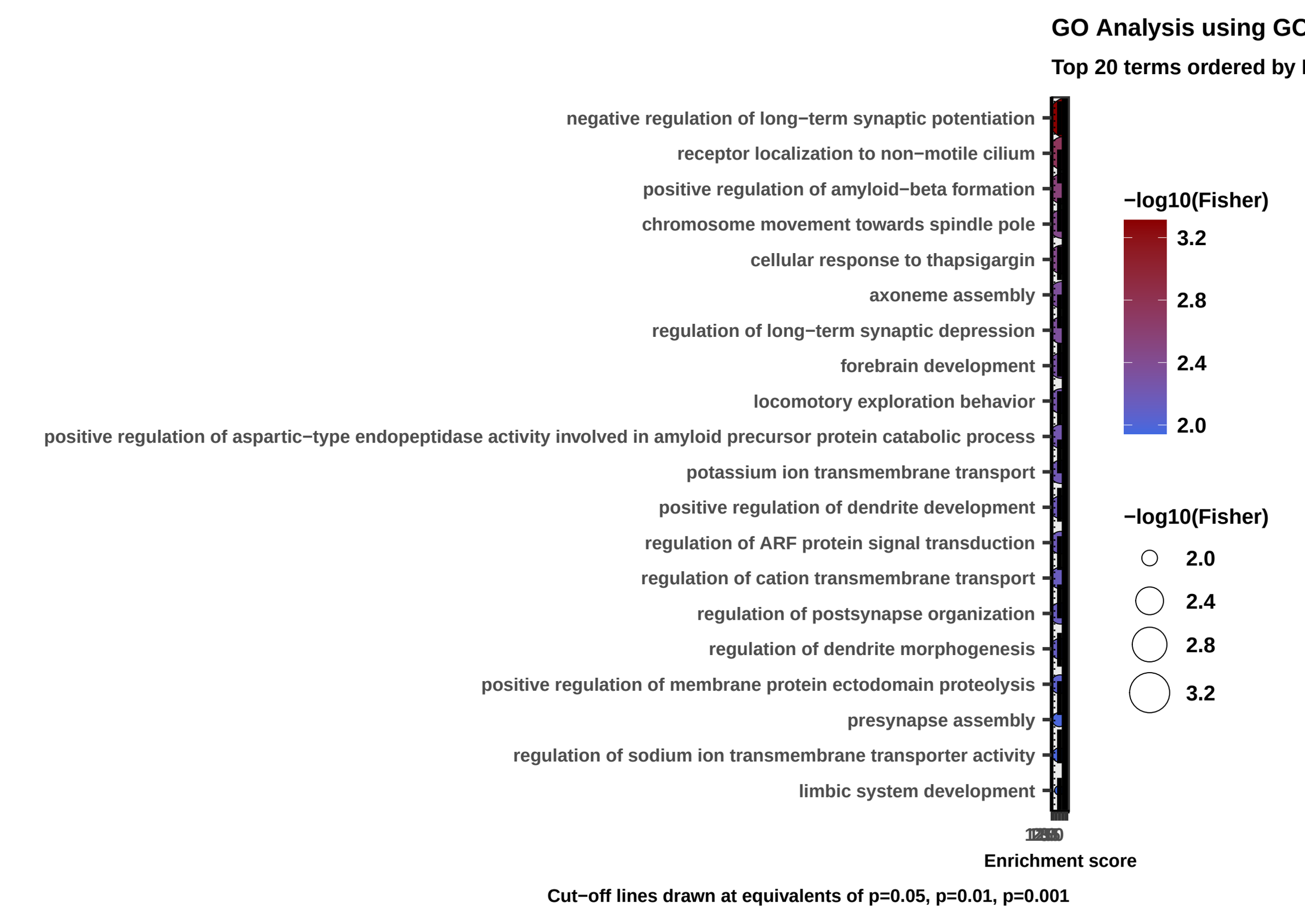
GO Analysis using GO
Top 20 terms ordered by
negative regulation of long−term synaptic potentiation
receptor localization to non−motile cilium
positive regulation of amyloid−beta formation
chromosome movement towards spindle pole
cellular response to thapsigargin
axoneme assembly
regulation of long−term synaptic depression
forebrain development
locomotory exploration behavior
positive regulation of aspartic−type endopeptidase activity involved in amyloid precursor protein catabolic process
potassium ion transmembrane transport
positive regulation of dendrite development
regulation of ARF protein signal transduction
regulation of cation transmembrane transport
regulation of postsynapse organization
regulation of dendrite morphogenesis
positive regulation of membrane protein ectodomain proteolysis
presynapse assembly
regulation of sodium ion transmembrane transporter activity
limbic system development
Enrichment score
−log10(Fisher)
3.2
2.8
2.4
2.0
−log10(Fisher)
2.0
2.4
2.8
3.2
Cut−off lines drawn at equivalents of p=0.05, p=0.01, p=0.001