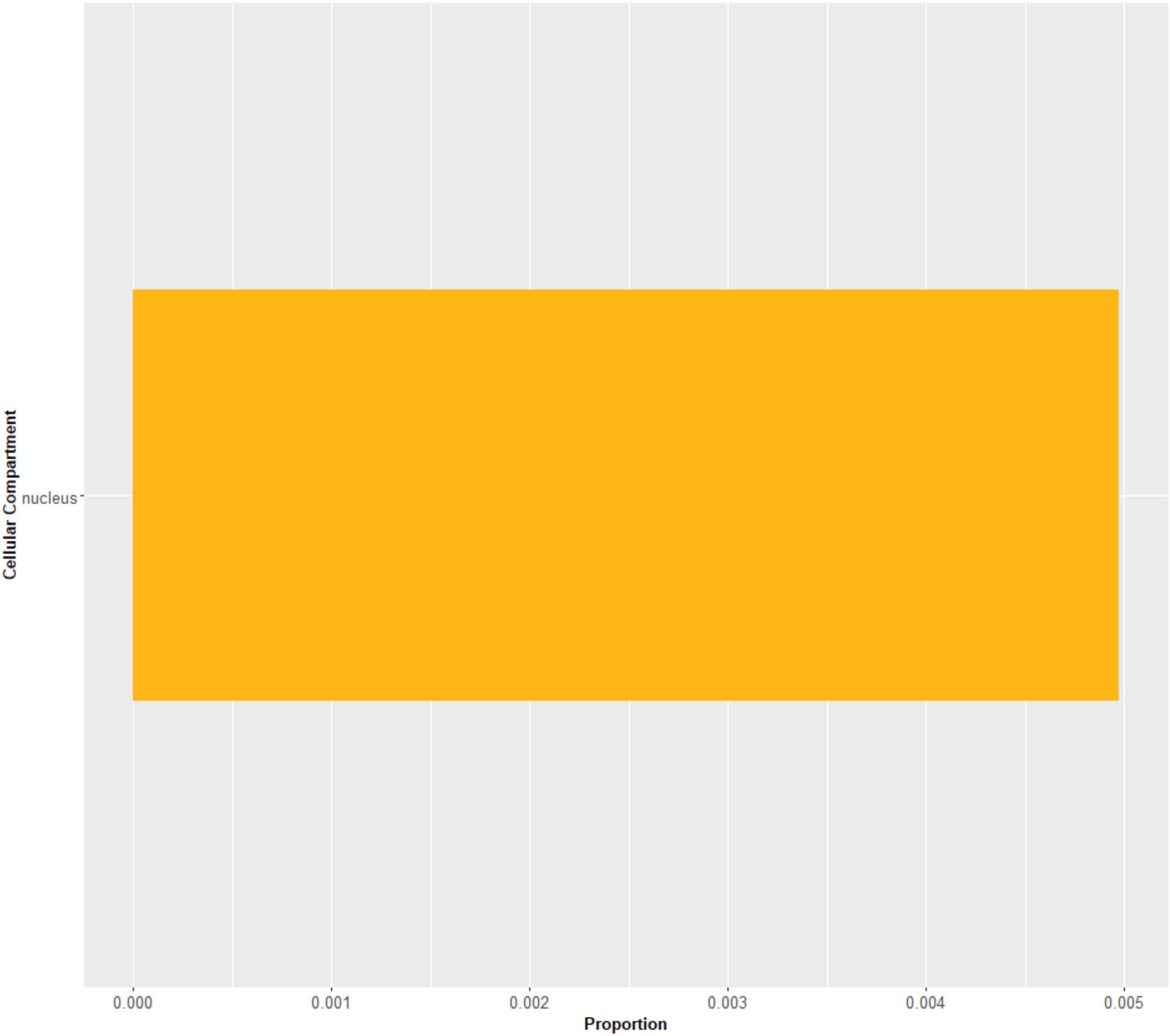
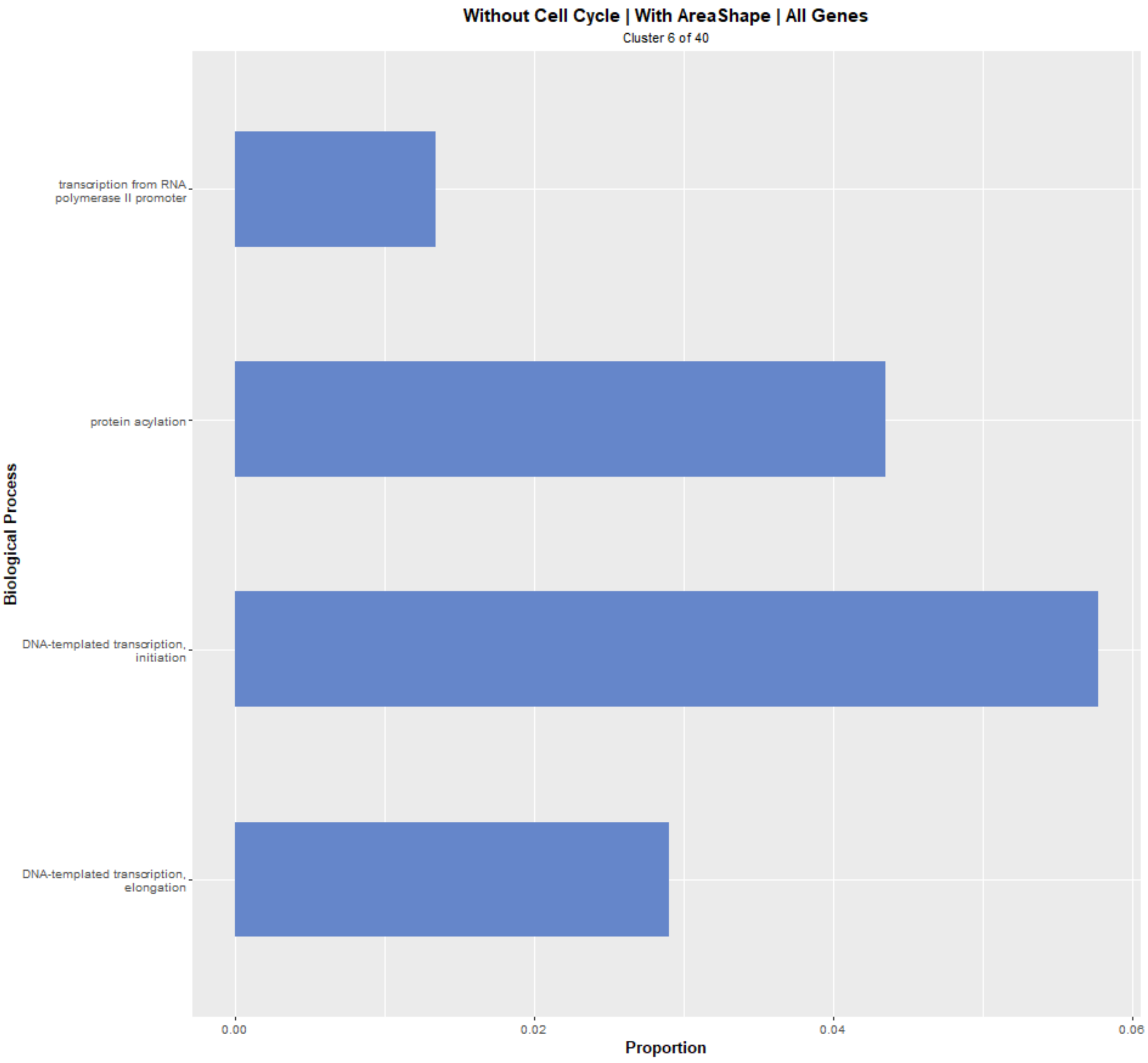
Without Cell Cycle | With AreaShape | All Genes

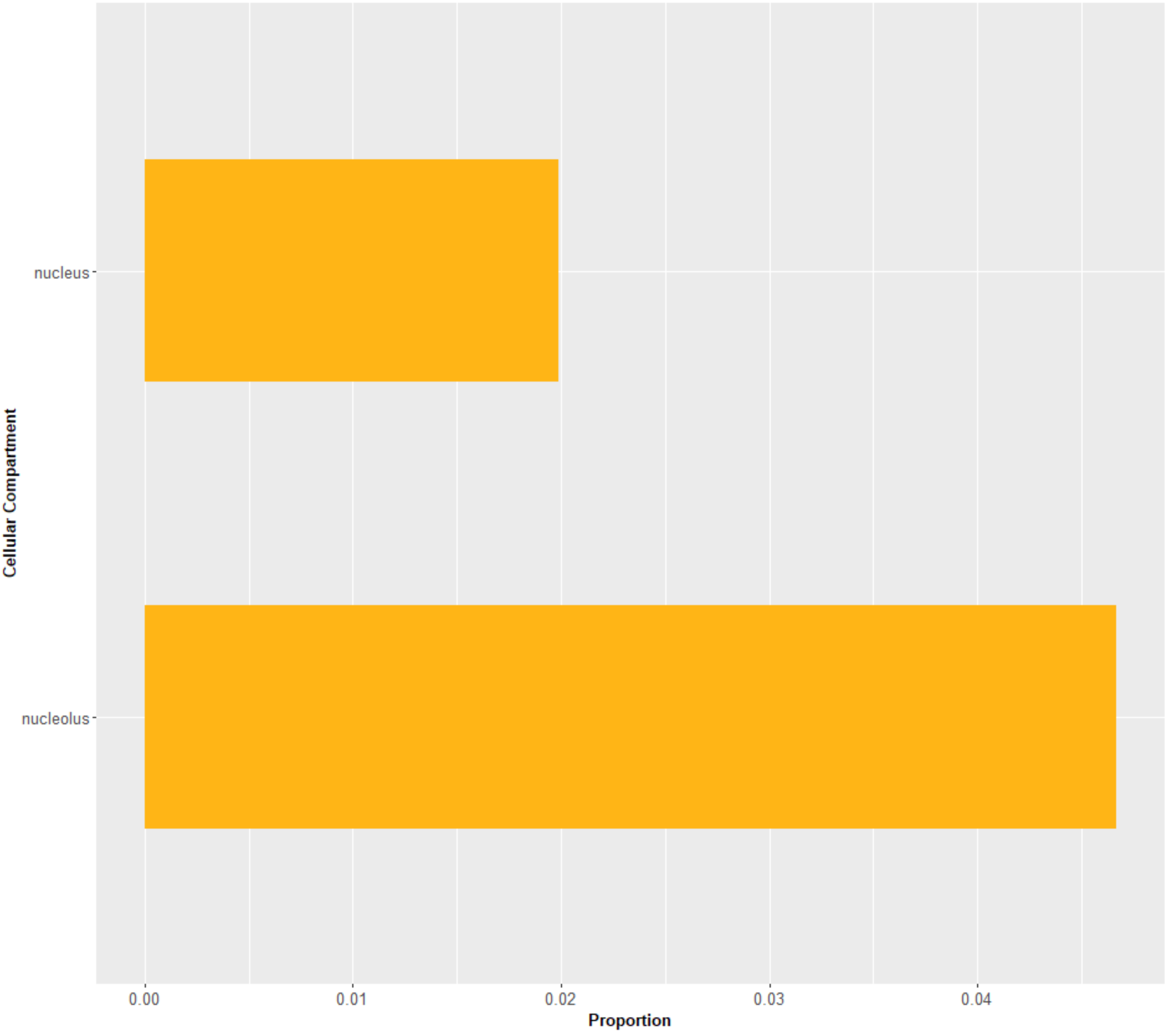
Cluster 6 of 40




Without Cell Cycle | With AreaShape | All Genes
Cluster 6 of 40
transcription from RNA polymerase II promoter
protein acylation
DNA-templated transcription, initiation
DNA-templated transcription, elongation
Biological Process
0.00
0.02
0.04
0.06
Proportion

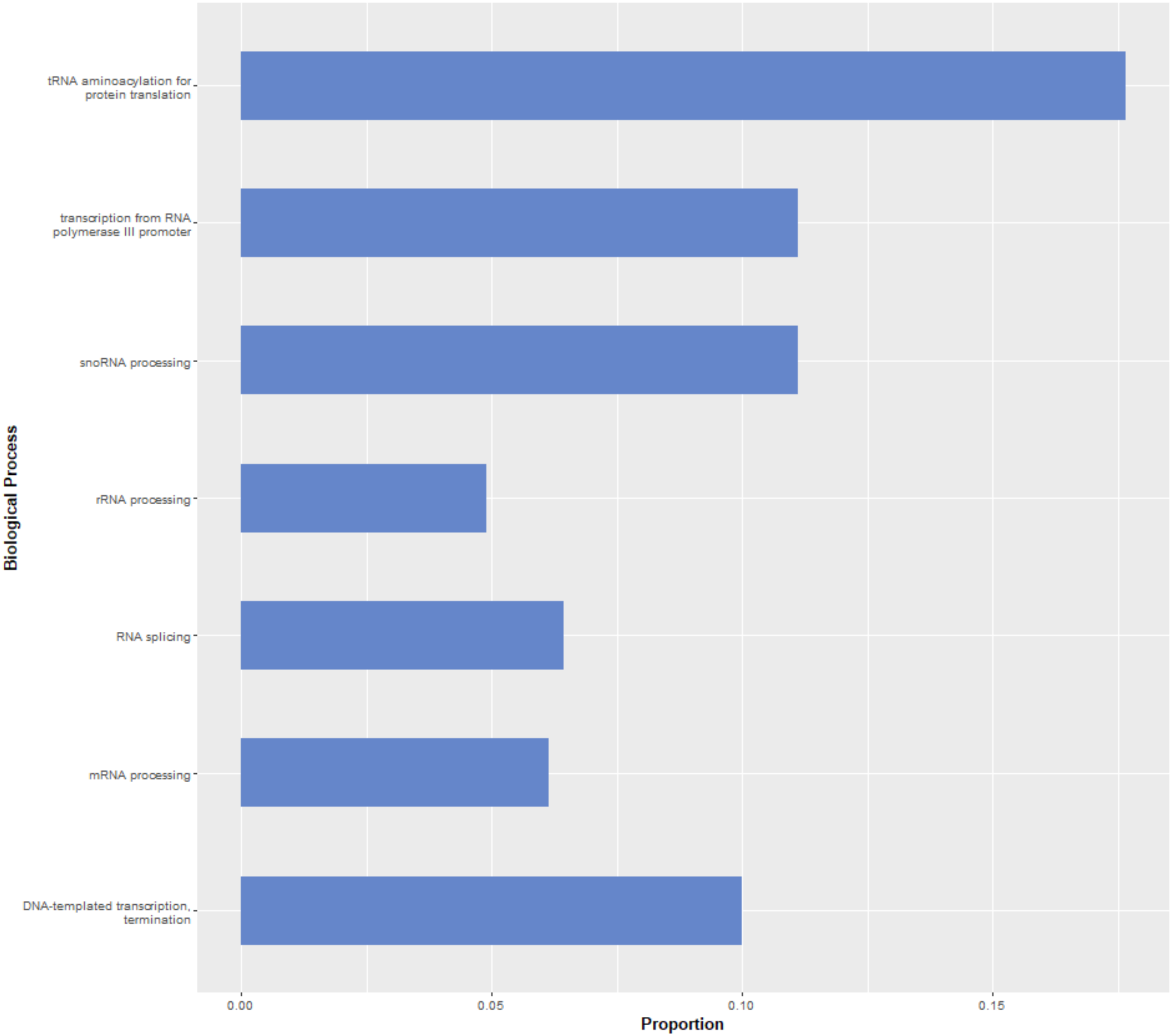
Without Cell Cycle | With AreaShape | All Genes

