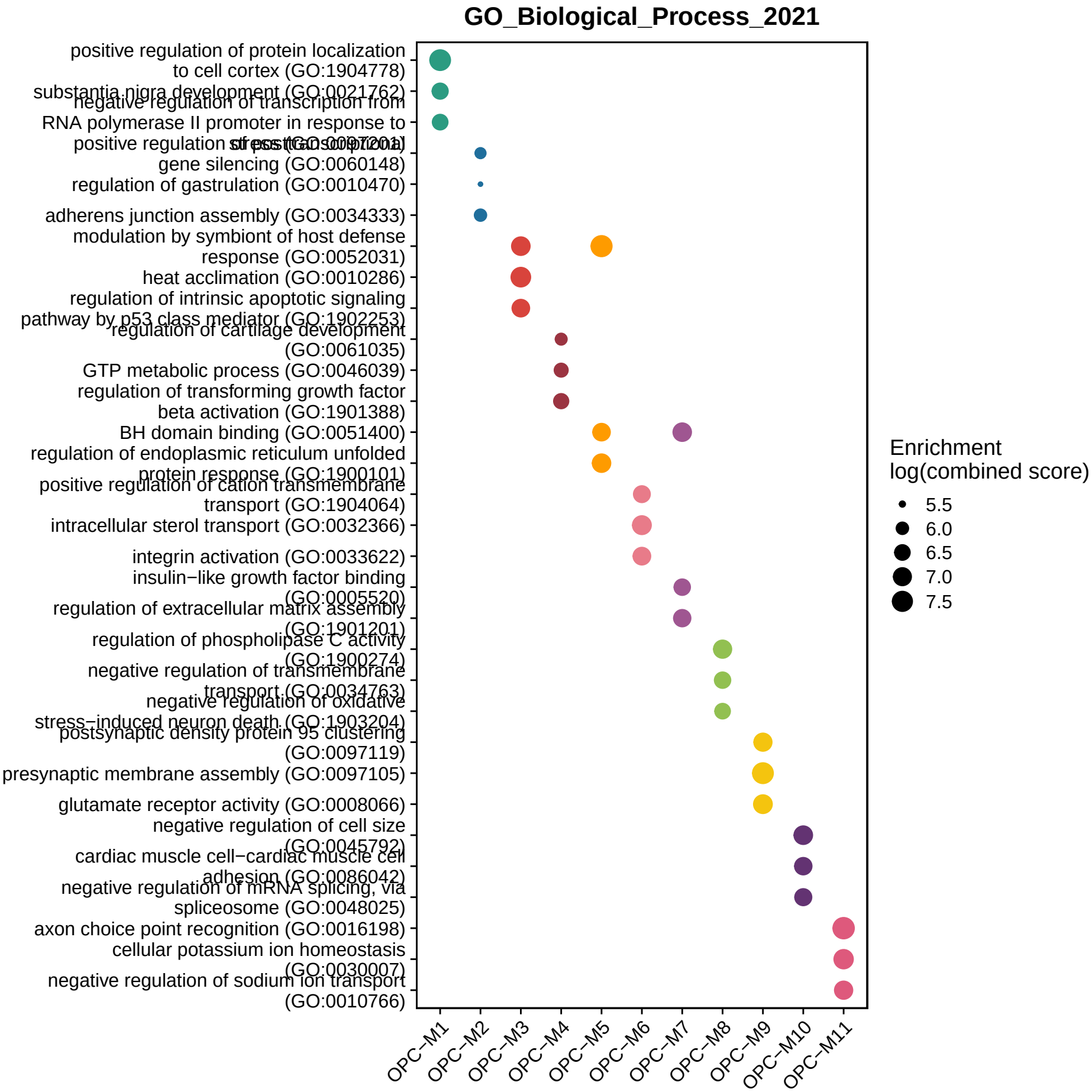
GO_Biological_Process_2021
positive regulation of protein localization to cell cortex (GO:1904778)
substantia nigra development (GO:0021762)
negative regulation of transcription from RNA polymerase II promoter in response to stress (GO:0097201)
positive regulation of [illegible] (GO:0097201)
gene silencing (GO:0060148)
regulation of gastrulation (GO:0010470)
adherens junction assembly (GO:0034333)
modulation by symbiont of host defense response (GO:0052031)
heat acclimation (GO:0010286)
regulation of intrinsic apoptotic signaling pathway by p53 class mediator (GO:1902253)
regulation of cartilage development (GO:0061035)
GTP metabolic process (GO:0046039)
regulation of transforming growth factor beta activation (GO:1901388)
BH domain binding (GO:0051400)
regulation of endoplasmic reticulum unfolded protein response (GO:1900101)
positive regulation of cation transmembrane transport (GO:1904064)
intracellular sterol transport (GO:0032366)
integrin activation (GO:0033622)
insulin−like growth factor binding (GO:0005520)
regulation of extracellular matrix assembly (GO:1901201)
regulation of phospholipase C activity (GO:1900274)
negative regulation of transmembrane transport (GO:0034763)
negative regulation of oxidative stress−induced neuron death (GO:1903204)
postsynaptic density protein 95 clustering (GO:0097119)
presynaptic membrane assembly (GO:0097105)
glutamate receptor activity (GO:0008066)
negative regulation of cell size (GO:0045792)
cardiac muscle cell−cardiac muscle cell adhesion (GO:0086042)
negative regulation of mRNA splicing, via spliceosome (GO:0048025)
axon choice point recognition (GO:0016198)
cellular potassium ion homeostasis (GO:0030007)
negative regulation of sodium ion transport (GO:0010766)
OPC−M1
OPC−M2
OPC−M3
OPC−M4
OPC−M5
OPC−M6
OPC−M7
OPC−M8
OPC−M9
OPC−M10
OPC−M11
Enrichment
log(combined score)
5.5
6.0
6.5
7.0
7.5