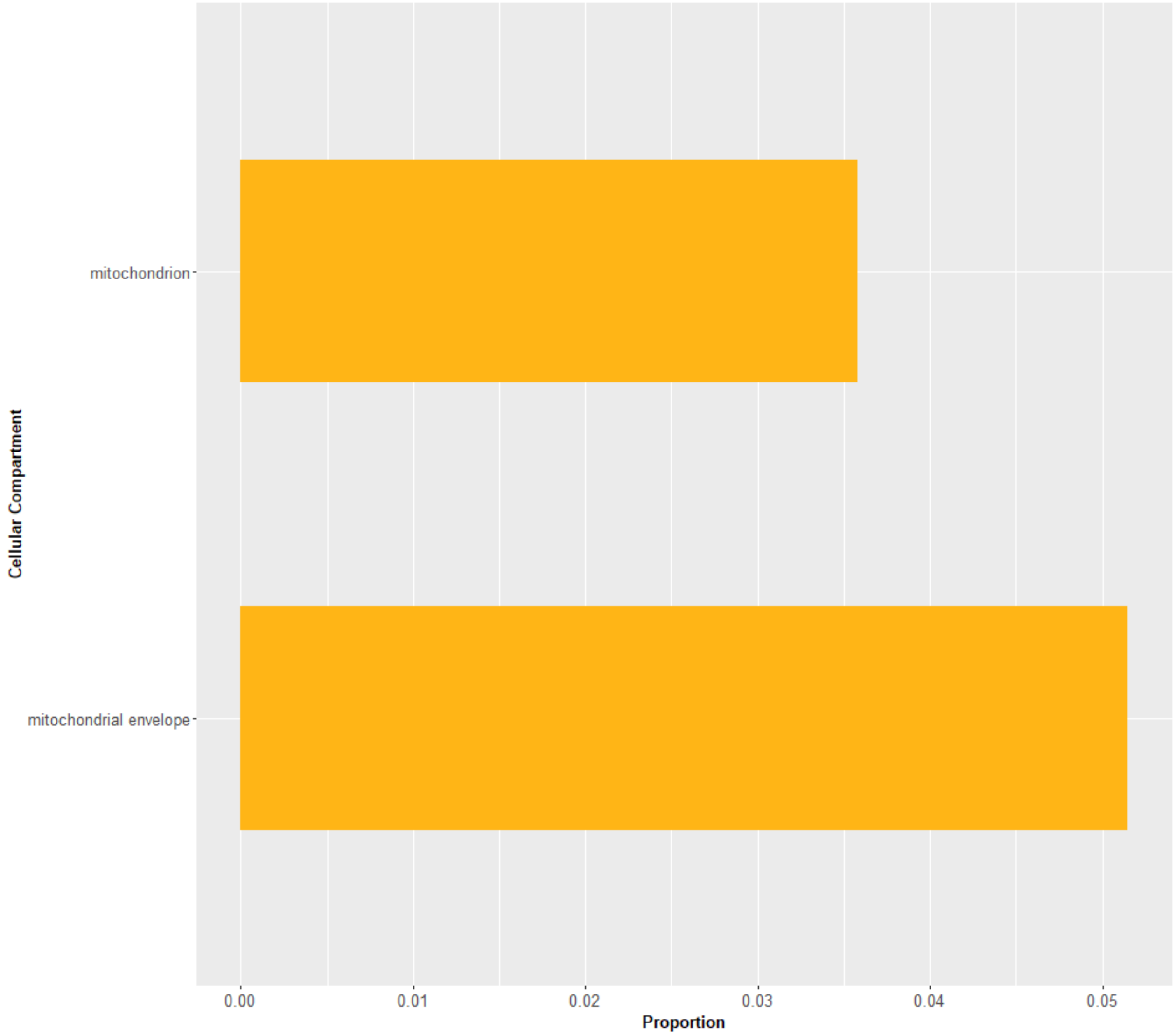
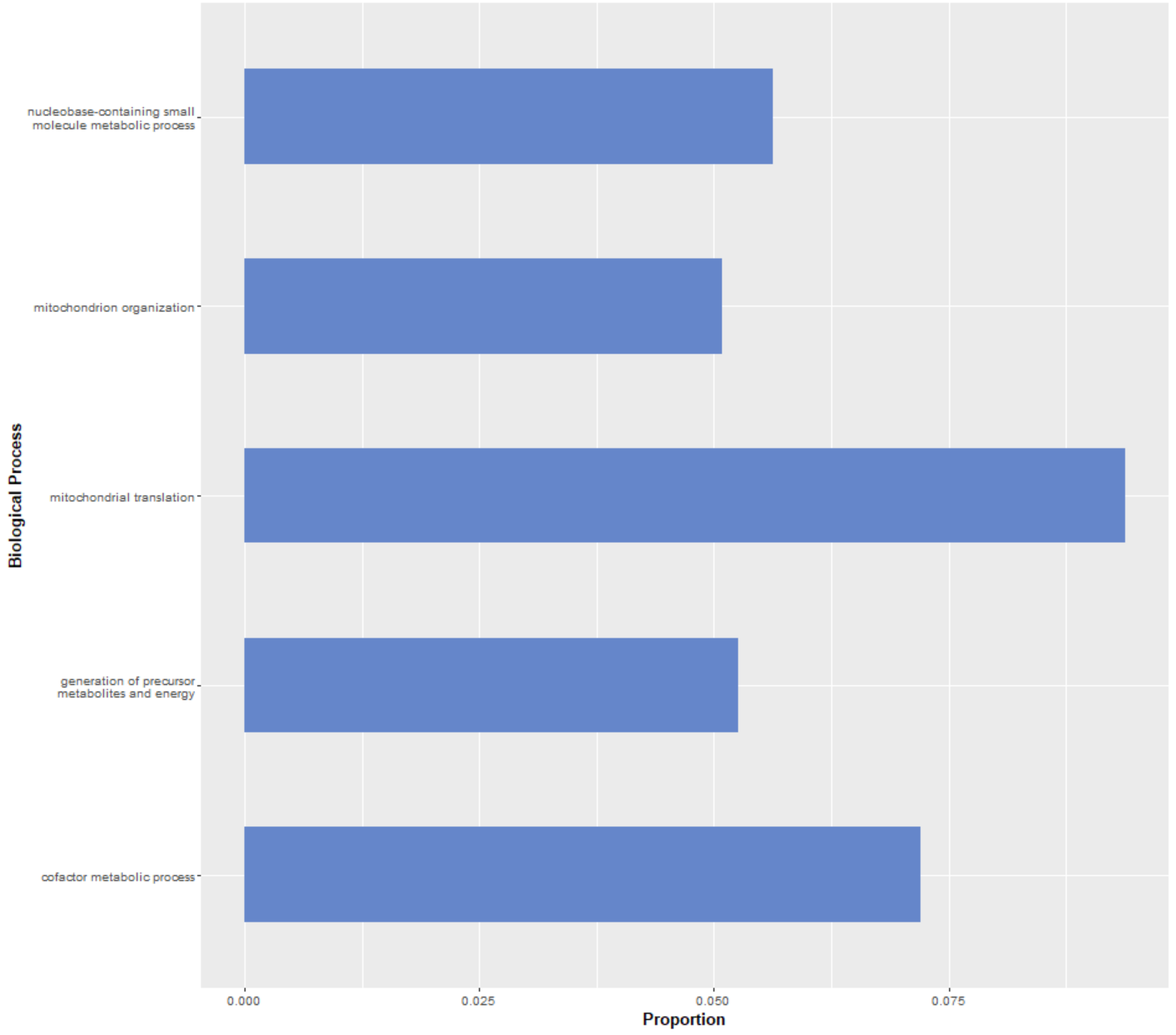
Without Cell Cycle | Without AreaShape | All Genes

Cluster 2 of 42



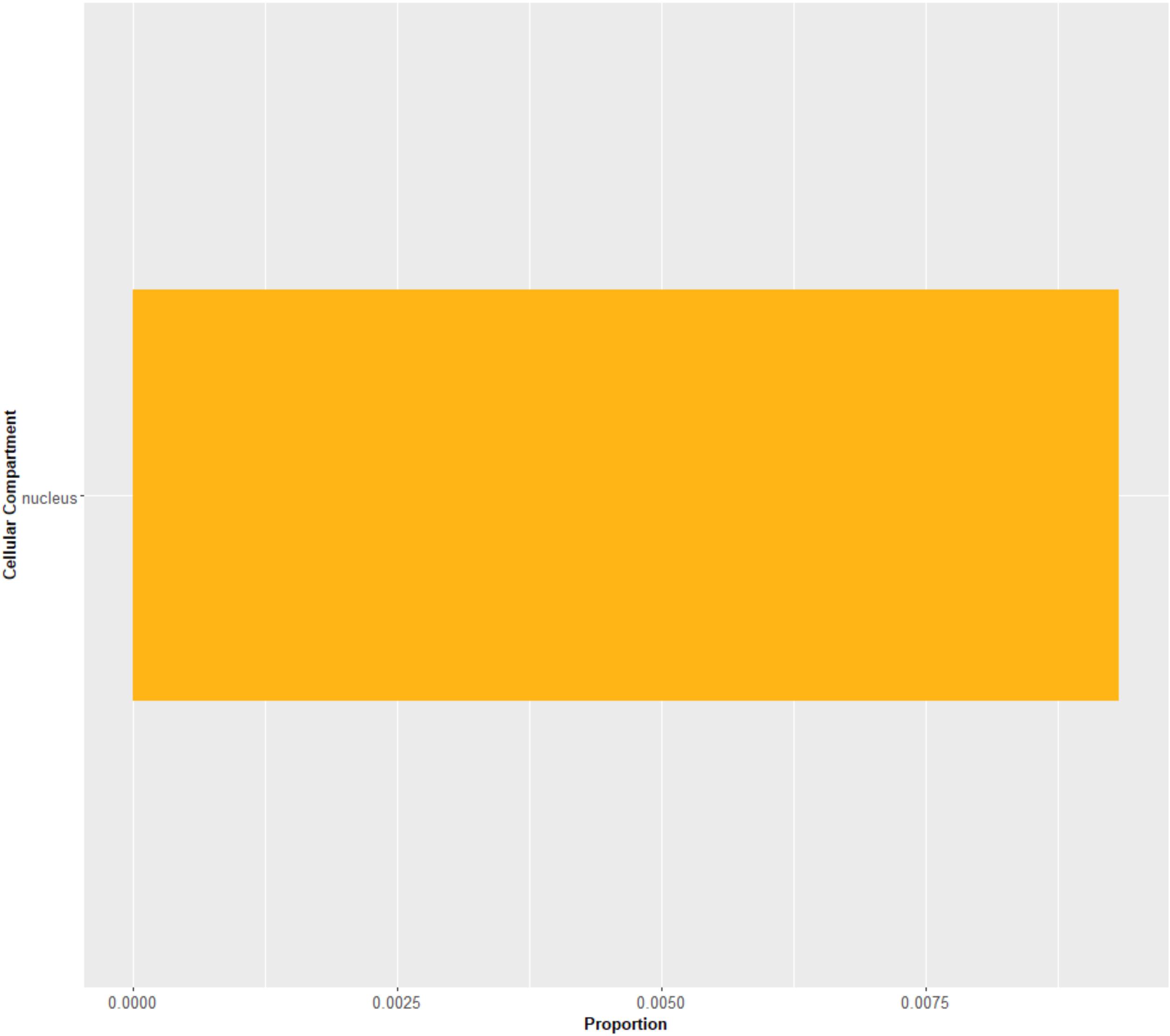
Without Cell Cycle | Without AreaShape | All Genes

Cluster 2 of 42



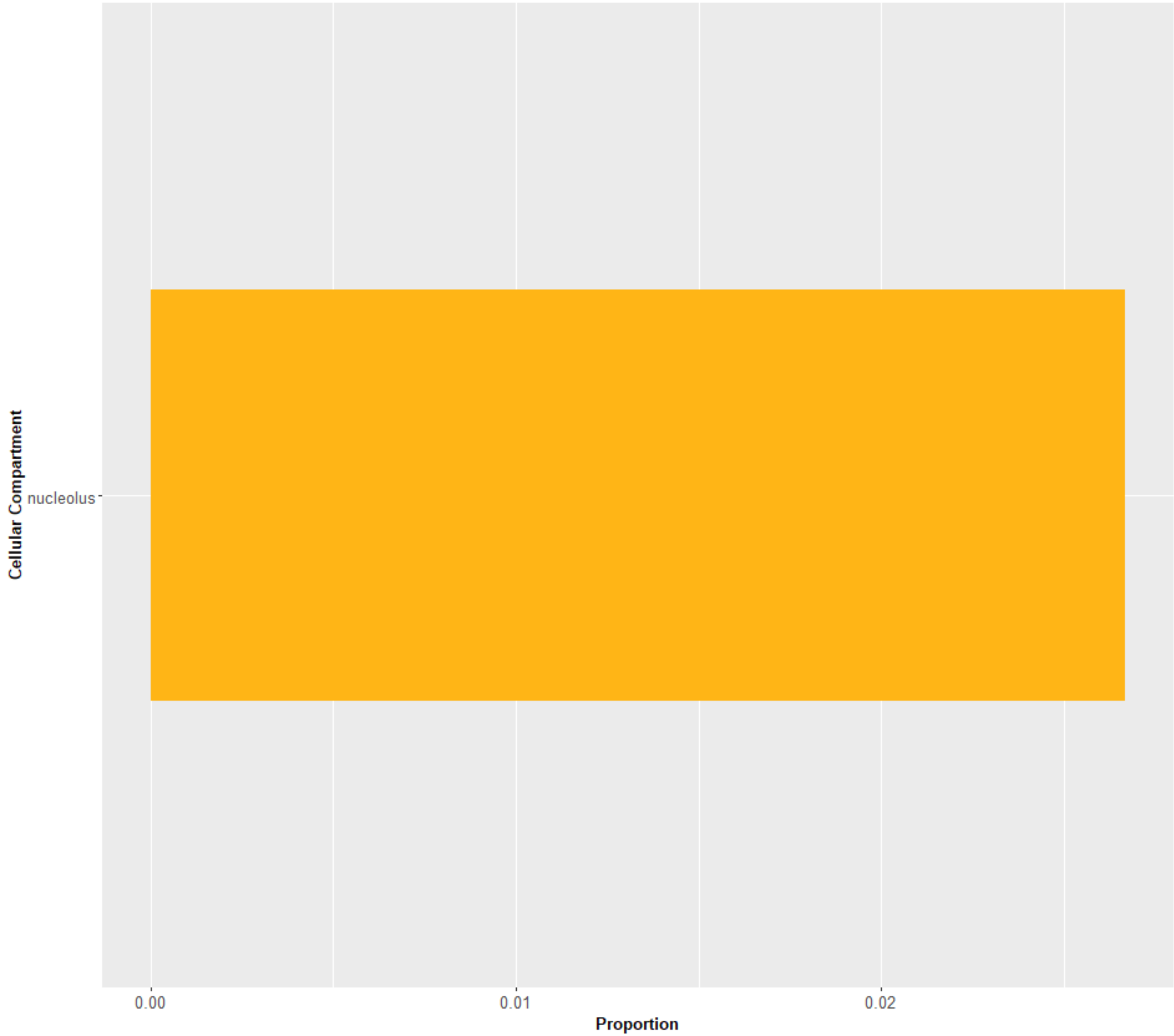
Without Cell Cycle | Without AreaShape | All Genes