Cluster 7 of 40



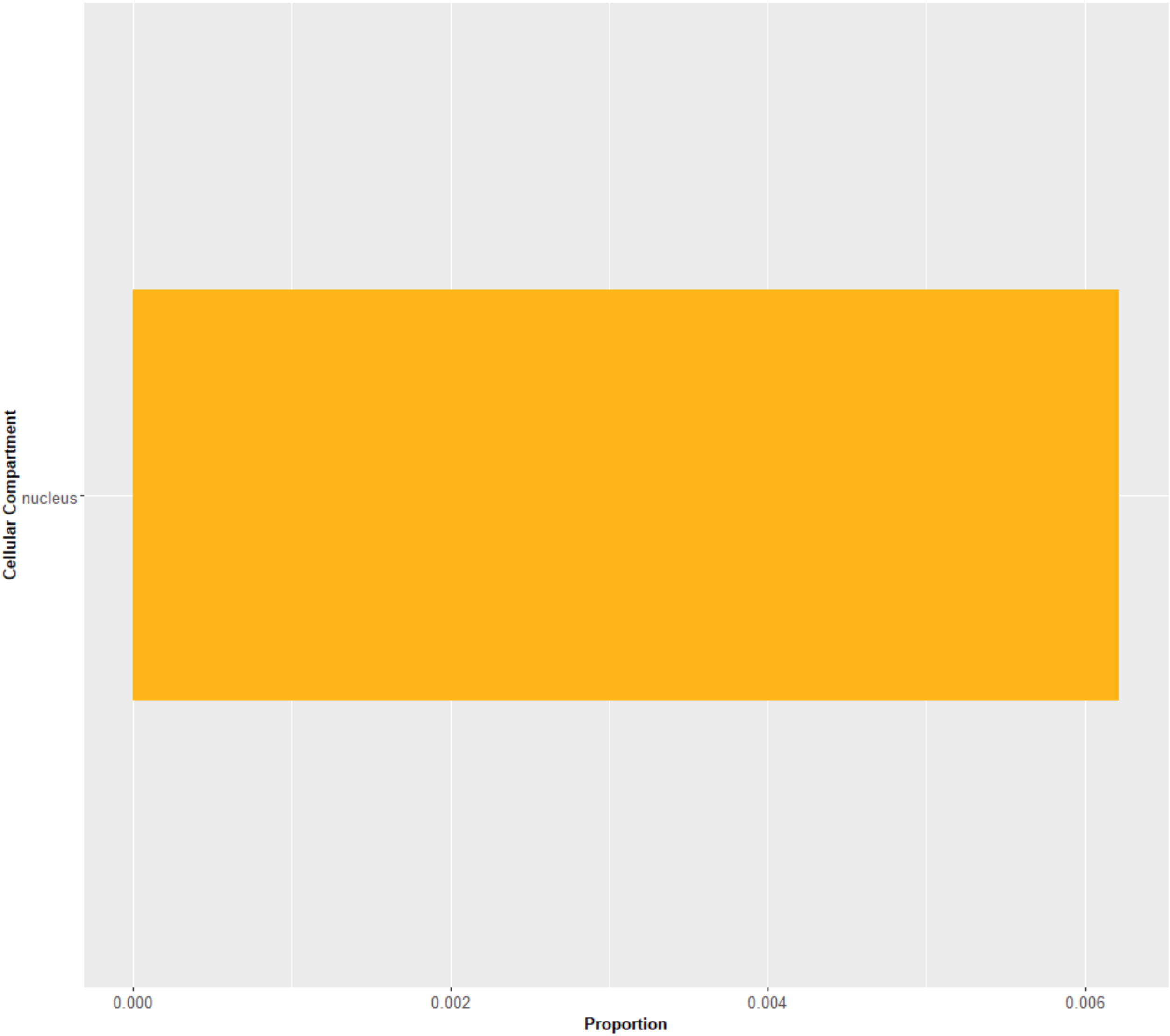
Without Cell Cycle | With AreaShape | All Genes

Cluster 7 of 40



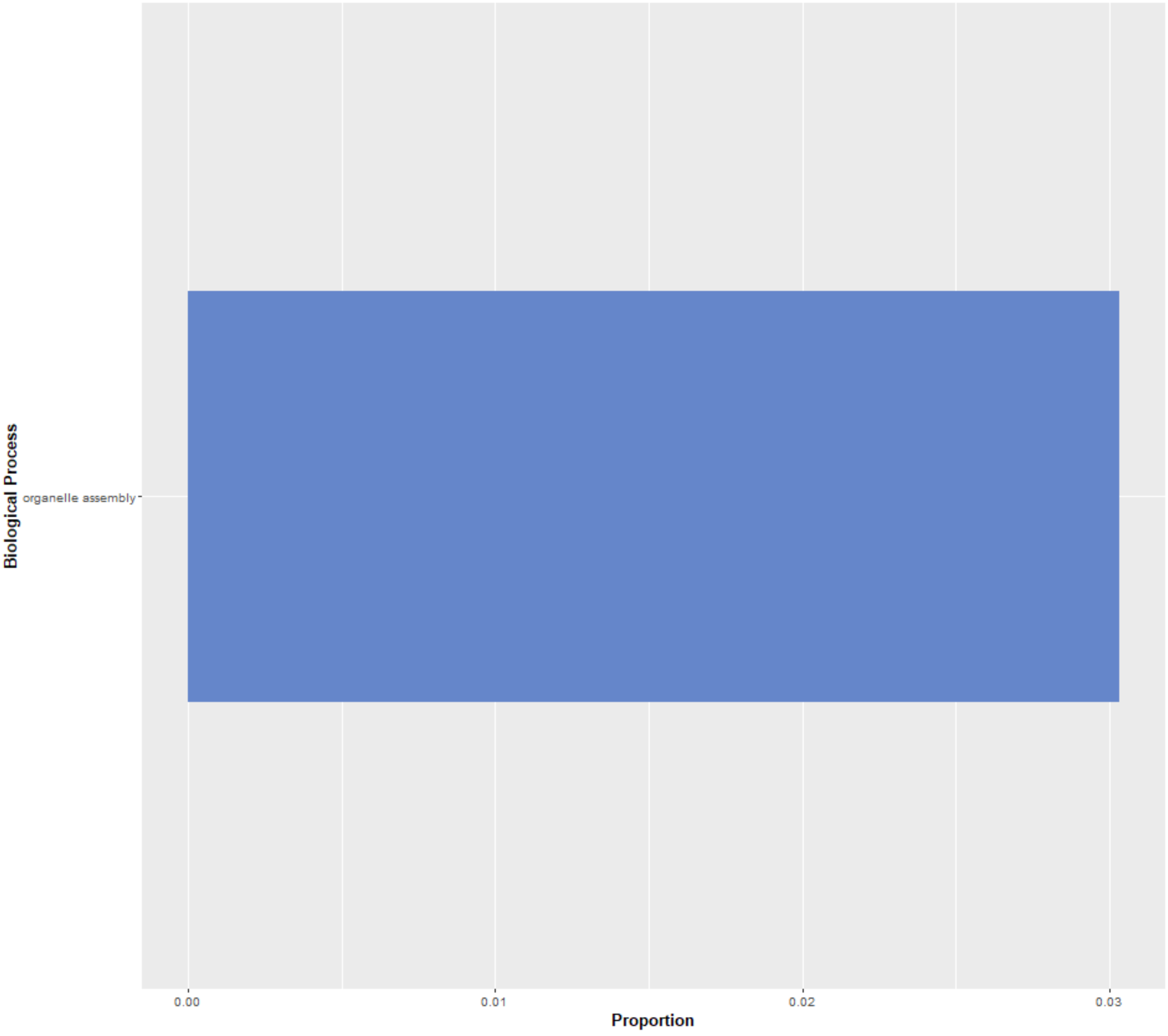
Without Cell Cycle | With AreaShape | All Genes

Cluster 8 of 40



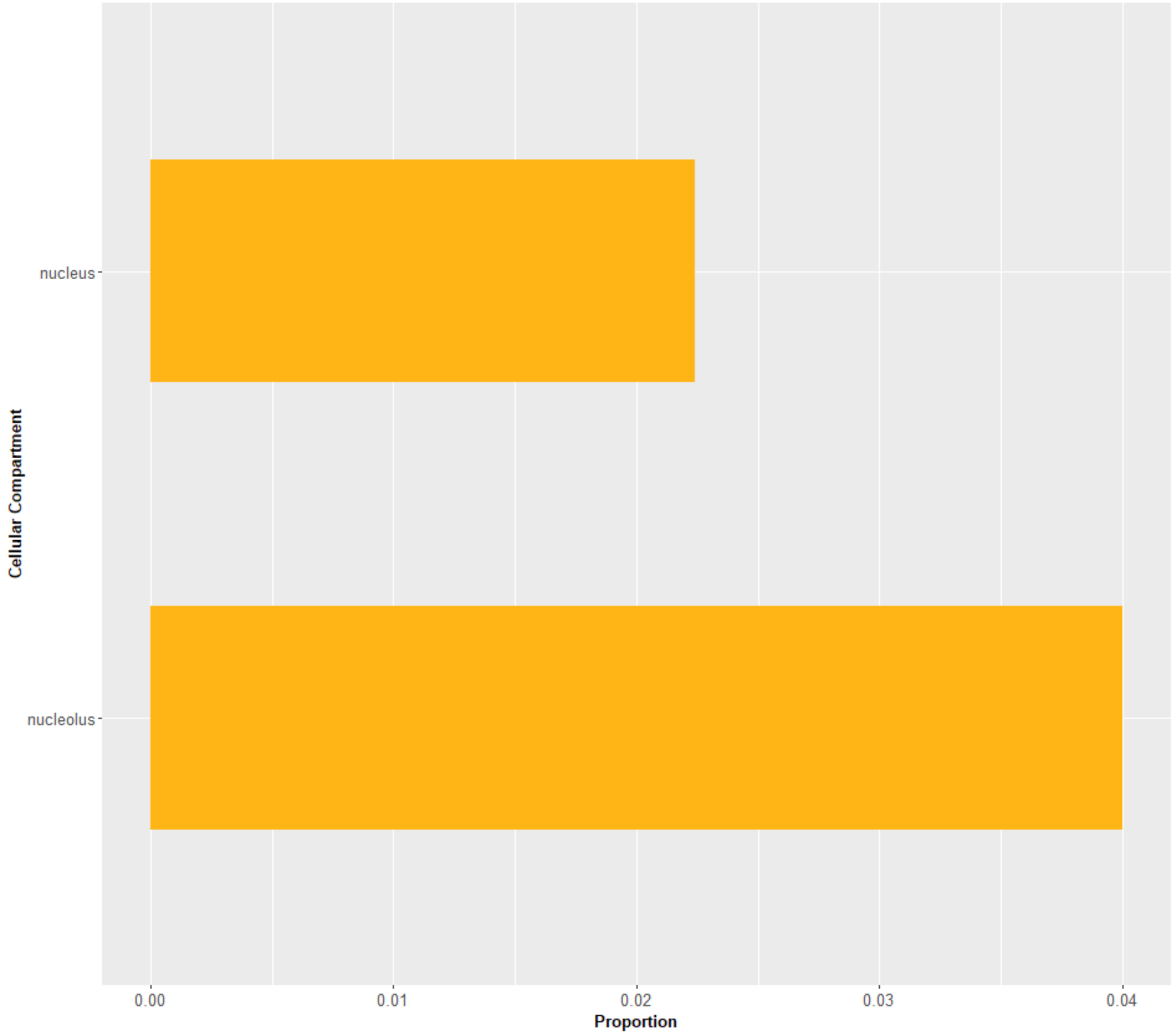
Without Cell Cycle | With AreaShape | All Genes

