

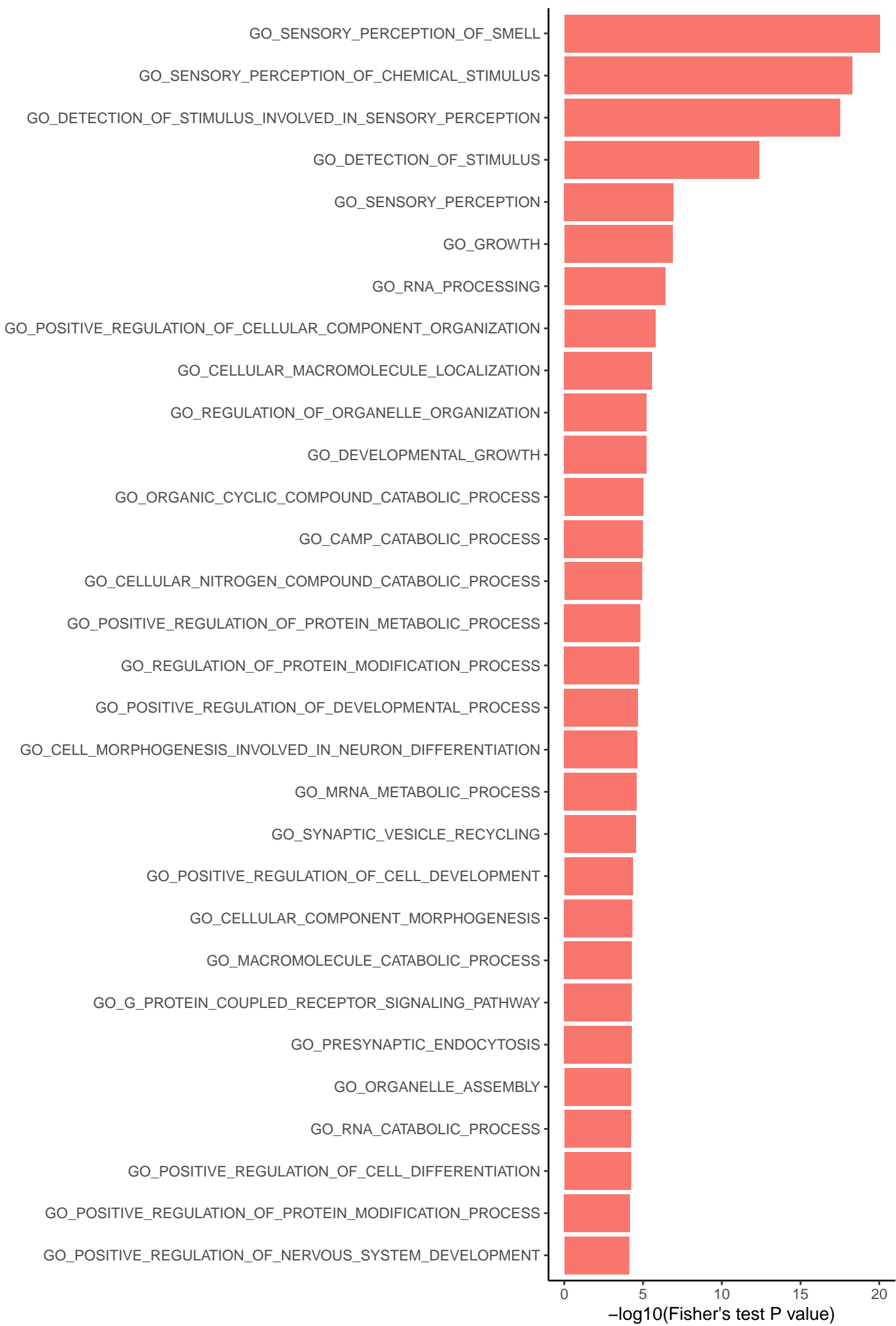
Top 30 enriched GO terms (p < 0.05 ) in All

