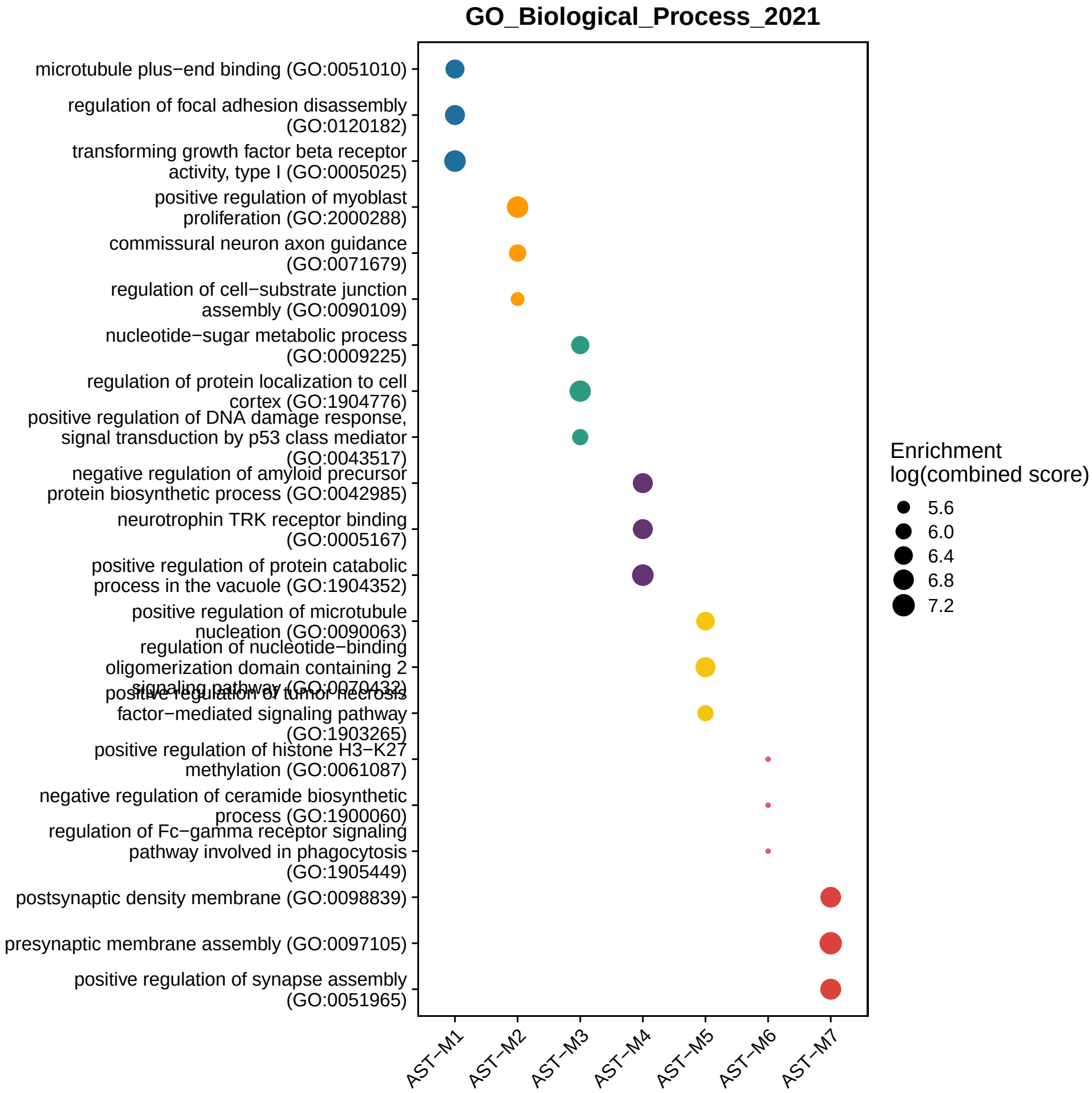
GO_Biological_Process_2021
microtubule plus−end binding (GO:0051010)
regulation of focal adhesion disassembly (GO:0120182)
transforming growth factor beta receptor activity, type I (GO:0005025)
positive regulation of myoblast proliferation (GO:2000288)
commissural neuron axon guidance (GO:0071679)
regulation of cell−substrate junction assembly (GO:0090109)
nucleotide−sugar metabolic process (GO:0009225)
regulation of protein localization to cell cortex (GO:1904776)
positive regulation of DNA damage response, signal transduction by p53 class mediator (GO:0043517)
negative regulation of amyloid precursor protein biosynthetic process (GO:0042985)
neurotrophin TRK receptor binding (GO:0005167)
positive regulation of protein catabolic process in the vacuole (GO:1904352)
positive regulation of microtubule nucleation (GO:0090063)
regulation of nucleotide−binding oligomerization domain containing 2 signaling pathway (GO:0070432)
positive regulation of tumor necrosis factor−mediated signaling pathway (GO:1903265)
positive regulation of histone H3−K27 methylation (GO:0061087)
negative regulation of ceramide biosynthetic process (GO:1900060)
regulation of Fc−gamma receptor signaling pathway involved in phagocytosis (GO:1905449)
postsynaptic density membrane (GO:0098839)
presynaptic membrane assembly (GO:0097105)
positive regulation of synapse assembly (GO:0051965)
AST−M1
AST−M2
AST−M3
AST−M4
AST−M5
AST−M6
AST−M7
Enrichment
log(combined score)
5.6
6.0
6.4
6.8
7.2