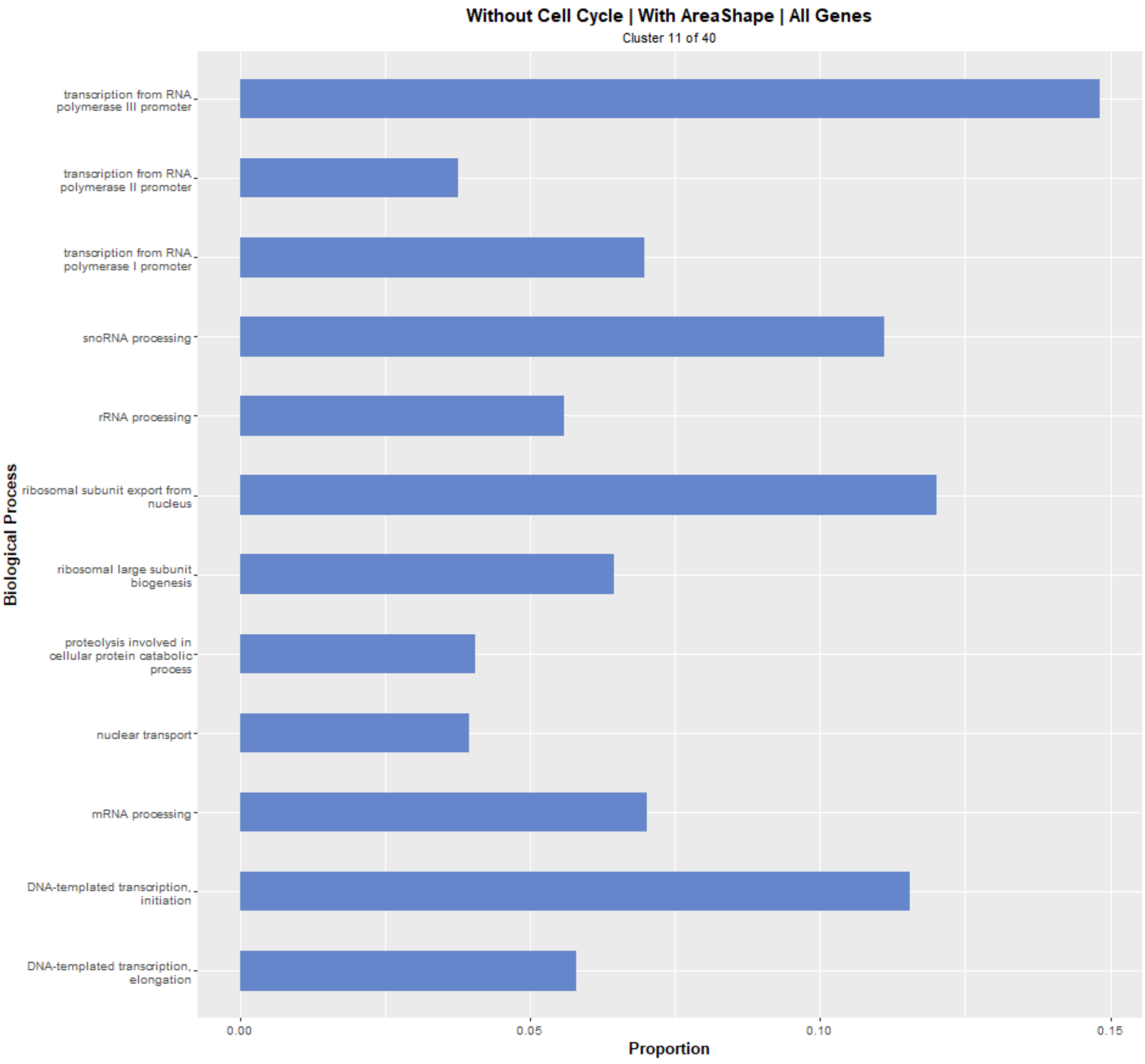
Cluster 8 of 40

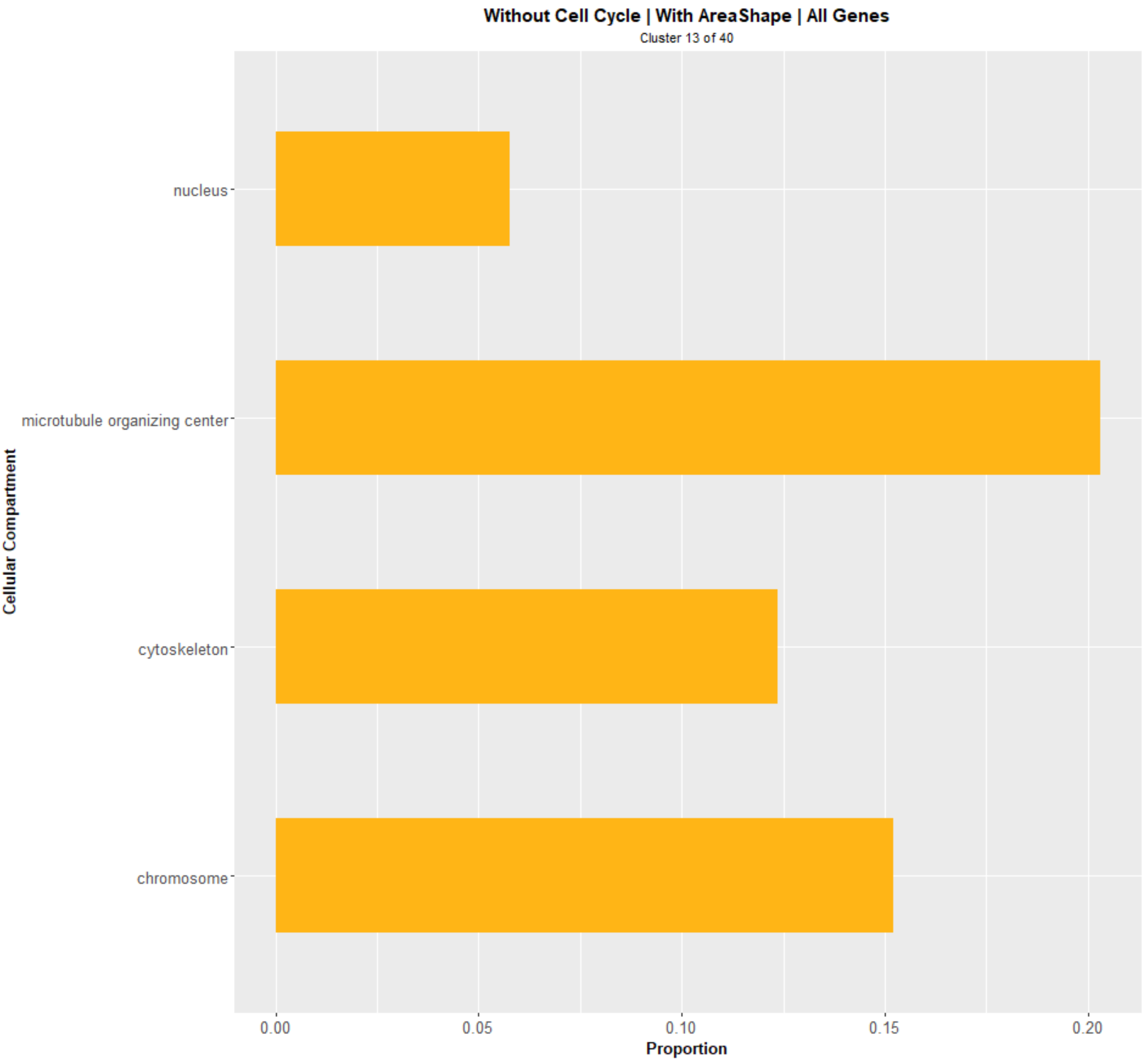


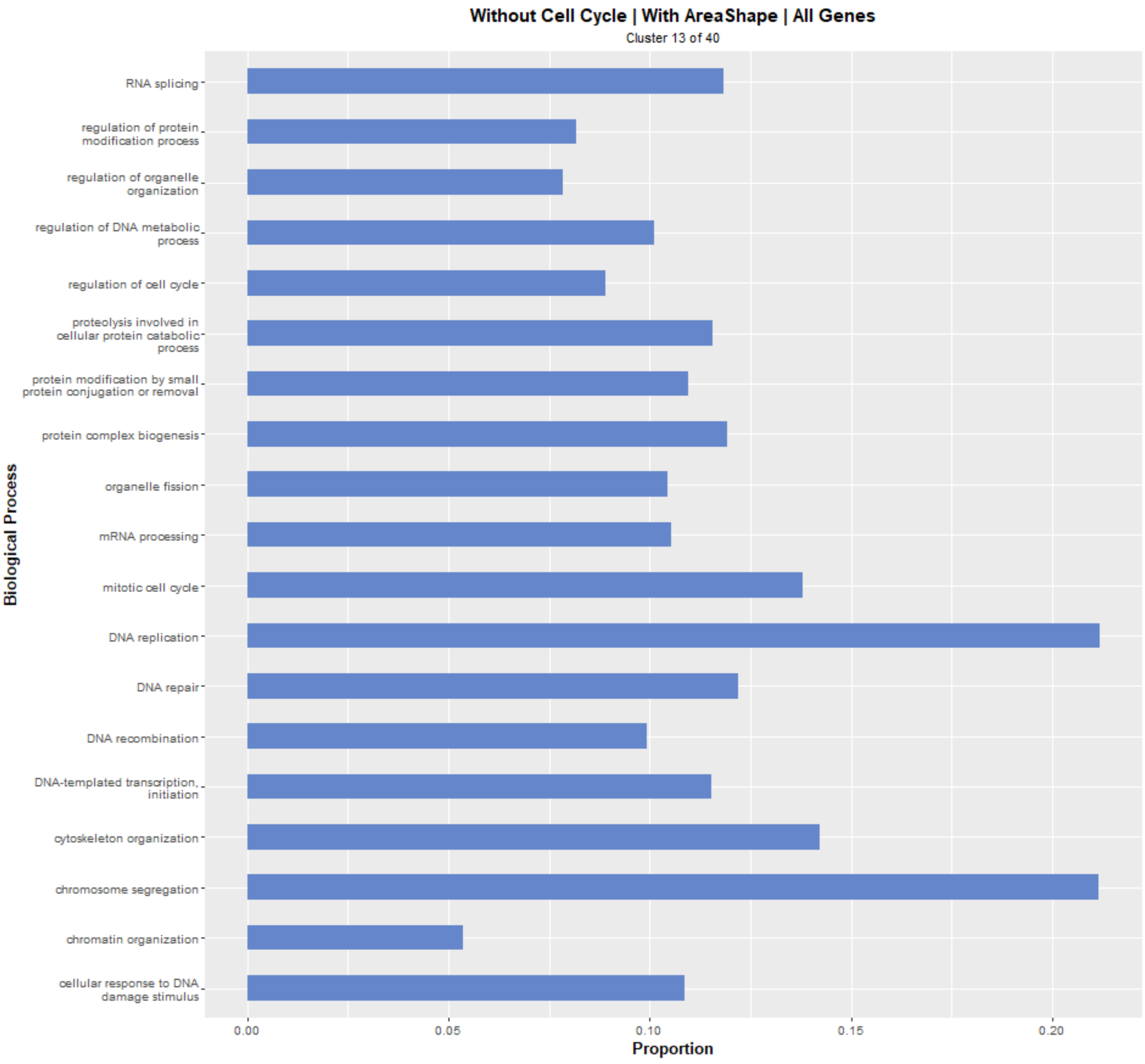
Without Cell Cycle | With AreaShape | All Genes

Cluster 11 of 40



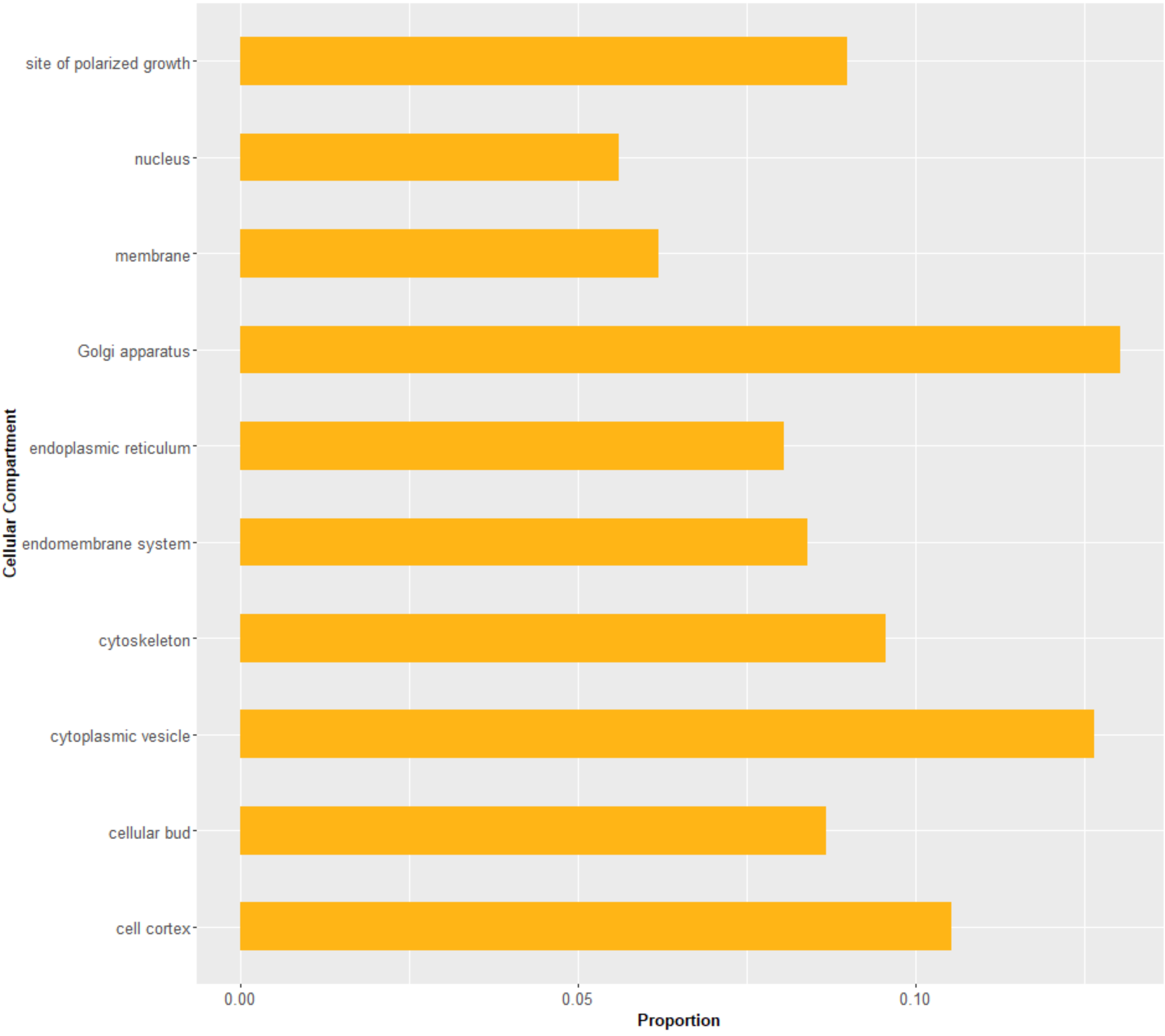

Without Cell Cycle | With AreaShape | All Genes
Cluster 11 of 40
transcription from RNA polymerase III promoter
transcription from RNA polymerase II promoter
transcription from RNA polymerase I promoter
snoRNA processing
rRNA processing
ribosomal subunit export from nucleus
ribosomal large subunit biogenesis
proteolysis involved in cellular protein catabolic process
nuclear transport
mRNA processing
DNA-templated transcription, initiation
DNA-templated transcription, elongation
Biological Process
0.00
0.05
0.10
0.15
Proportion

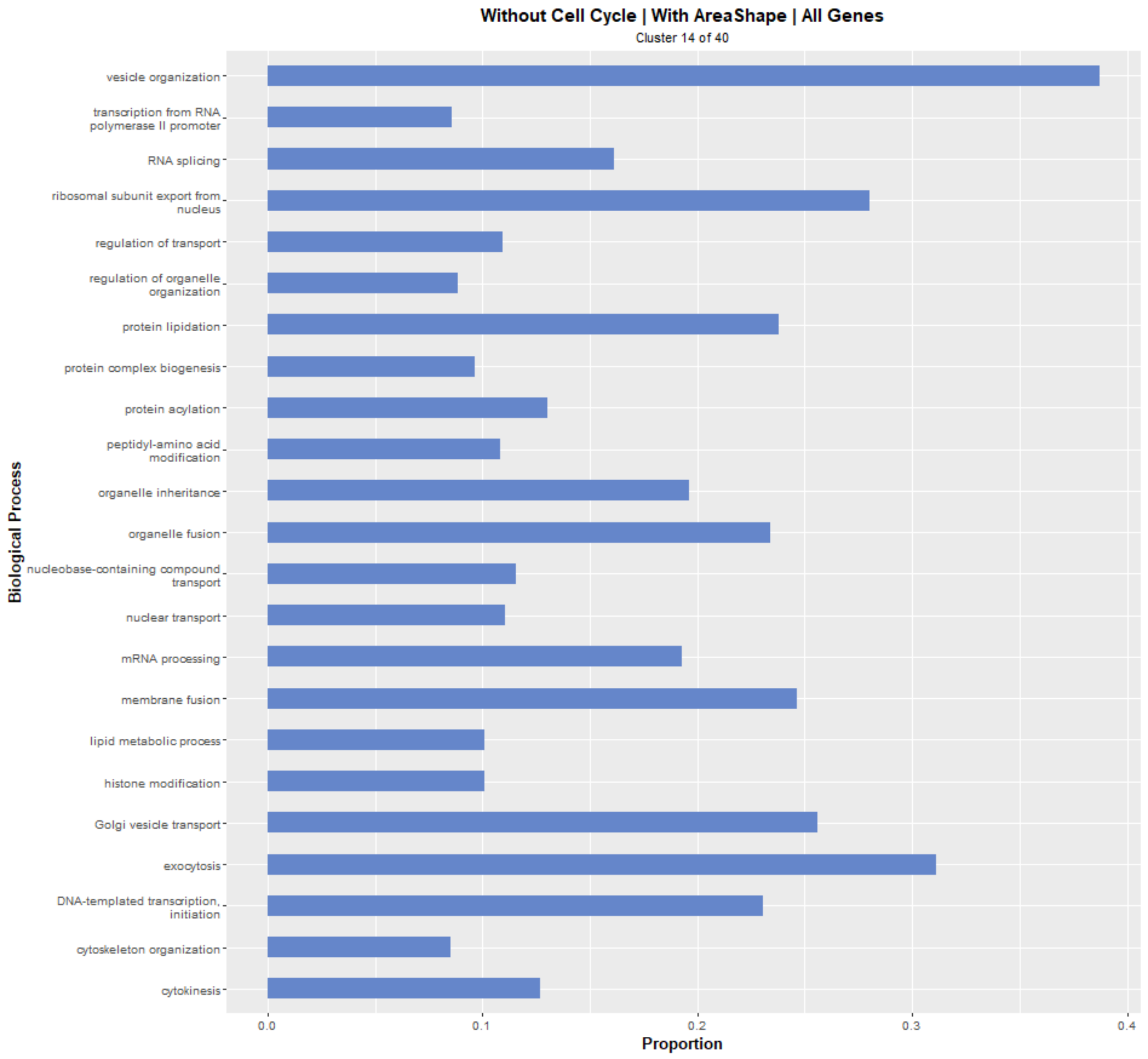

Without Cell Cycle | With AreaShape | All Genes
Cluster 13 of 40
Cellular Compartment
nucleus
microtubule organizing center
cytoskeleton
chromosome
0.00
0.05
0.10
0.15
0.20
Proportion