Cluster 3 of 42



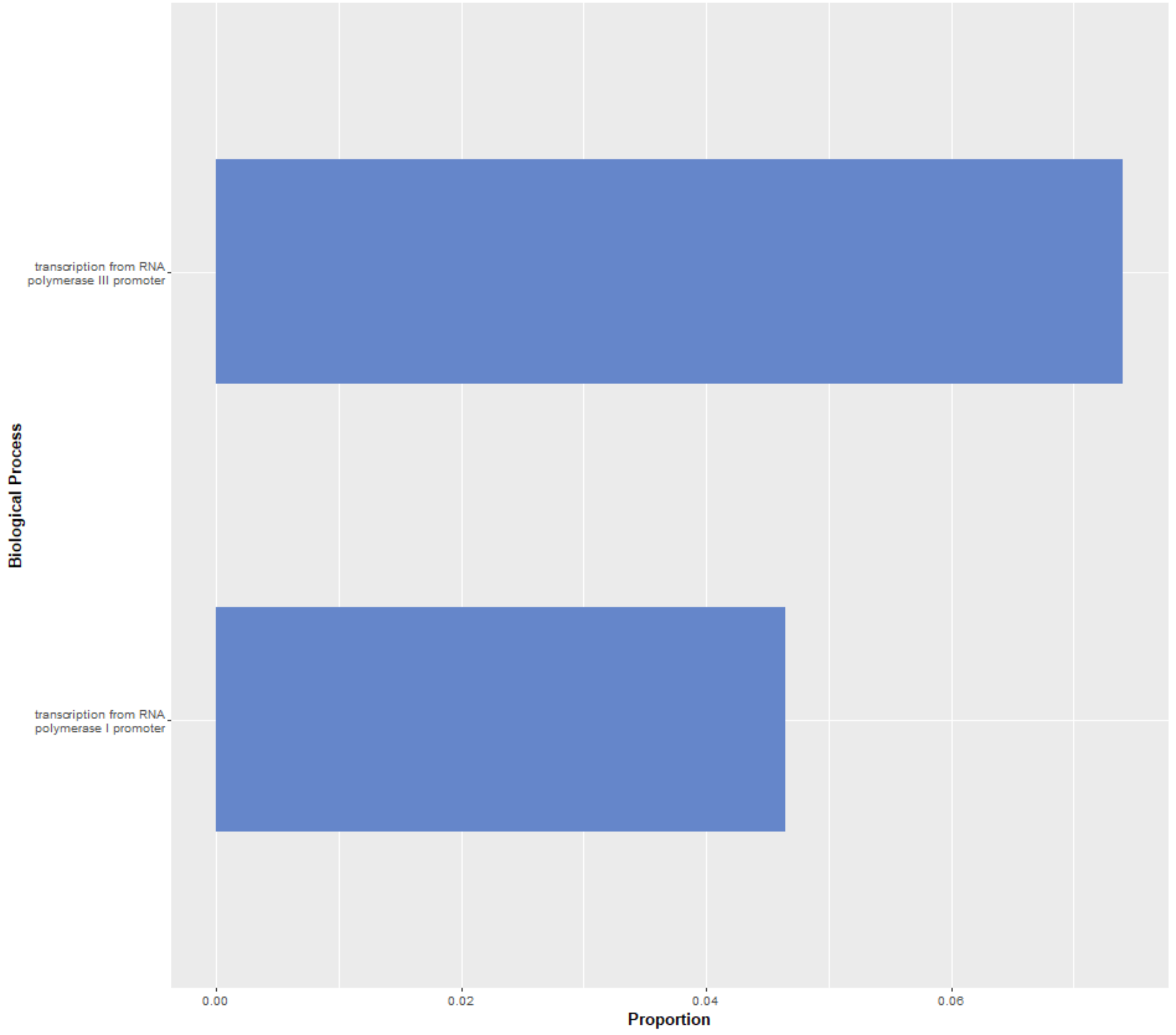
Without Cell Cycle | Without AreaShape | All Genes

Cluster 10 of 42



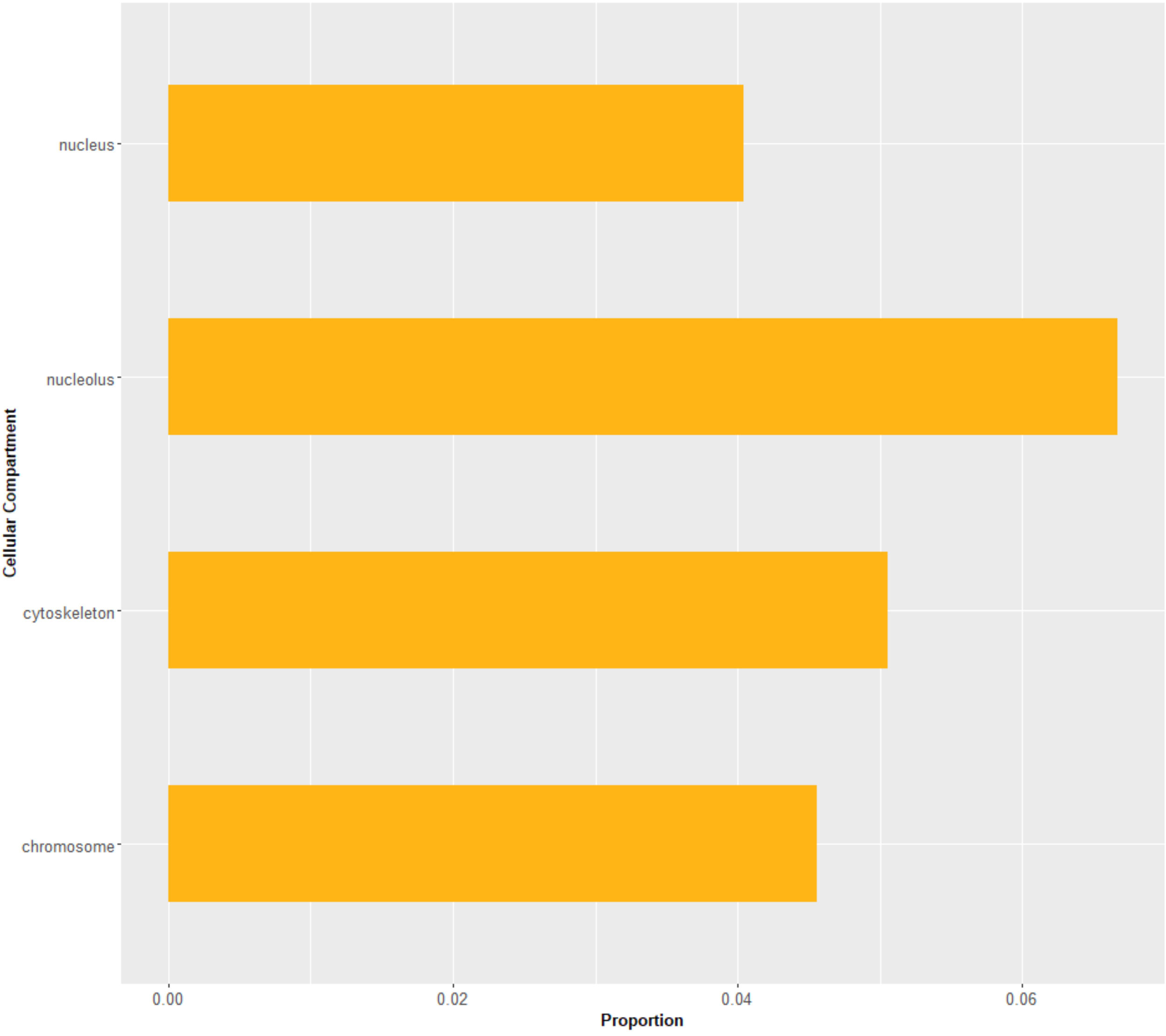
Without Cell Cycle | Without AreaShape | All Genes

