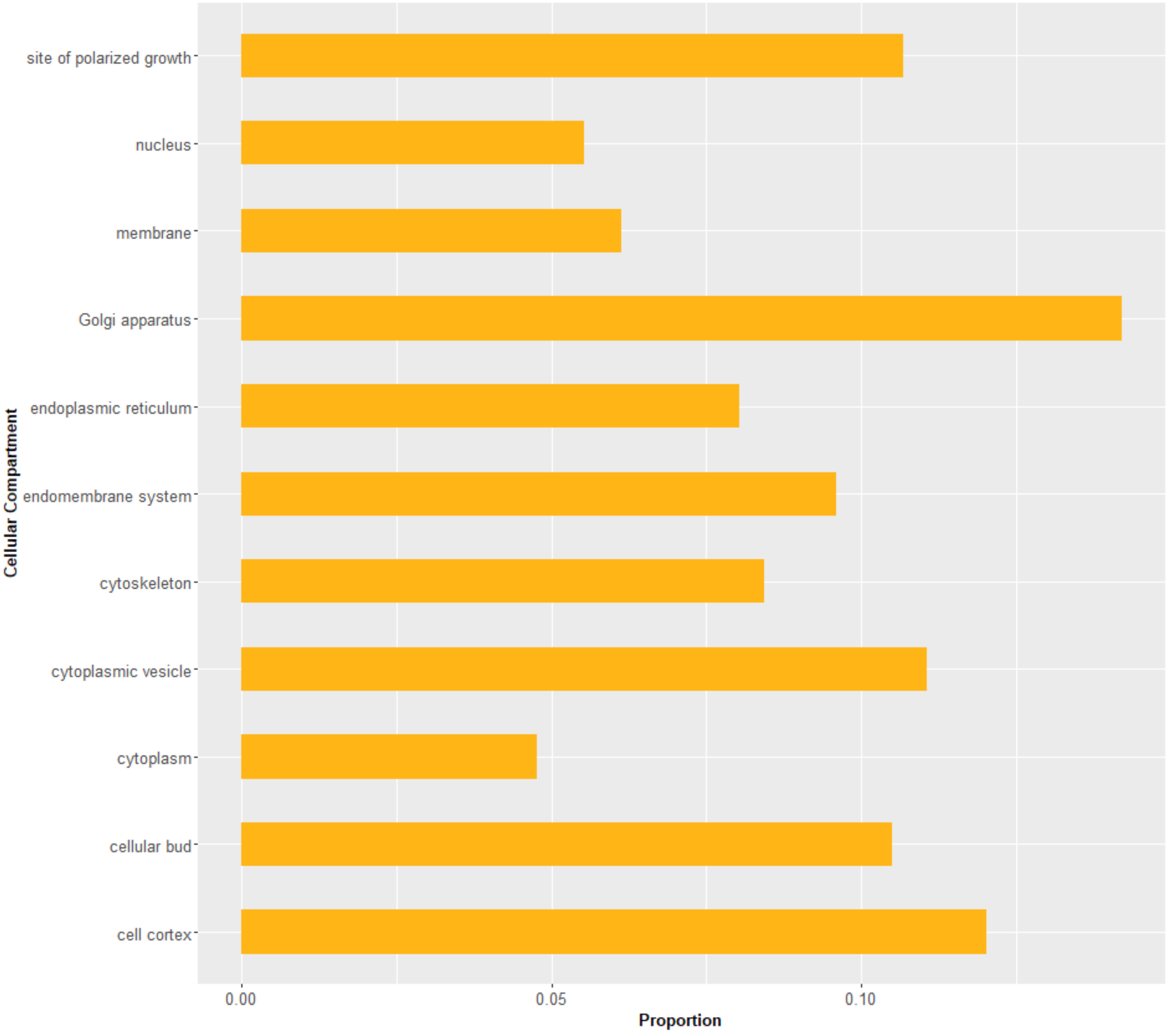