Without Cell Cycle | With AreaShape | All Genes
Cluster 13 of 40
RNA splicing
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 modification by small protein conjugation or removal
protein complex biogenesis
organelle fission
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
0.00
0.05
0.10
0.15
0.20
Proportion

Without Cell Cycle | With AreaShape | All Genes

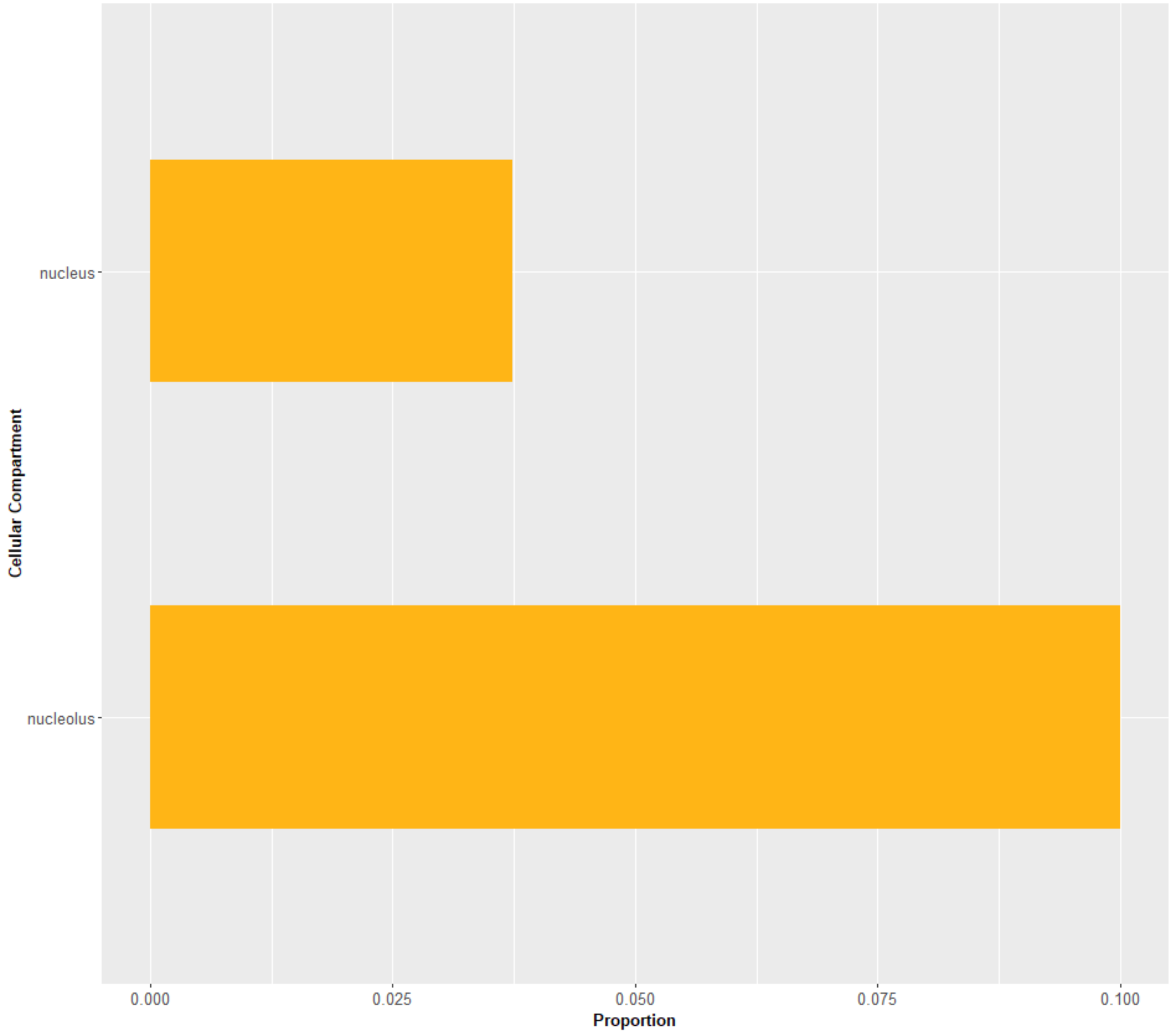
Cluster 14 of 40

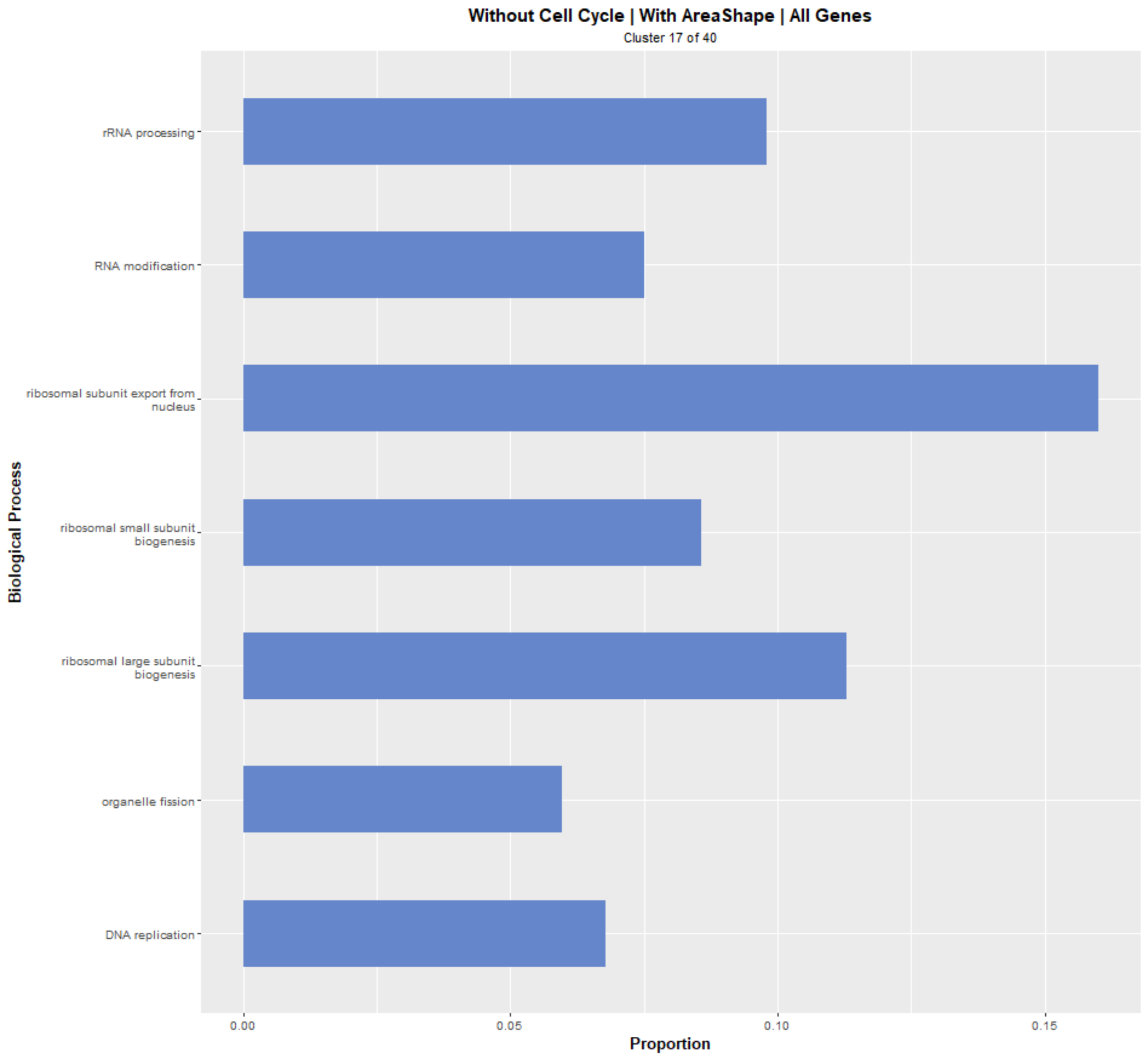



Without Cell Cycle | With AreaShape | All Genes
Cluster 14 of 40
Biological Process
vesicle organization
transcription from RNA polymerase II promoter
RNA splicing
ribosomal subunit export from nucleus
regulation of transport
regulation of organelle organization
protein lipidation
protein complex biogenesis
protein acylation
peptidyl-amino acid modification
organelle inheritance
organelle fusion
nucleobase-containing compound transport
nuclear transport
mRNA processing
membrane fusion
lipid metabolic process
histone modification
Golgi vesicle transport
exocytosis
DNA-templated transcription, initiation
cytoskeleton organization
cytokinesis
0.0
0.1
0.2
0.3
0.4
Proportion

Without Cell Cycle | With AreaShape | All Genes

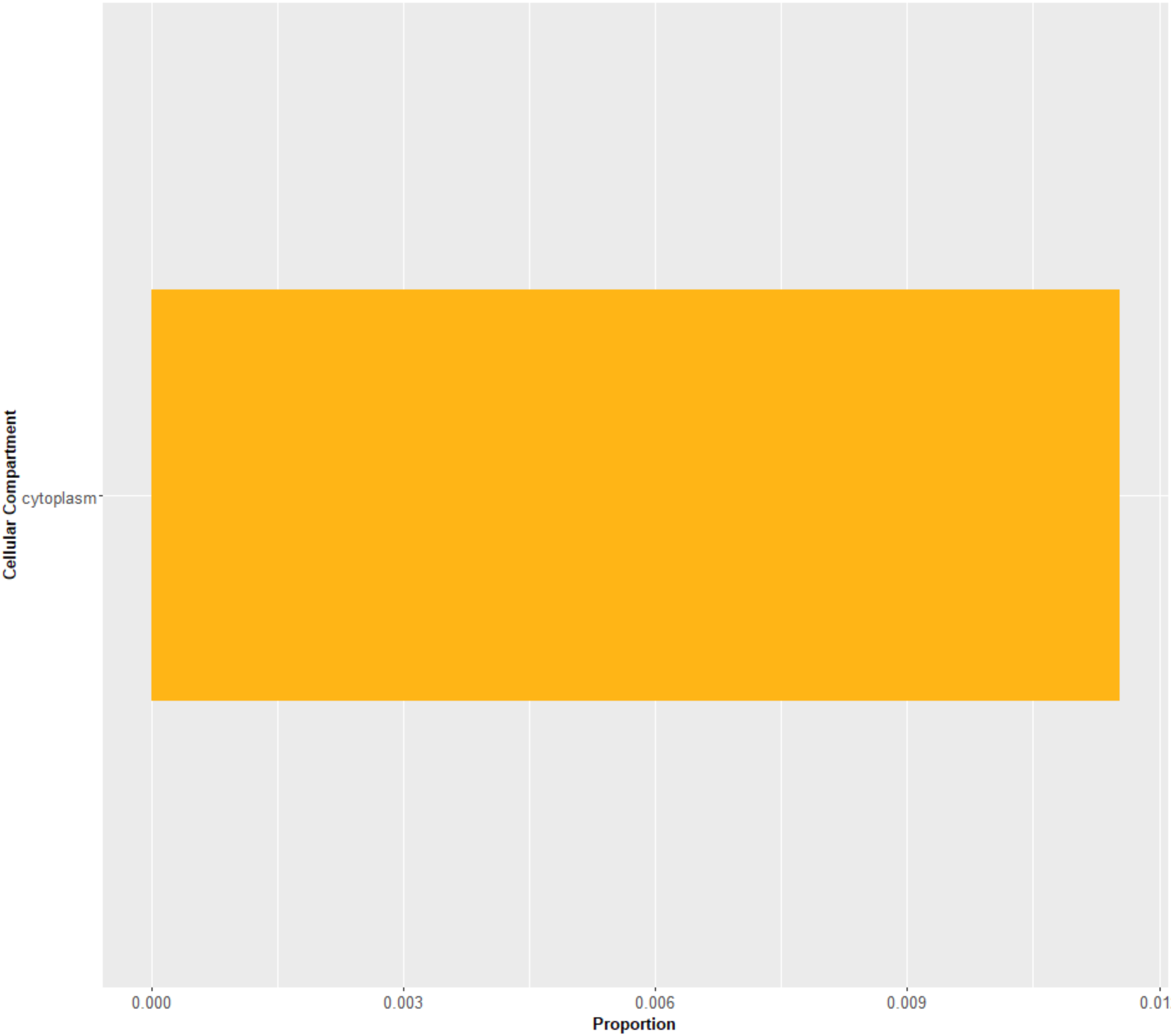
Cluster 17 of 40




Without Cell Cycle | With AreaShape | All Genes
Cluster 17 of 40
rRNA processing
RNA modification
ribosomal subunit export from nucleus
ribosomal small subunit biogenesis
ribosomal large subunit biogenesis
organelle fission
DNA replication
Biological Process
0.00
0.05
0.10
0.15
Proportion

