

With Cell Cycle | Without AreaShape | All Genes

Cluster 1 of 44

Cellular Compartment

nucleus

microtubule organizing center

cytoskeleton

chromosome

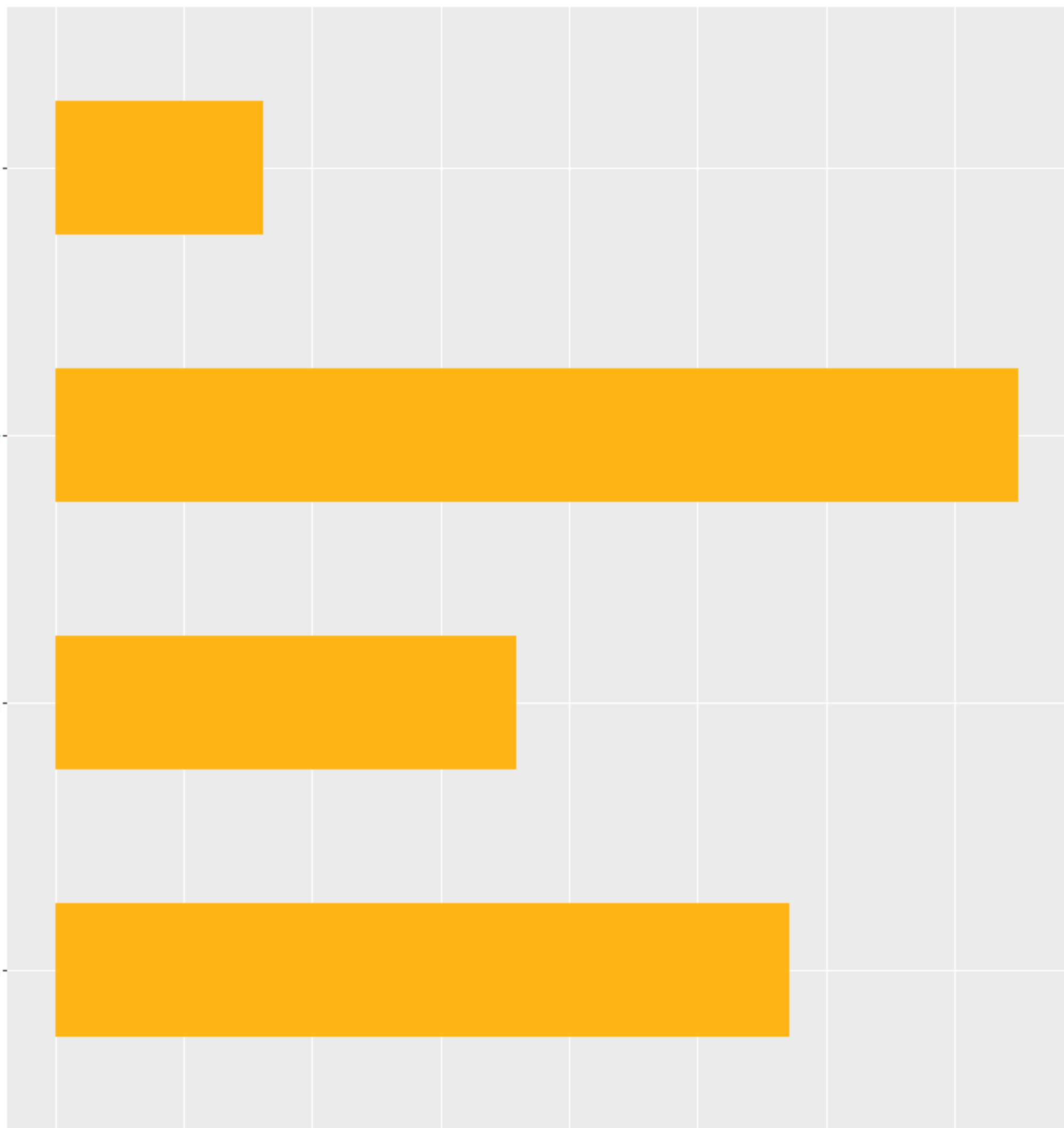
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0.05

0.10

0.15

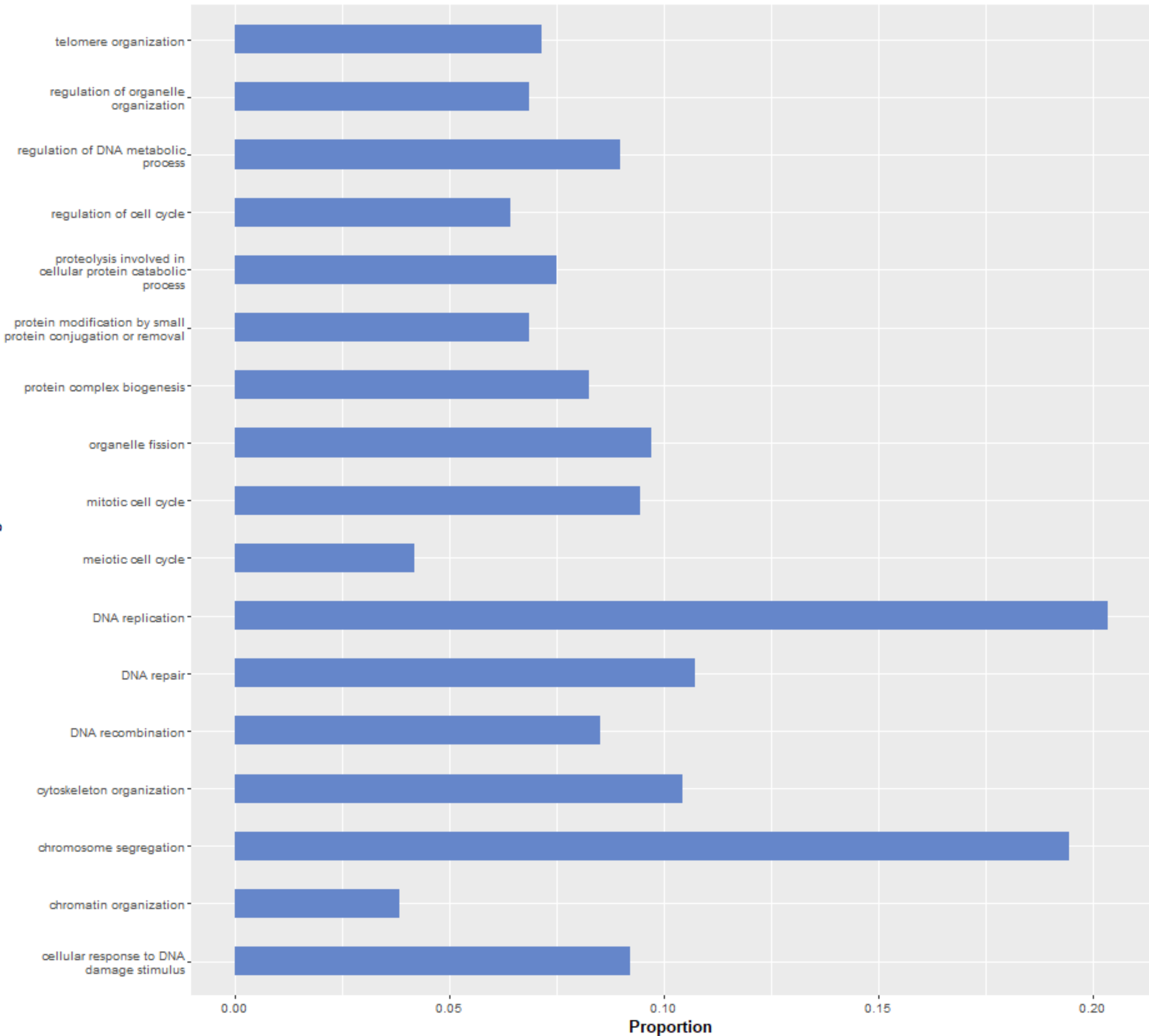
Proportion



With Cell Cycle | Without AreaShape | All Genes

Cluster 1 of 44

Biological Process



With Cell Cycle | Without AreaShape | All Genes

Cluster 3 of 44

Cellular Compartment

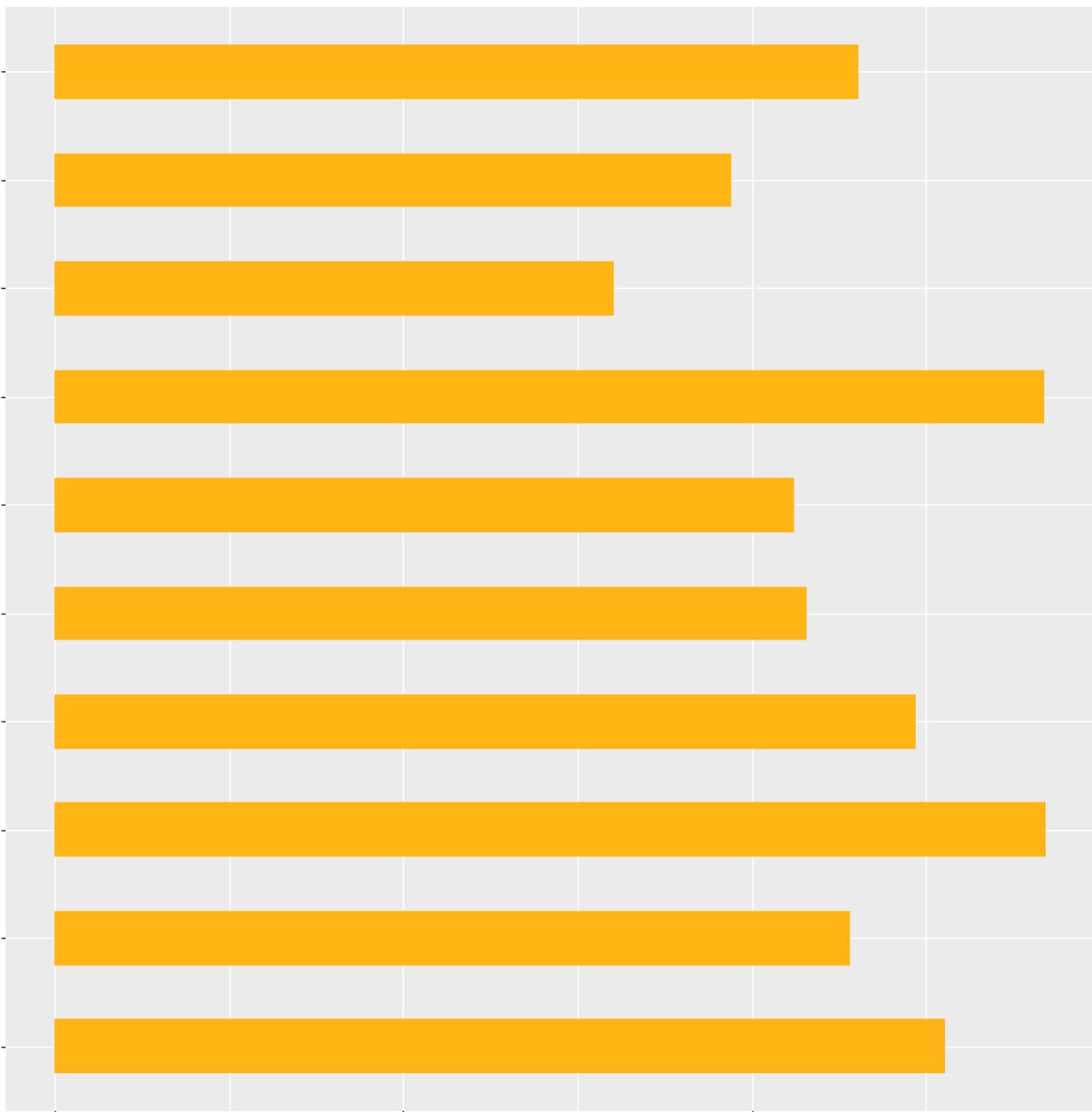
site of polarized growth
nucleus
membrane
Golgi apparatus
endoplasmic reticulum
endomembrane system
cytoskeleton
cytoplasmic vesicle
cellular bud
cell cortex

