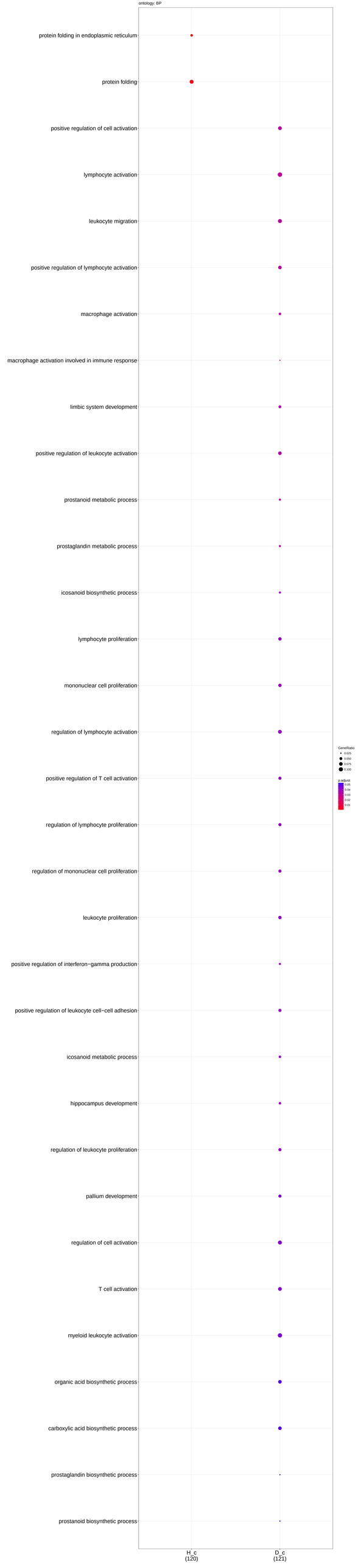
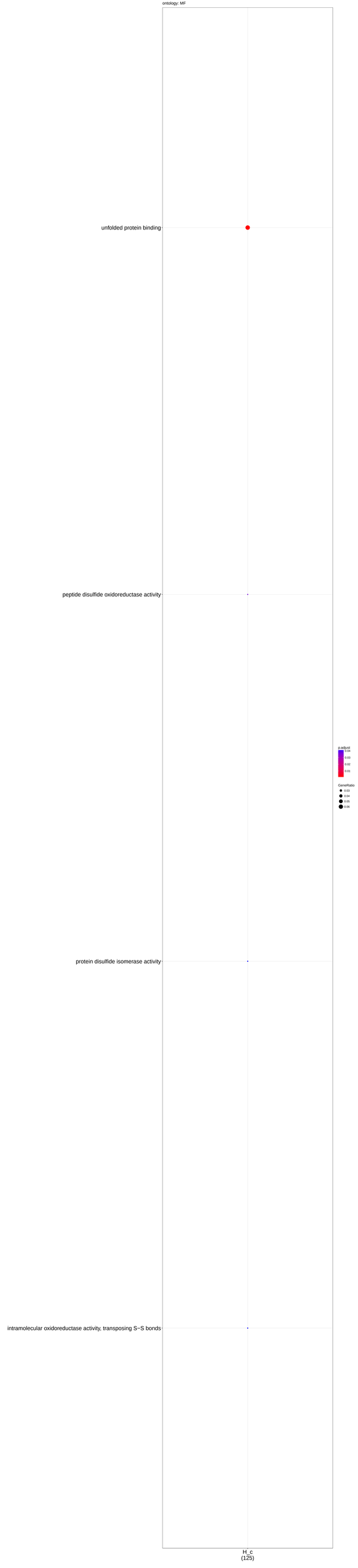
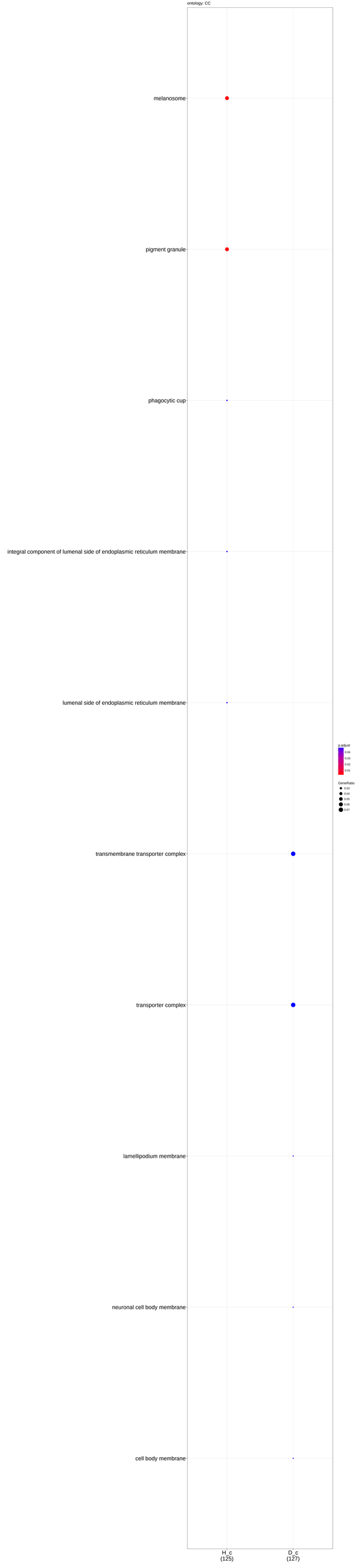

ontology: BP
protein folding in endoplasmic reticulum
protein folding
positive regulation of cell activation
lymphocyte activation
leukocyte migration
positive regulation of lymphocyte activation
macrophage activation
macrophage activation involved in immune response
limbic system development
positive regulation of leukocyte activation
prostanoid metabolic process
prostaglandin metabolic process
icosanoid biosynthetic process
lymphocyte proliferation
mononuclear cell proliferation
regulation of lymphocyte activation
positive regulation of T cell activation
regulation of lymphocyte proliferation
regulation of mononuclear cell proliferation
leukocyte proliferation
positive regulation of interferon−gamma production
positive regulation of leukocyte cell−cell adhesion
icosanoid metabolic process
hippocampus development
regulation of leukocyte proliferation
pallium development
regulation of cell activation
T cell activation
myeloid leukocyte activation
organic acid biosynthetic process
carboxylic acid biosynthetic process
prostaglandin biosynthetic process
prostanoid biosynthetic process
H_c
(120)
D_c
(121)
GeneRatio
0.025
0.050
0.075
0.100
p.adjust
0.05
0.04
0.03
0.02
0.01


ontology: MF
unfolded protein binding
peptide disulfide oxidoreductase activity
protein disulfide isomerase activity
intramolecular oxidoreductase activity, transposing S−S bonds
p.adjust
0.04
0.03
0.02
0.01
GeneRatio
0.03
0.04
0.05
0.06
H_c
(125)


ontology: CC
melanosome
pigment granule
phagocytic cup
integral component of lumenal side of endoplasmic reticulum membrane
lumenal side of endoplasmic reticulum membrane
transmembrane transporter complex
transporter complex
lamellipodium membrane
neuronal cell body membrane
cell body membrane
p.adjust
0.04
0.03
0.02
0.01
GeneRatio
0.03
0.04
0.05
0.06
0.07
H_c
(125)
D_c
(127)