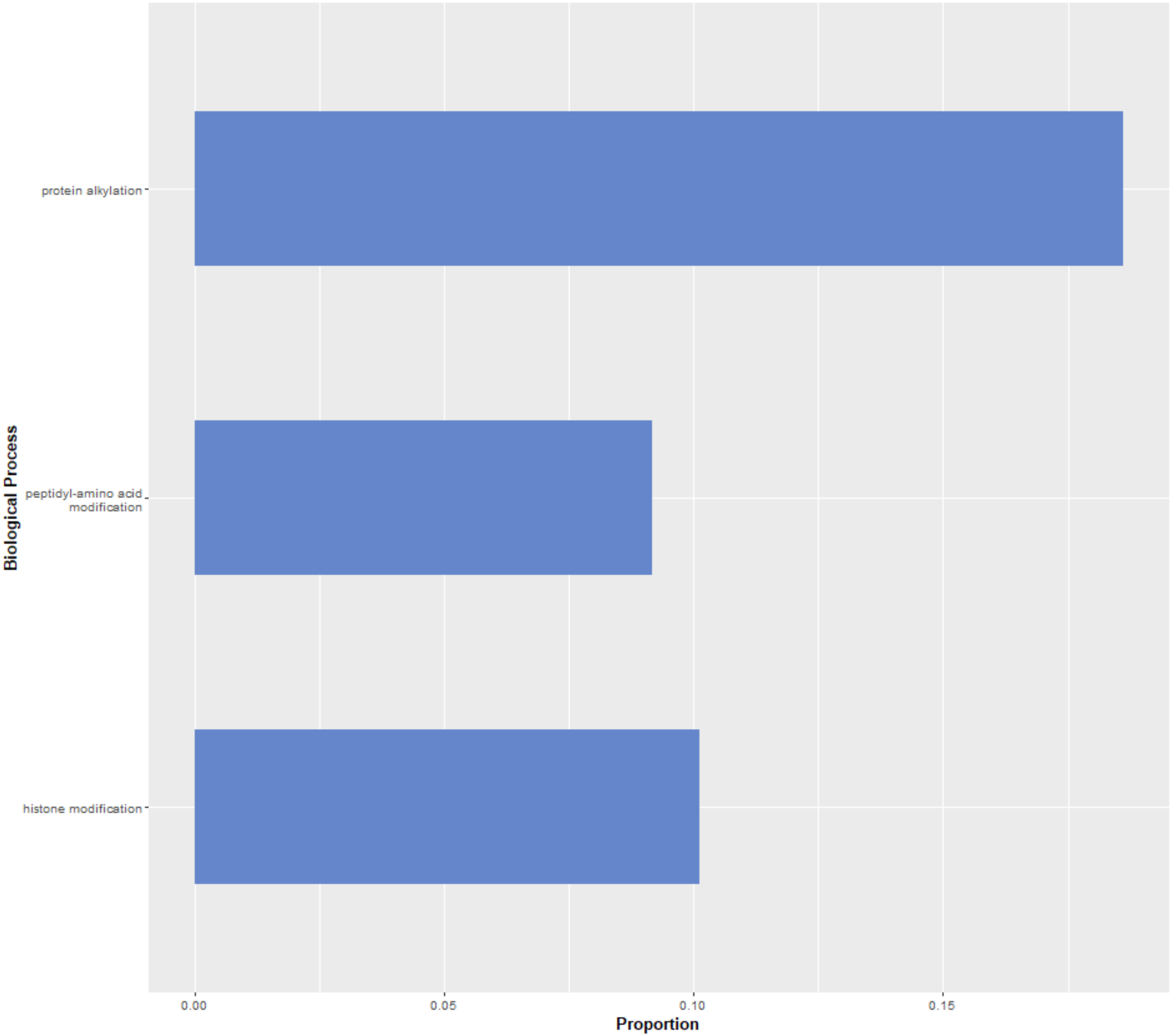
Without Cell Cycle | With AreaShape | All Genes

Cluster 18 of 40



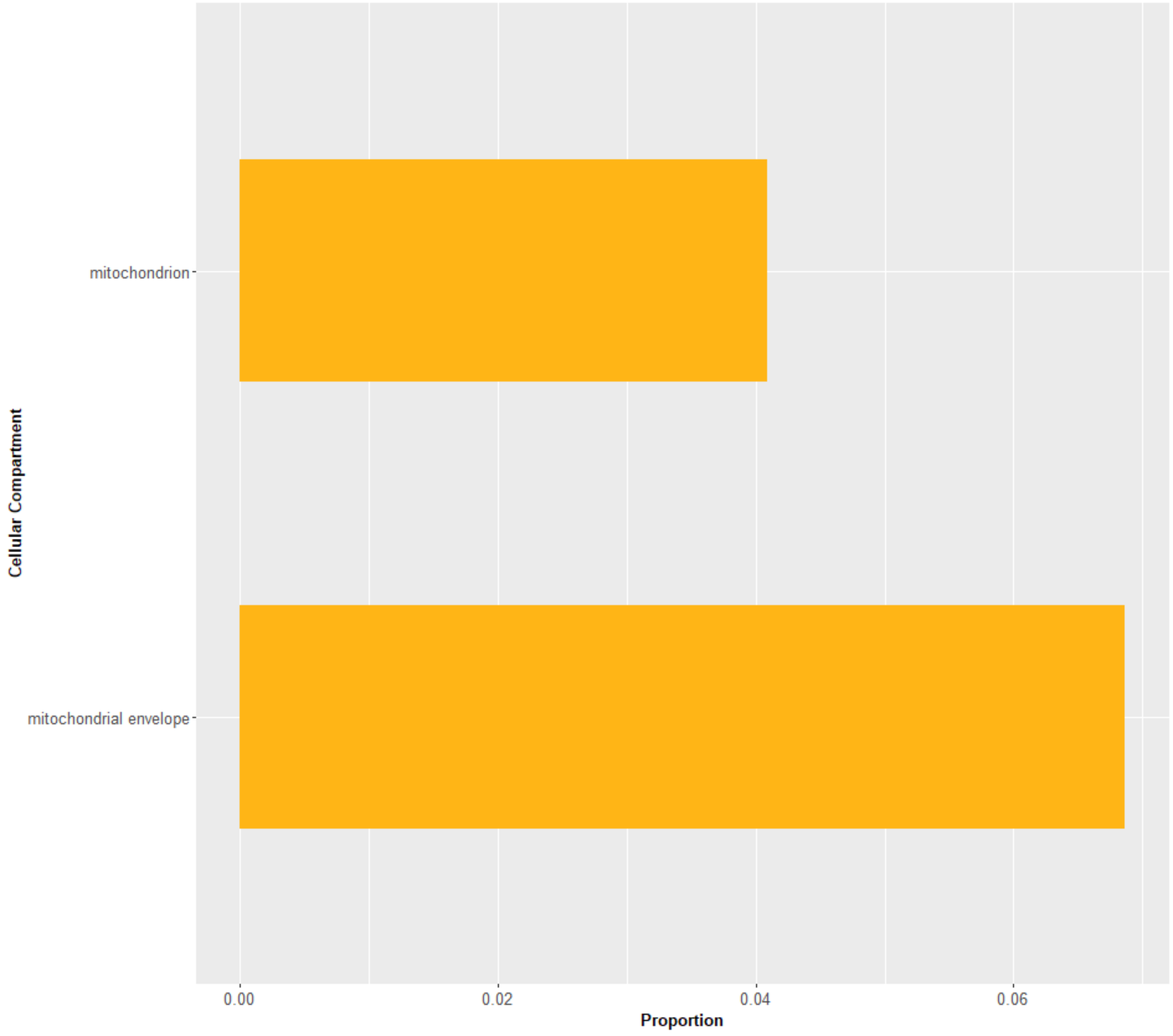
Without Cell Cycle | With AreaShape | All Genes

Cluster 19 of 40



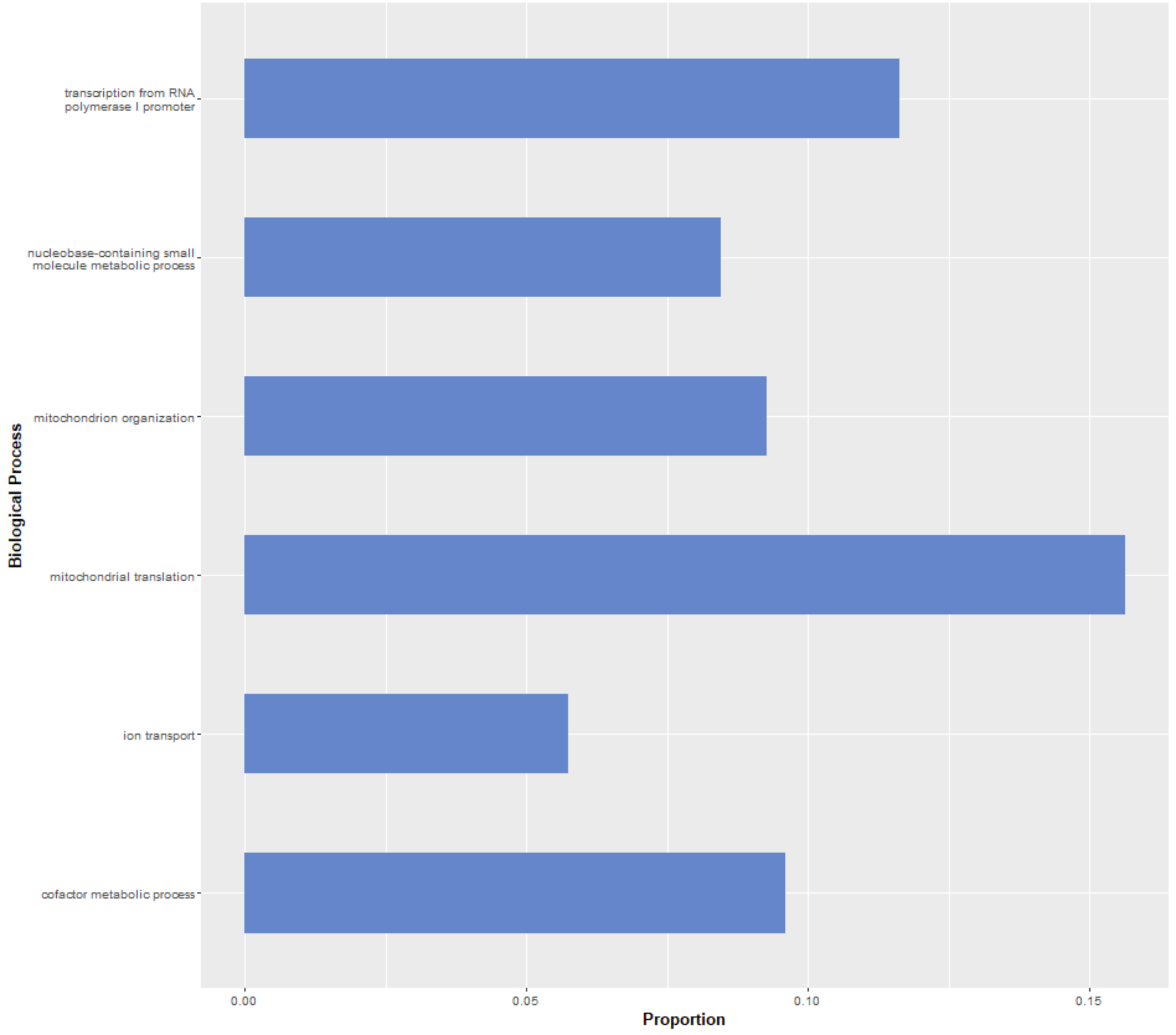
Without Cell Cycle | With AreaShape | All Genes

Cluster 20 of 40



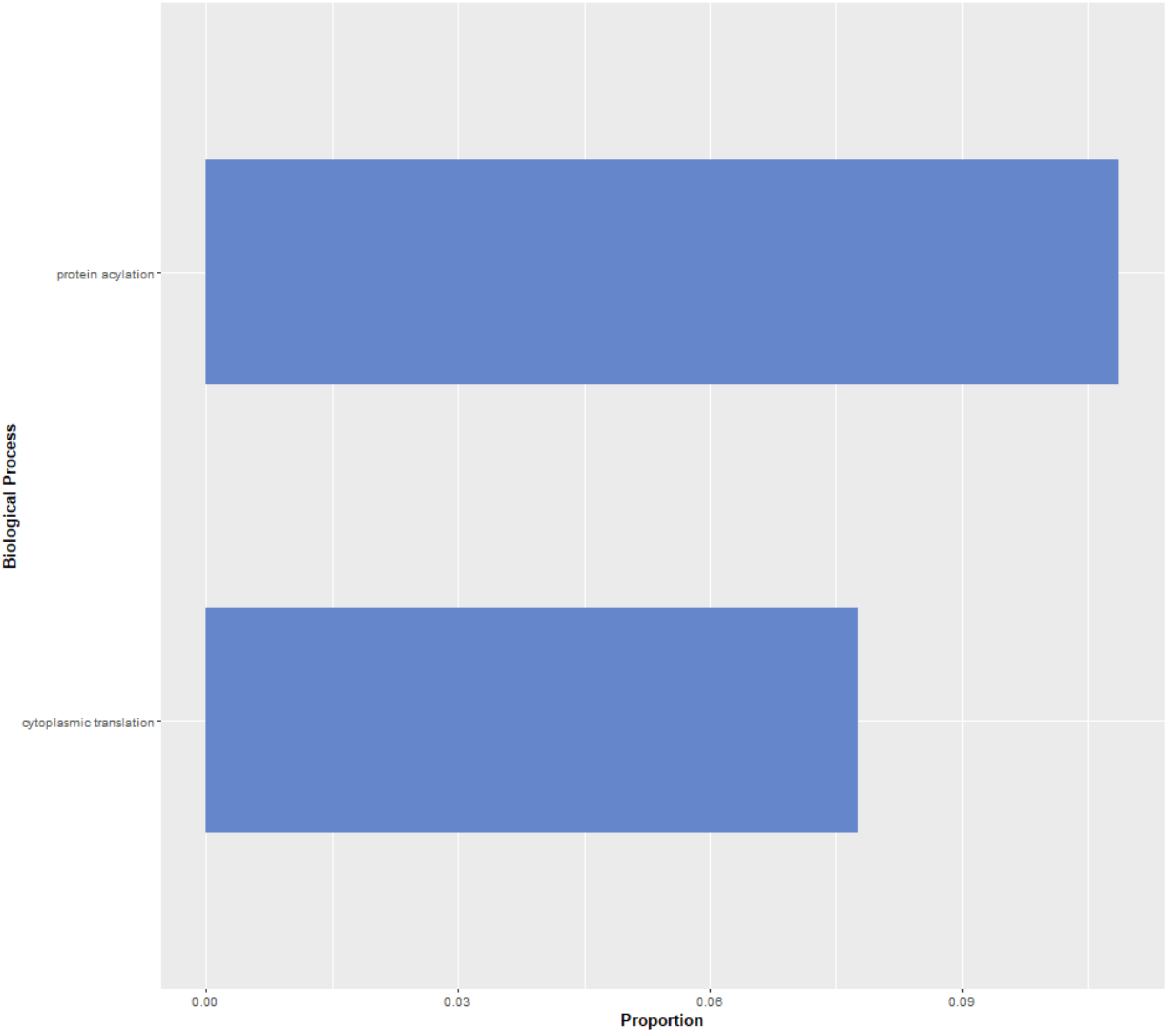
Without Cell Cycle | With AreaShape | All Genes

