
EnrichmentGO
adenylate cyclase binding
activation of adenylate cyclase activity
positive regulation of adenylate cyclase activity
cortical actin cytoskeleton
positive regulation of lyase activity
positive regulation of cyclase activity
regulation of adenylate cyclase activity
positive regulation of cAMP biosynthetic process
cortical cytoskeleton
regulation of cyclase activity
positive regulation of cAMP metabolic process
regulation of lyase activity
positive regulation of cyclic nucleotide biosynthetic process
positive regulation of purine nucleotide biosynthetic process
positive regulation of nucleotide biosynthetic process
positive regulation of cyclic nucleotide metabolic process
regulation of cAMP biosynthetic process
cell cortex part
cAMP biosynthetic process
regulation of cAMP metabolic process
positive regulation of purine nucleotide metabolic process
positive regulation of nucleotide metabolic process
regulation of cyclic nucleotide biosynthetic process
regulation of purine nucleotide biosynthetic process
regulation of nucleotide biosynthetic process
cAMP metabolic process
cyclic purine nucleotide metabolic process
cyclic nucleotide biosynthetic process
regulation of cyclic nucleotide metabolic process
establishment or maintenance of cell polarity
cyclic nucleotide metabolic process
actin polymerization or depolymerization
regulation of purine nucleotide metabolic process
regulation of nucleotide metabolic process
cell cortex
purine ribonucleotide biosynthetic process