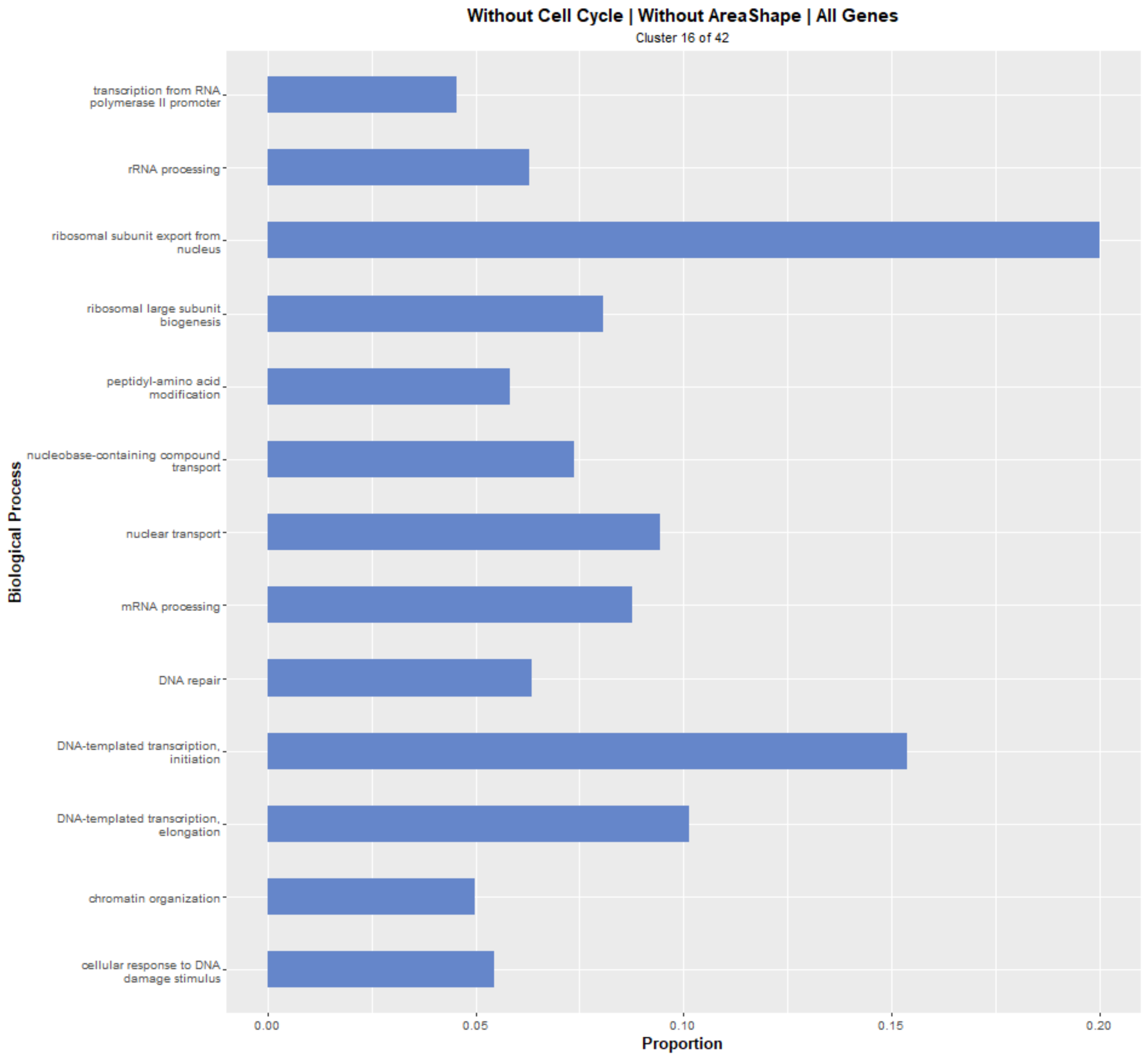
Cluster 10 of 42

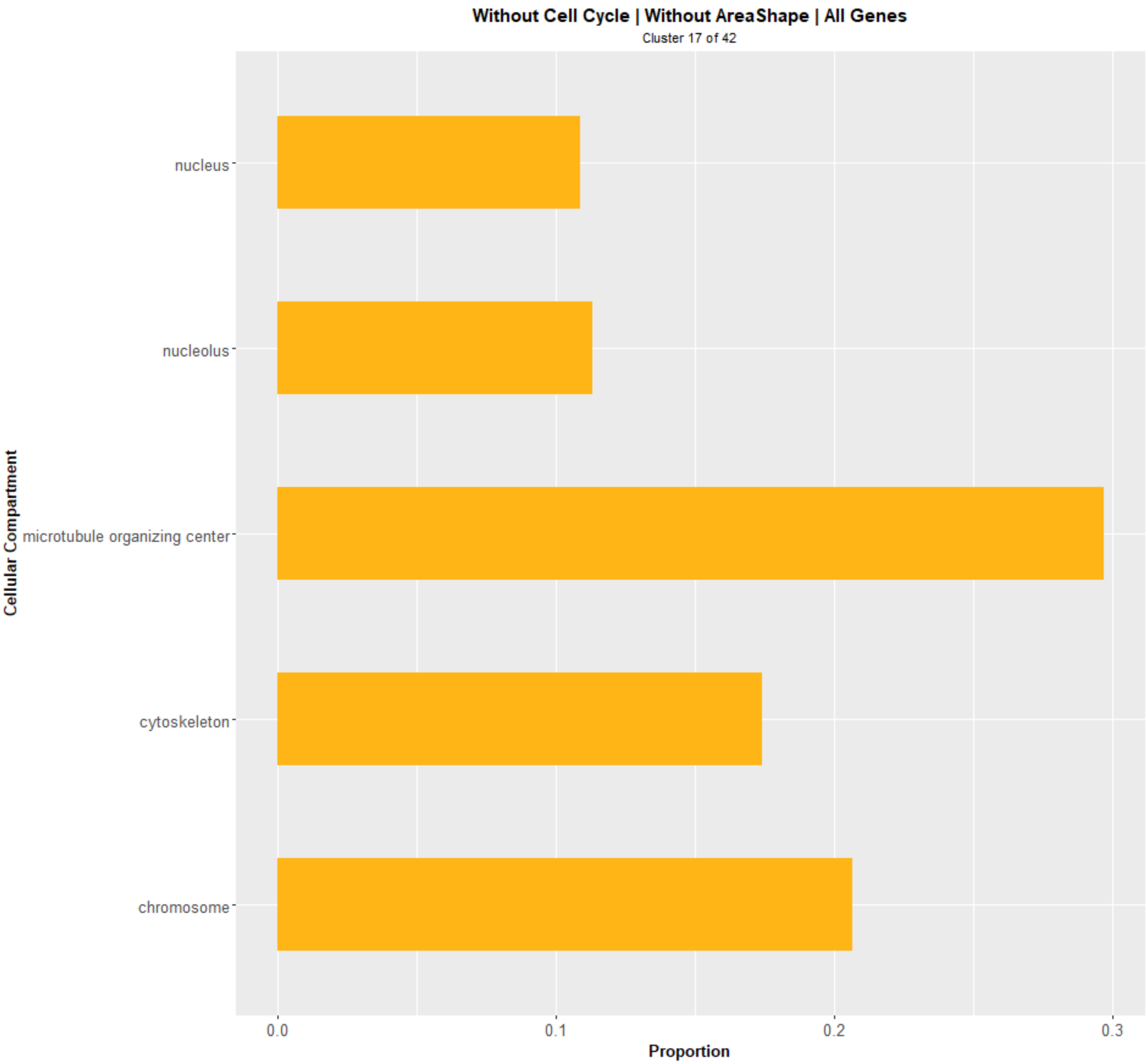


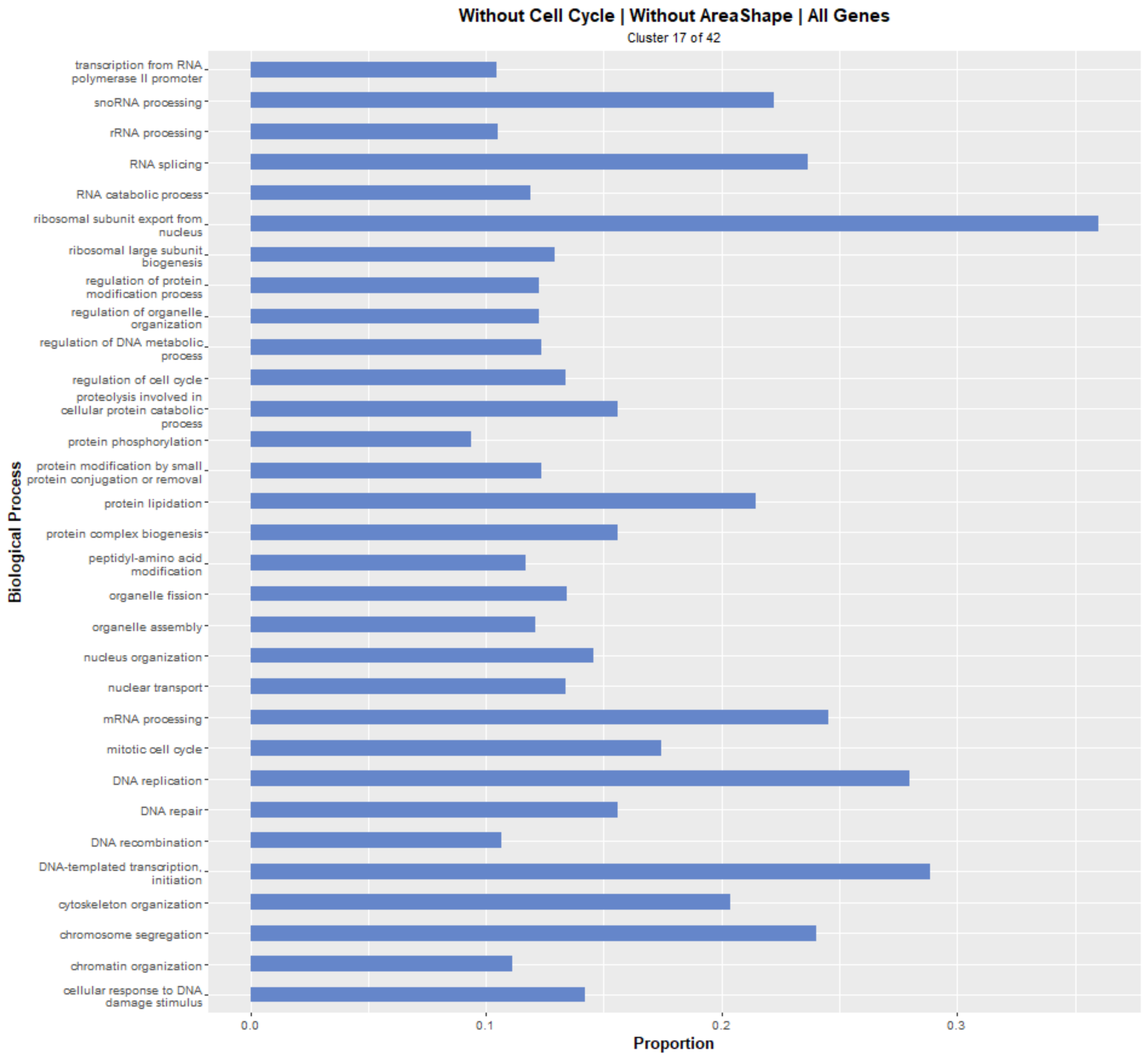
Without Cell Cycle | Without AreaShape | All Genes

Cluster 16 of 42



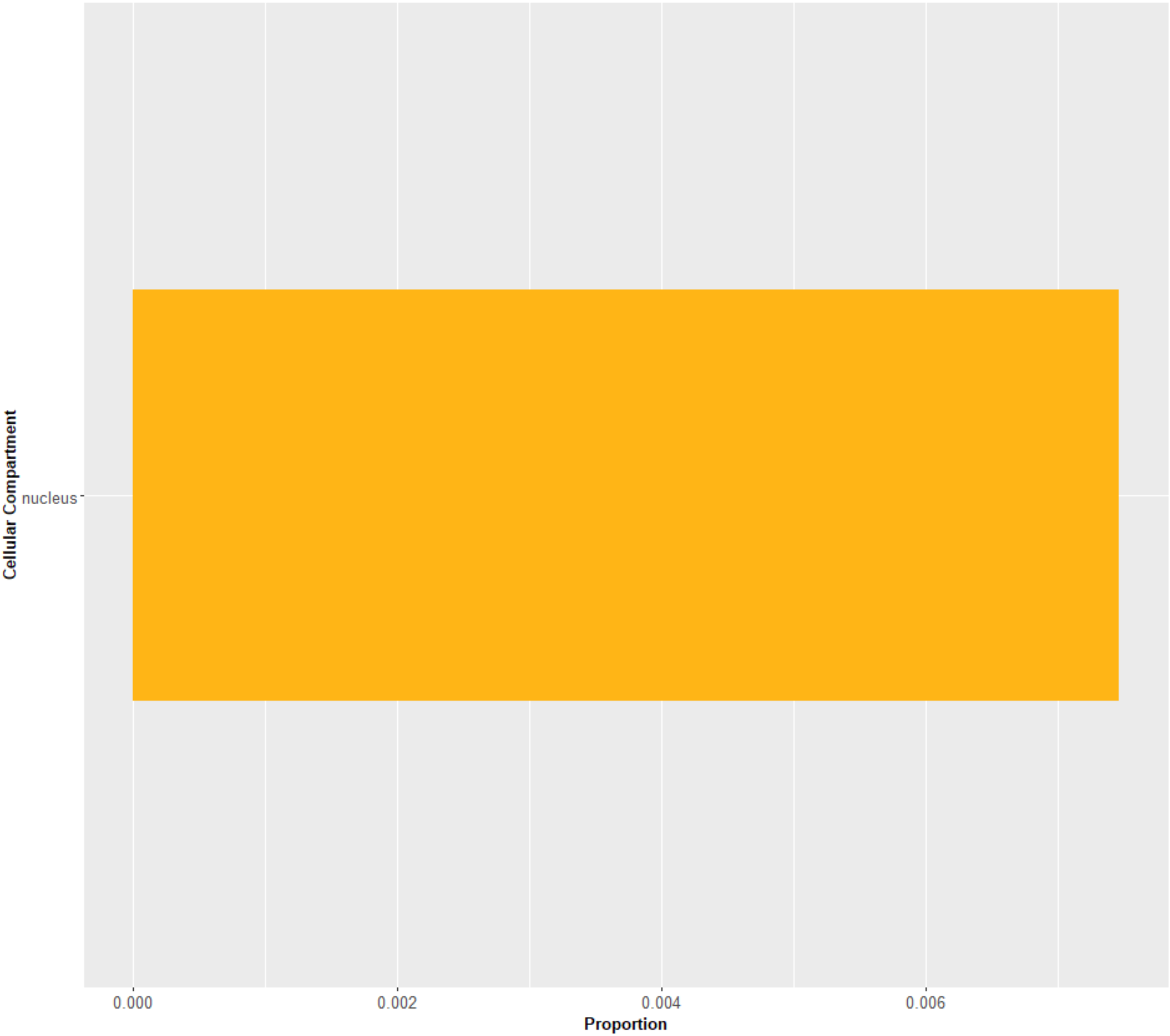

Without Cell Cycle | Without AreaShape | All Genes
Cluster 16 of 42
transcription from RNA polymerase II promoter
rRNA processing
ribosomal subunit export from nucleus
ribosomal large subunit biogenesis
peptidyl-amino acid modification
nucleobase-containing compound transport
nuclear transport
mRNA processing
DNA repair
DNA-templated transcription, initiation
DNA-templated transcription, elongation
chromatin organization
cellular response to DNA damage stimulus
Biological Process
0.00
0.05
0.10
0.15
0.20
Proportion


Without Cell Cycle | Without AreaShape | All Genes
Cluster 17 of 42
Cellular Compartment
nucleus
nucleolus
microtubule organizing center
cytoskeleton
chromosome
0.0
0.1
0.2
0.3
Proportion


Without Cell Cycle | Without AreaShape | All Genes
Cluster 17 of 42
transcription from RNA polymerase II promoter
snoRNA processing
rRNA processing
RNA splicing
RNA catabolic process
ribosomal subunit export from nucleus
ribosomal large subunit biogenesis
regulation of protein modification process
regulation of organelle organization
regulation of DNA metabolic process
regulation of cell cycle
proteolysis involved in cellular protein catabolic process
protein phosphorylation
protein modification by small protein conjugation or removal
protein lipidation
protein complex biogenesis
peptidyl-amino acid modification
organelle fission
organelle assembly
nucleus organization
nuclear transport
mRNA processing
mitotic cell cycle
DNA replication
DNA repair
DNA recombination
DNA-templated transcription, initiation
cytoskeleton organization
chromosome segregation
chromatin organization
cellular response to DNA damage stimulus
Biological Process
0.0
0.1
0.2
0.3
Proportion