Cluster 20 of 40



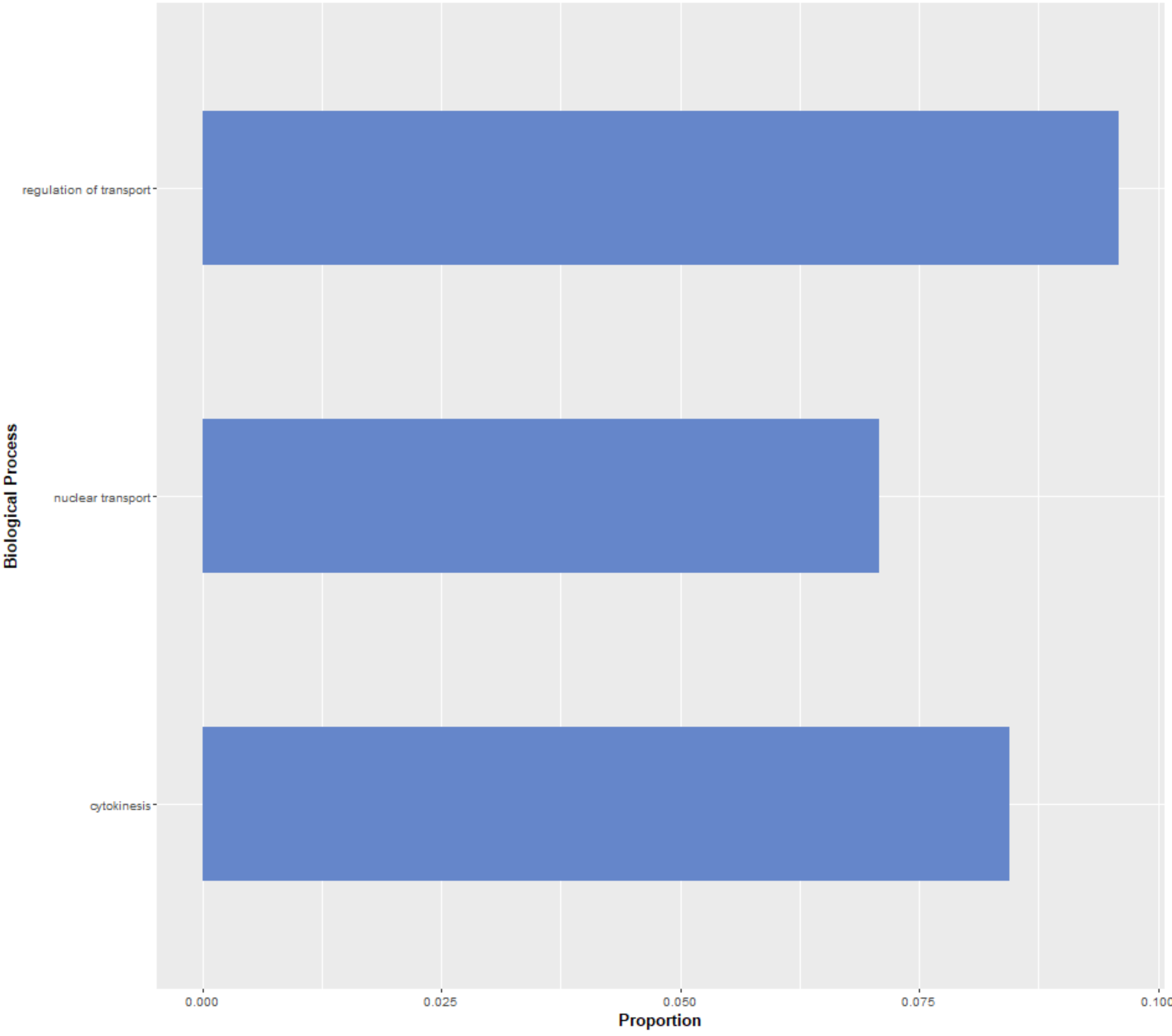
Without Cell Cycle | With AreaShape | All Genes

Cluster 22 of 40



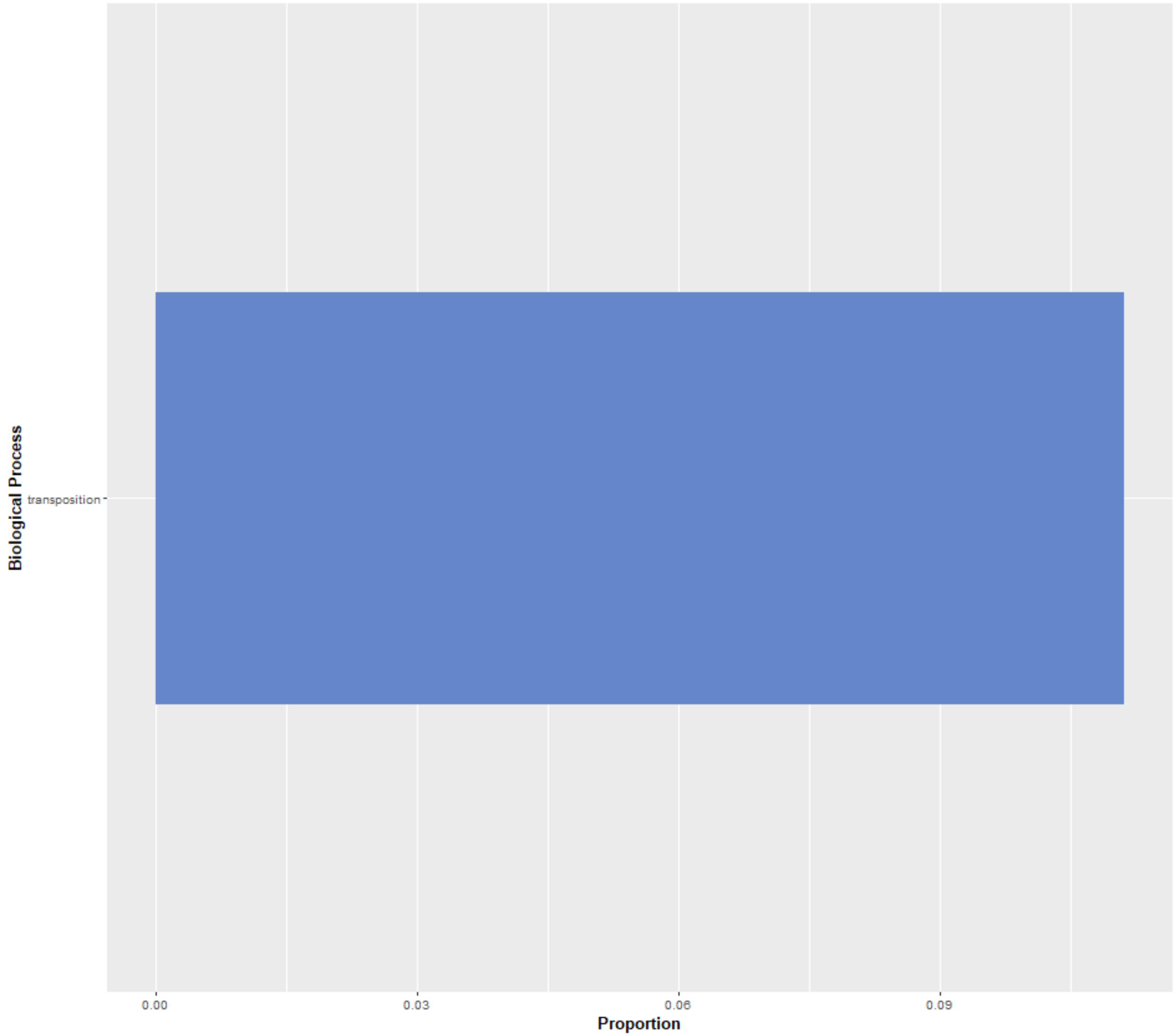
Without Cell Cycle | With AreaShape | All Genes

Cluster 23 of 40



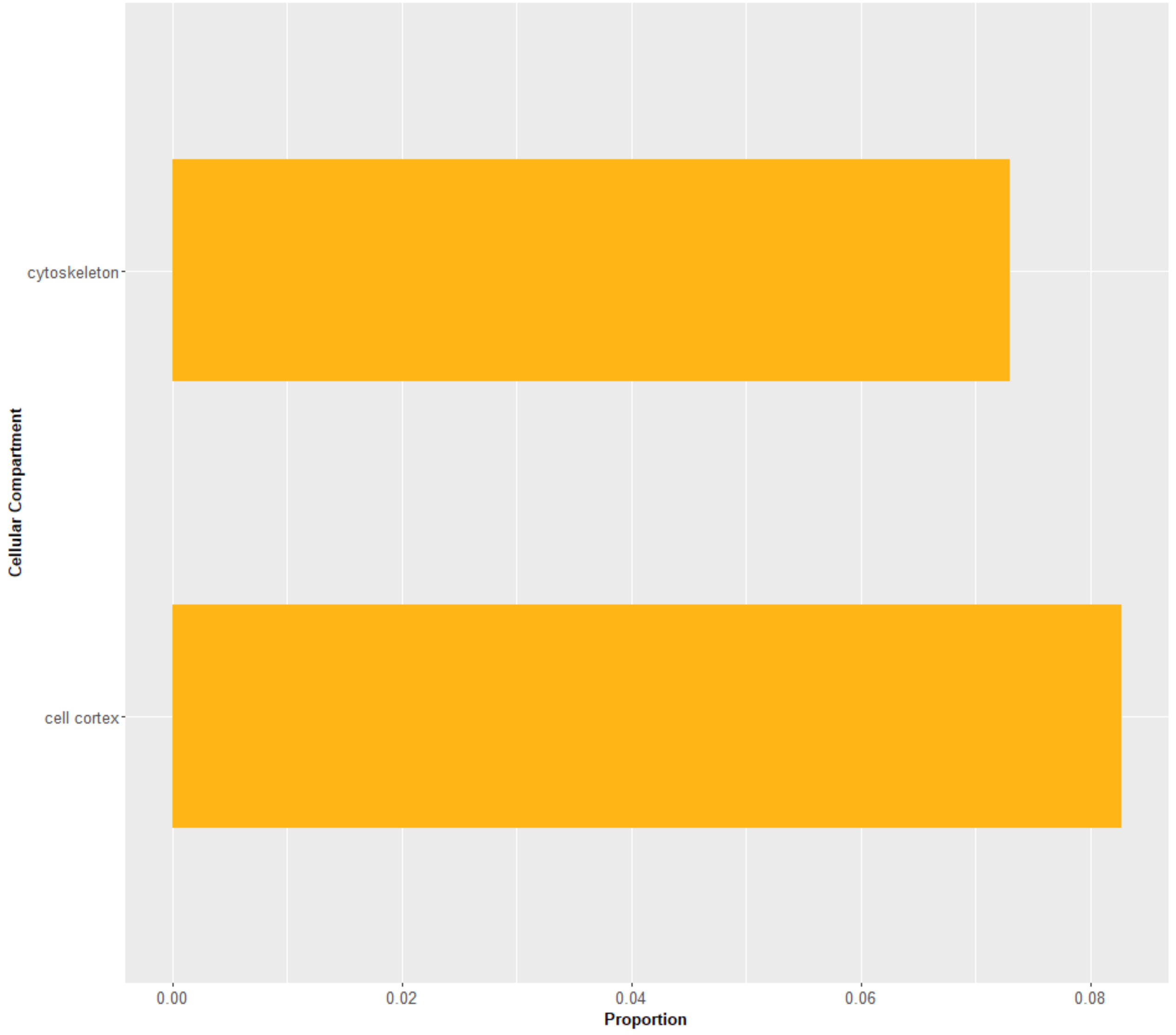
Without Cell Cycle | With AreaShape | All Genes

Cluster 25 of 40



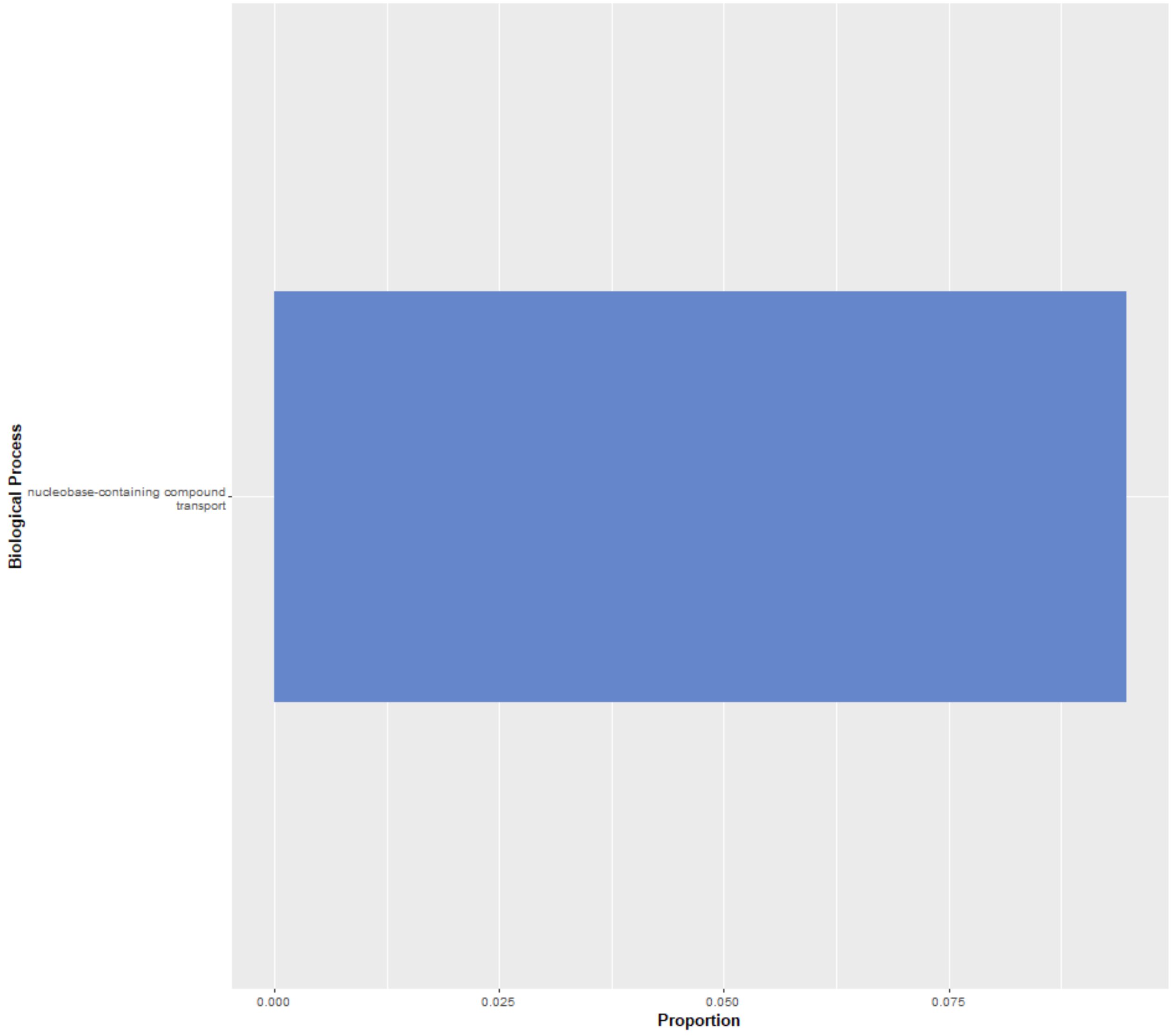
Without Cell Cycle | With AreaShape | All Genes

