the force of skeletal muscle contraction
regulation of slow−twitch skeletal muscle fiber contraction
regulation of myosin−light−chain−phosphatase activity
regulation of meiotic nuclear division
regulation of mast cell activation
regulation of hydrogen peroxide metabolic process
regulation of Golgi inheritance
regulation of fibroblast migration
regulation of female gonad development
regulation of DNA endoreduplication
regulation of adenylate cyclase−activating G protein−coupled receptor signaling pathway
regulation of acetylcholine secretion, neurotransmission
radial spoke assembly
putrescine acetylation
proximal/distal pattern formation involved in pronephric nephron development
protein transport along microtubule
protein repair
protein localization to mitotic spindle pole body
protein adenylylation
proline catabolic process
positive regulation of transcription from RNA polymerase II promoter in response to heat stress
positive regulation of TORC2 signaling
positive regulation of testosterone secretion
positive regulation of synaptic plasticity
positive regulation of synapse maturation
positive regulation of skeletal muscle tissue growth
positive regulation of RNA biosynthetic process
positive regulation of respiratory burst
positive regulation of protein−containing complex disassembly
positive regulation of protein localization to nucleolus
positive regulation of protein insertion into mitochondrial outer membrane
positive regulation of positive chemotaxis to cAMP
positive regulation of mRNA polyadenylation
positive regulation of meiotic cell cycle process involved in oocyte maturation
positive regulation of meiotic cell cycle
positive regulation of metaphase/anaphase transition of cell cycle
positive regulation of lipase activity
positive regulation of immature T cell proliferation in thymus
positive regulation of glycoprotein biosynthetic process
positive regulation of glycogen biosynthetic process
positive regulation of fatty acid oxidation
positive regulation of chemorepellent activity
positive regulation of cell cycle phase transition
positive regulation of amacrine cell differentiation
pore formation in membrane of other organism
polyphosphate−mediated signaling
phloem or xylem histogenesis
peroxisome fission
ornithine biosynthetic process
oocyte dorsal/ventral axis specification
notochord cell differentiation
Notch signaling pathway involved in arterial endothelial cell fate commitment
nor−spermidine metabolic process
neuron projection maintenance
neural nucleus development
negative regulation of transporter activity
negative regulation of transcription from RNA polymerase II promoter involved in smooth muscle cell differentiation
negative regulation of regulated secretory pathway
negative regulation of post−embryonic development
negative regulation of interleukin−10 production
negative regulation of hydrogen peroxide−mediated programmed cell death
negative regulation of glycogen biosynthetic process
negative regulation of glucocorticoid secretion
negative regulation of glucocorticoid receptor signaling pathway
negative regulation of glucagon secretion
negative regulation of G protein−coupled receptor signaling pathway
negative regulation of feeding behavior
negative regulation of double−strand break repair via nonhomologous end joining
negative regulation of chaperone−mediated protein folding
negative regulation of cellular response to transforming growth factor beta stimulus
negative regulation of cell division
negative regulation of cardiac muscle cell differentiation
negative regulation of appetite
N−acylphosphatidylethanolamine metabolic process
muscle filament sliding
methionyl−tRNA aminoacylation
methionine catabolic process
membrane depolarization
manchette disassembly
male sex determination
maintenance of ciliary planar beating movement pattern
leukocyte migration involved in inflammatory response
intercellular transport
hematopoietic stem cell migration
hemangioblast cell differentiation
green leaf volatile biosynthetic process
glucosinolate biosynthetic process
glucose 6−phosphate metabolic process
glomus development
GABAergic neuron differentiation in basal ganglia
forebrain neuron differentiation
fatty acid derivative biosynthetic process
epithelial cell proliferation involved in lung morphogenesis
endothelial tip cell fate specification
embryonic liver development
DNA double−strand break processing involved in repair via single−strand annealing
cytokinin transport
cytokinesis
cotyledon vascular tissue pattern formation
complement activation, classical pathway
citrulline biosynthetic process
chondrocyte intercalation involved in growth plate cartilage morphogenesis
chemotaxis to folate
cerebral cortex GABAergic interneuron fate commitment
cerebellar Purkinje cell layer structural organization
cerebellar molecular layer formation
cerebellar cortex development
cellular response to sodium arsenite
cellular response to gonadotropin−releasing hormone
cellular response to diamide
cellular response to cAMP
cardiolipin acyl−chain remodeling
carbohydrate transmembrane transport
blastoderm segmentation
asexual reproduction
anterograde neuronal dense core vesicle transport
activation of transmembrane receptor protein tyrosine kinase activity
activation of phospholipase D activity
abscisic acid−activated signaling pathway
abscisic acid transport
4−hydroxyproline catabolic process
0
10
20
30
40
ontology
BP
CC
MF