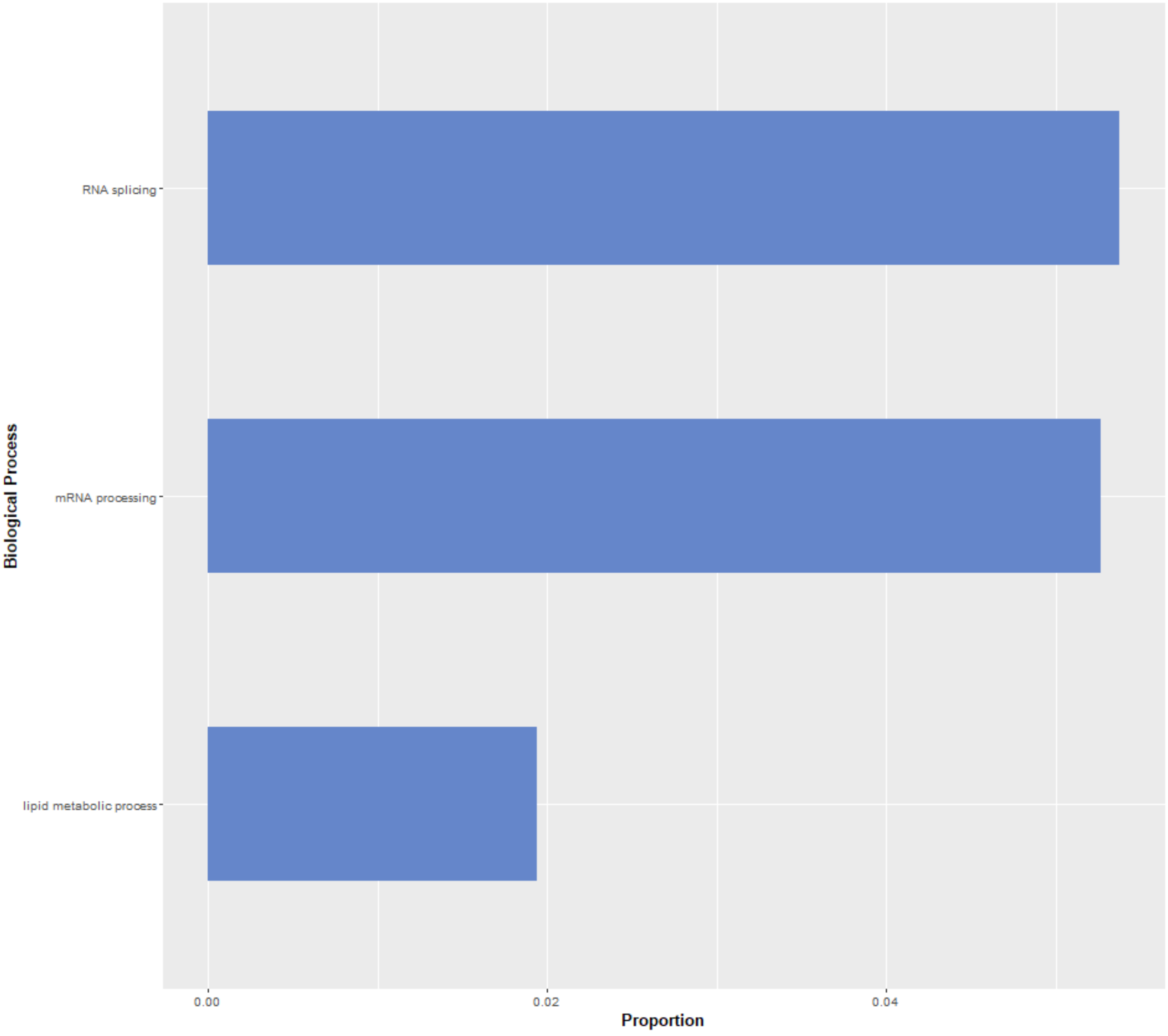
Without Cell Cycle | Without AreaShape | All Genes

Cluster 18 of 42



Without Cell Cycle | Without AreaShape | All Genes

Cluster 18 of 42



Without Cell Cycle | Without AreaShape | All Genes

Cluster 19 of 42

Cellular Compartment

site of polarized growth

membrane

Golgi apparatus

endoplasmic reticulum

endomembrane system

cytoplasmic vesicle

cytoplasm

cellular bud

cell cortex

0.00

0.04

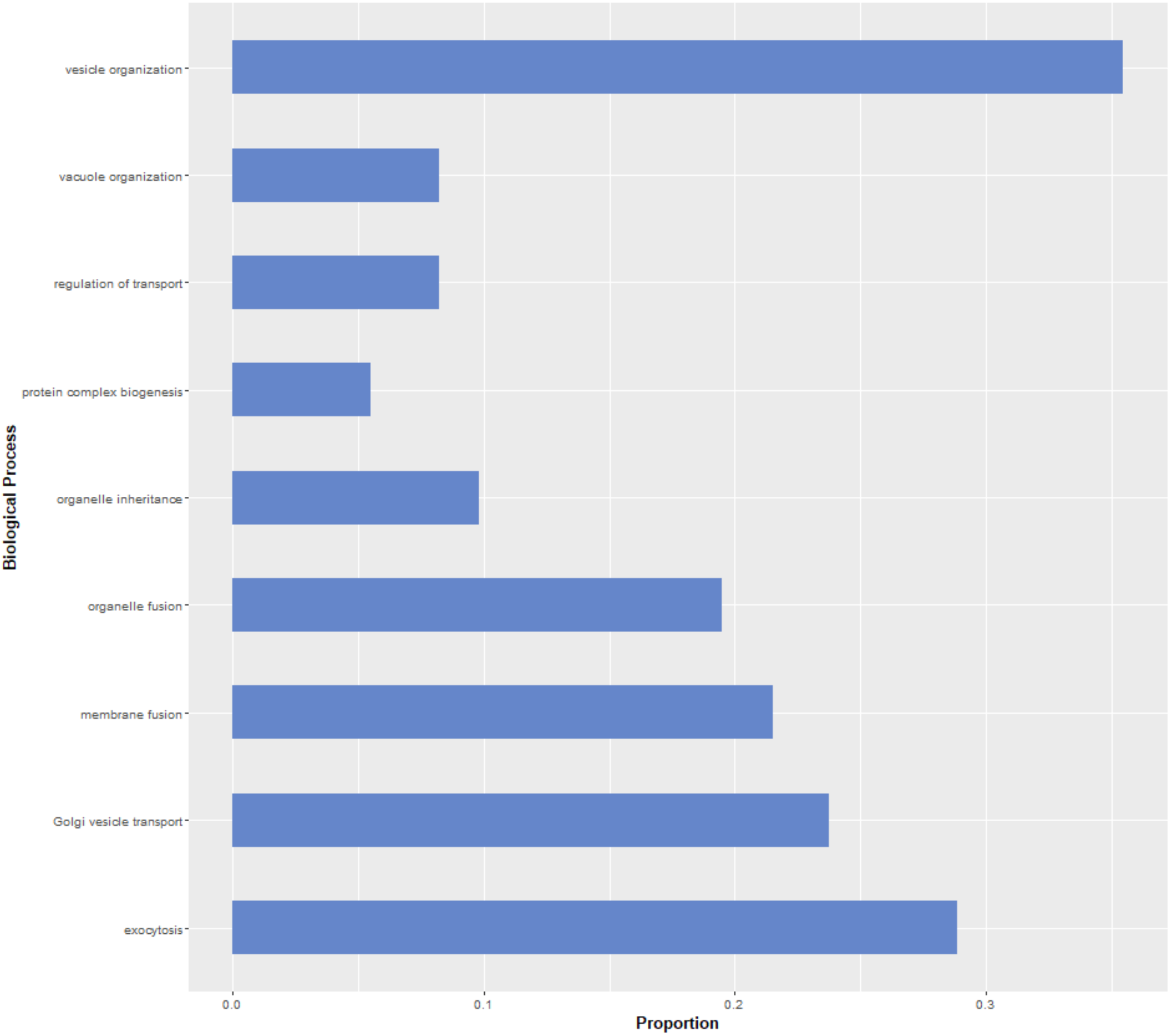
0.08

0.12

Proportion

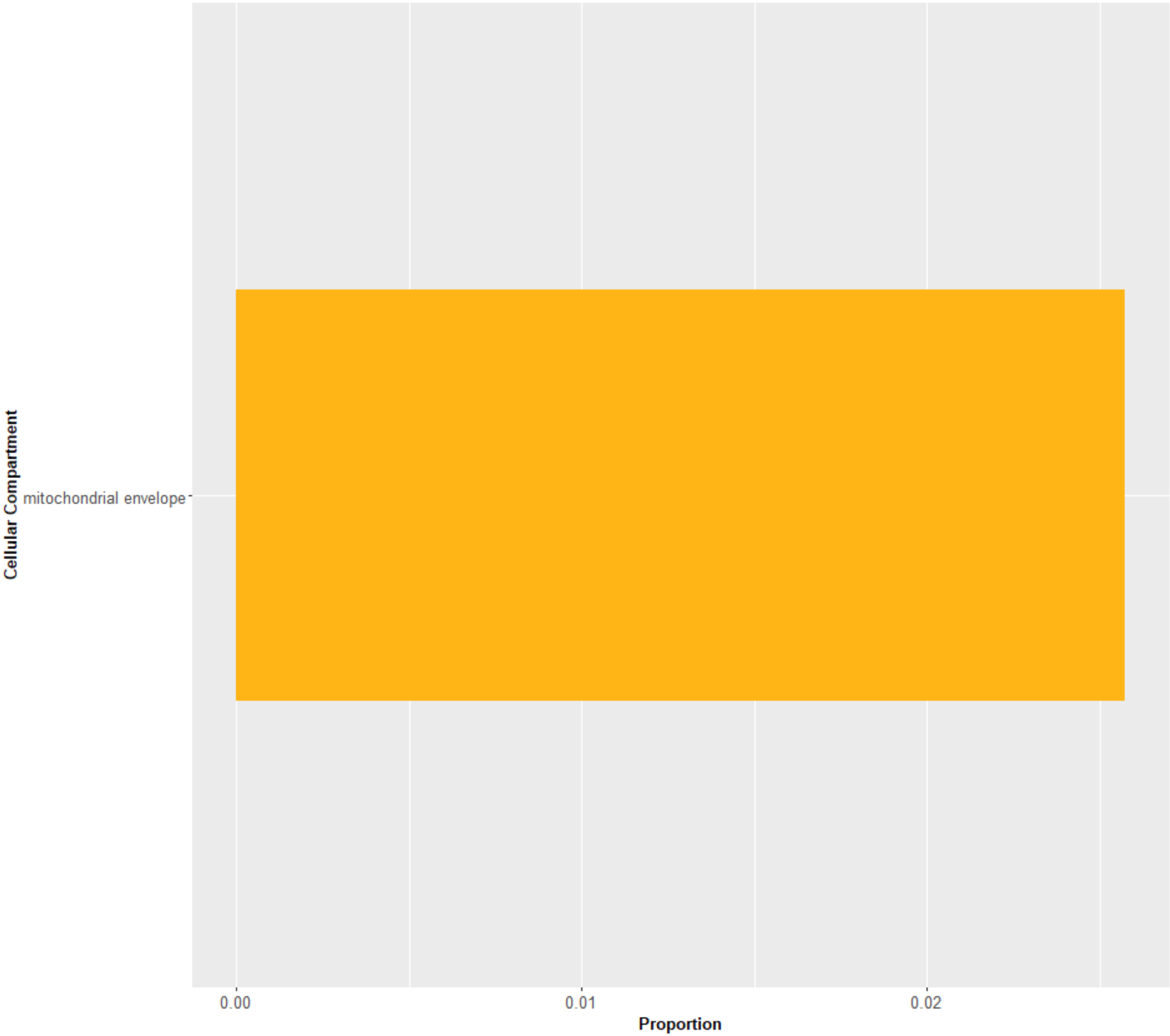
Without Cell Cycle | Without AreaShape | All Genes

Cluster 19 of 42



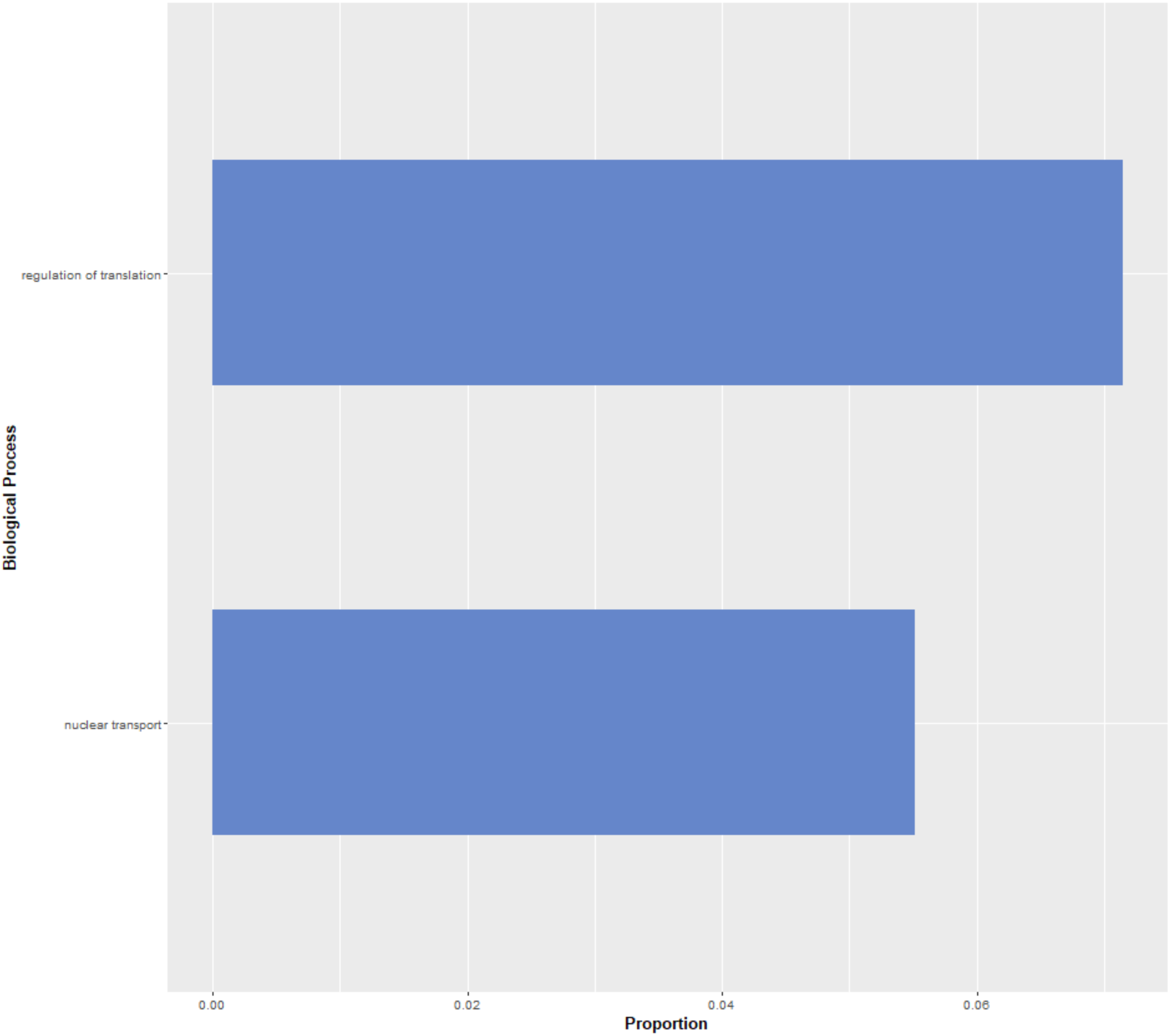
Without Cell Cycle | Without AreaShape | All Genes