Cluster 27 of 40



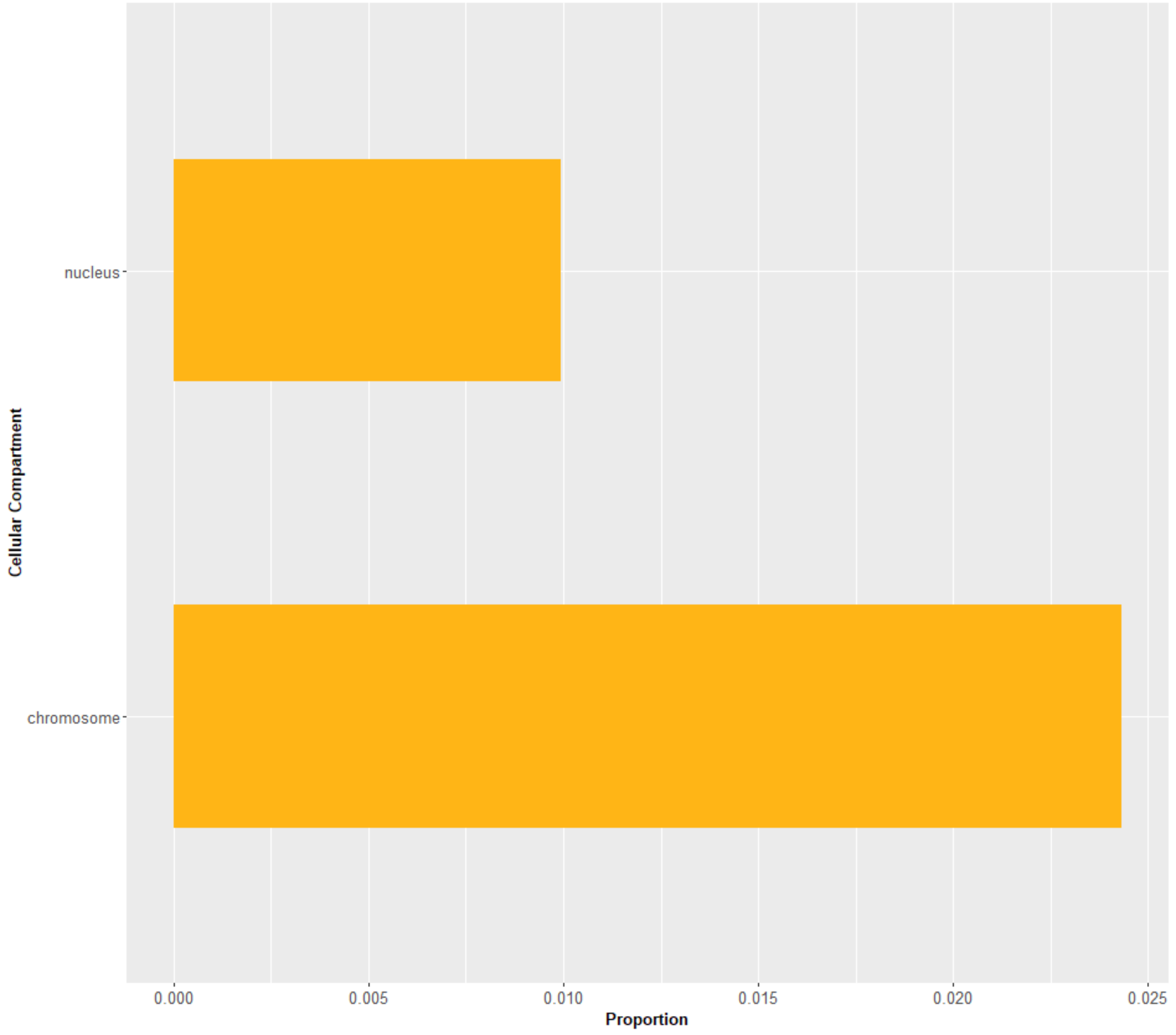
Without Cell Cycle | With AreaShape | All Genes

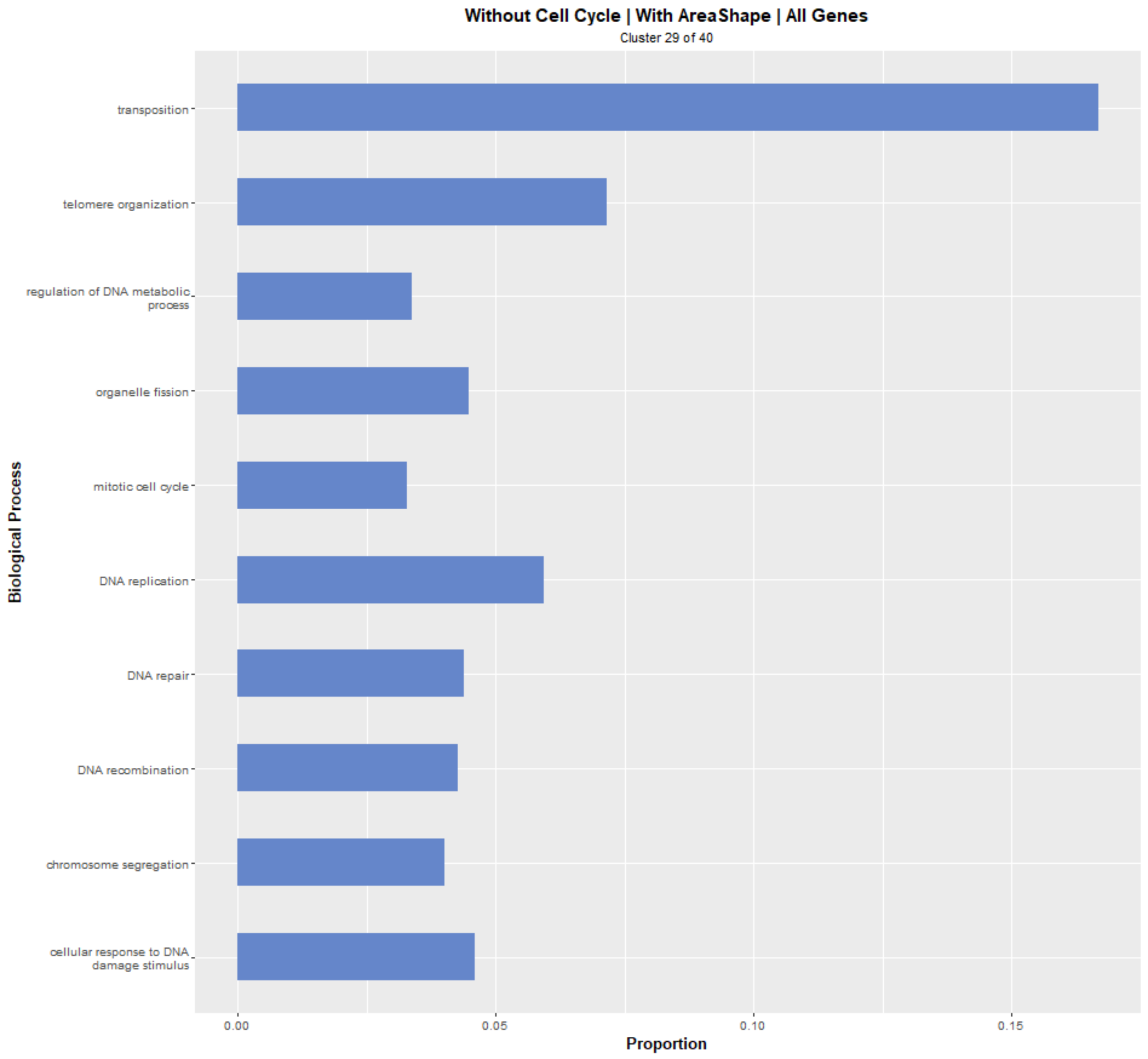
Cluster 27 of 40



Without Cell Cycle | With AreaShape | All Genes

Cluster 29 of 40




Without Cell Cycle | With AreaShape | All Genes
Cluster 29 of 40
transposition
telomere organization
regulation of DNA metabolic process
organelle fission
mitotic cell cycle
DNA replication
DNA repair
DNA recombination
chromosome segregation
cellular response to DNA damage stimulus
Biological Process
0.00
0.05
0.10
0.15
Proportion

Without Cell Cycle | With AreaShape | All Genes

Cluster 30 of 40

Cellular Compartment

site of polarized growth

cellular bud

cell cortex

0.00

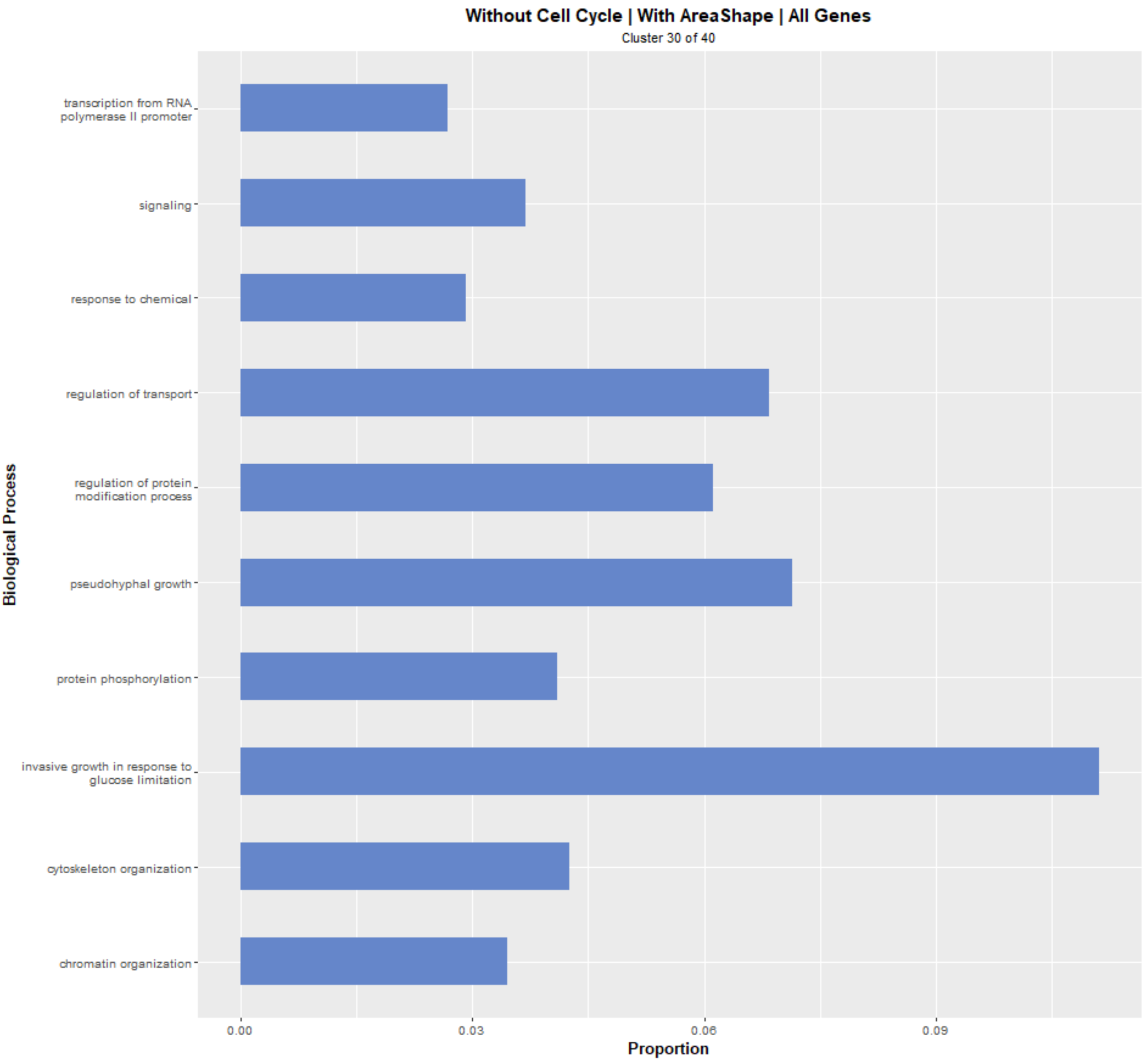
0.01

0.02

0.03

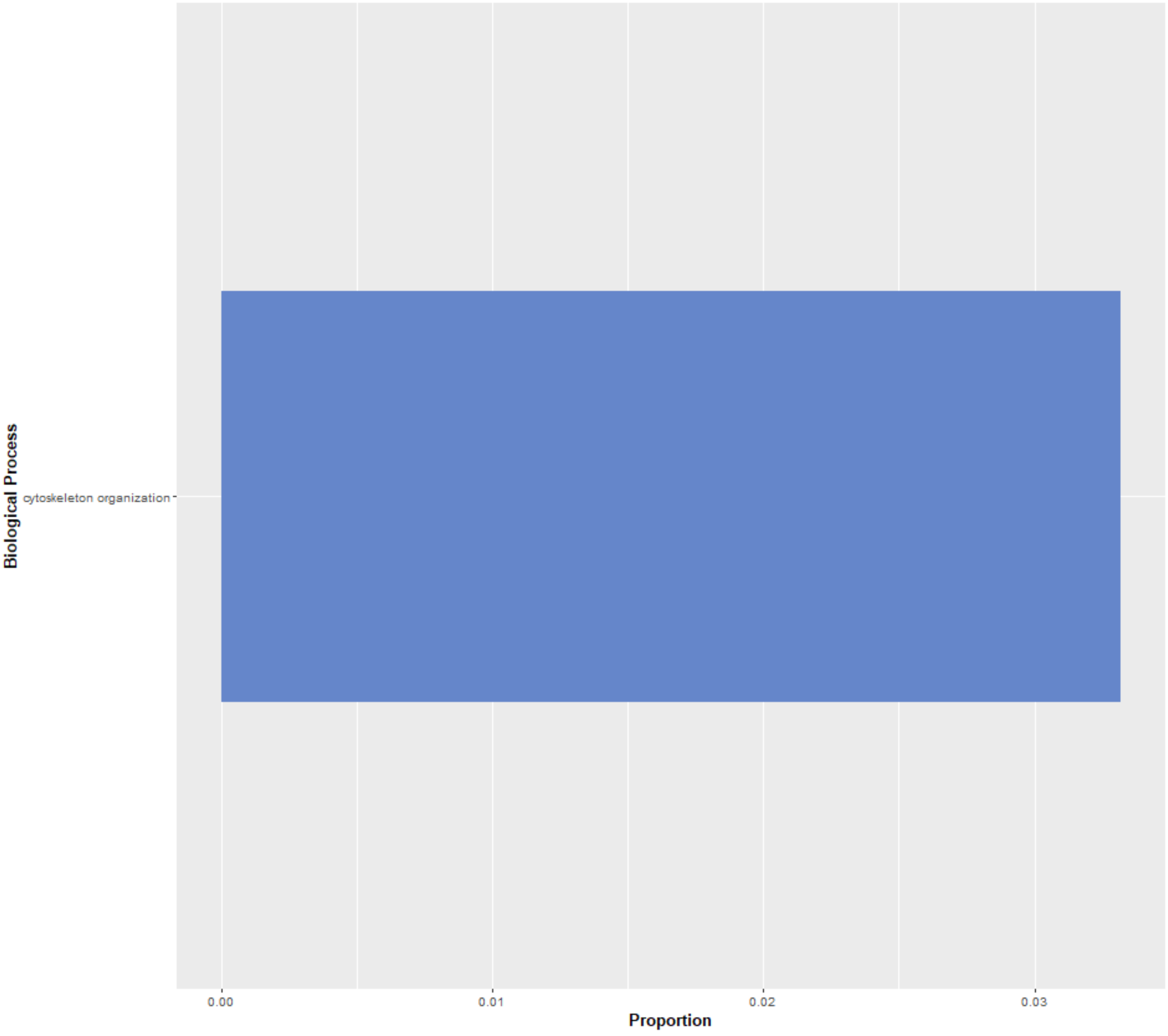
0.04

Proportion


Without Cell Cycle | With AreaShape | All Genes
Cluster 30 of 40
transcription from RNA polymerase II promoter
signaling
response to chemical
regulation of transport
regulation of protein modification process
pseudohyphal growth
protein phosphorylation
invasive growth in response to glucose limitation
cytoskeleton organization
chromatin organization
Biological Process
0.00
0.03
0.06
0.09
Proportion