0.00

0.05

0.10

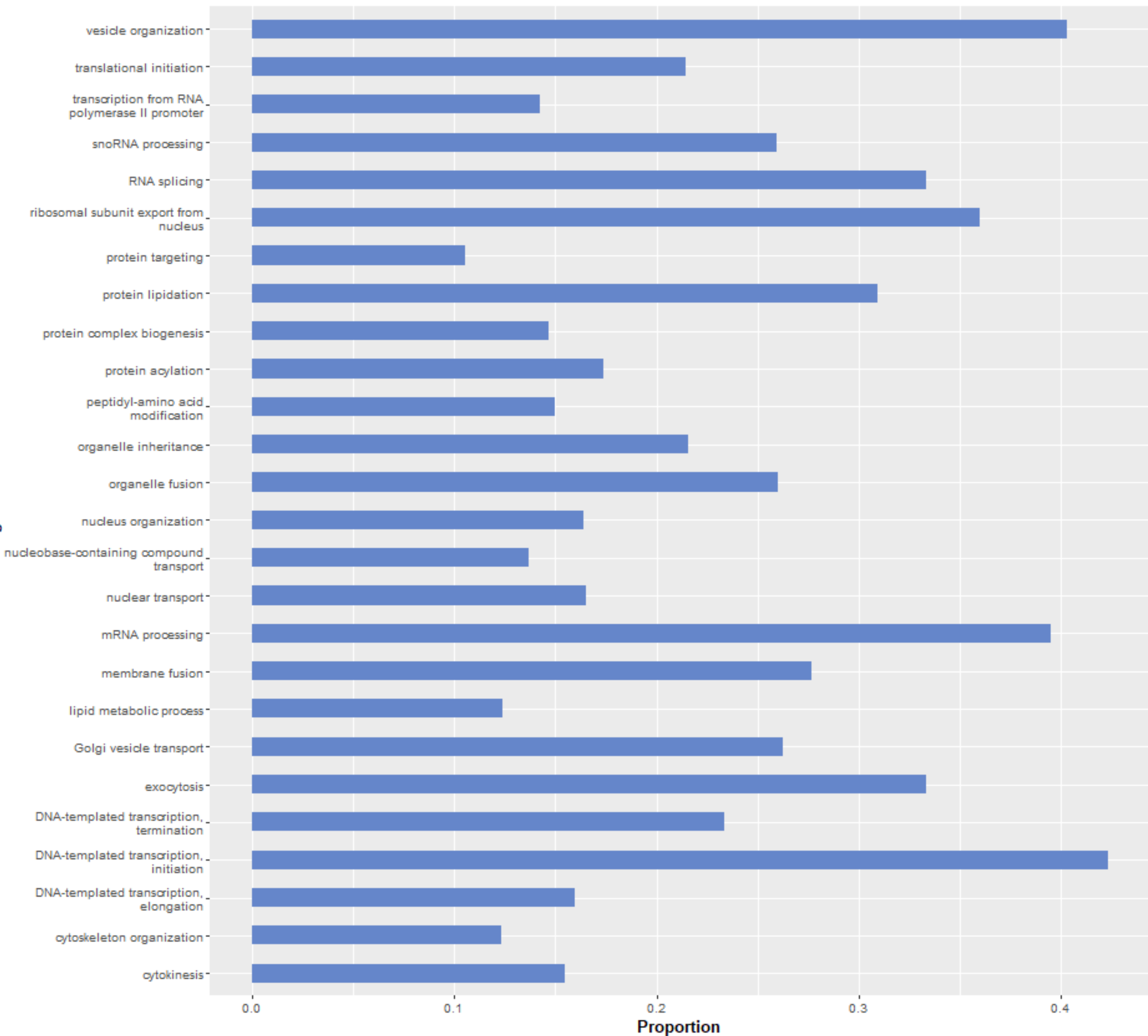
Proportion



With Cell Cycle | Without AreaShape | All Genes

Cluster 3 of 44

Biological Process



Cellular Compartment

nucleus

nucleolus

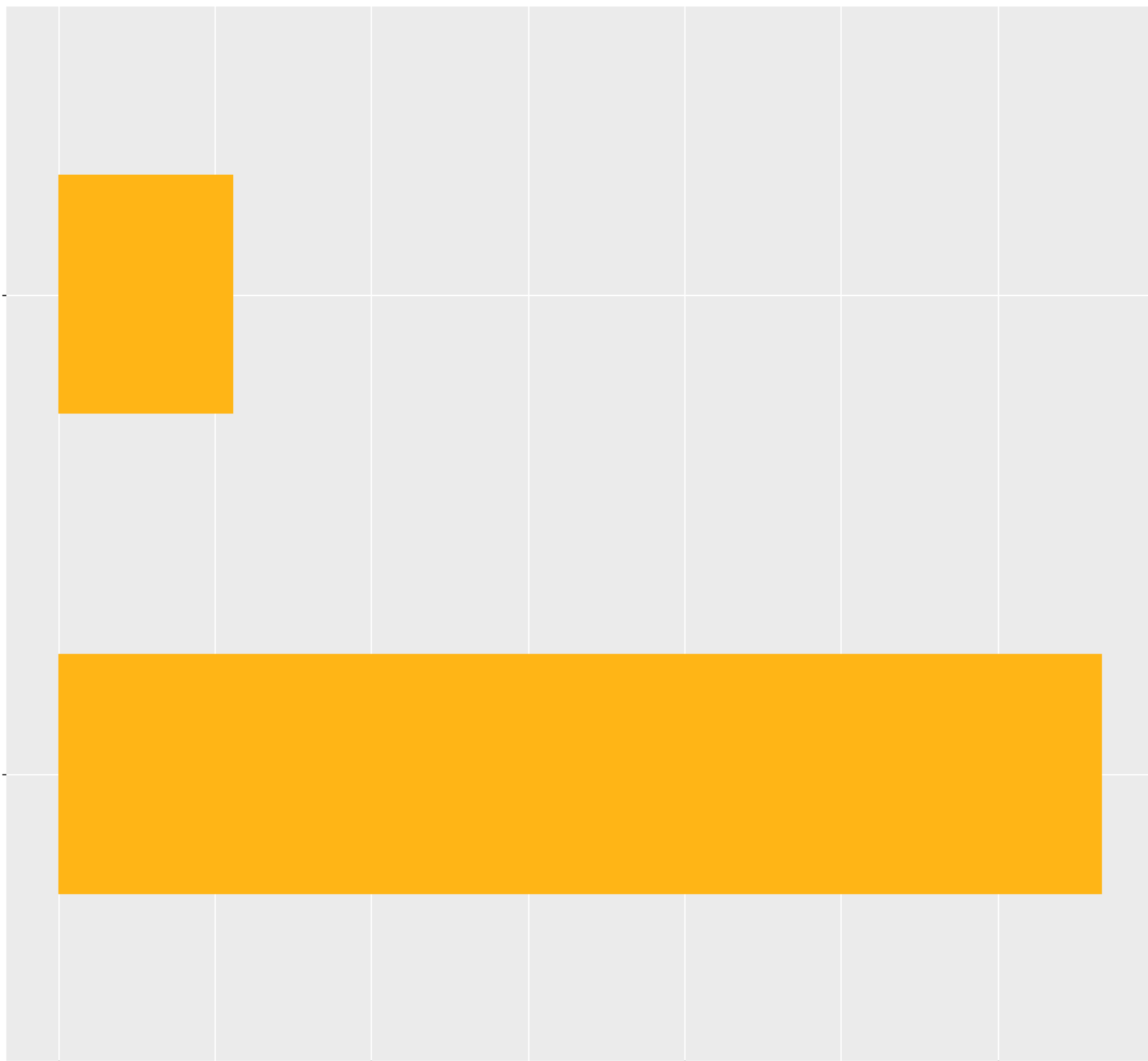
0.00

0.01

0.02

0.03

Proportion



With Cell Cycle | Without AreaShape | All Genes

Cluster 4 of 44

Biological Process

rRNA processing

RNA catabolic process

ribosome assembly

ribosomal large subunit
biogenesis

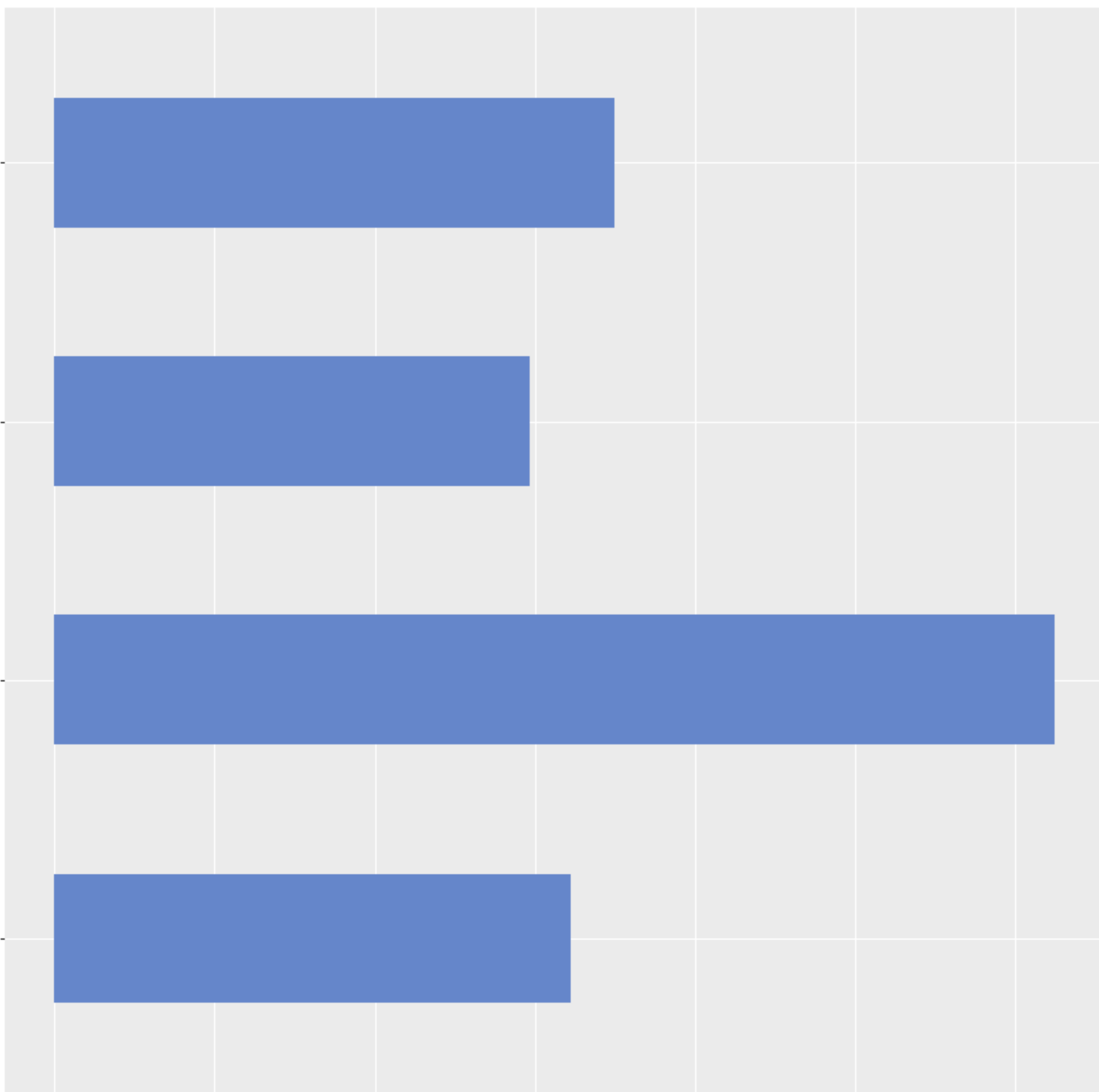
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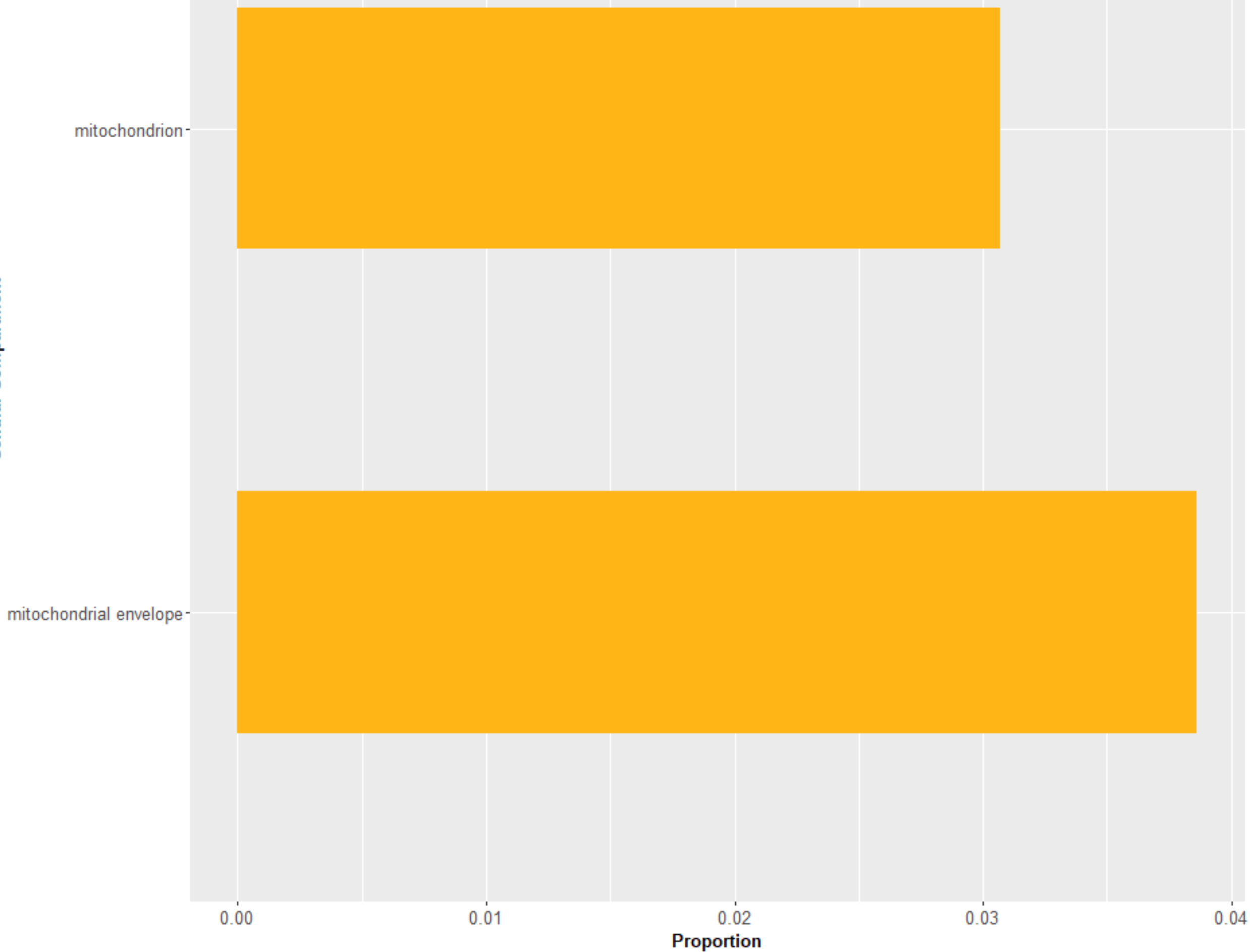
0.04