GO terms



Top 30 enriched GO terms (p < 0.05 ) in  
bp

GO terms



gene\_set  
c5.go.bp

Top 30 enriched GO terms (p < 0.05 ) in

cc

GO terms

GO\_SYNAPSE

GO\_PERINUCLEAR\_REGION\_OF\_CYTOPLASM

GO\_GLUTAMATERGIC\_SYNAPSE

GO\_NEURON\_PROJECTION

GO\_MICROTUBULE\_CYTOSKELETON

GO\_SYNAPTIC\_MEMBRANE

GO\_ORGANELLE\_SUBCOMPARTMENT

GO\_ACTIN\_FILAMENT\_BUNDLE

GO\_NEURON\_TO\_NEURON\_SYNAPSE

GO\_TRANS\_GOLGI\_NETWORK

GO\_PLASMA\_MEMBRANE\_REGION

GO\_FILOPODIUM\_MEMBRANE

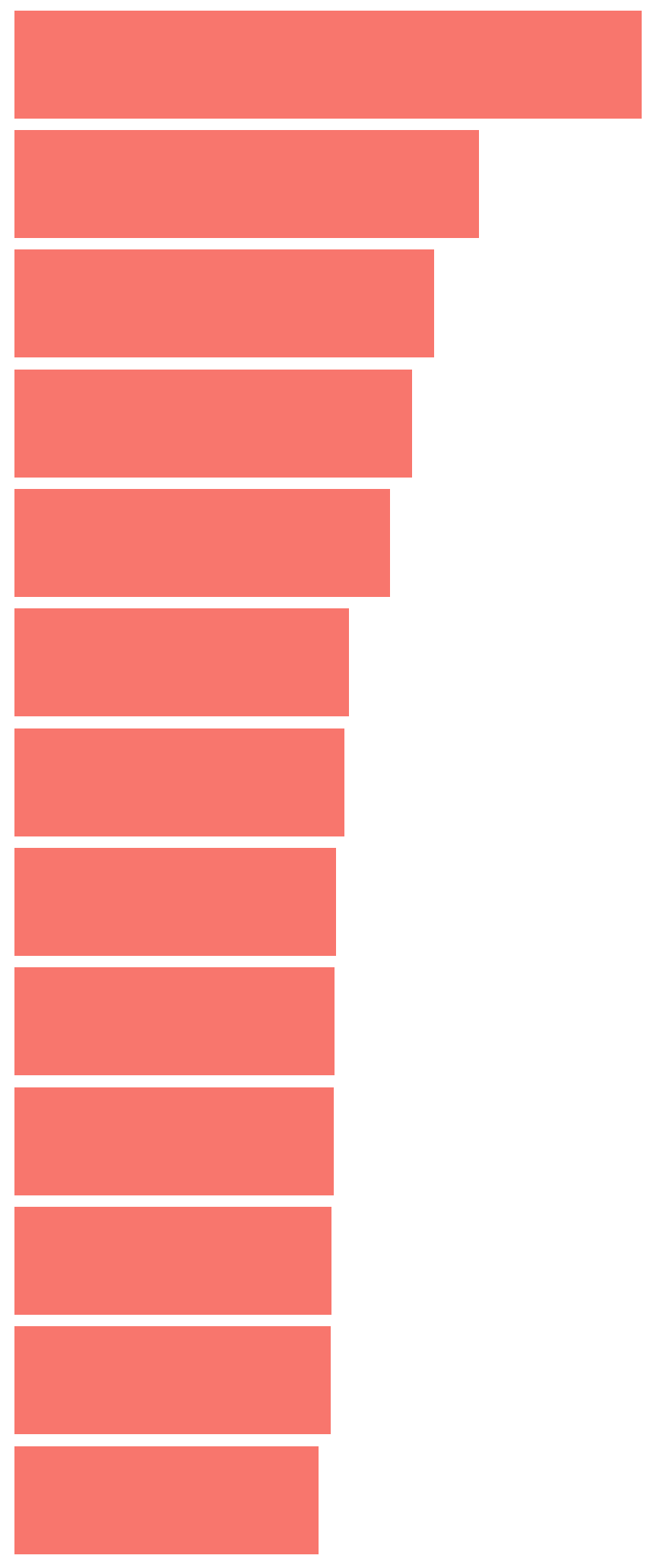
GO\_NUCLEAR\_BODY

gene\_set

c5.go.cc

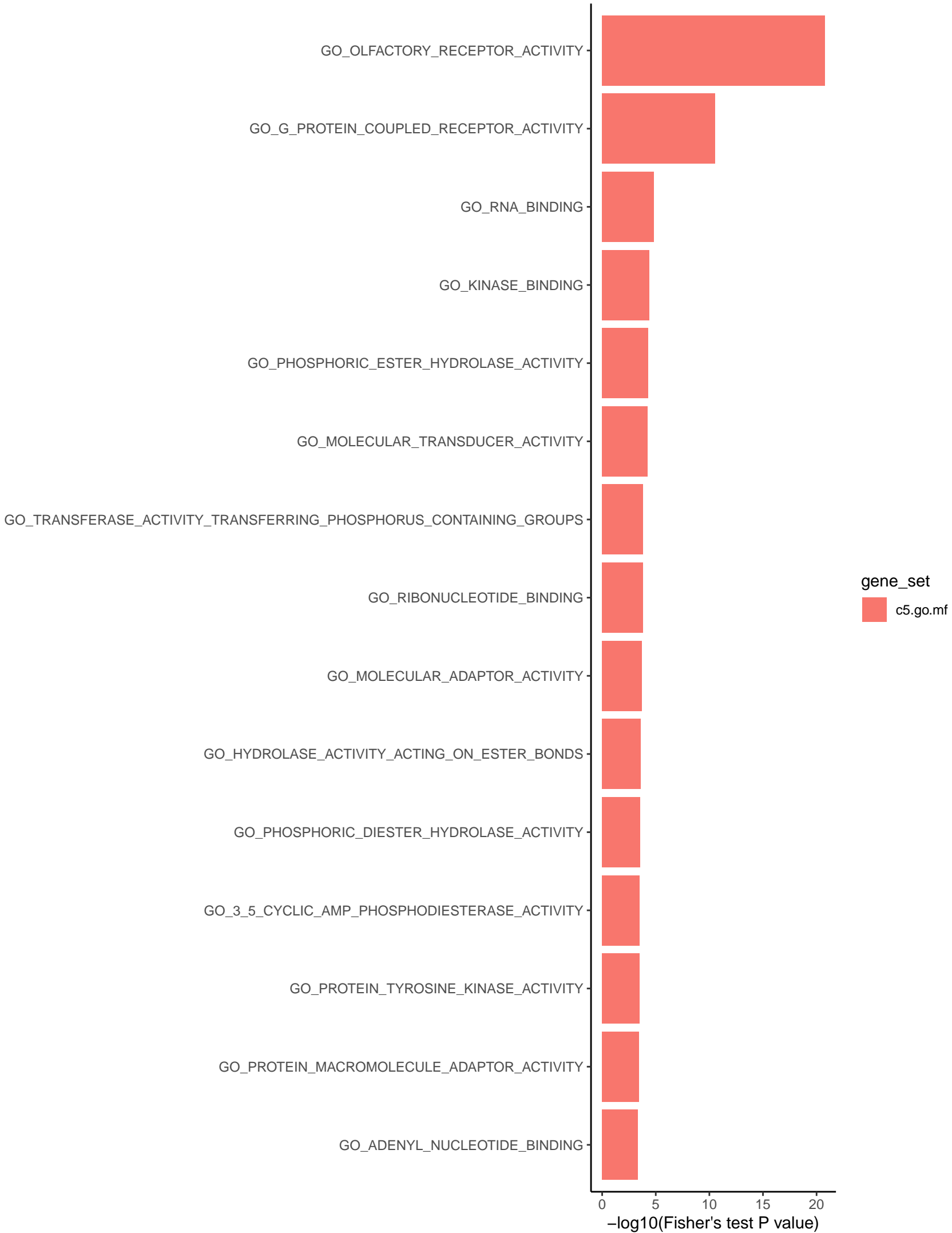
0 2 4 6

$-\log_{10}(\text{Fisher's test P value})$



Top 30 enriched GO terms (p < 0.05 ) in mf

GO terms



Top 30 enriched pathways (p < 0.05 ) in KEGG

pathway

KEGG\_OLFACTORY\_TRANSDUCTION

gene\_set

c2.cp.kegg

$-\log_{10}(\text{Fisher's test P value})$

0

5

10

15