purine nucleotide biosynthetic process
ribonucleotide biosynthetic process
purine−containing compound biosynthetic process
ribose phosphate biosynthetic process
cytoplasmic region
nucleotide biosynthetic process
nucleoside phosphate biosynthetic process
actin filament organization
actin binding
actin cytoskeleton
purine ribonucleotide metabolic process
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autophagy	2.98
regulation of homeostatic process	2.97
negative regulation of intracellular signal transduction	2.94
blood vessel morphogenesis	2.85
purine ribonucleotide metabolic process	2.84

GO Terms

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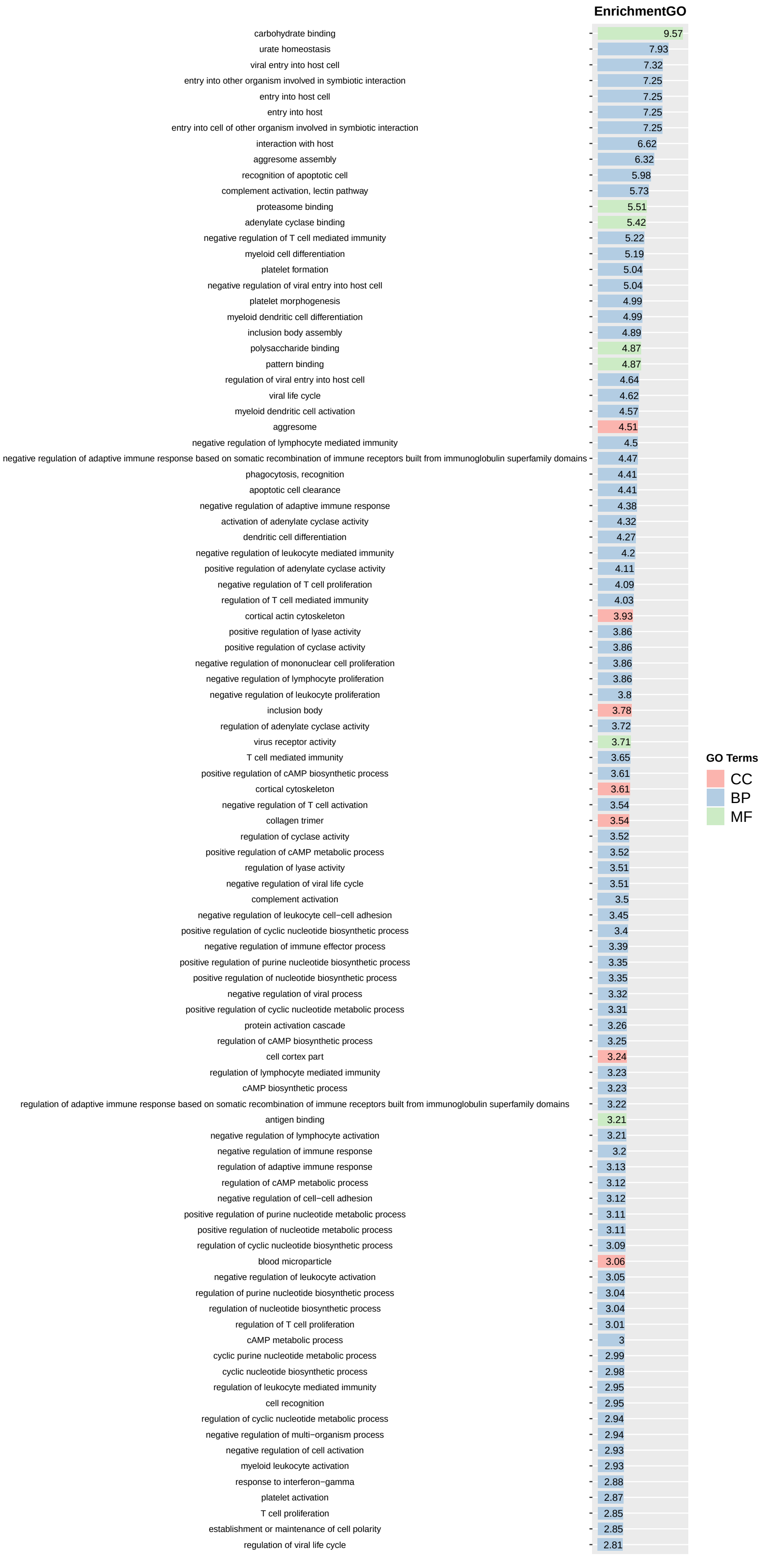
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carbohydrate binding 9.57
urate homeostasis 7.93
viral entry into host cell 7.32
entry into other organism involved in symbiotic interaction 7.25
entry into host cell 7.25
entry into host 7.25
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positive regulation of nucleotide biosynthetic process 3.35
negative regulation of viral process 3.32
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regulation of lymphocyte mediated immunity 3.23
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negative regulation of immune response 3.2
regulation of adaptive immune response 3.13
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negative regulation of cell−cell adhesion 3.12
positive regulation of purine nucleotide metabolic process 3.11
