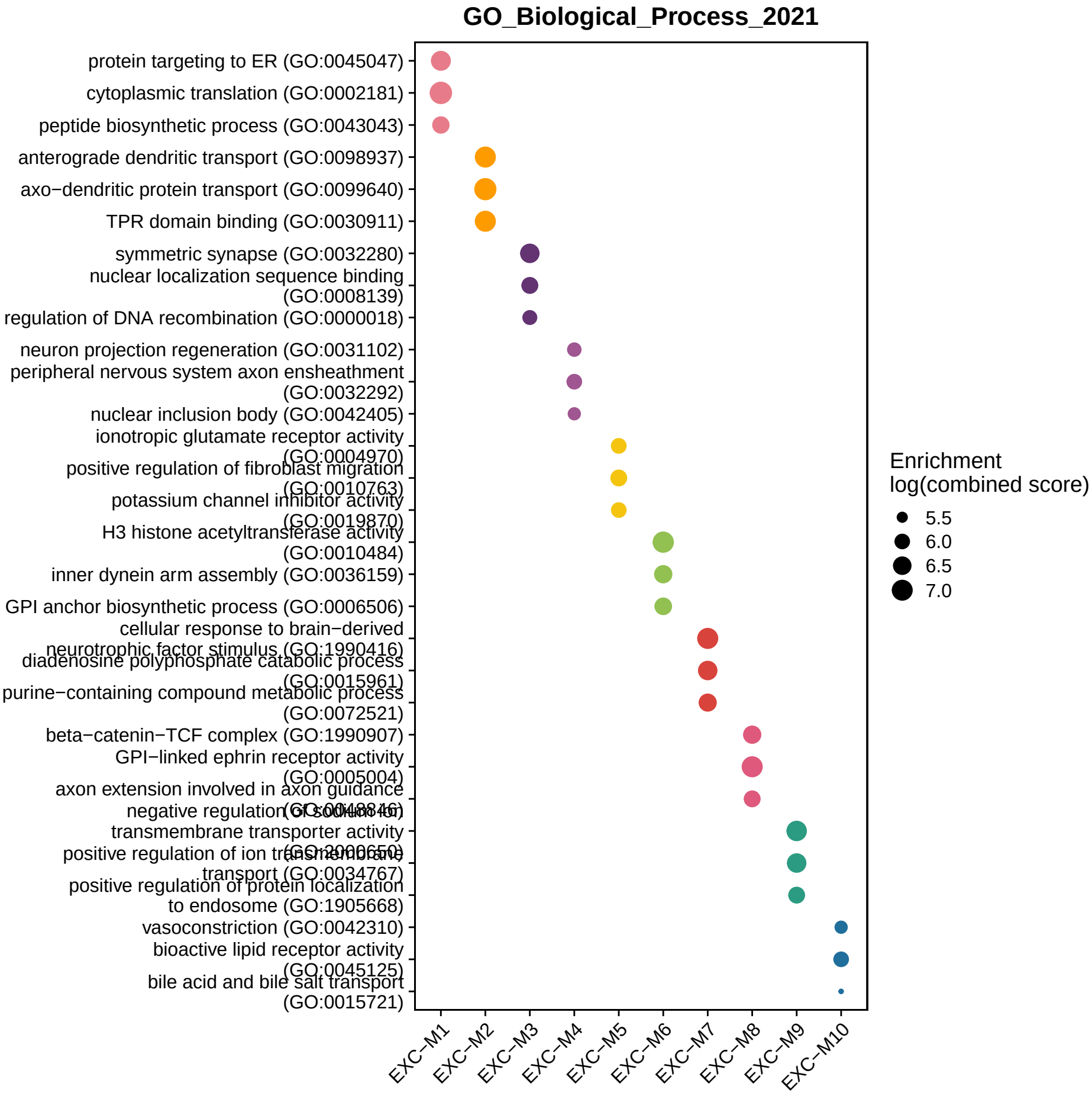
GO_Biological_Process_2021
protein targeting to ER (GO:0045047)
cytoplasmic translation (GO:0002181)
peptide biosynthetic process (GO:0043043)
anterograde dendritic transport (GO:0098937)
axo−dendritic protein transport (GO:0099640)
TPR domain binding (GO:0030911)
symmetric synapse (GO:0032280)
nuclear localization sequence binding (GO:0008139)
regulation of DNA recombination (GO:0000018)
neuron projection regeneration (GO:0031102)
peripheral nervous system axon ensheathment (GO:0032292)
nuclear inclusion body (GO:0042405)
ionotropic glutamate receptor activity (GO:0004970)
positive regulation of fibroblast migration (GO:0010763)
potassium channel inhibitor activity (GO:0019870)
H3 histone acetyltransferase activity (GO:0010484)
inner dynein arm assembly (GO:0036159)
GPI anchor biosynthetic process (GO:0006506)
cellular response to brain−derived neurotrophic factor stimulus (GO:1990416)
diadenosine polyphosphate catabolic process (GO:0015961)
purine−containing compound metabolic process (GO:0072521)
beta−catenin−TCF complex (GO:1990907)
GPI−linked ephrin receptor activity (GO:0005004)
axon extension involved in axon guidance (GO:0048846)
negative regulation of sodium ion transmembrane transporter activity (GO:2000650)
positive regulation of ion transmembrane transport (GO:0034767)
positive regulation of protein localization to endosome (GO:1905668)
vasoconstriction (GO:0042310)
bioactive lipid receptor activity (GO:0045125)
bile acid and bile salt transport (GO:0015721)
EXC−M1
EXC−M2
EXC−M3
EXC−M4
EXC−M5
EXC−M6
EXC−M7
EXC−M8
EXC−M9
EXC−M10
Enrichment
log(combined score)
5.5
6.0
6.5
7.0