Cluster 20 of 42



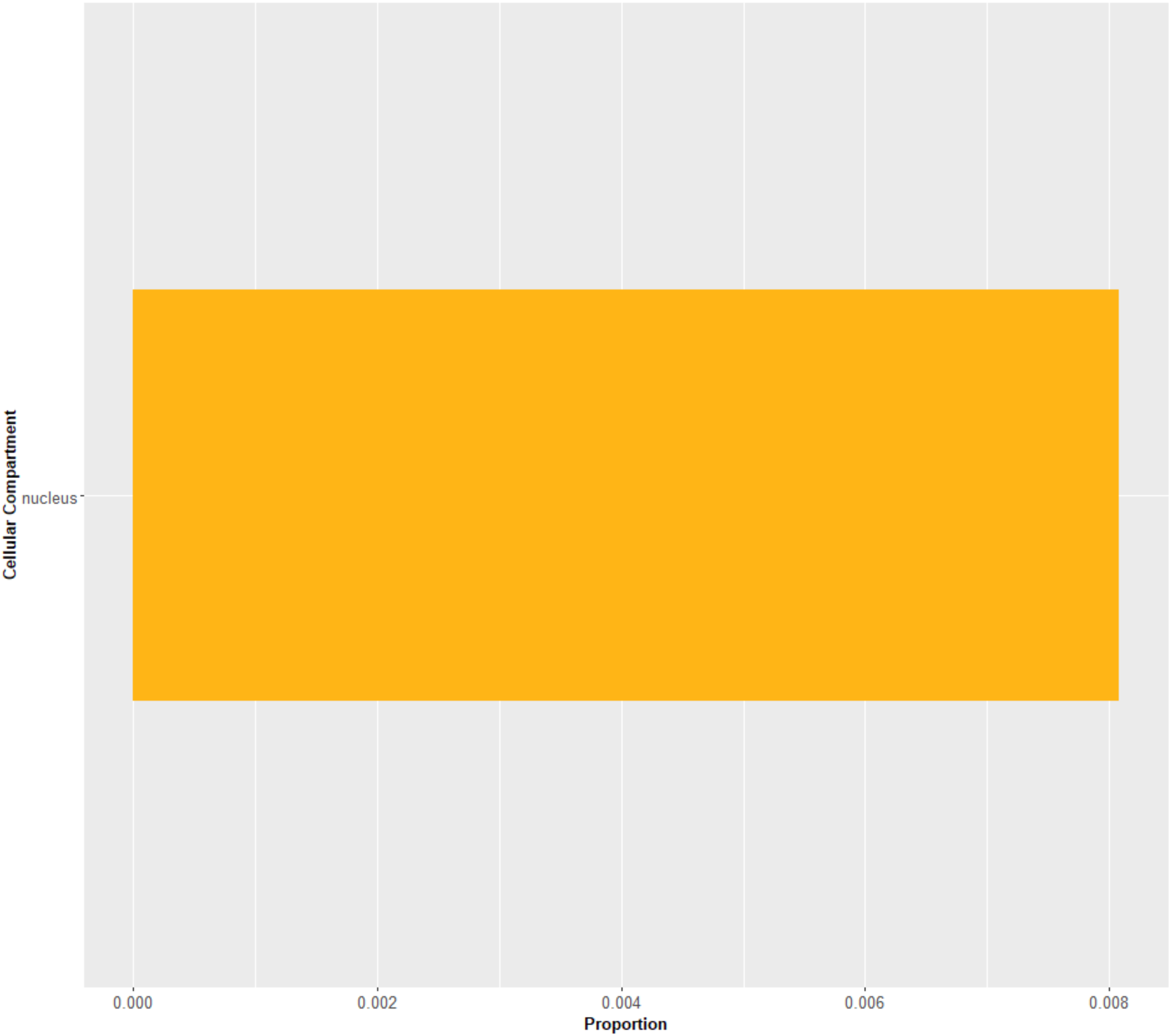
Without Cell Cycle | Without AreaShape | All Genes

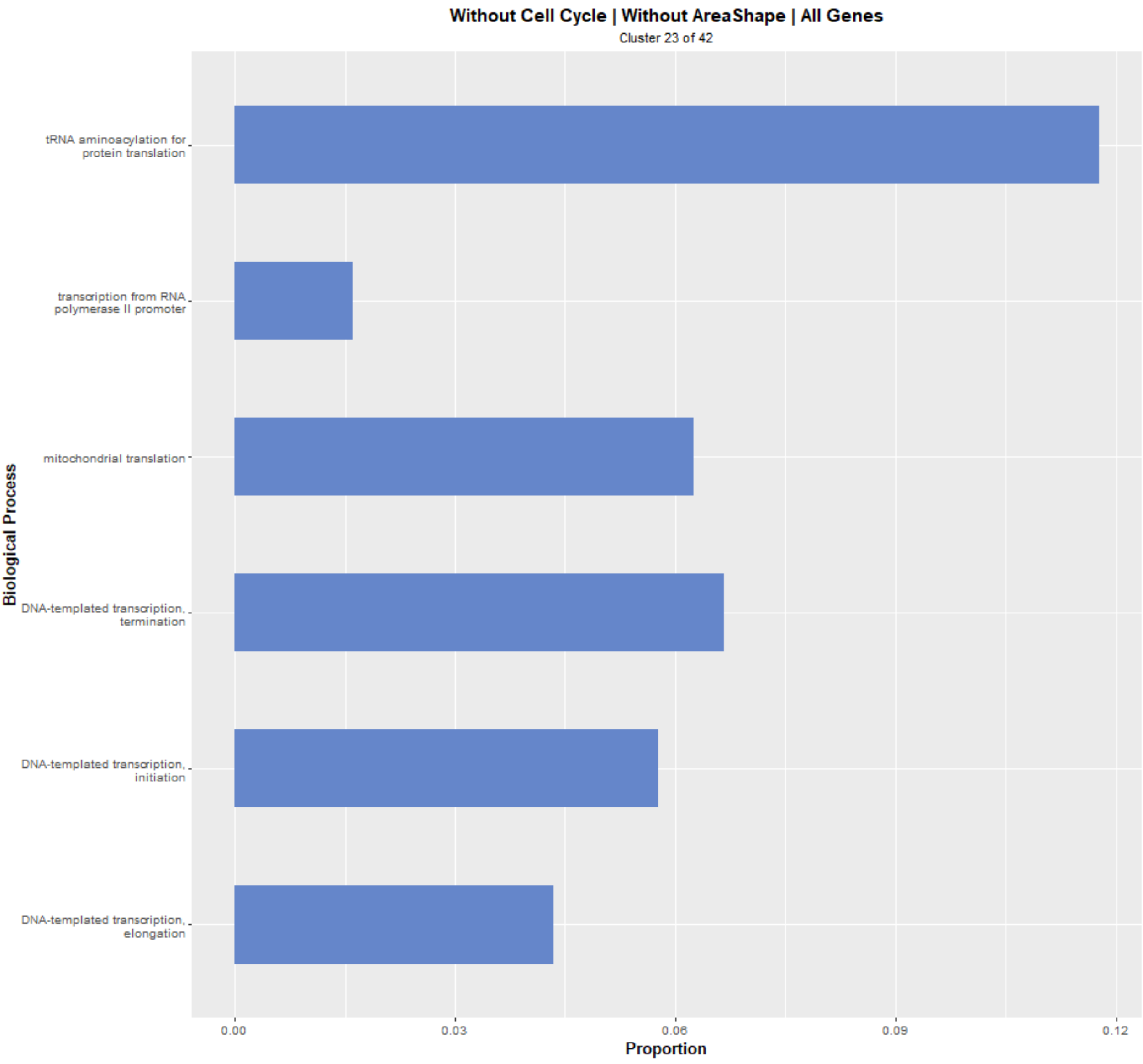
Cluster 21 of 42

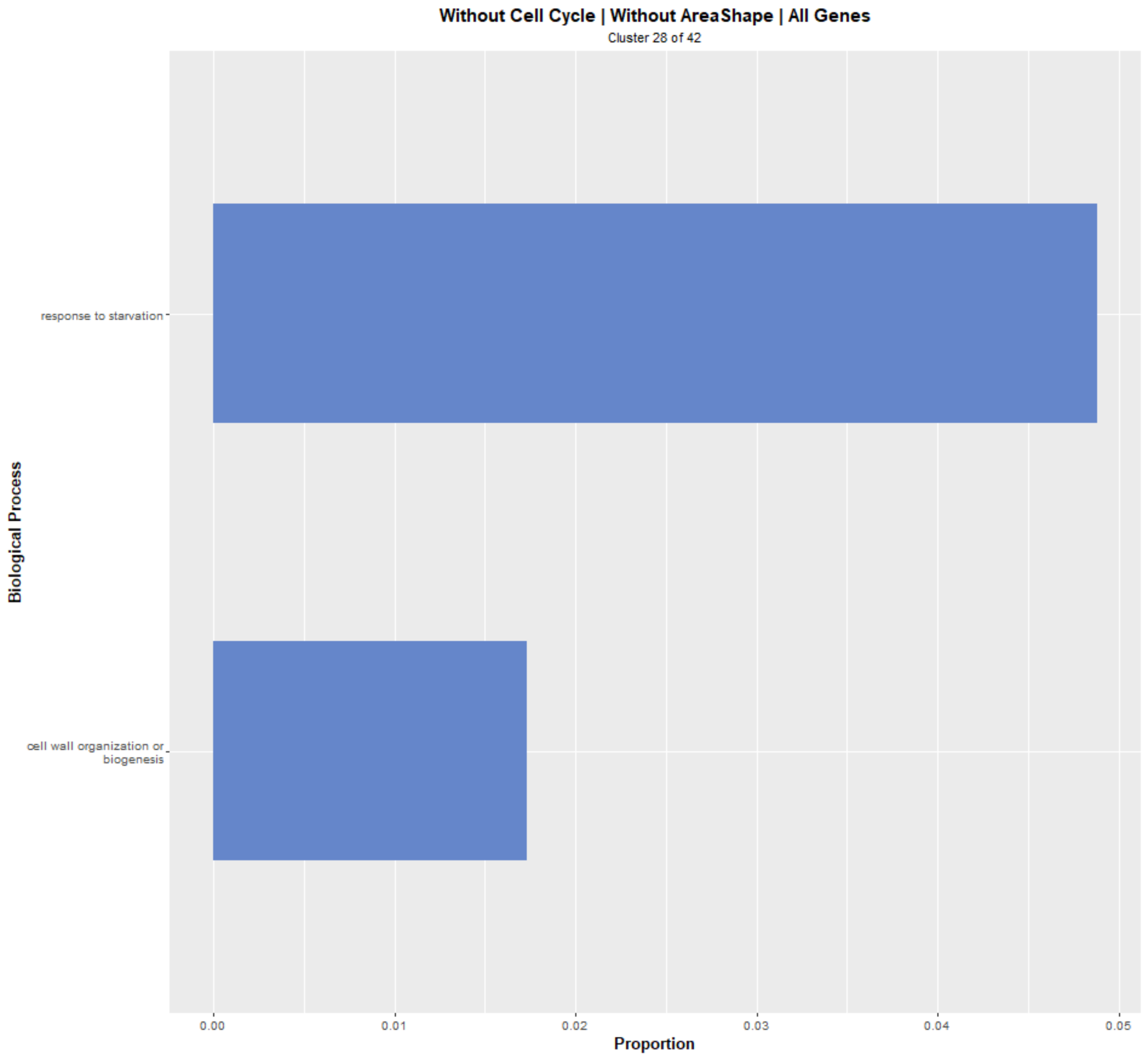


Without Cell Cycle | Without AreaShape | All Genes

Cluster 23 of 42

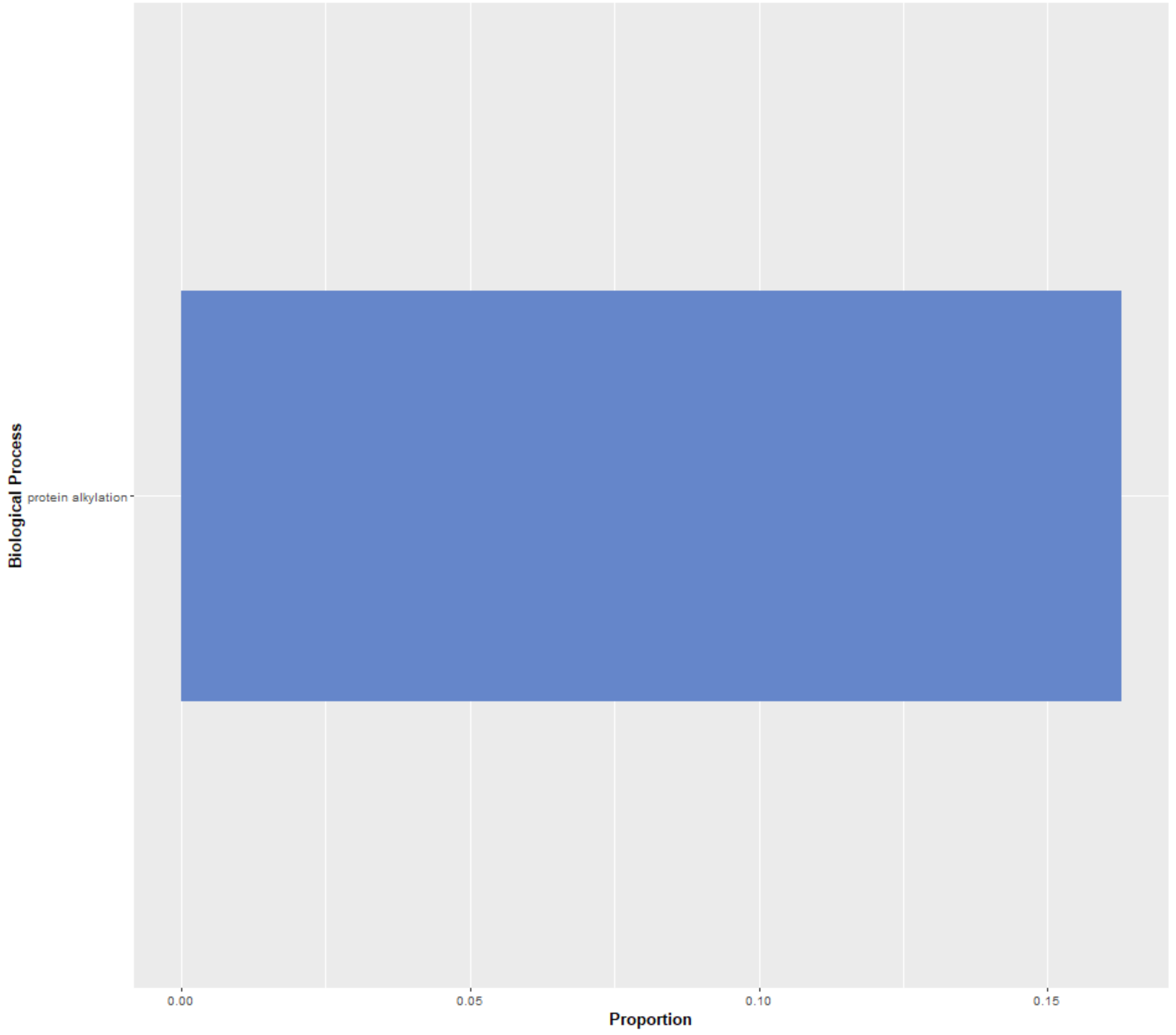



Without Cell Cycle | Without AreaShape | All Genes
Cluster 23 of 42
tRNA aminoacylation for protein translation
transcription from RNA polymerase II promoter
mitochondrial translation
DNA-templated transcription, termination
DNA-templated transcription, initiation
DNA-templated transcription, elongation
Biological Process
0.00
0.03
0.06
0.09
0.12
Proportion