positive regulation of nucleotide metabolic process 3.11
regulation of cyclic nucleotide biosynthetic process 3.09
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negative regulation of leukocyte activation 3.05
regulation of purine nucleotide biosynthetic process 3.04
regulation of nucleotide biosynthetic process 3.04
regulation of T cell proliferation 3.01
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cyclic purine nucleotide metabolic process 2.99
cyclic nucleotide biosynthetic process 2.98
regulation of leukocyte mediated immunity 2.95
cell recognition 2.95
regulation of cyclic nucleotide metabolic process 2.94
negative regulation of multi−organism process 2.94
negative regulation of cell activation 2.93
myeloid leukocyte activation 2.93
response to interferon−gamma 2.88
platelet activation 2.87
T cell proliferation 2.85
establishment or maintenance of cell polarity 2.85
regulation of viral life cycle 2.81
GO Terms
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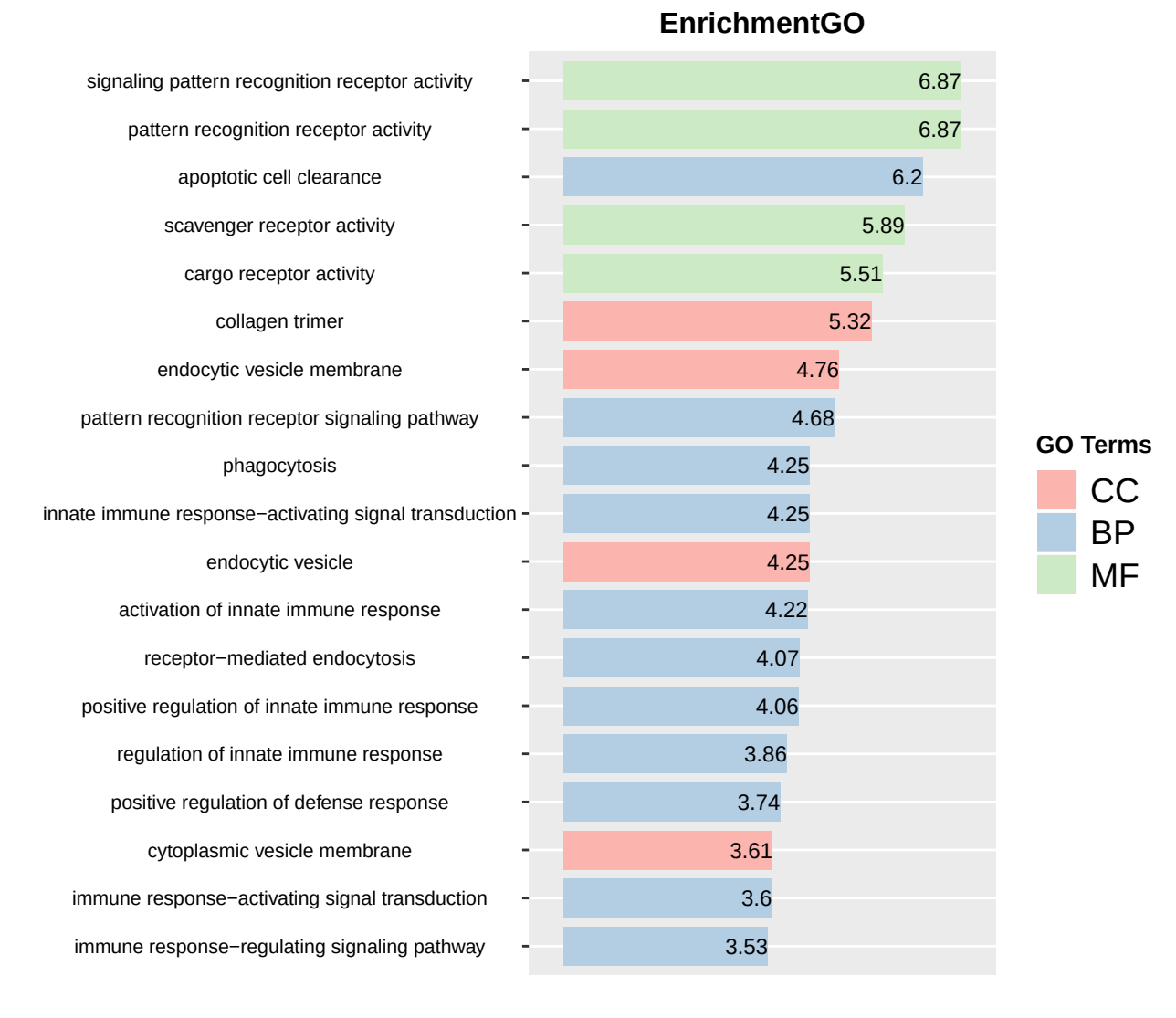
EnrichmentGO
signaling pattern recognition receptor activity
pattern recognition receptor activity
apoptotic cell clearance
scavenger receptor activity
cargo receptor activity
collagen trimer
endocytic vesicle membrane
pattern recognition receptor signaling pathway
phagocytosis
innate immune response−activating signal transduction
endocytic vesicle
activation of innate immune response
receptor−mediated endocytosis
positive regulation of innate immune response