0.06

Proportion



Cellular Compartment



With Cell Cycle | Without AreaShape | All Genes

Cluster 9 of 44

Biological Process

nucleobase-containing small
molecule metabolic process

mitochondrion organization

mitochondrial translation

generation of precursor
metabolites and energy

cofactor metabolic process

carbohydrate metabolic process

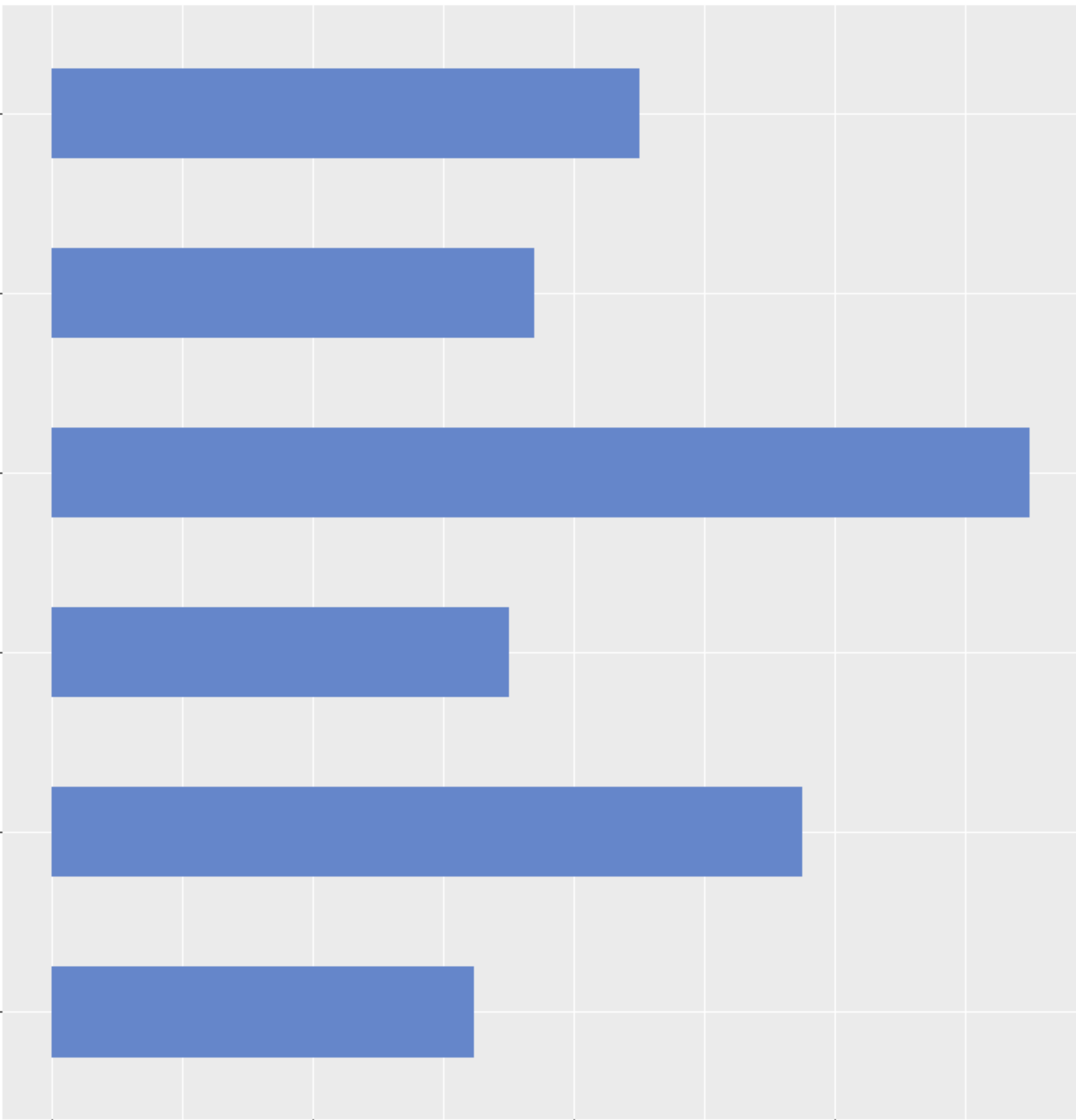
0.000

0.025

0.050

0.075

Proportion



With Cell Cycle | Without AreaShape | All Genes

Cluster 11 of 44

Cellular Compartment

nucleus

0.000

0.005

0.010

0.015

Proportion



With Cell Cycle | Without AreaShape | All Genes

Cluster 11 of 44

Biological Process

tRNA aminoacylation for
protein translation

transcription from RNA
polymerase II promoter

RNA splicing

mRNA processing

DNA-templated transcription,
initiation

chromosome segregation

0.00

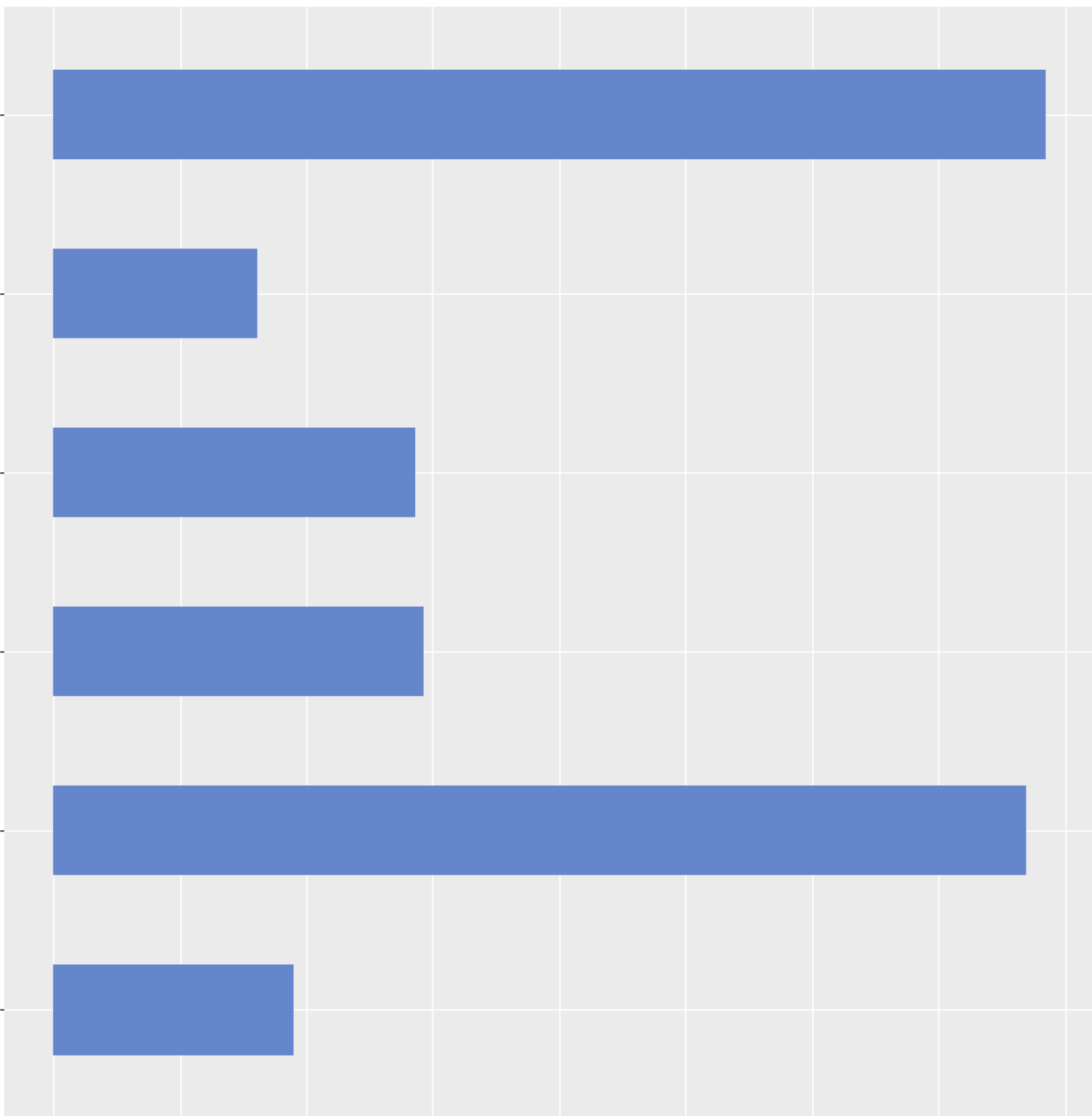
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0.06