Without Cell Cycle | Without AreaShape | All Genes
Cluster 28 of 42
response to starvation
cell wall organization or biogenesis
Biological Process
0.00
0.01
0.02
0.03
0.04
0.05
Proportion

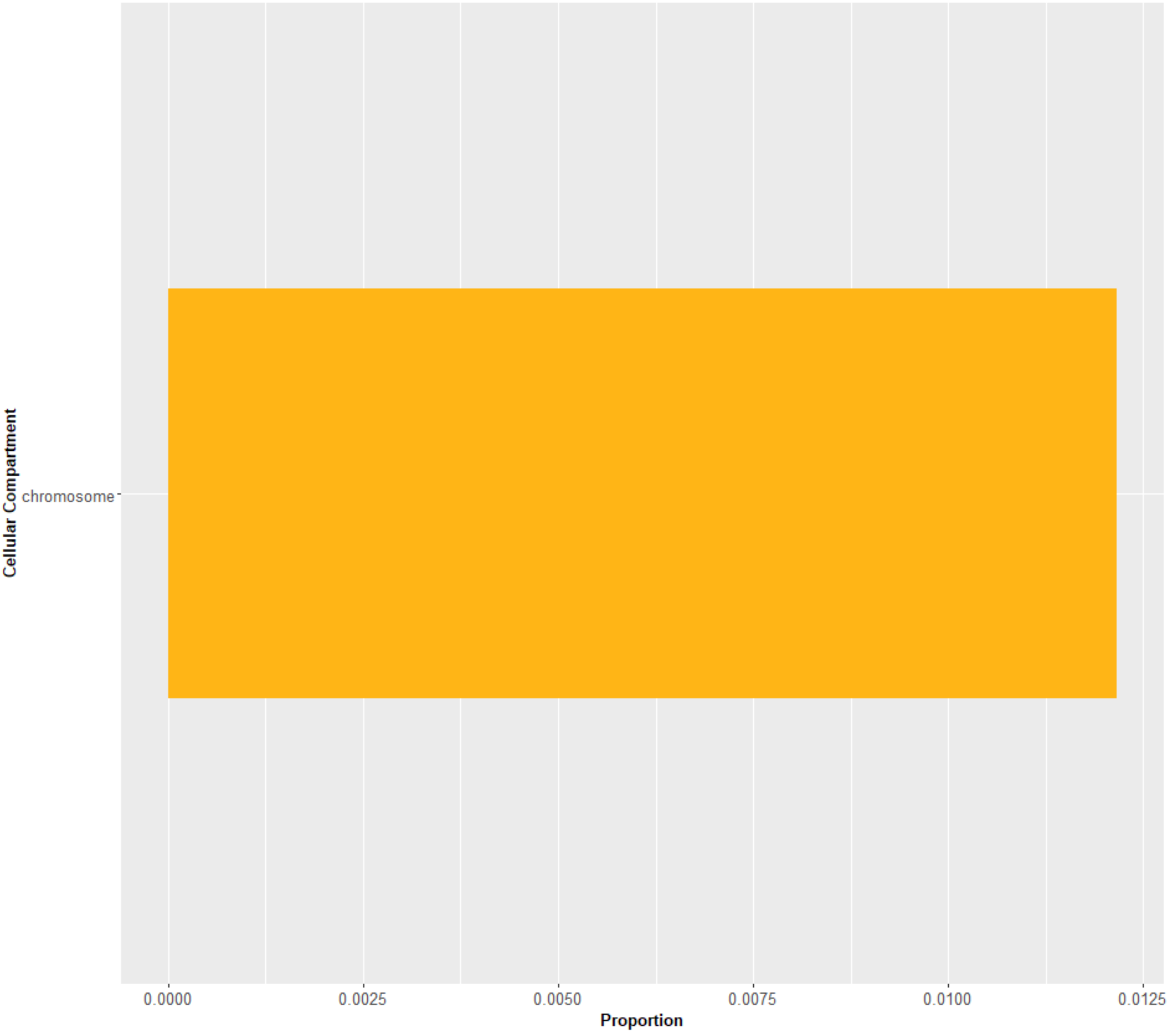
Without Cell Cycle | Without AreaShape | All Genes

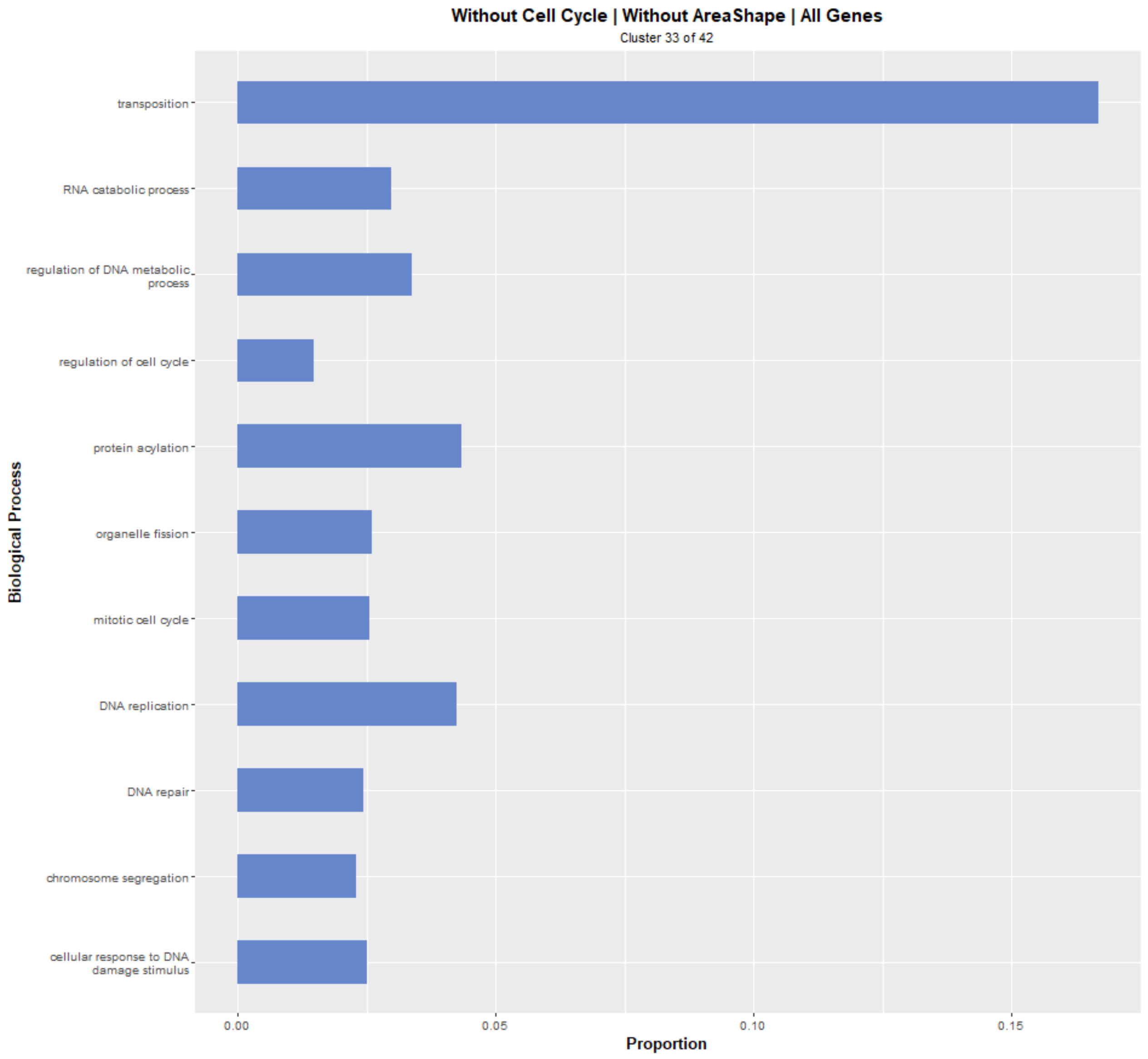
Cluster 30 of 42



Without Cell Cycle | Without AreaShape | All Genes

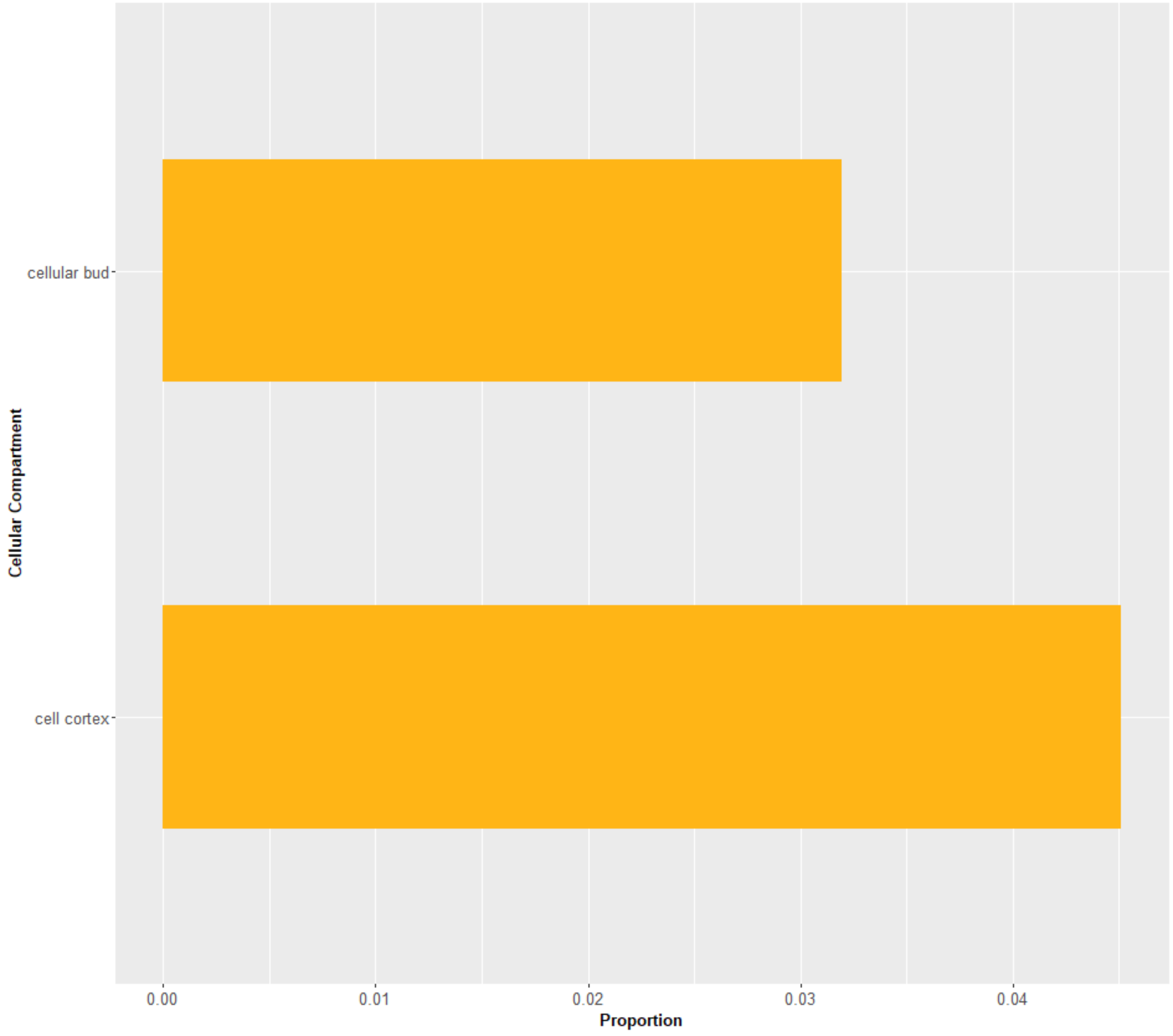
Cluster 33 of 42




Without Cell Cycle | Without AreaShape | All Genes
Cluster 33 of 42
transposition
RNA catabolic process
regulation of DNA metabolic process
regulation of cell cycle
protein acylation
organelle fission
mitotic cell cycle
DNA replication
DNA repair
chromosome segregation
cellular response to DNA damage stimulus
Biological Process
0.00
0.05
0.10
0.15
Proportion