regulation of innate immune response
positive regulation of defense response
cytoplasmic vesicle membrane
immune response−activating signal transduction
immune response−regulating signaling pathway
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GO Terms
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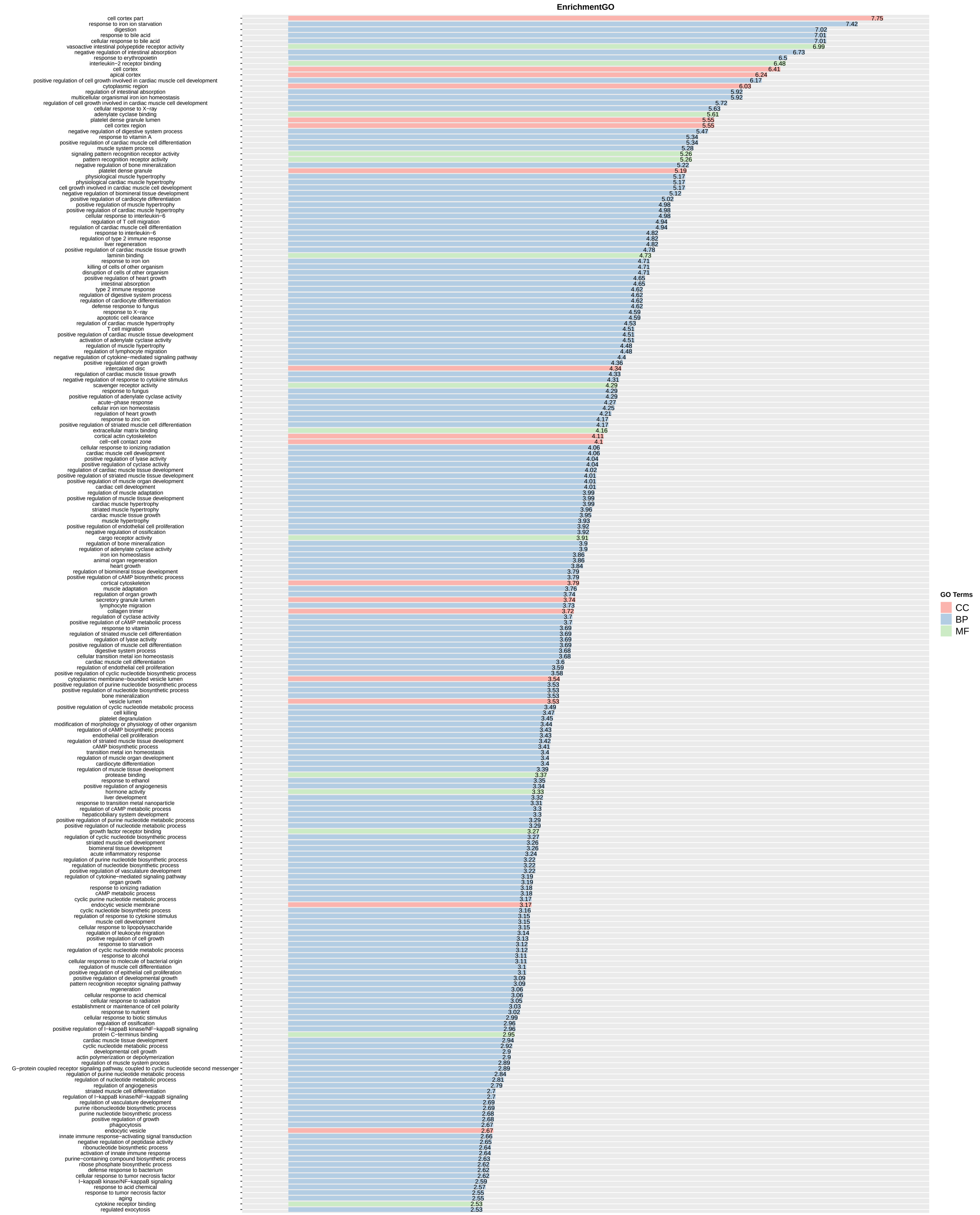
EnrichmentGO
cell cortex part 7.75
response to iron ion starvation 7.42
digestion 7.02
response to bile acid 7.01
cellular response to bile acid 7.01
vasoactive intestinal polypeptide receptor activity 6.99
negative regulation of intestinal absorption 6.73
response to erythropoietin 6.5
interleukin−2 receptor binding 6.48
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regulation of intestinal absorption 5.92
multicellular organismal iron ion homeostasis 5.92
regulation of cell growth involved in cardiac muscle cell development 5.72
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regulation of cardiac muscle cell differentiation 4.94
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regulation of type 2 immune response 4.82
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positive regulation of cardiac muscle tissue growth 4.78
laminin binding 4.73
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killing of cells of other organism 4.71
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regulation of digestive system process 4.62
regulation of cardiocyte differentiation 4.62