0.09

0.12

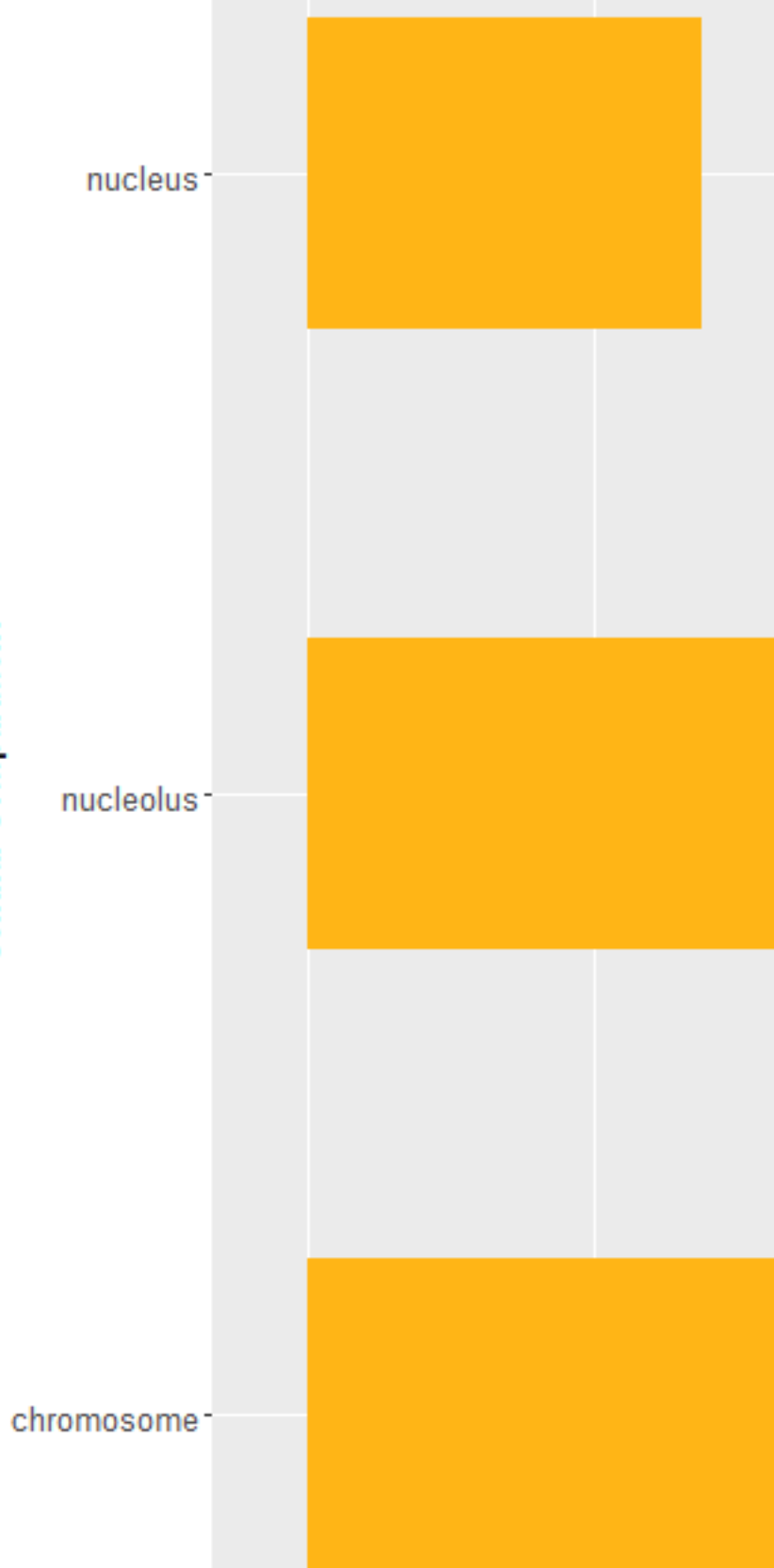
Proportion



With Cell Cycle | Without AreaShape | All Genes

Cluster 15 of 44

Cellular Compartment



With Cell Cycle | Without AreaShape | All Genes

Cluster 15 of 44

Biological Process

snoRNA processing

rRNA processing

DNA replication

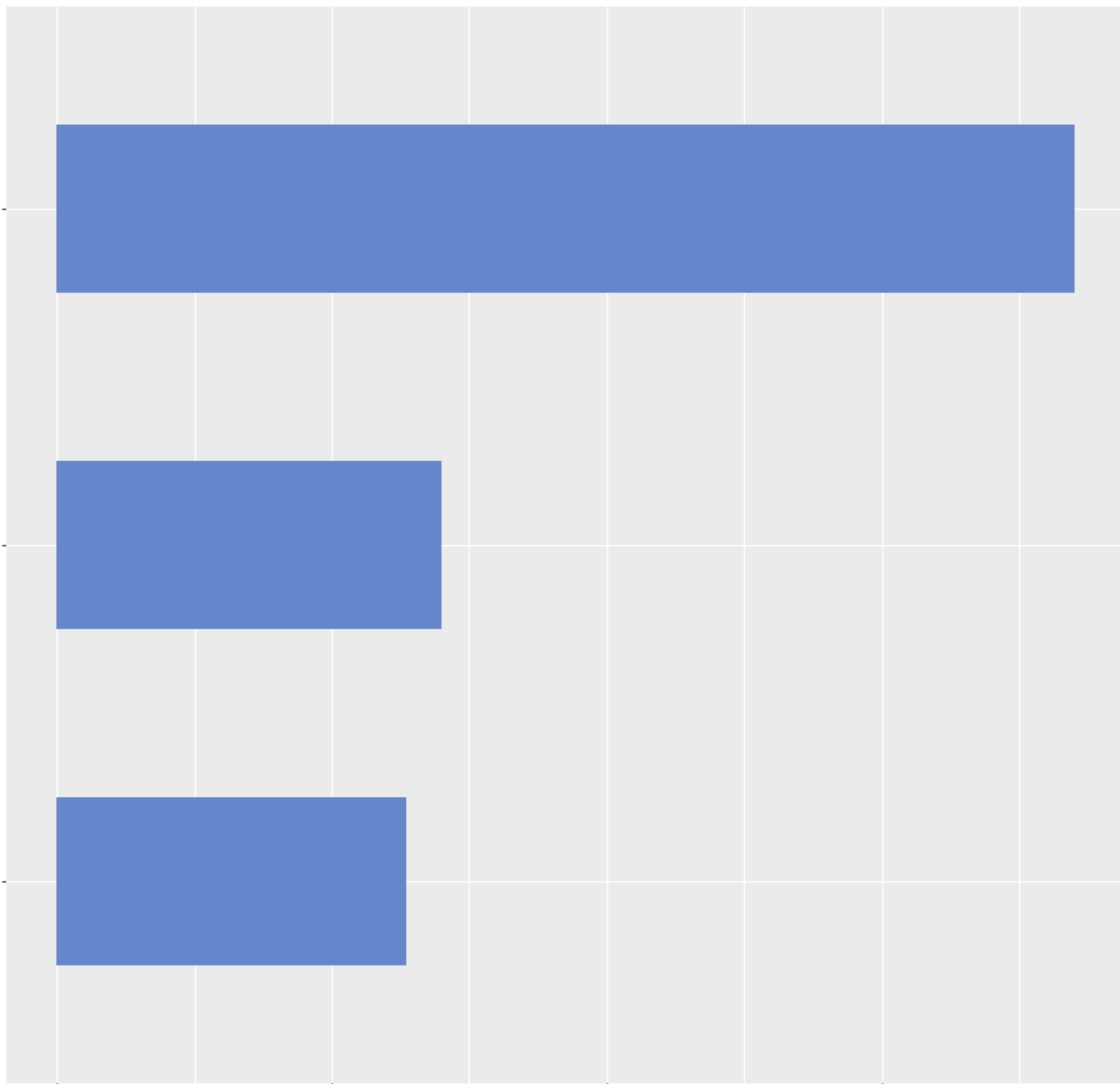
0.00

0.02

0.04

0.06

Proportion



With Cell Cycle | Without AreaShape | All Genes

Cluster 16 of 44

Cellular Compartment

nucleus

0.000

0.005

0.010

0.015

0.020

Proportion



With Cell Cycle | Without AreaShape | All Genes

Cluster 16 of 44

Biological Process

rRNA processing

ribosomal large subunit
biogenesis

nuclear transport

DNA-templated transcription,
termination

cytoskeleton organization

0.000

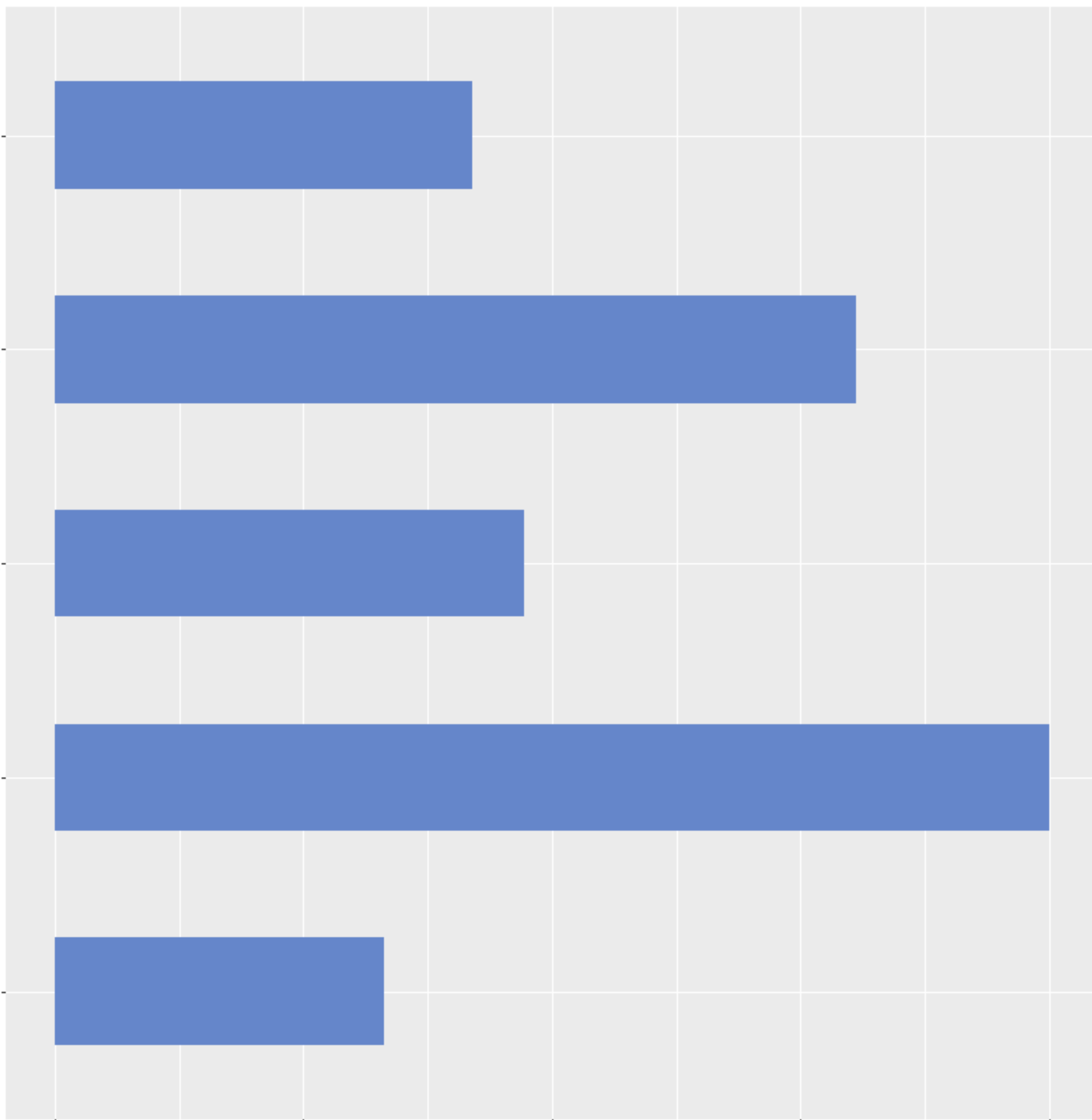
0.025

0.050

0.075

0.100

Proportion



Biological Process

response to starvation