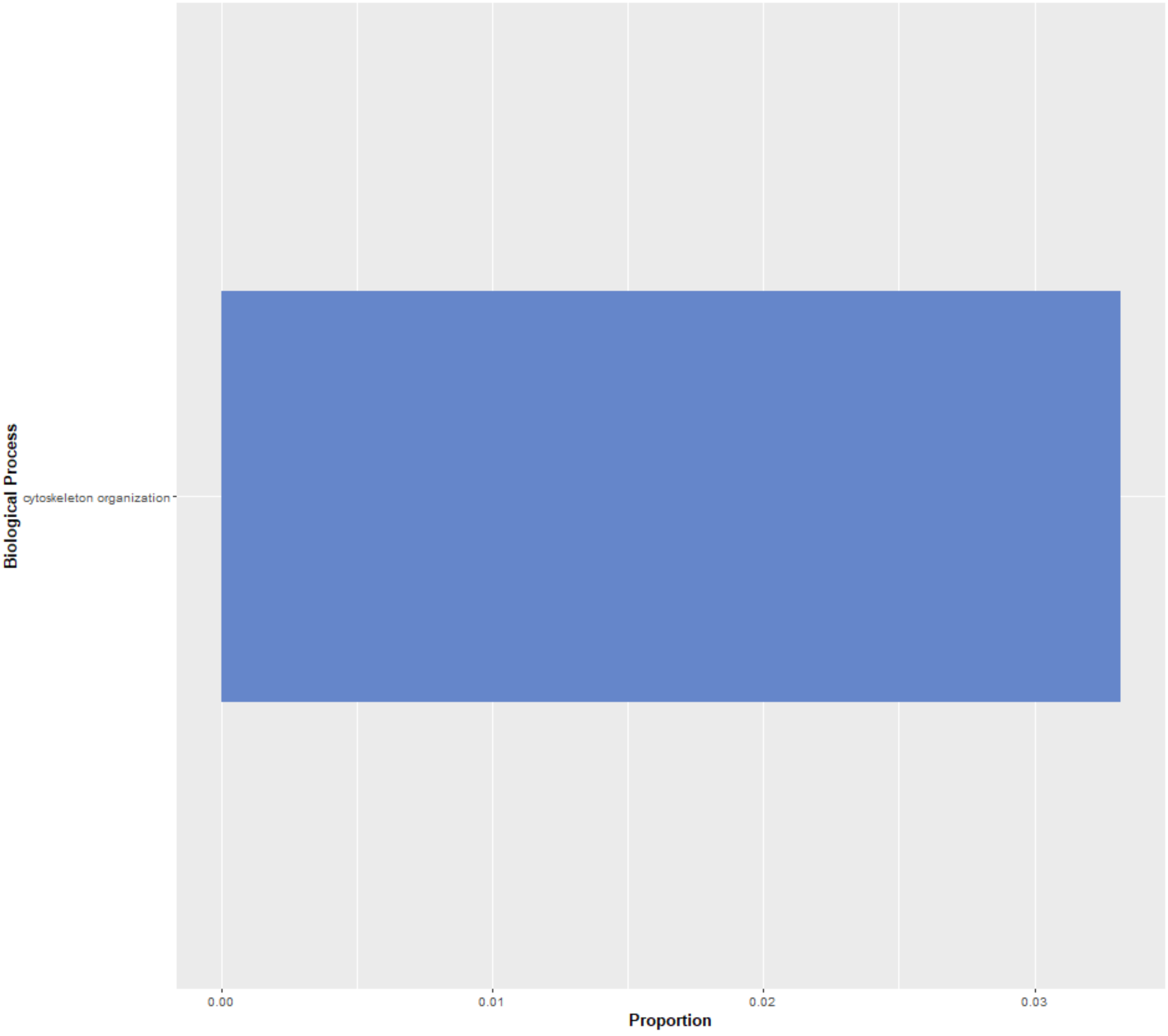
Without Cell Cycle | Without AreaShape | All Genes

Cluster 34 of 42



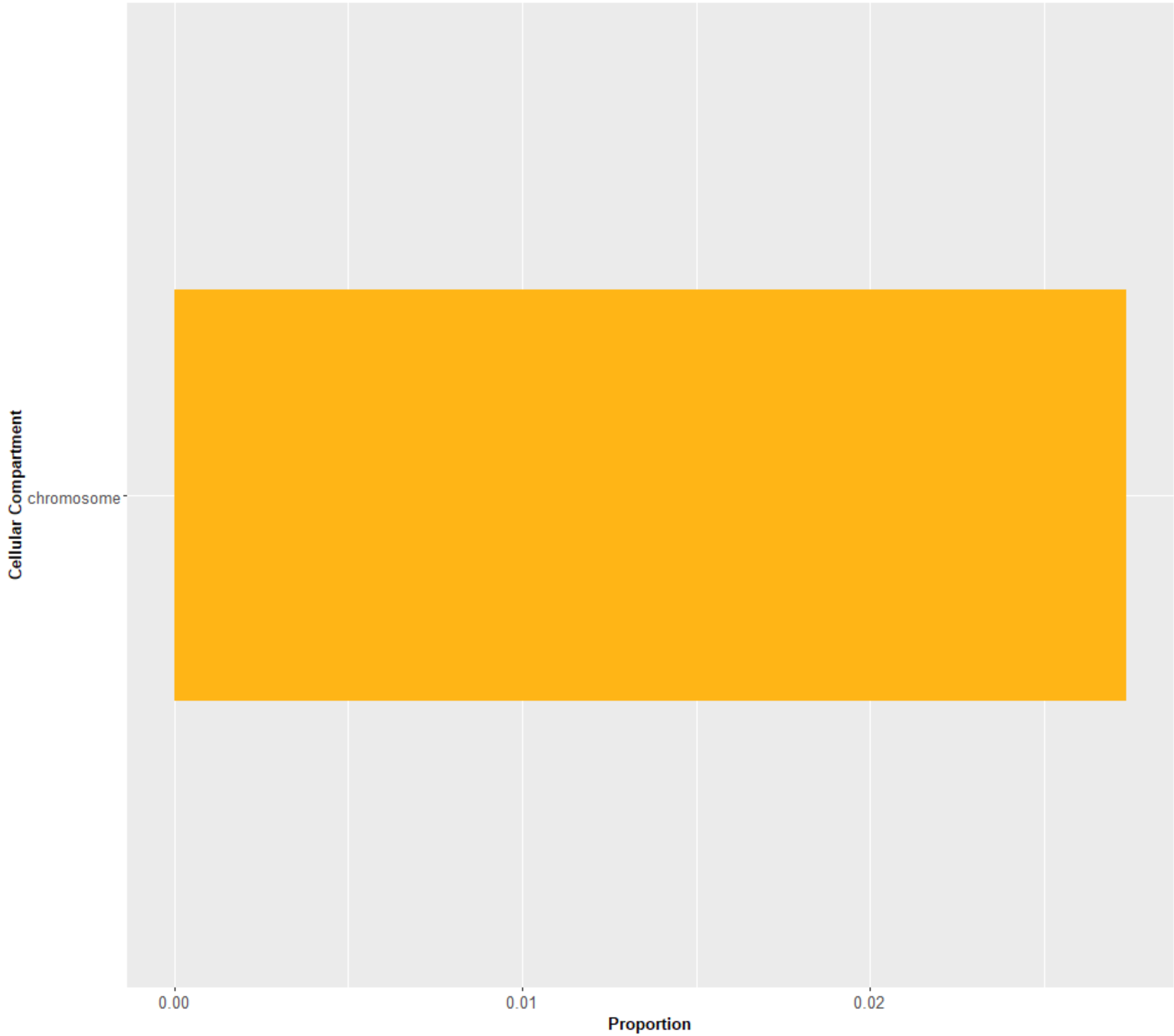
Without Cell Cycle | Without AreaShape | All Genes

Cluster 34 of 42



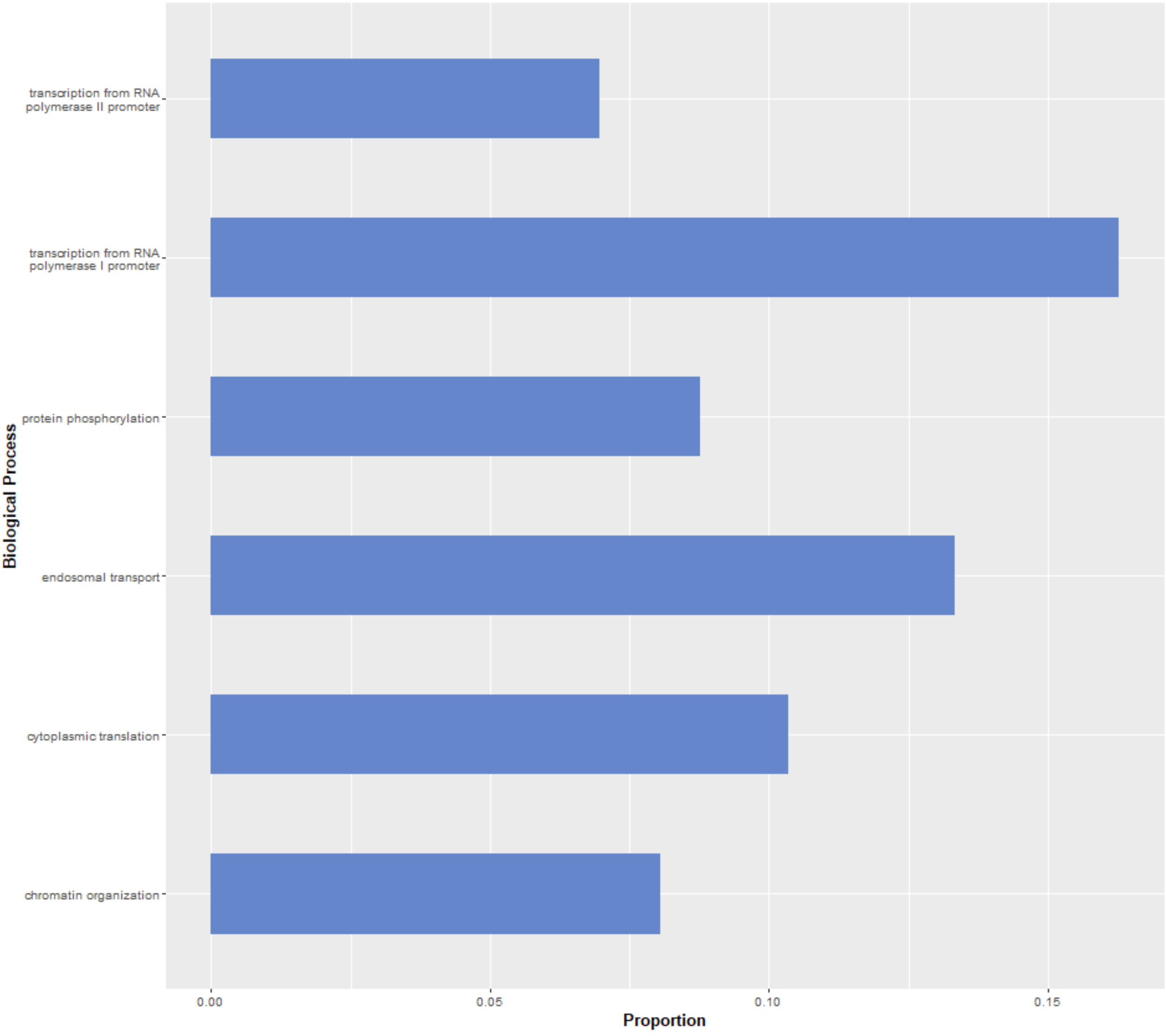
Without Cell Cycle | Without AreaShape | All Genes

Cluster 36 of 42



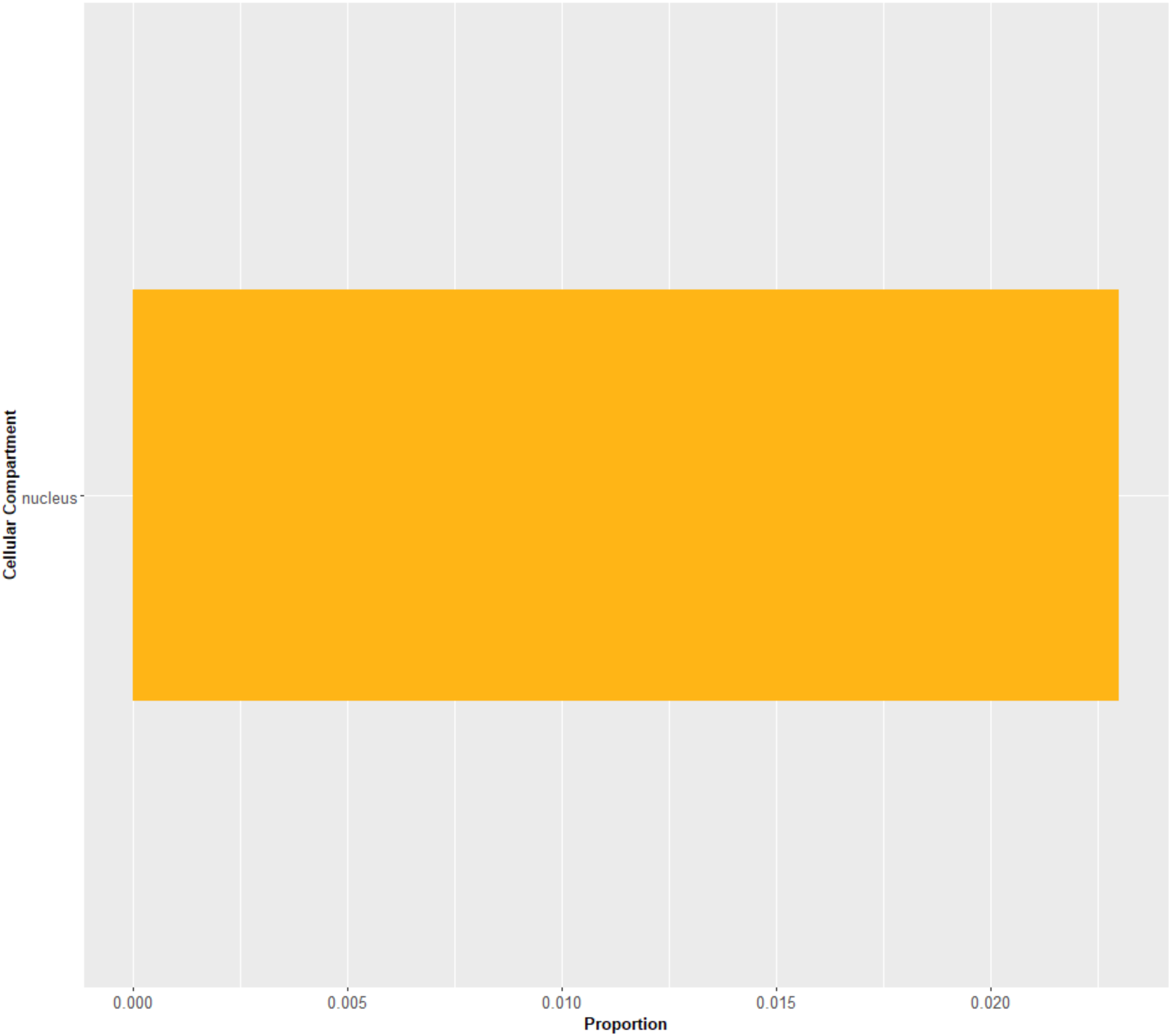
Without Cell Cycle | Without AreaShape | All Genes

