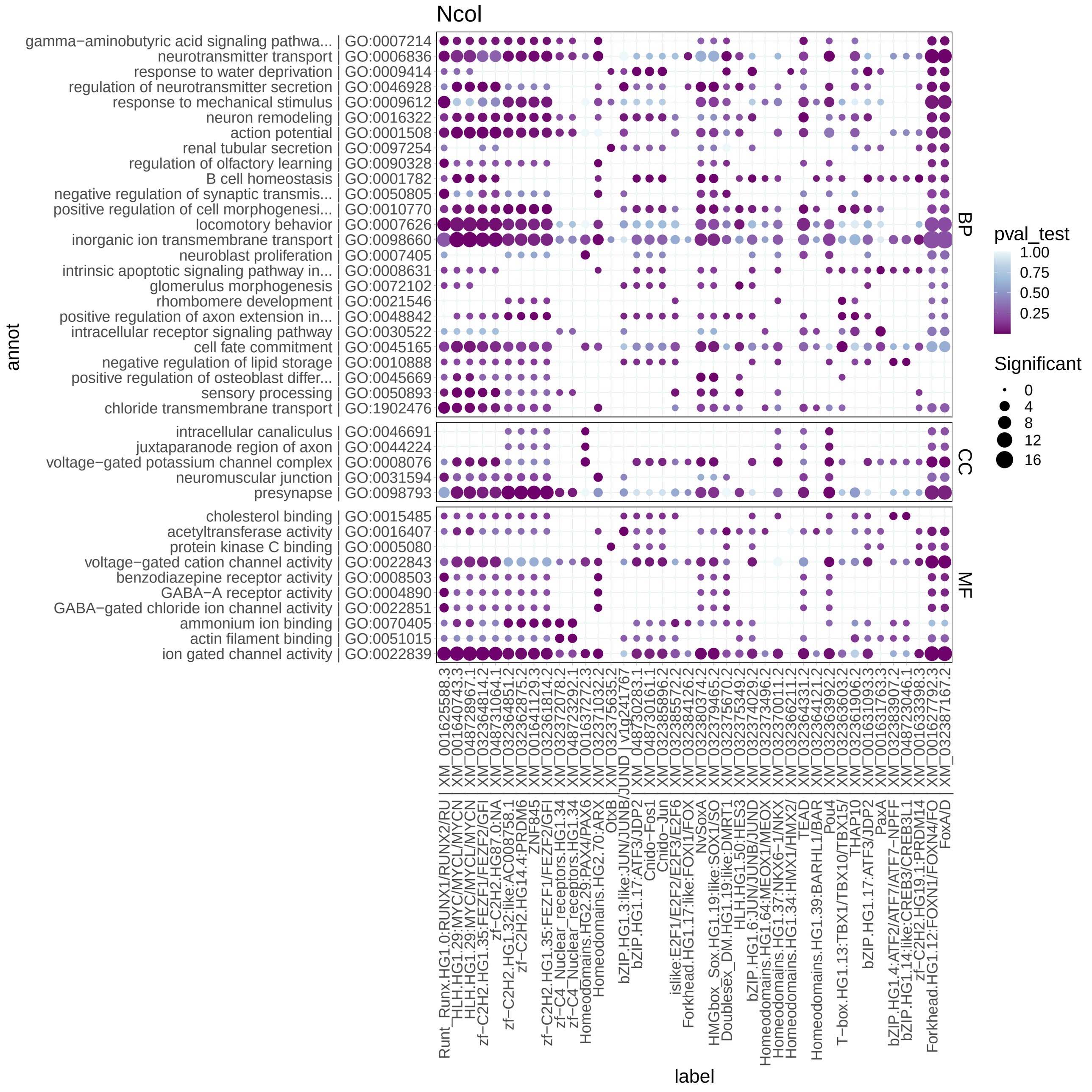
Ncol
gamma−aminobutyric acid signaling pathwa... | GO:0007214
neurotransmitter transport | GO:0006836
response to water deprivation | GO:0009414
regulation of neurotransmitter secretion | GO:0046928
response to mechanical stimulus | GO:0009612
neuron remodeling | GO:0016322
action potential | GO:0001508
renal tubular secretion | GO:0097254
regulation of olfactory learning | GO:0090328
B cell homeostasis | GO:0001782
negative regulation of synaptic transmis... | GO:0050805
positive regulation of cell morphogenesi... | GO:0010770
locomotory behavior | GO:0007626
inorganic ion transmembrane transport | GO:0098660
neuroblast proliferation | GO:0007405
intrinsic apoptotic signaling pathway in... | GO:0008631
glomerulus morphogenesis | GO:0072102
rhombomere development | GO:0021546
positive regulation of axon extension in... | GO:0048842
intracellular receptor signaling pathway | GO:0030522
cell fate commitment | GO:0045165
negative regulation of lipid storage | GO:0010888
positive regulation of osteoblast differ... | GO:0045669
sensory processing | GO:0050893
chloride transmembrane transport | GO:1902476
annot
BP
intracellular canaliculus | GO:0046691
juxtaparanode region of axon | GO:0044224
voltage−gated potassium channel complex | GO:0008076
neuromuscular junction | GO:0031594
presynapse | GO:0098793
CC
cholesterol binding | GO:0015485
acetyltransferase activity | GO:0016407
protein kinase C binding | GO:0005080
voltage−gated cation channel activity | GO:0022843
benzodiazepine receptor activity | GO:0008503
GABA−A receptor activity | GO:0004890