0.00

0.01

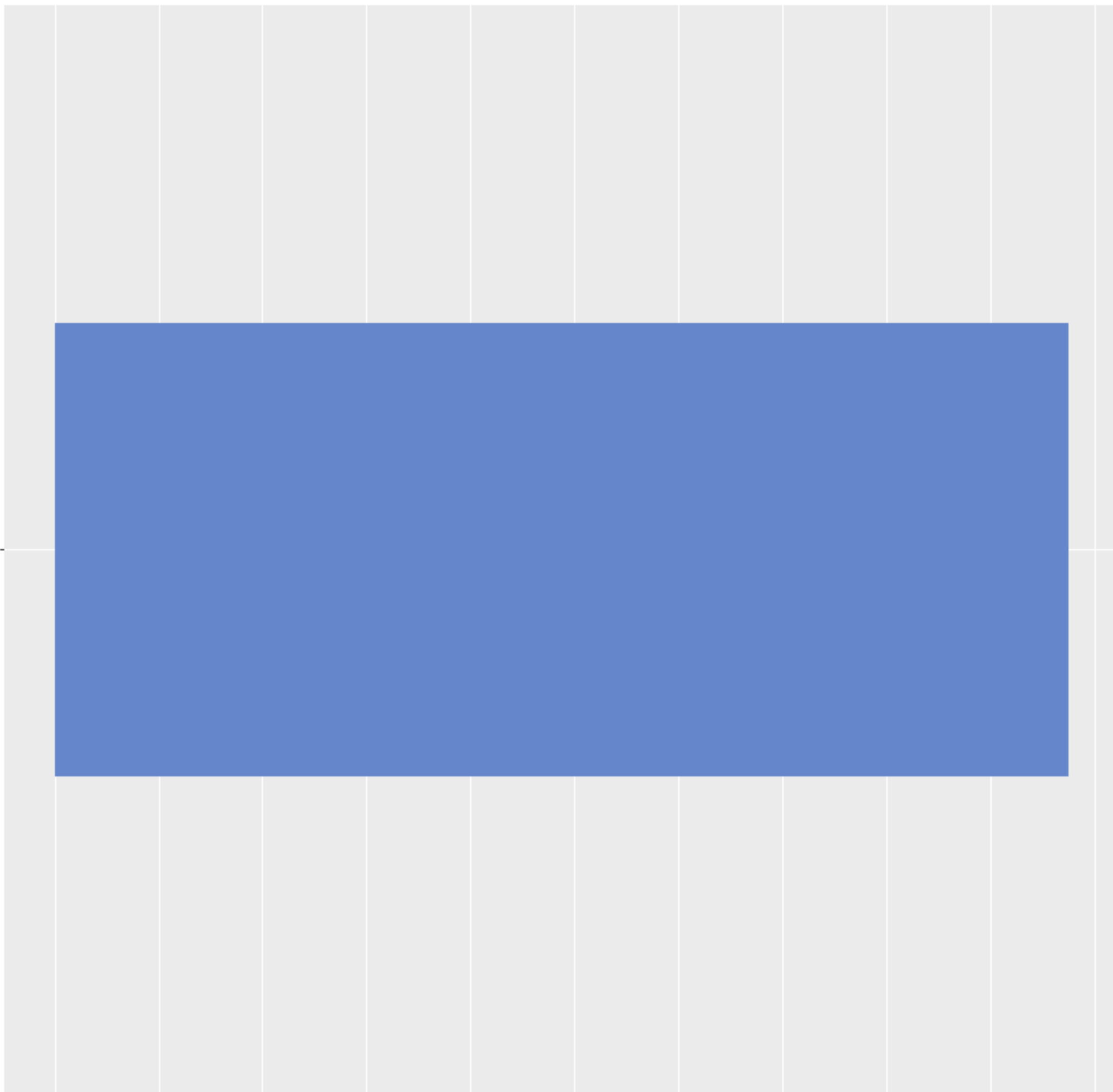
0.02

0.03

0.04

0.05

Proportion



Biological Process

response to chemical

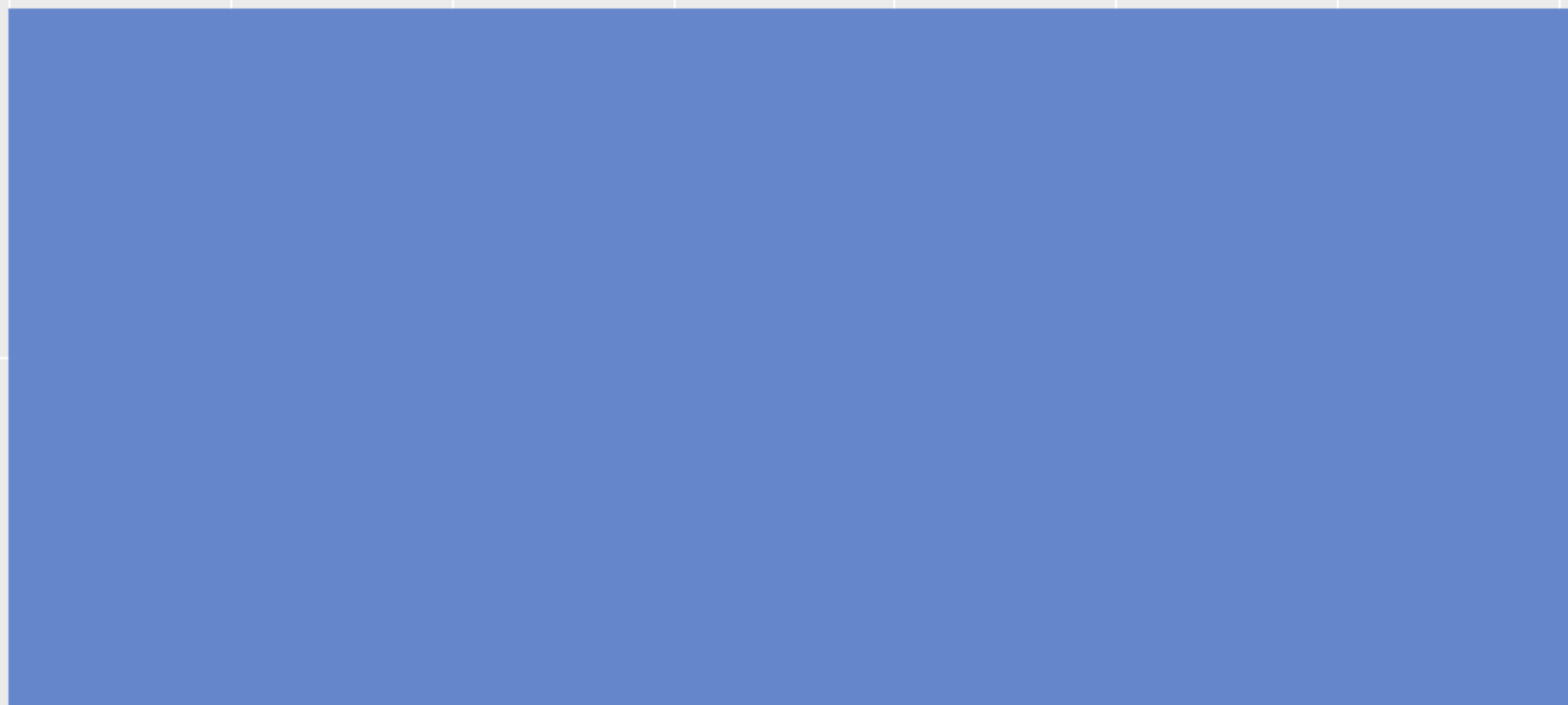
0.000

0.003

0.006

0.009

Proportion



With Cell Cycle | Without AreaShape | All Genes

Cluster 22 of 44

Biological Process

protein modification by small
protein conjugation or removal

peptidyl-amino acid
modification

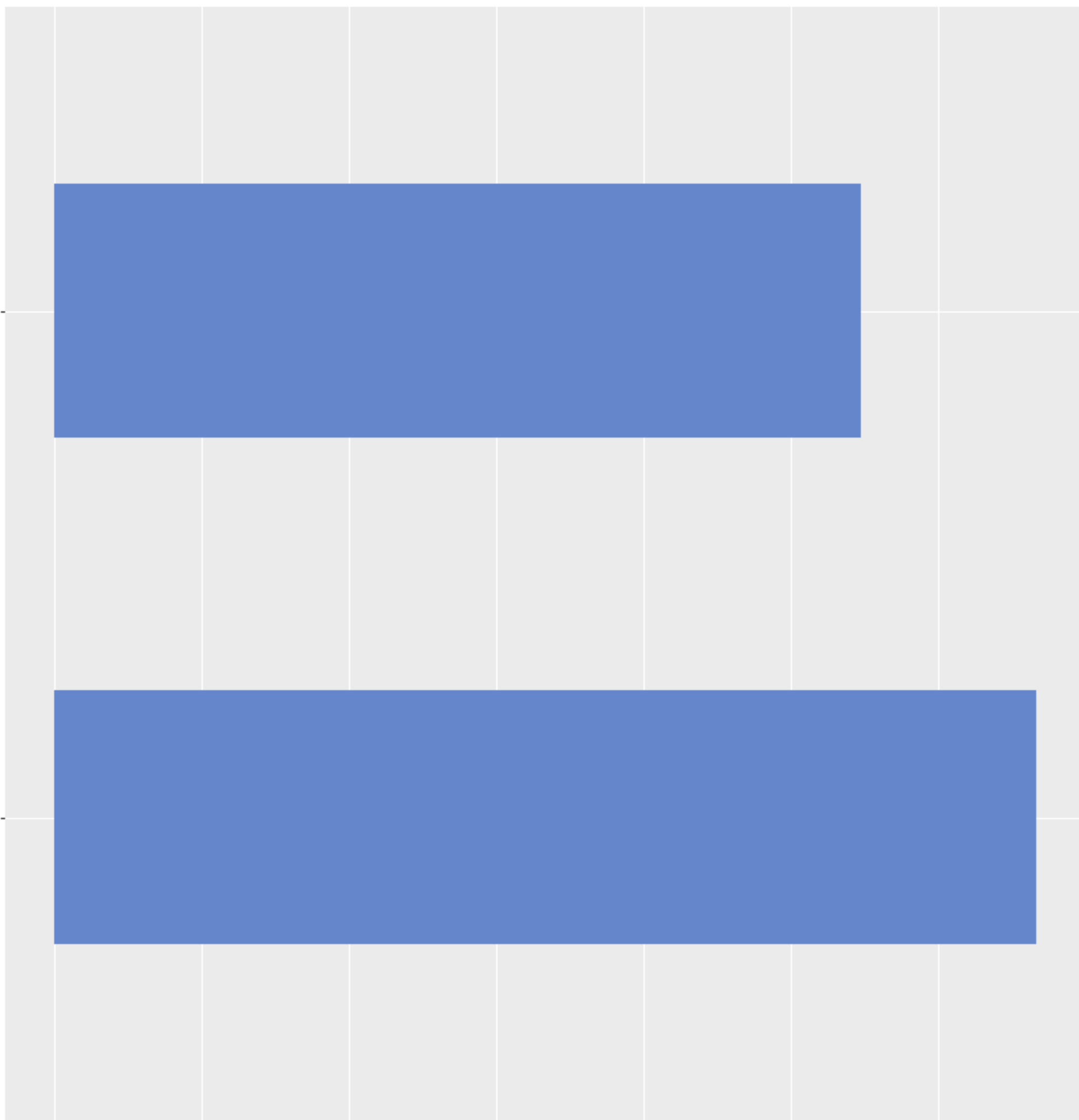
0.00

0.01

0.02

0.03

Proportion



With Cell Cycle | Without AreaShape | All Genes

Cluster 23 of 44

Cellular Compartment

cell cortex

0.00

0.01

0.02

0.03

0.04

Proportion



With Cell Cycle | Without AreaShape | All Genes

Cluster 24 of 44

Cellular Compartment

vacuole

site of polarized growth

cellular bud

0.000

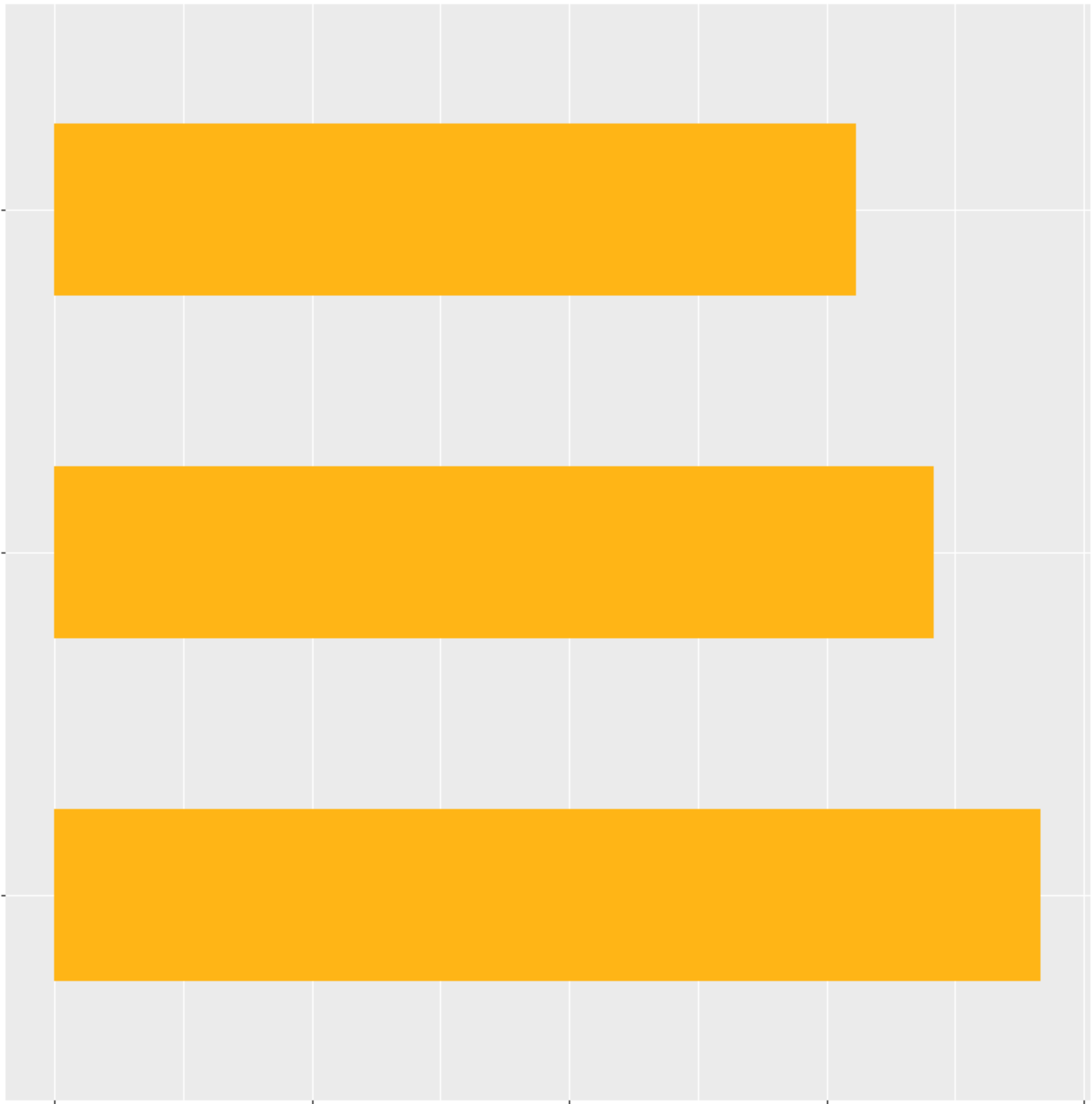
0.025

0.050

0.075

0.100

Proportion



With Cell Cycle | Without AreaShape | All Genes