Cluster 37 of 42



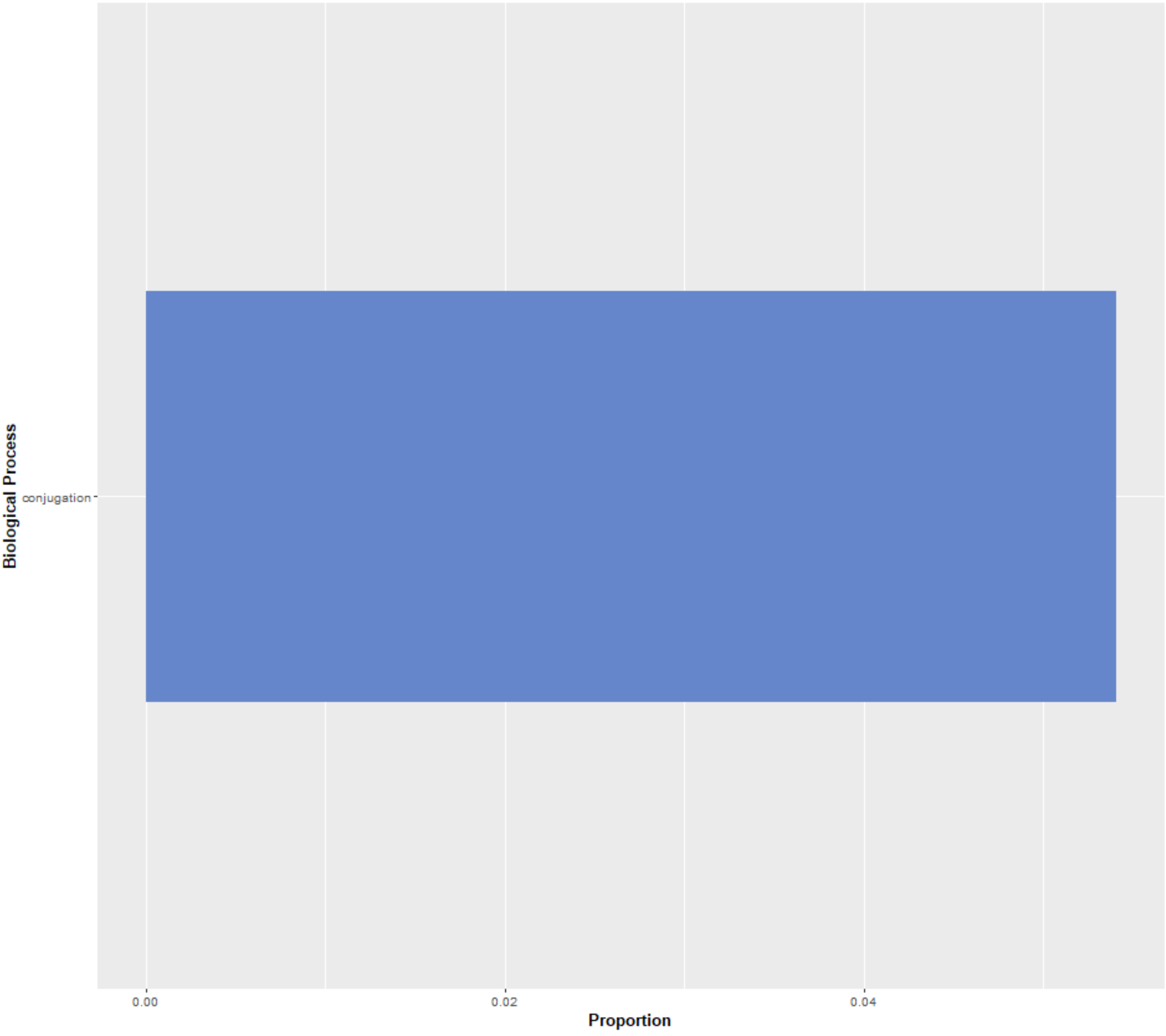
Without Cell Cycle | Without AreaShape | All Genes

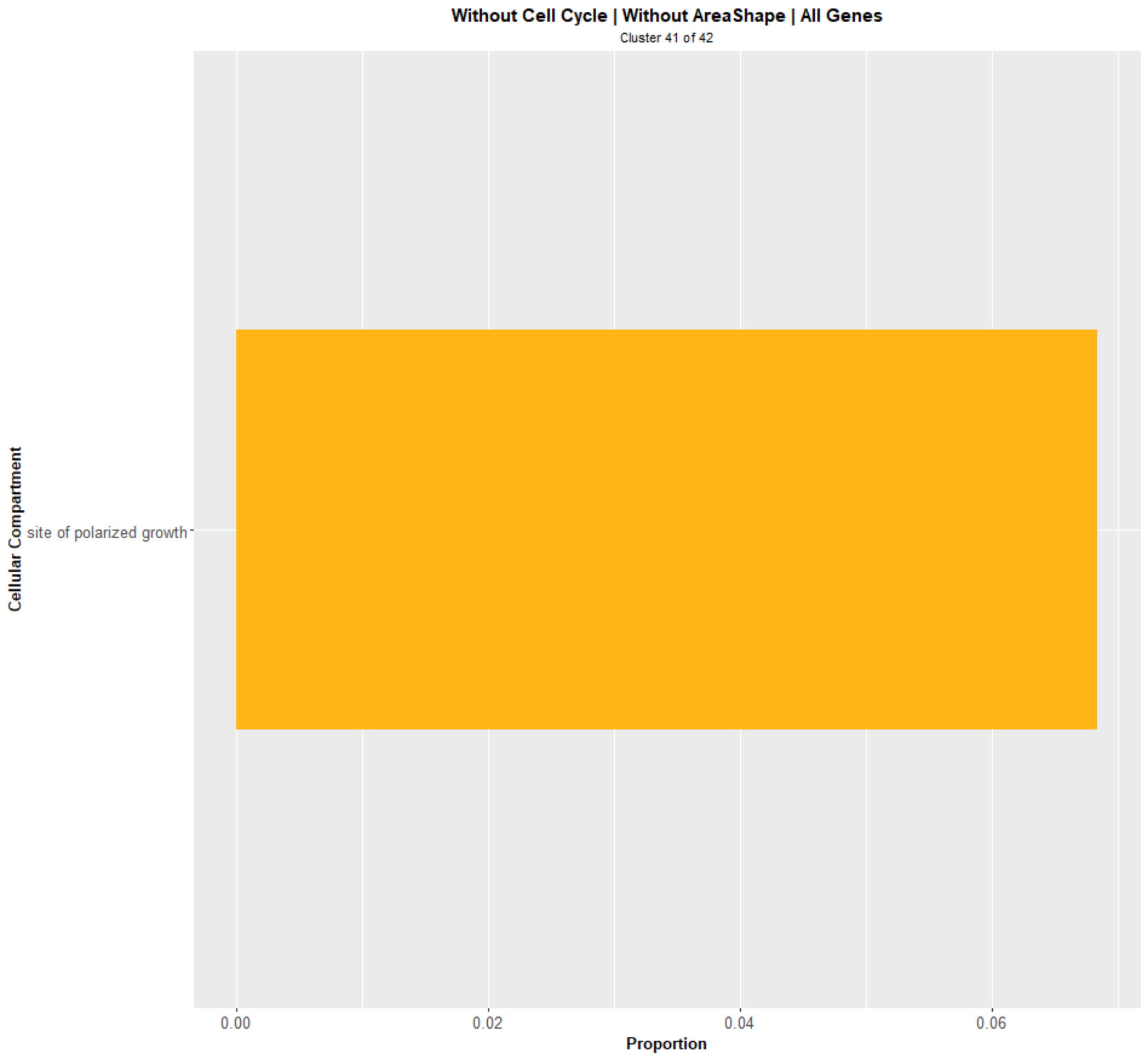
Cluster 40 of 42



Without Cell Cycle | Without AreaShape | All Genes

Cluster 40 of 42




Without Cell Cycle | Without AreaShape | All Genes
Cluster 41 of 42
Cellular Compartment
site of polarized growth
0.00
0.02
0.04
0.06
Proportion

Without Cell Cycle | Without AreaShape | All Genes

Cluster 41 of 42

vacuole organization

regulation of transport

organelle fission

mitotic cell cycle

cytokinesis

Biological Process

0.00

0.03

0.06

0.09

Proportion

Without Cell Cycle | Without AreaShape | All Genes

Cluster 42 of 42

Cellular Compartment

site of polarized growth

cytoskeleton

cell cortex

0.00

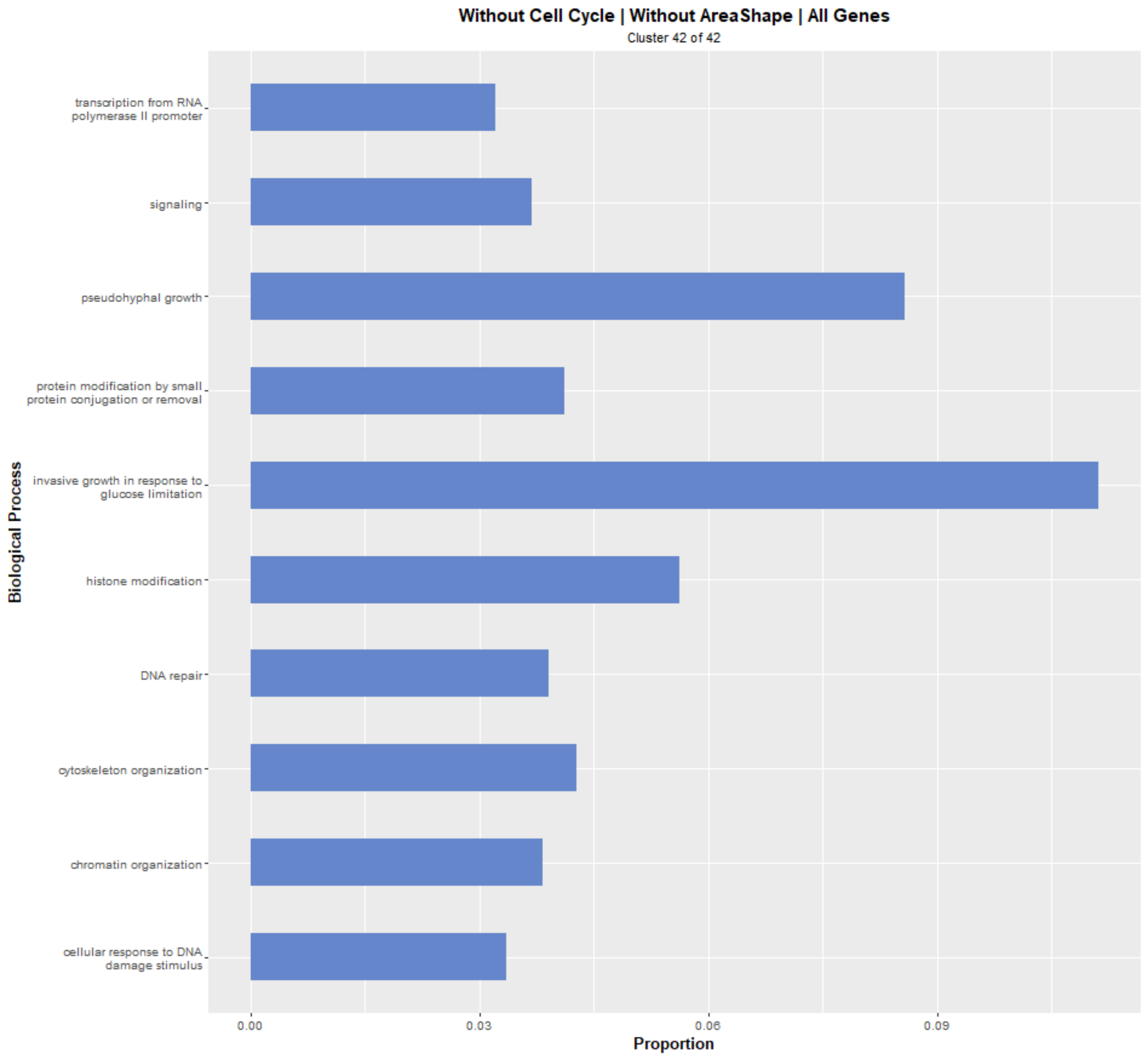
0.01

0.02

0.03

0.04

Proportion


Without Cell Cycle | Without AreaShape | All Genes
Cluster 42 of 42
transcription from RNA polymerase II promoter
signaling
pseudohyphal growth
protein modification by small protein conjugation or removal
invasive growth in response to glucose limitation
histone modification
DNA repair
cytoskeleton organization
chromatin organization
cellular response to DNA damage stimulus
Biological Process
0.00
0.03
0.06
0.09
Proportion