GABA−gated chloride ion channel activity | GO:0022851
ammonium ion binding | GO:0070405
actin filament binding | GO:0051015
ion gated channel activity | GO:0022839
MF
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XM_032364814.2 | zf−C2H2.HG1.35:FEZF1/FEZF2/GFI
XM_048731064.1 | zf−C2H2.HG87.0:NA
XM_032364851.2 | zf−C2H2.HG1.32:like:AC008758.1
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XM_001637272.3 | Homeodomains.HG2.29:PAX4/PAX6
XM_032371032.2 | Homeodomains.HG2.70:ARX
XM_032375635.2 | OtxB
v1g241767 | bZIP.HG1.3:like:JUN/JUNB/JUND
XM_048730283.1 | bZIP.HG1.17:ATF3/JDP2
XM_048730161.1 | Cnido−Fos1
XM_032385896.2 | Cnido−Jun
XM_032385572.2 | islike:E2F1/E2F2/E2F3/E2F6
XM_032384126.2 | Forkhead.HG1.17:like:FOXI1/FOX
XM_032380374.2 | NvSoxA
XM_032379465.2 | HMGbox_Sox.HG1.19:like:SOX1/SO
XM_032375670.2 | Doublesex_DM.HG1.19:like:DMRT1
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XM_032374029.2 | bZIP.HG1.6:JUN/JUNB/JUND
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XM_032366211.2 | Homeodomains.HG1.34:HMX1/HMX2/
XM_032364331.2 | TEAD
XM_032364121.2 | Homeodomains.HG1.39:BARHL1/BAR
XM_032363992.2 | Pou4
XM_032363603.2 | T−box.HG1.13:TBX1/TBX10/TBX15/
XM_032361906.2 | THAP10
XM_001631093.3 | bZIP.HG1.17:ATF3/JDP2
XM_001631763.3 | PaxA
XM_032383907.2 | bZIP.HG1.4:ATF2/ATF7/ATF7−NPFF
XM_048723046.1 | bZIP.HG1.14:like:CREB3/CREB3L1
XM_001633398.3 | zf−C2H2.HG19.1:PRDM14
XM_001627792.3 | Forkhead.HG1.12:FOXN1/FOXN4/FO
XM_032387167.2 | FoxA/D
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0.50
0.25
Significant
0
4
8
12
16