Cluster 24 of 44

Biological Process

regulation of transport

chromatin organization

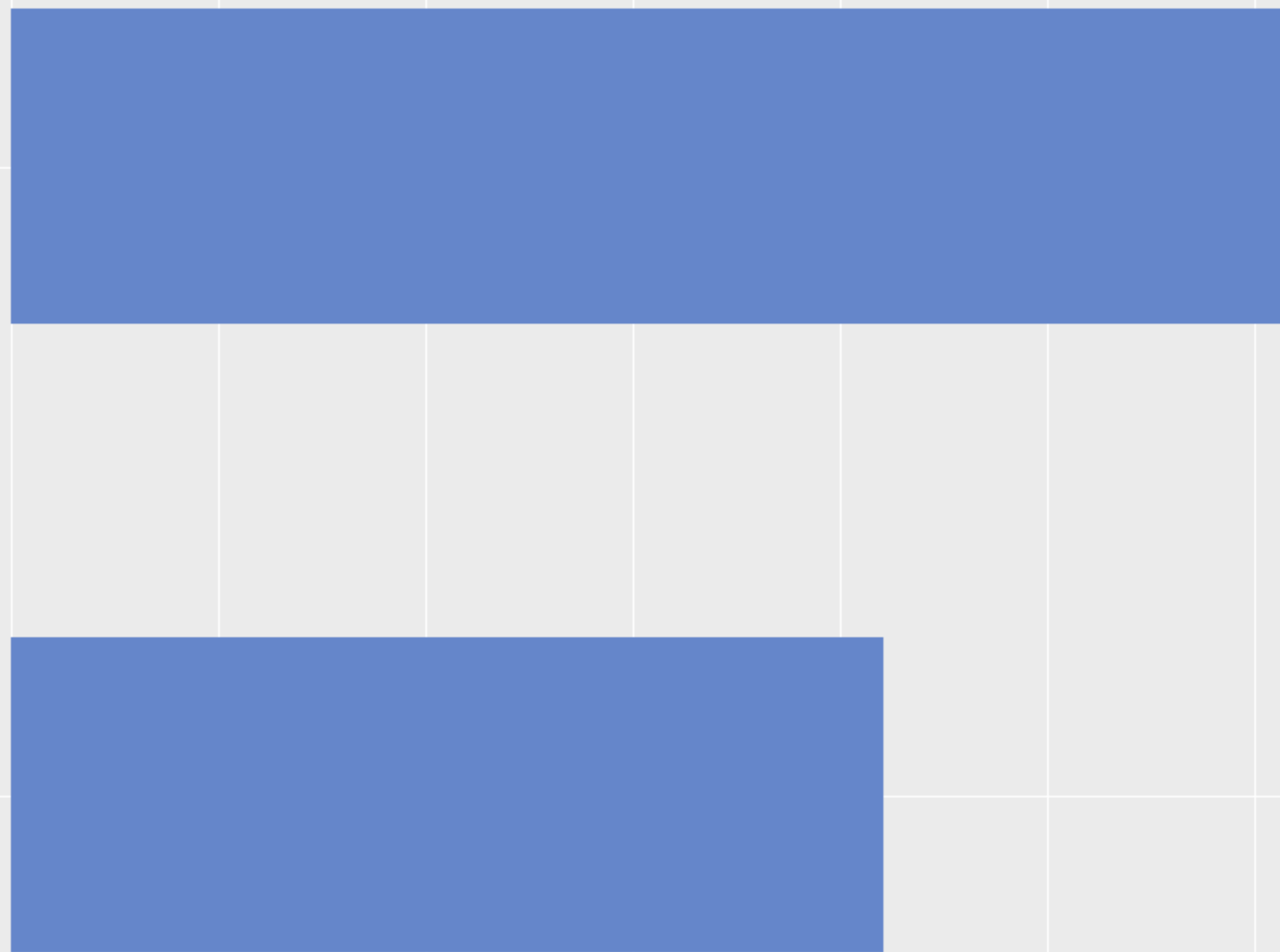
0.00

0.04

0.08

0.12

Proportion



With Cell Cycle | Without AreaShape | All Genes

Cluster 27 of 44

Cellular Compartment

nucleus

0.000

0.005

0.010

0.015

0.020

0.025

Proportion



With Cell Cycle | Without AreaShape | All Genes

Cluster 27 of 44

Biological Process

ribosomal small subunit
biogenesis

protein maturation

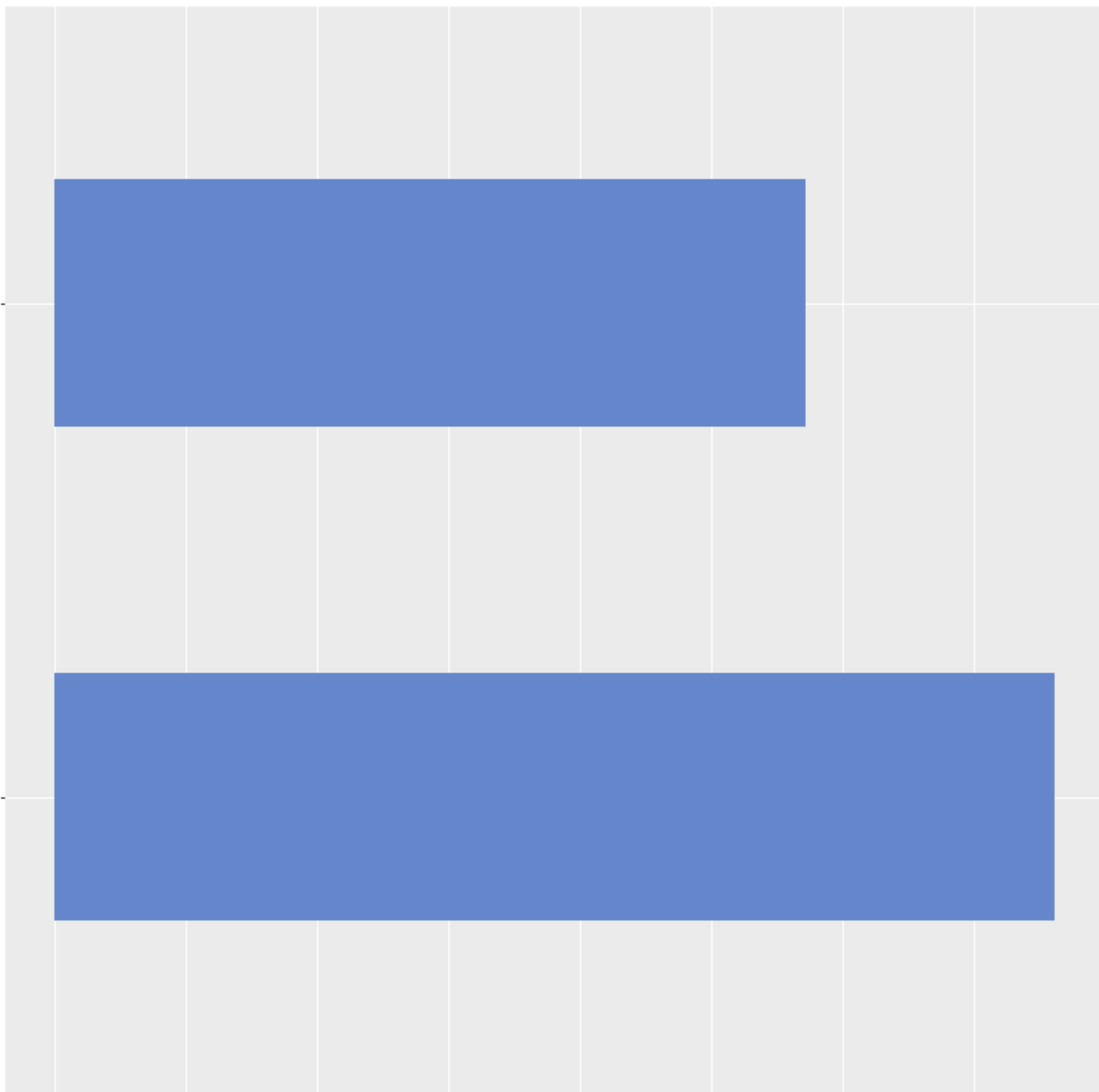
0.000

0.025

0.050

0.075

Proportion



With Cell Cycle | Without AreaShape | All Genes

Cluster 28 of 44

Biological Process

organelle fission

cytoplasmic translation

0.00

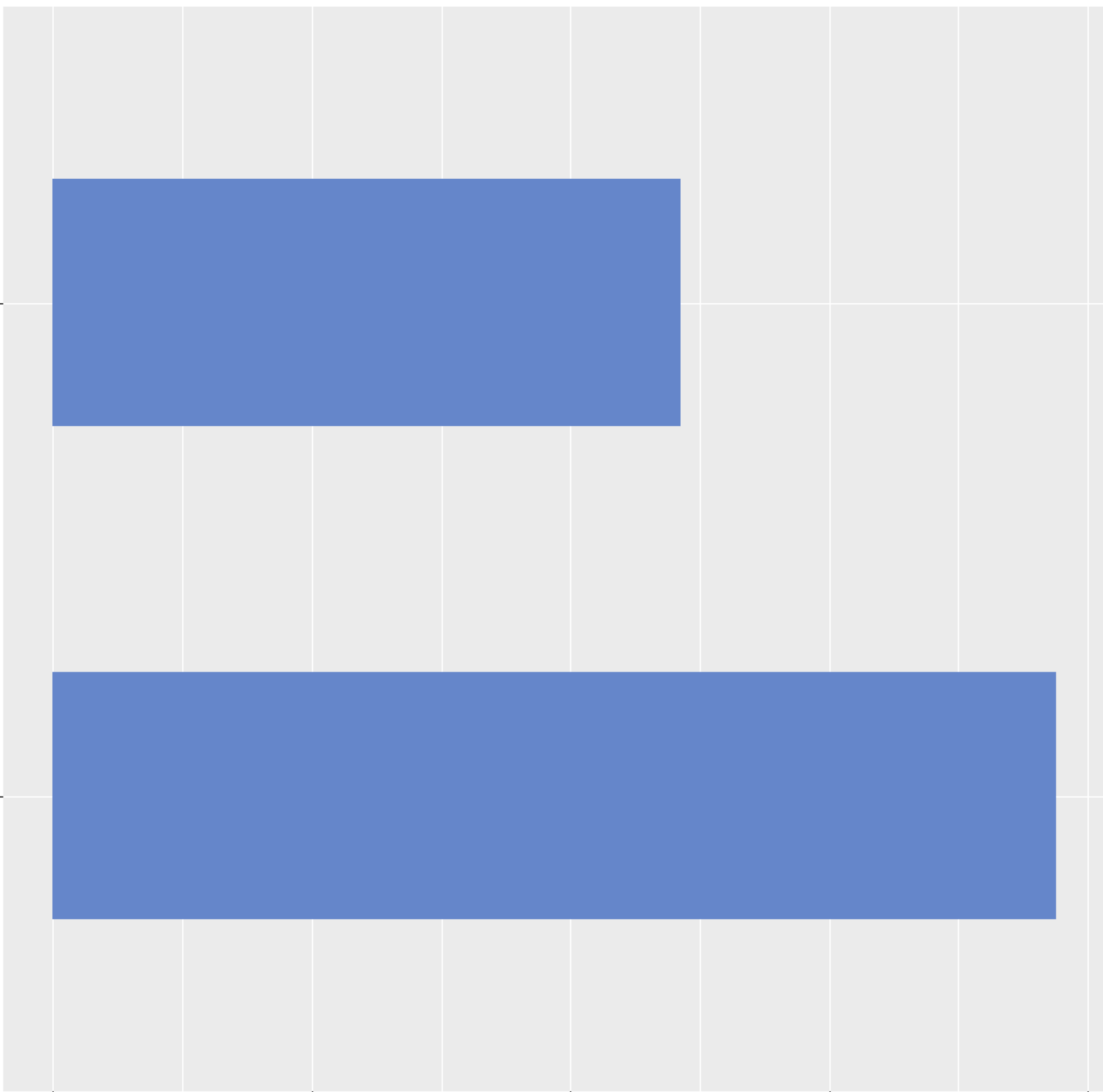
0.02

0.04

0.06

0.08

Proportion



With Cell Cycle | Without AreaShape | All Genes

Cluster 30 of 44

Cellular Compartment

site of polarized growth

