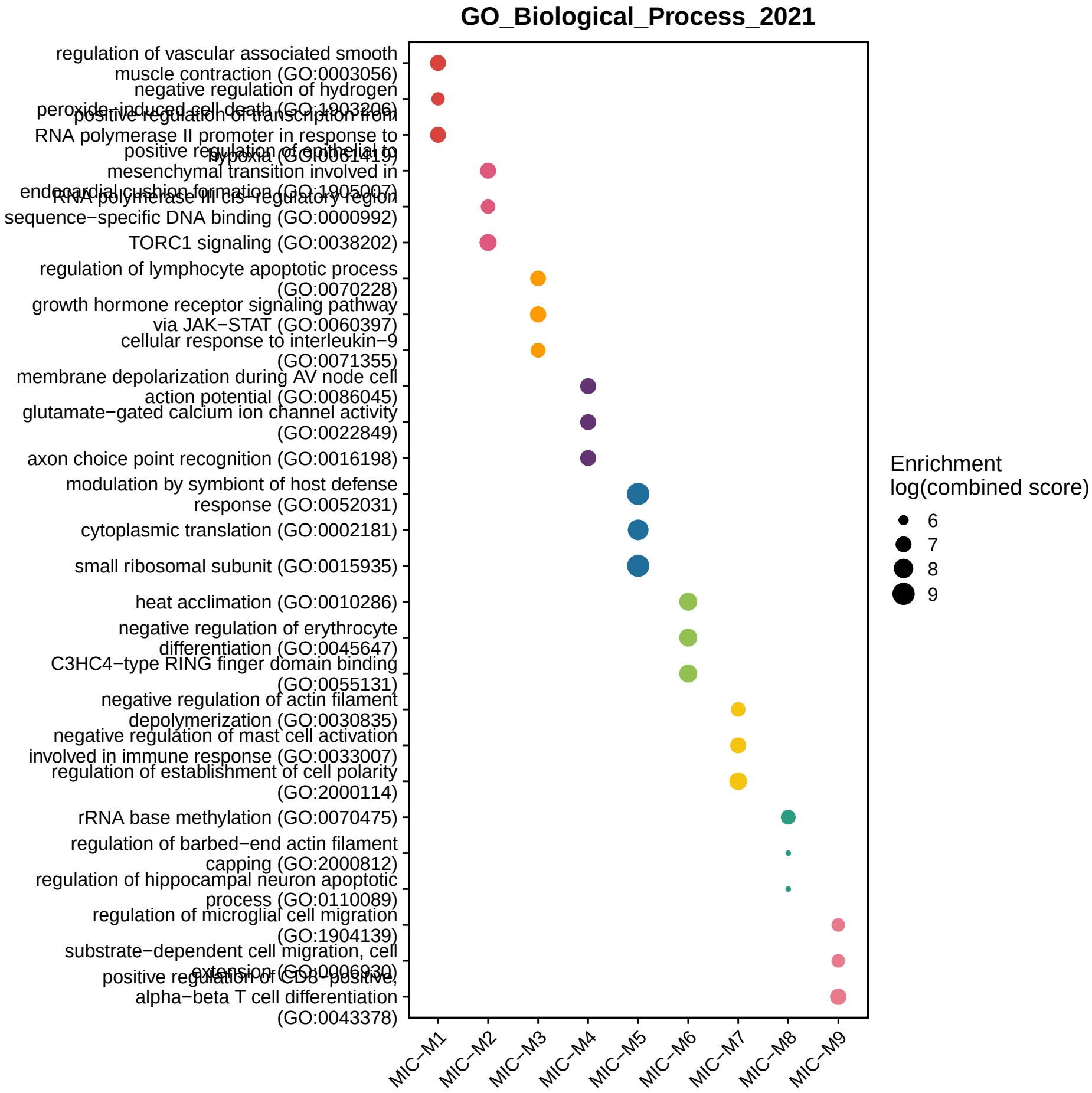
GO_Biological_Process_2021
regulation of vascular associated smooth muscle contraction (GO:0003056)
negative regulation of hydrogen peroxide-induced cell death (GO:1903206)
positive regulation of transcription from RNA polymerase II promoter in response to hypoxia (GO:0061419)
positive regulation of epithelial to mesenchymal transition involved in endocardial cushion formation (GO:1905007)
RNA polymerase III cis-regulatory region sequence-specific DNA binding (GO:0000992)
TORC1 signaling (GO:0038202)
regulation of lymphocyte apoptotic process (GO:0070228)
growth hormone receptor signaling pathway via JAK-STAT (GO:0060397)
cellular response to interleukin-9 (GO:0071355)
membrane depolarization during AV node cell action potential (GO:0086045)
glutamate-gated calcium ion channel activity (GO:0022849)
axon choice point recognition (GO:0016198)
modulation by symbiont of host defense response (GO:0052031)
cytoplasmic translation (GO:0002181)
small ribosomal subunit (GO:0015935)
heat acclimation (GO:0010286)
negative regulation of erythrocyte differentiation (GO:0045647)
C3HC4-type RING finger domain binding (GO:0055131)
negative regulation of actin filament depolymerization (GO:0030835)
negative regulation of mast cell activation involved in immune response (GO:0033007)
regulation of establishment of cell polarity (GO:2000114)
rRNA base methylation (GO:0070475)
regulation of barbed-end actin filament capping (GO:2000812)
regulation of hippocampal neuron apoptotic process (GO:0110089)
regulation of microglial cell migration (GO:1904139)
substrate-dependent cell migration, cell extension (GO:0006930)
positive regulation of CD8-positive, alpha-beta T cell differentiation (GO:0043378)
MIC-M1
MIC-M2
MIC-M3
MIC-M4
MIC-M5
MIC-M6
MIC-M7
MIC-M8
MIC-M9
Enrichment
log(combined score)
6
7
8
9