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response to X−ray 4.59
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negative regulation of ossification 3.92
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response to acid chemical 2.57
response to tumor necrosis factor 2.55
aging 2.55
cytokine receptor binding 2.53
regulated exocytosis 2.53
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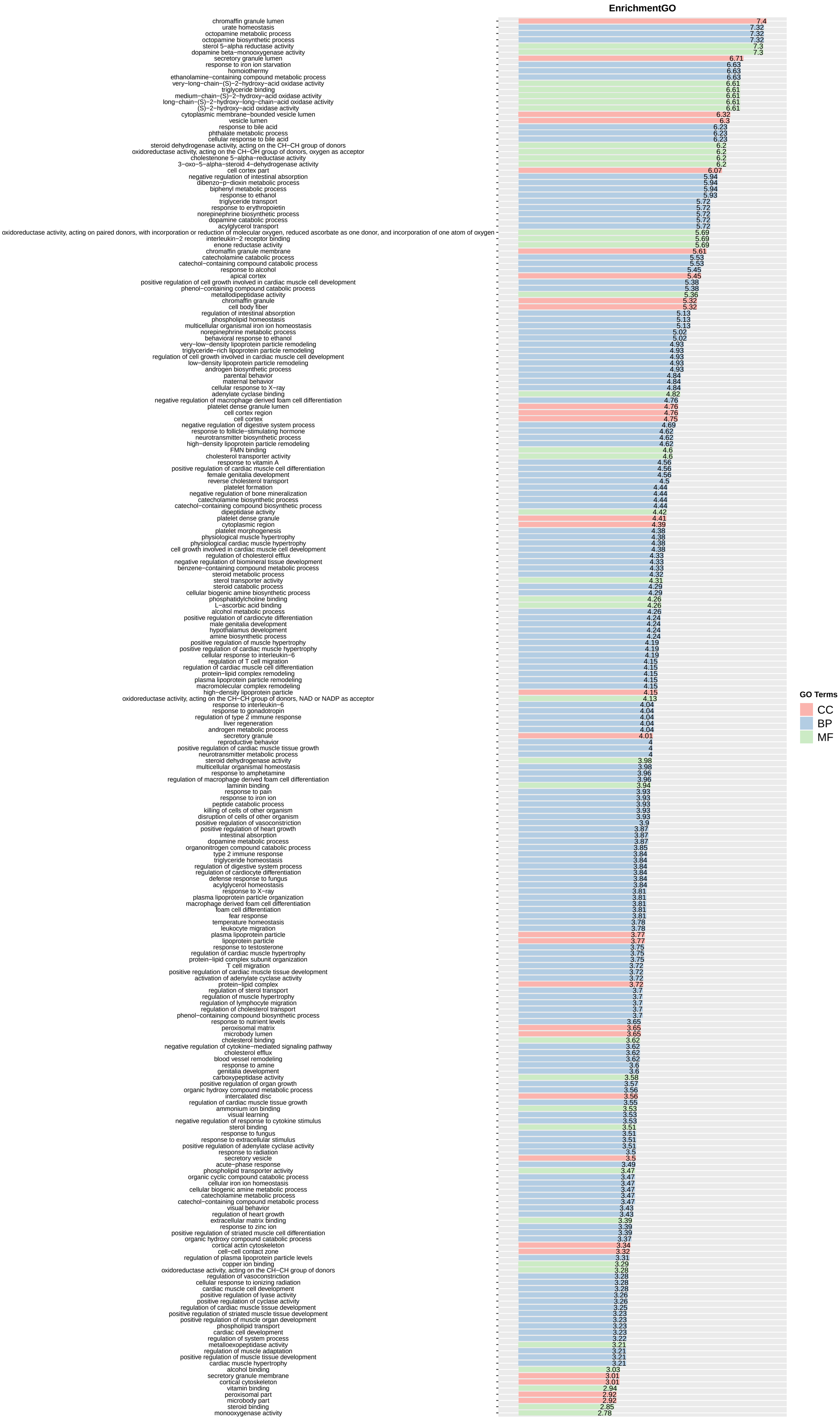
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chromaffin granule lumen 7.4
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octopamine metabolic process 7.32
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sterol 5−alpha reductase activity 7.3
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multicellular organismal iron ion homeostasis 5.13
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parental behavior 4.84
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positive regulation of cardiac muscle tissue growth 4
neurotransmitter metabolic process 4
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regulation of macrophage derived foam cell differentiation 3.96
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response to pain 3.93
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killing of cells of other organism 3.93
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positive regulation of vasoconstriction 3.9
positive regulation of heart growth 3.87
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