microtubule organizing center

cytoskeleton

cellular bud

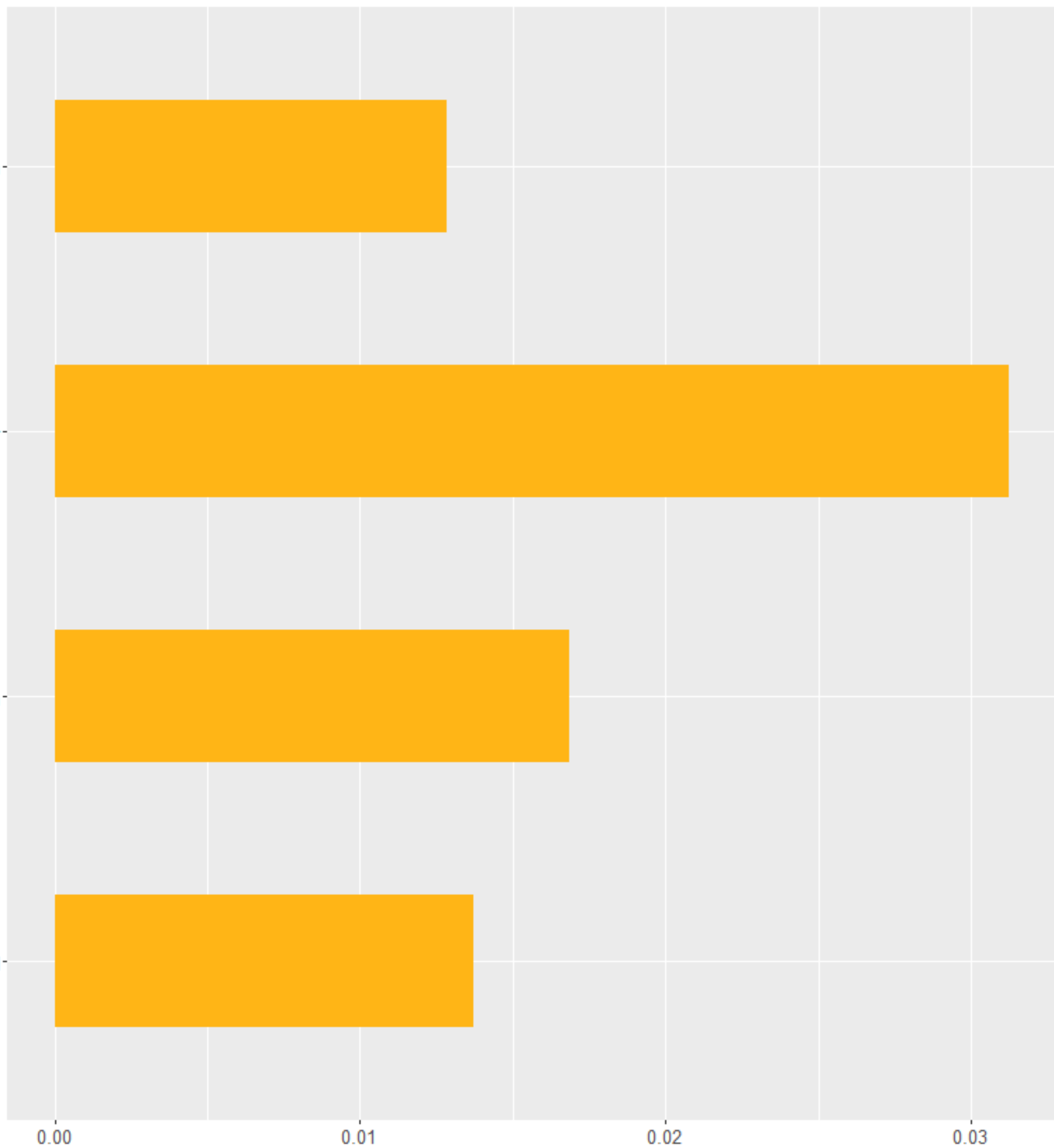
0.00

0.01

0.02

0.03

Proportion



With Cell Cycle | Without AreaShape | All Genes

Cluster 30 of 44

Biological Process

regulation of protein
modification process

regulation of cell cycle

protein phosphorylation

cytoskeleton organization

cytokinesis

cell budding

0.00

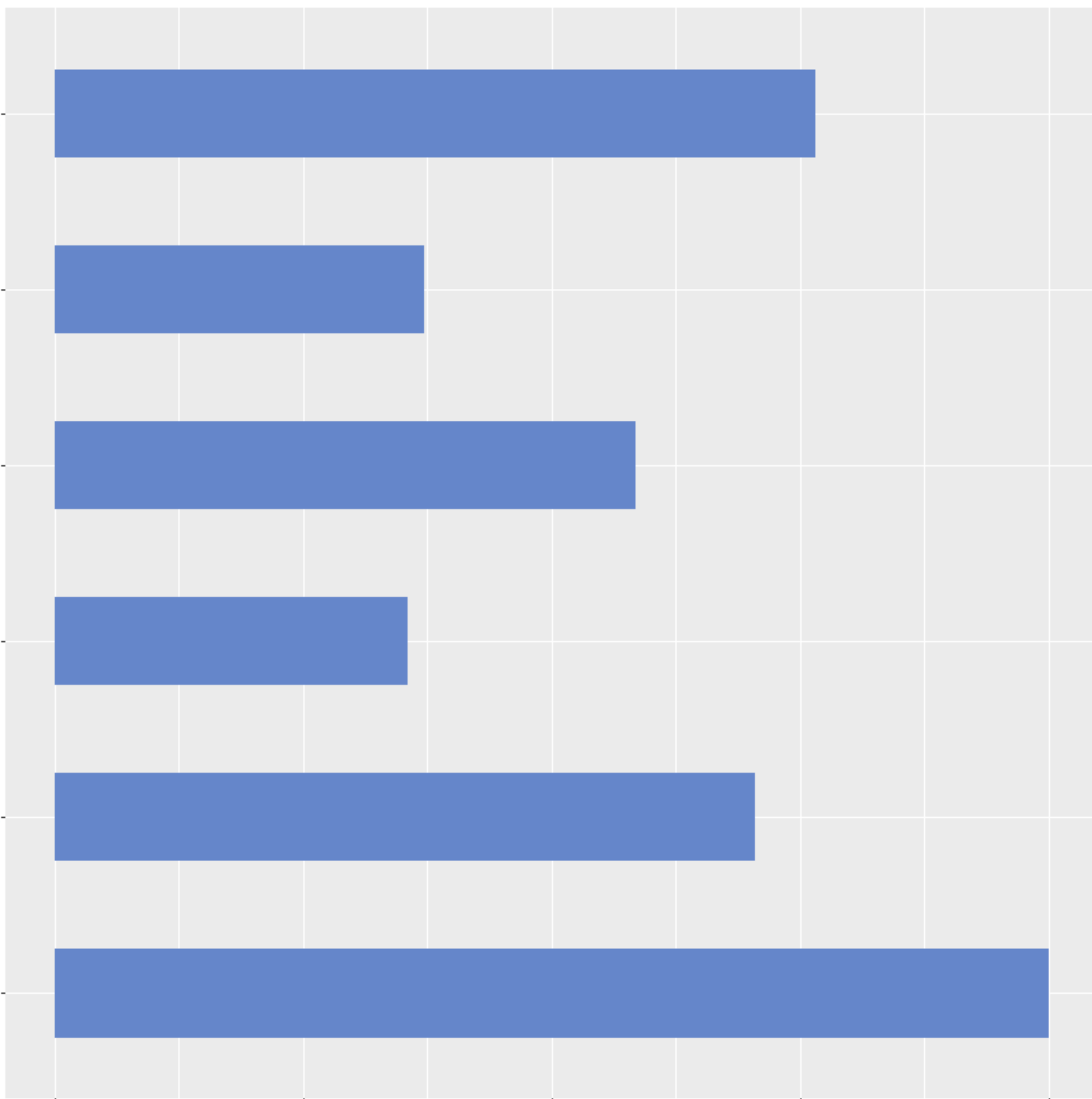
0.01

0.02

0.03

0.04

Proportion



With Cell Cycle | Without AreaShape | All Genes

Cluster 33 of 44

Biological Process

translational elongation

transcription from RNA
polymerase II promoter

transcription from RNA
polymerase I promoter

response to heat

protein acylation

endosomal transport

cytoplasmic translation

chromatin organization

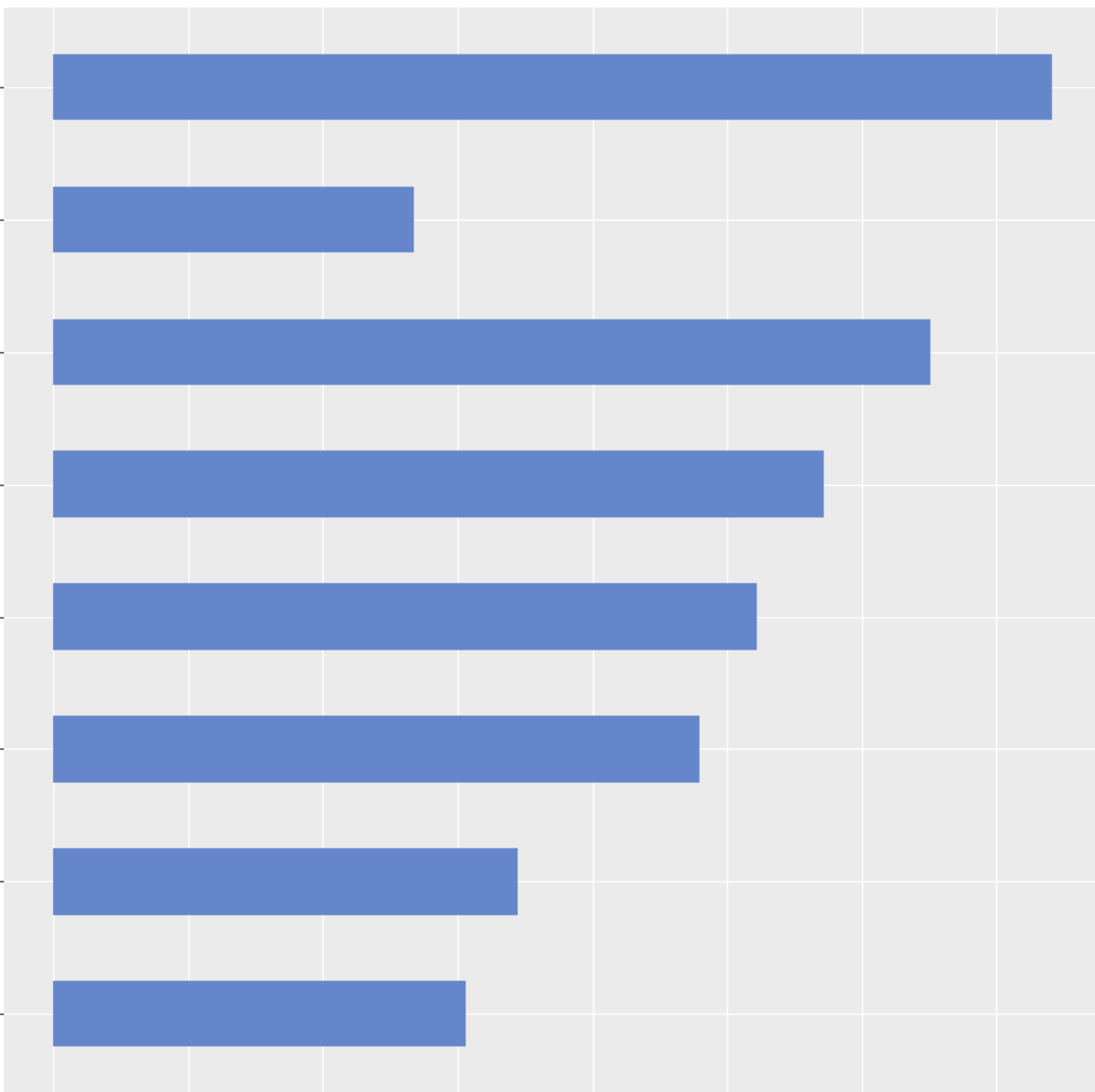
0.00

0.05

0.10

0.15

Proportion



With Cell Cycle | Without AreaShape | All Genes