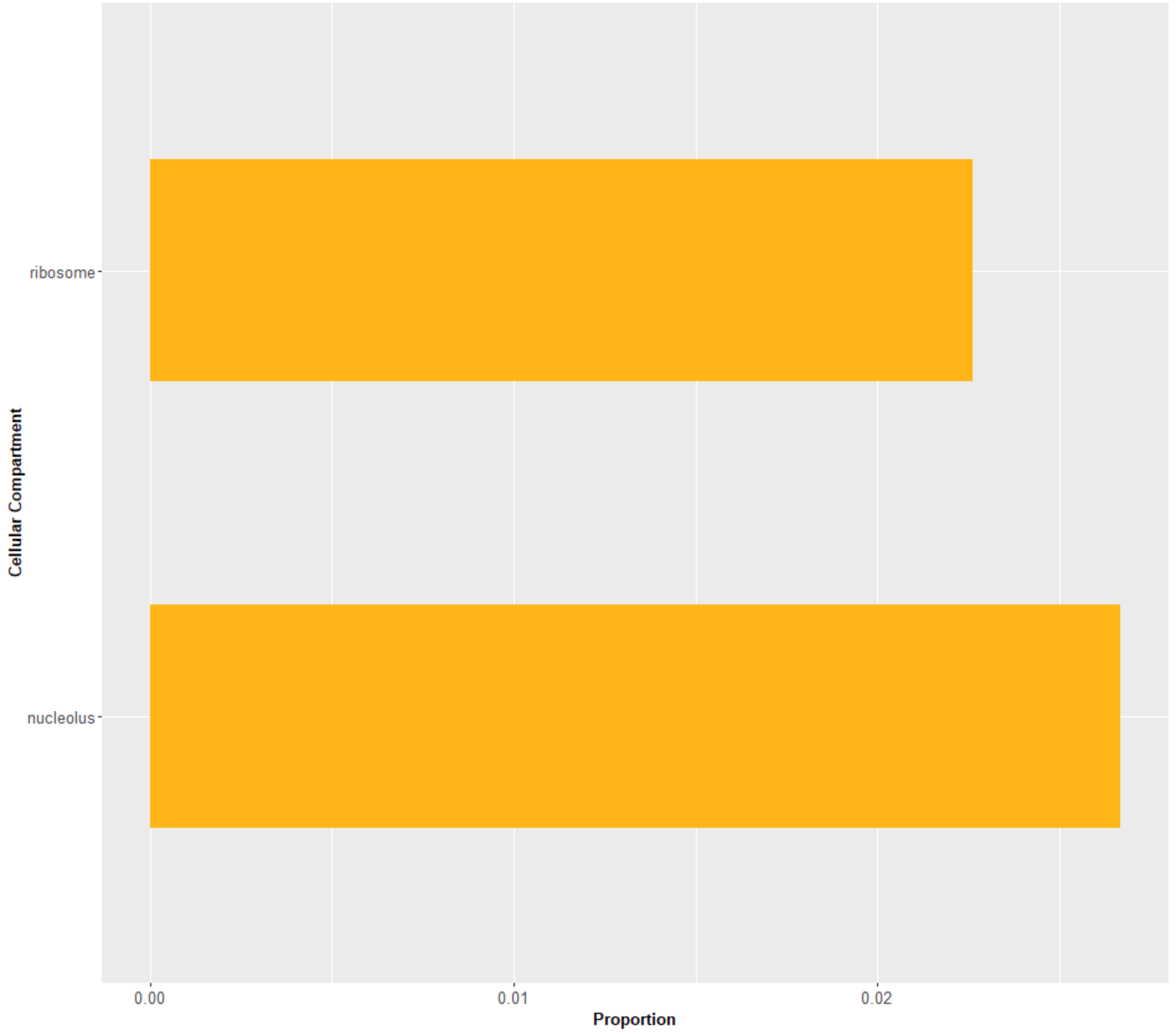
Without Cell Cycle | With AreaShape | All Genes

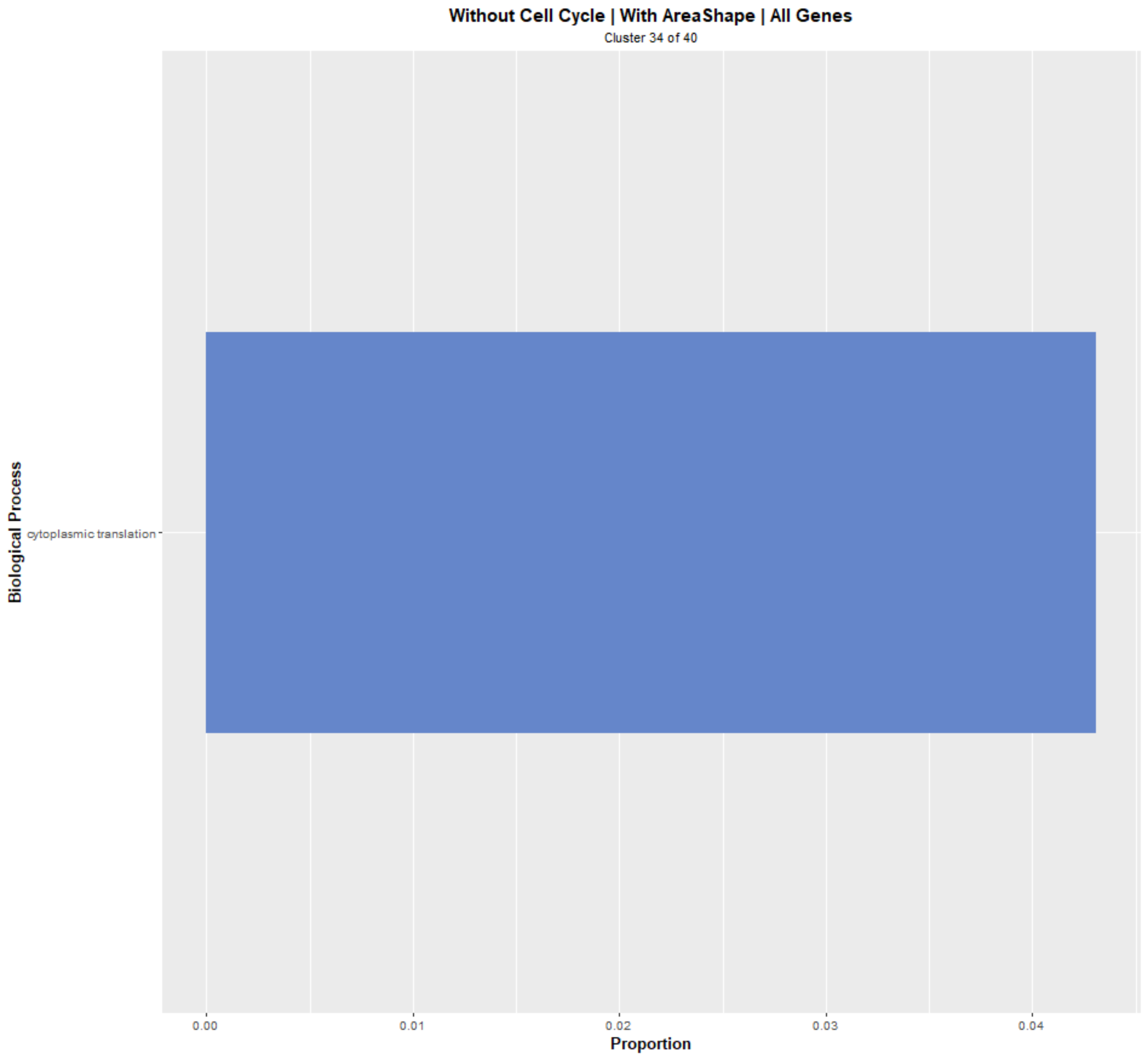
Cluster 33 of 40



Without Cell Cycle | With AreaShape | All Genes

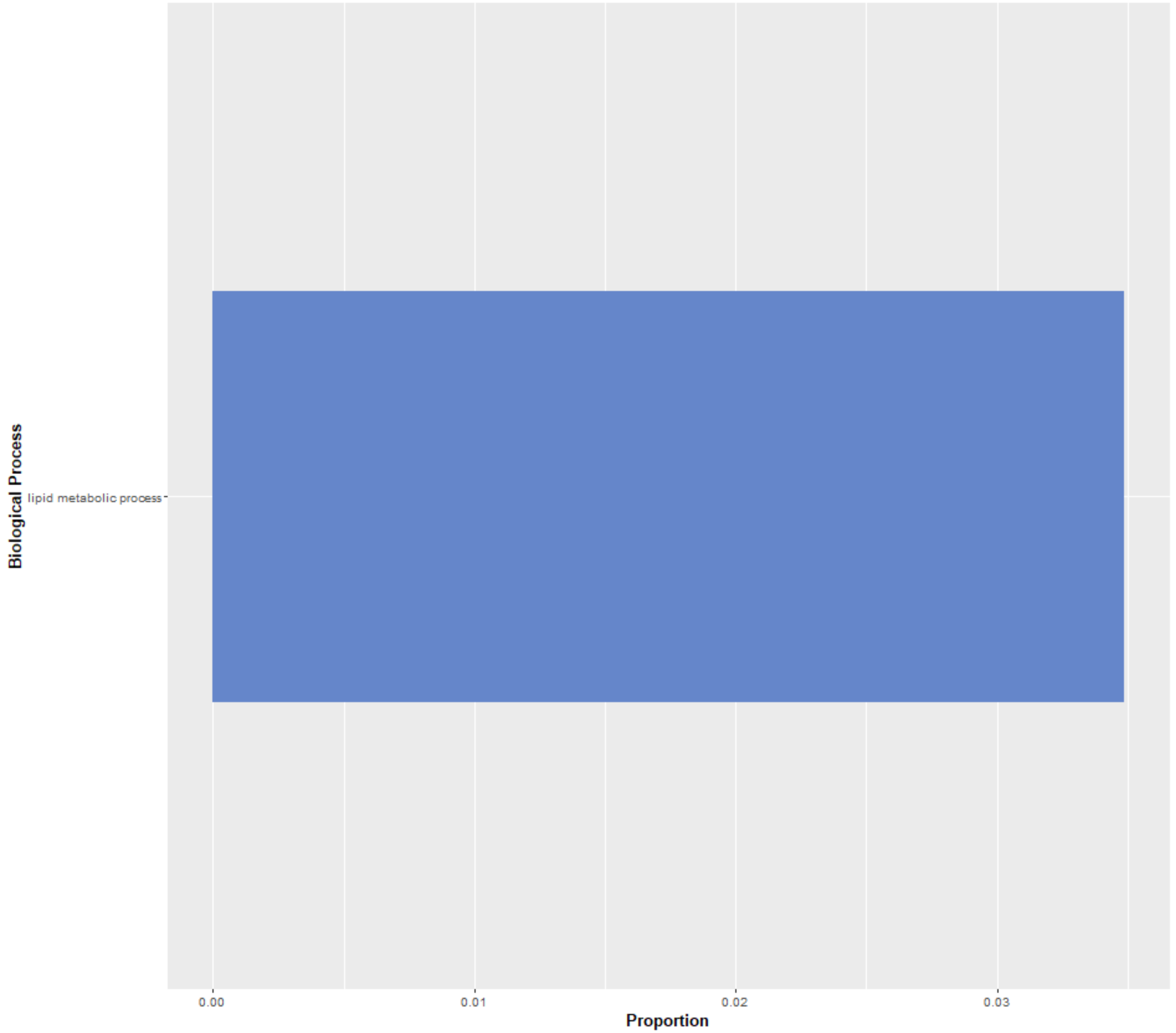
Cluster 34 of 40

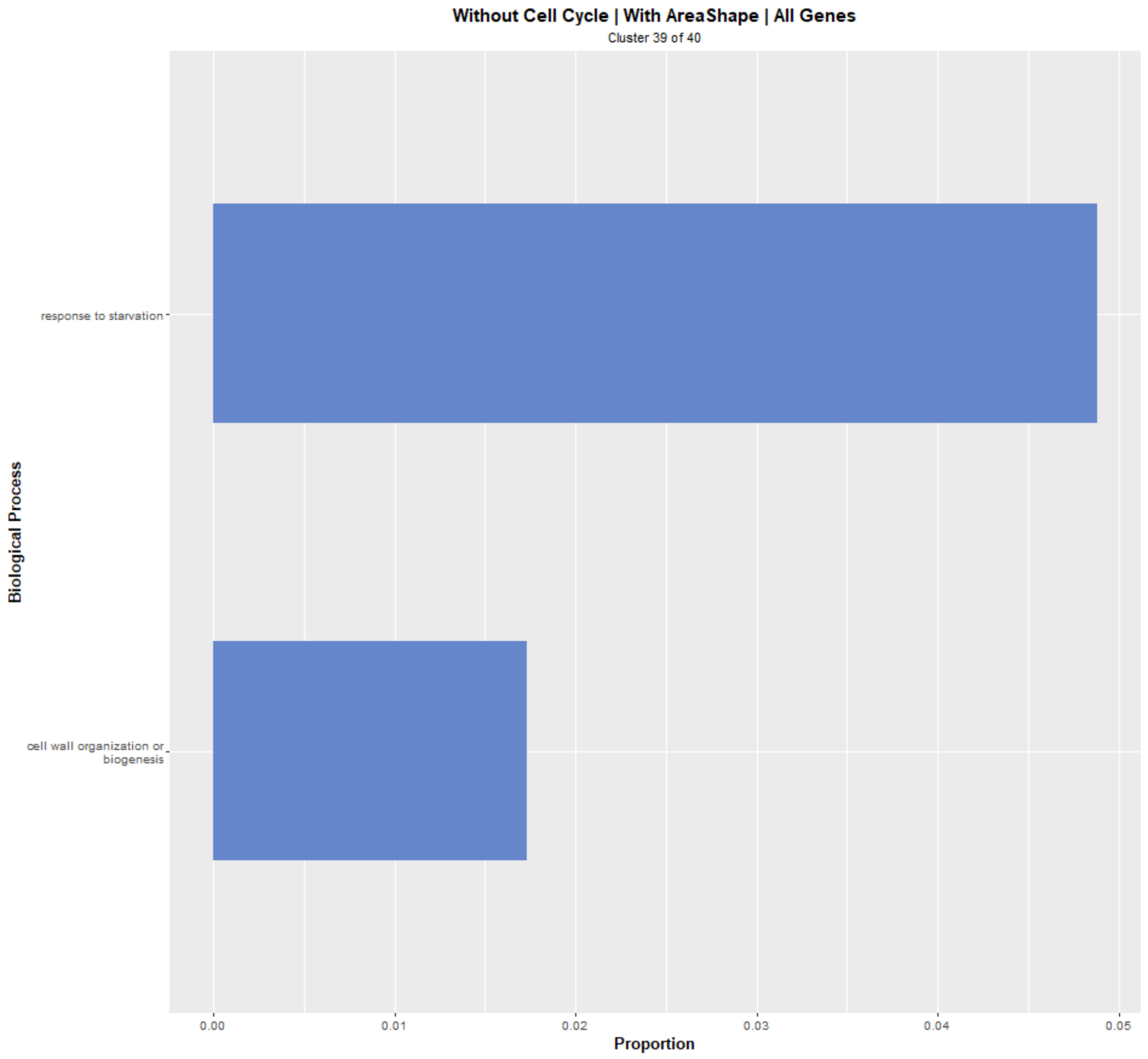



Without Cell Cycle | With AreaShape | All Genes
Cluster 34 of 40
Biological Process
cytoplasmic translation
0.00
0.01
0.02
0.03
0.04
Proportion

Without Cell Cycle | With AreaShape | All Genes

Cluster 36 of 40




Without Cell Cycle | With AreaShape | All Genes
Cluster 39 of 40
response to starvation
cell wall organization or biogenesis
Biological Process
0.00
0.01
0.02
0.03
0.04
0.05
Proportion