Cluster 34 of 44

Biological Process

protein modification by small
protein conjugation or removal

histone modification

0.00

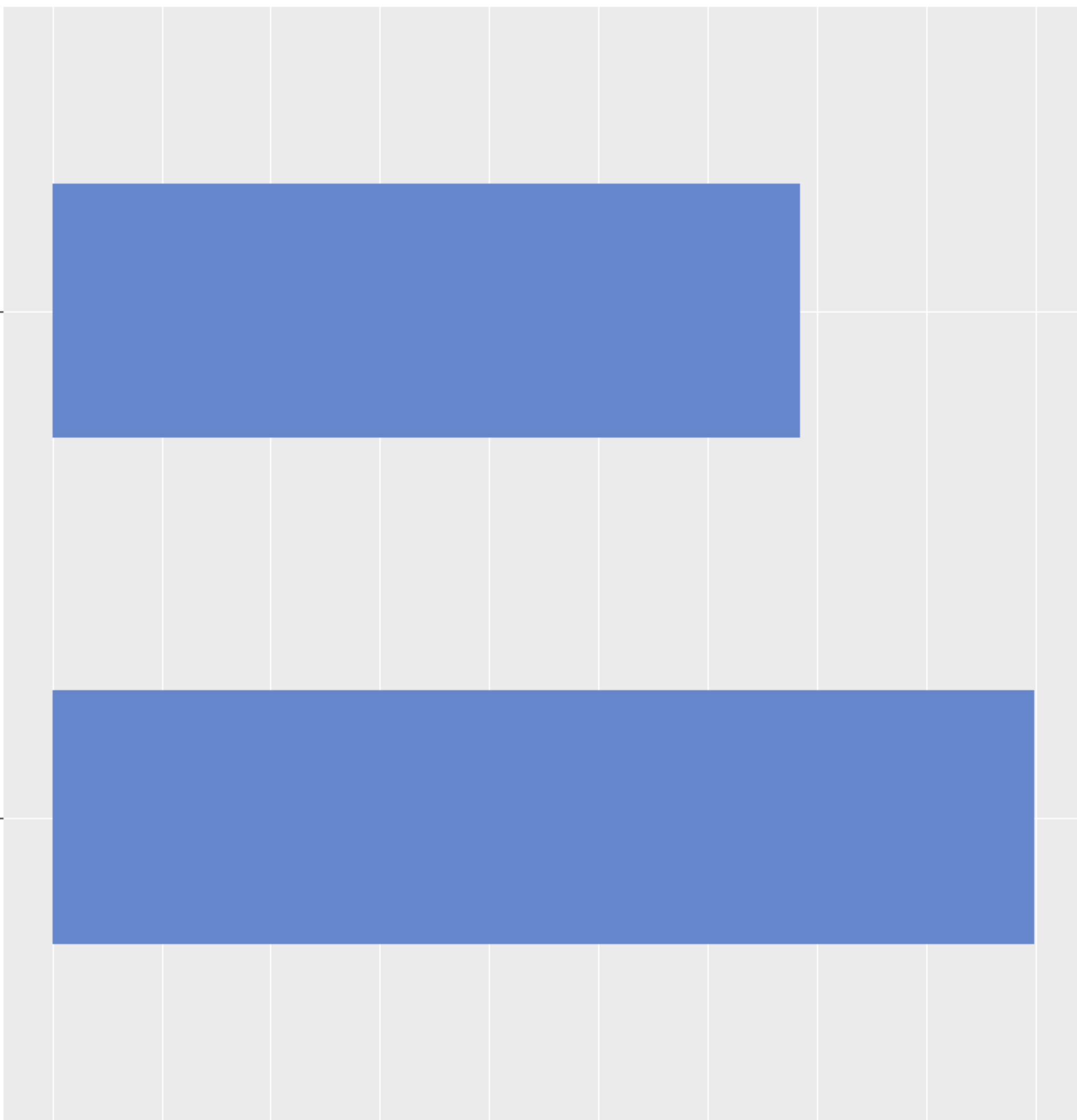
0.01

0.02

0.03

0.04

Proportion



With Cell Cycle | Without AreaShape | All Genes

Cluster 35 of 44

Cellular Compartment

nucleus

chromosome

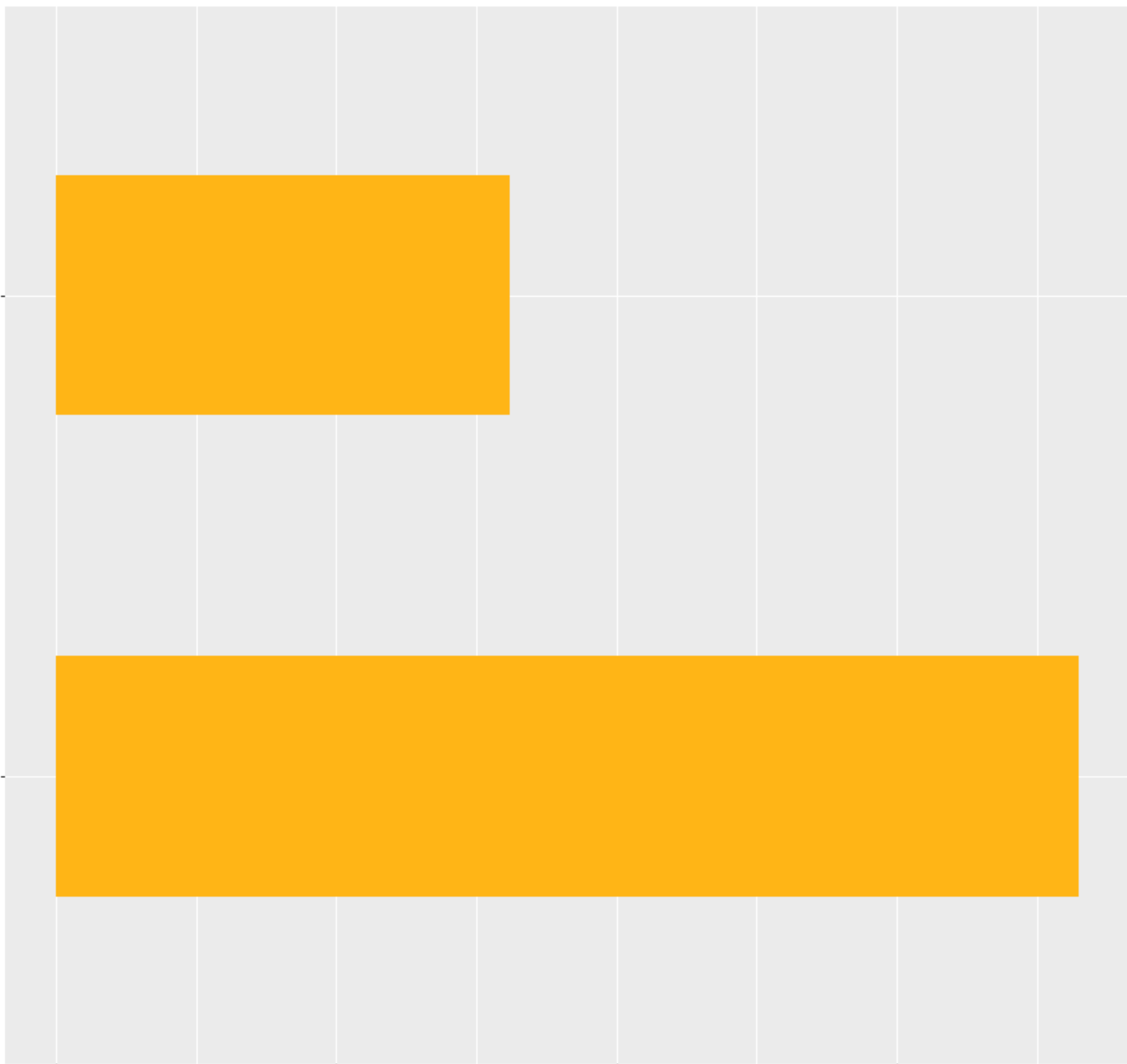
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0.010

0.015

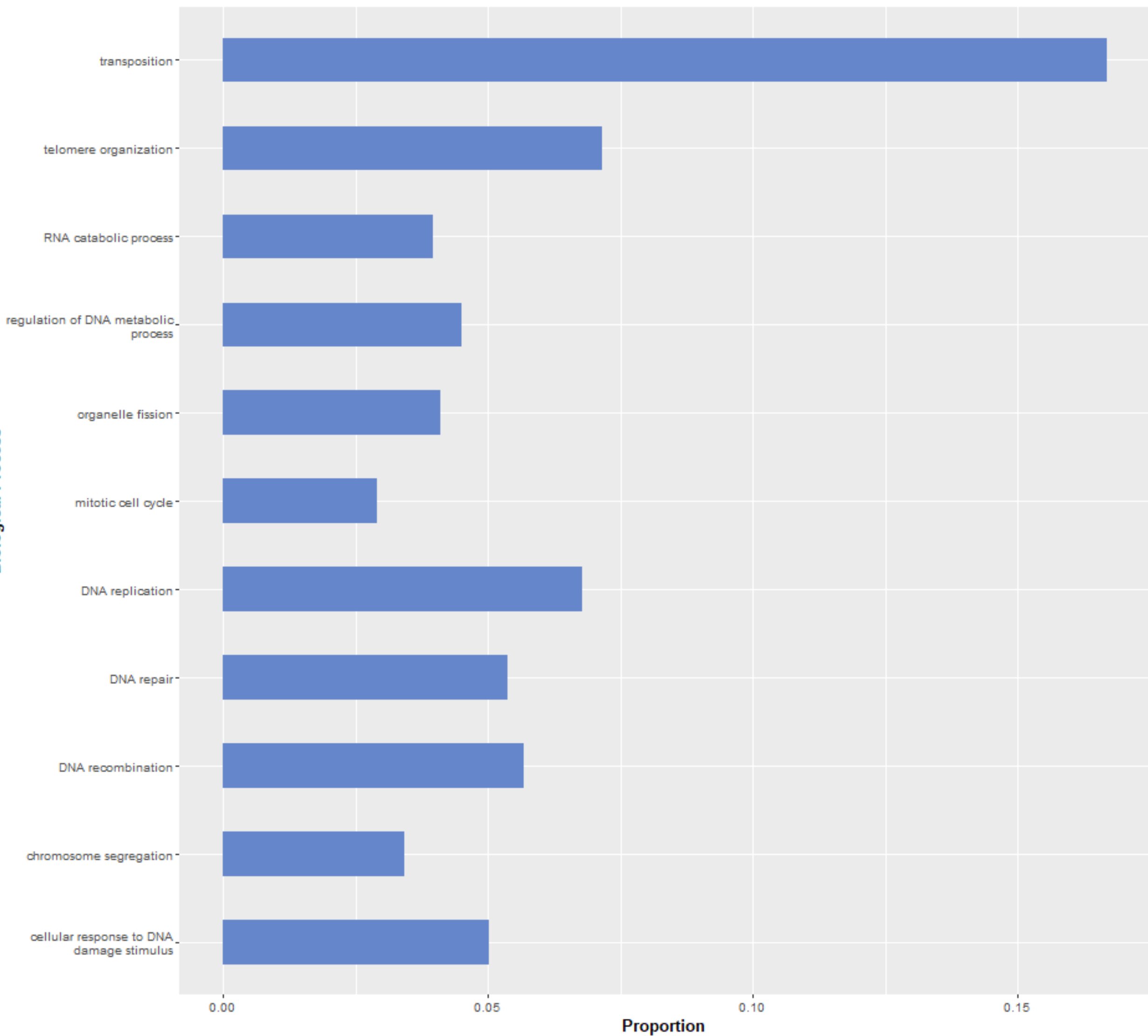
Proportion



With Cell Cycle | Without AreaShape | All Genes

Cluster 35 of 44

Biological Process



With Cell Cycle | Without AreaShape | All Genes

Cluster 42 of 44

Biological Process

cellular amino acid metabolic
process

0.00

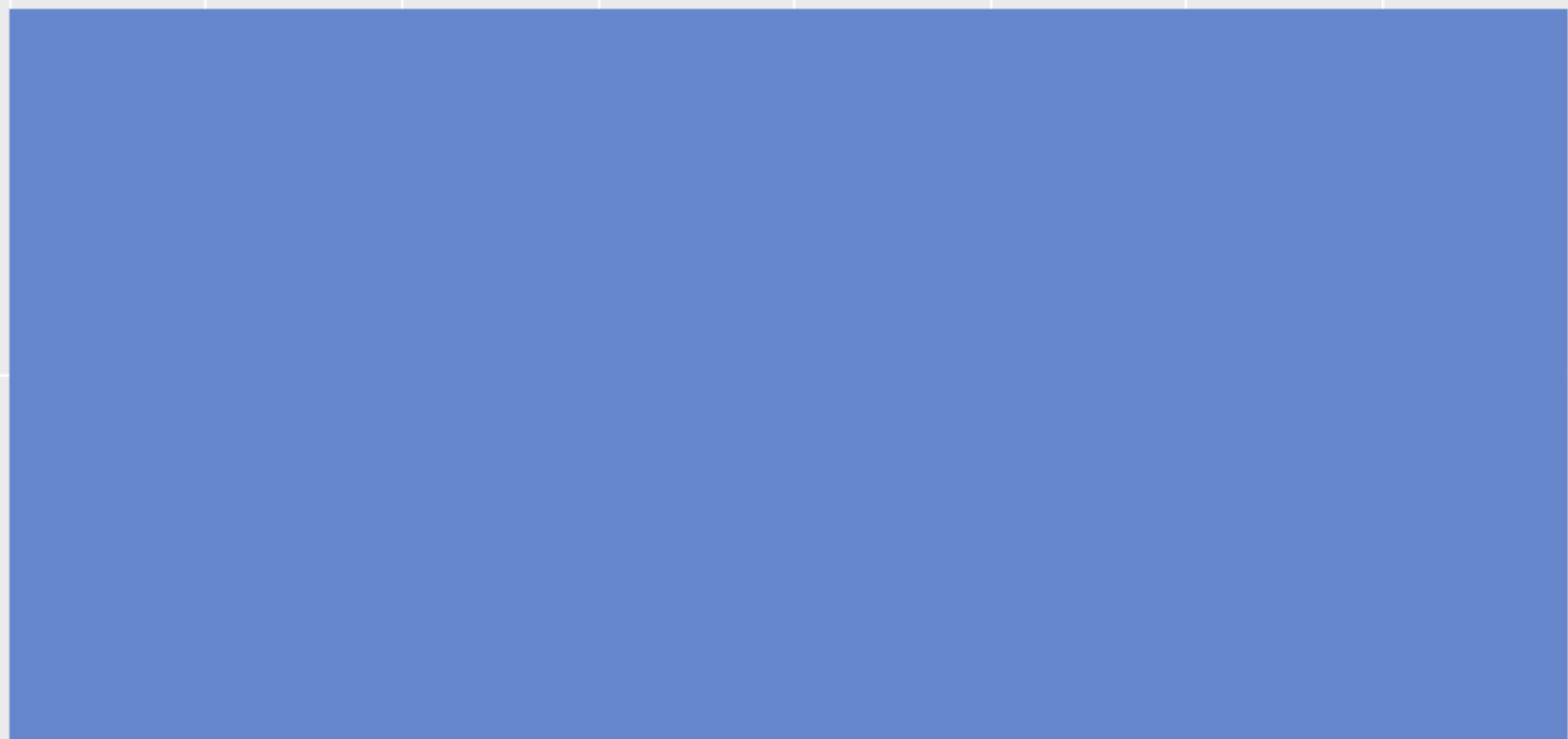
0.01

0.02

0.03

0.04

Proportion



Cellular Compartment

mitochondrial envelope

0.00

0.01

0.02

0.03

Proportion