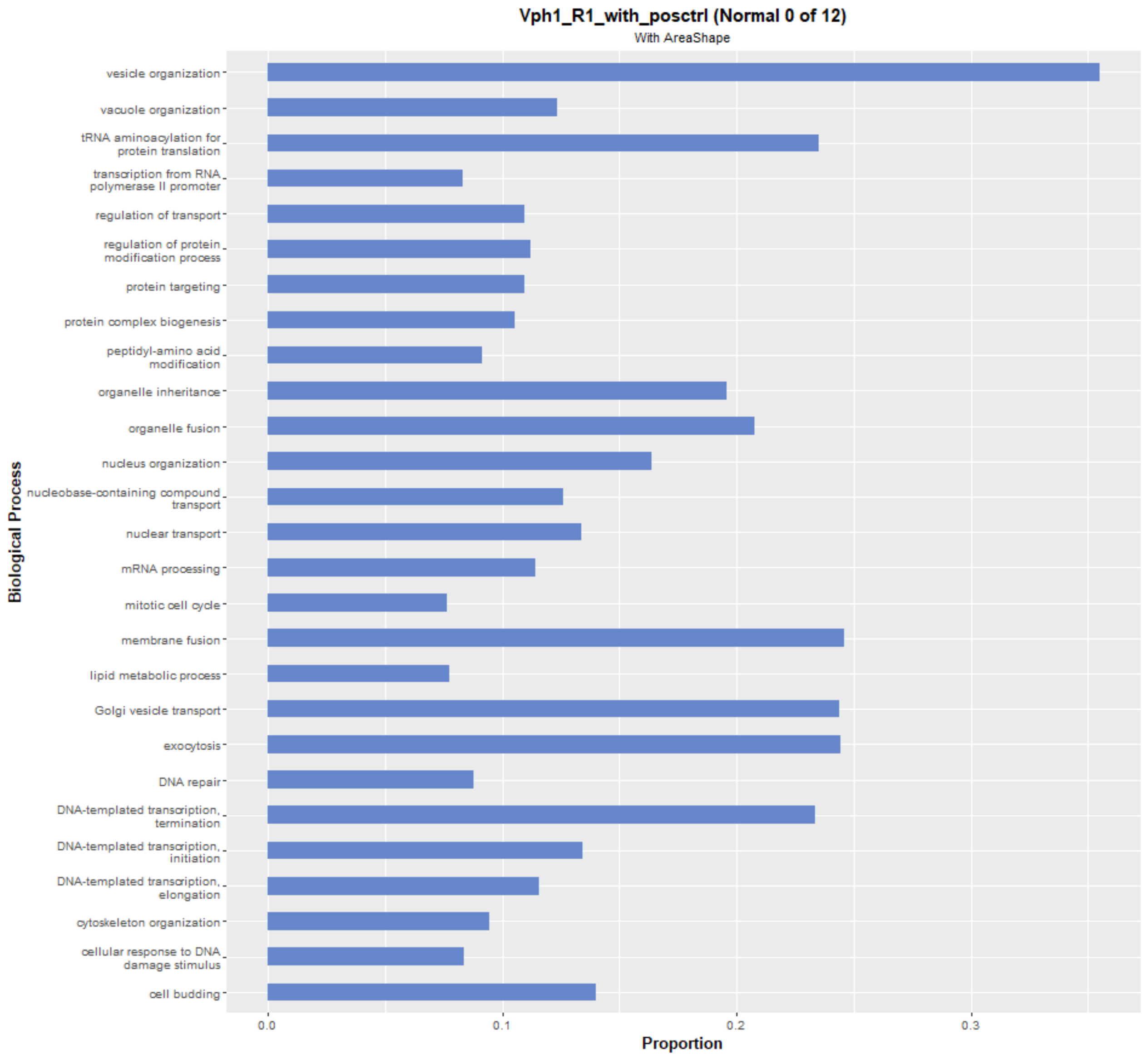
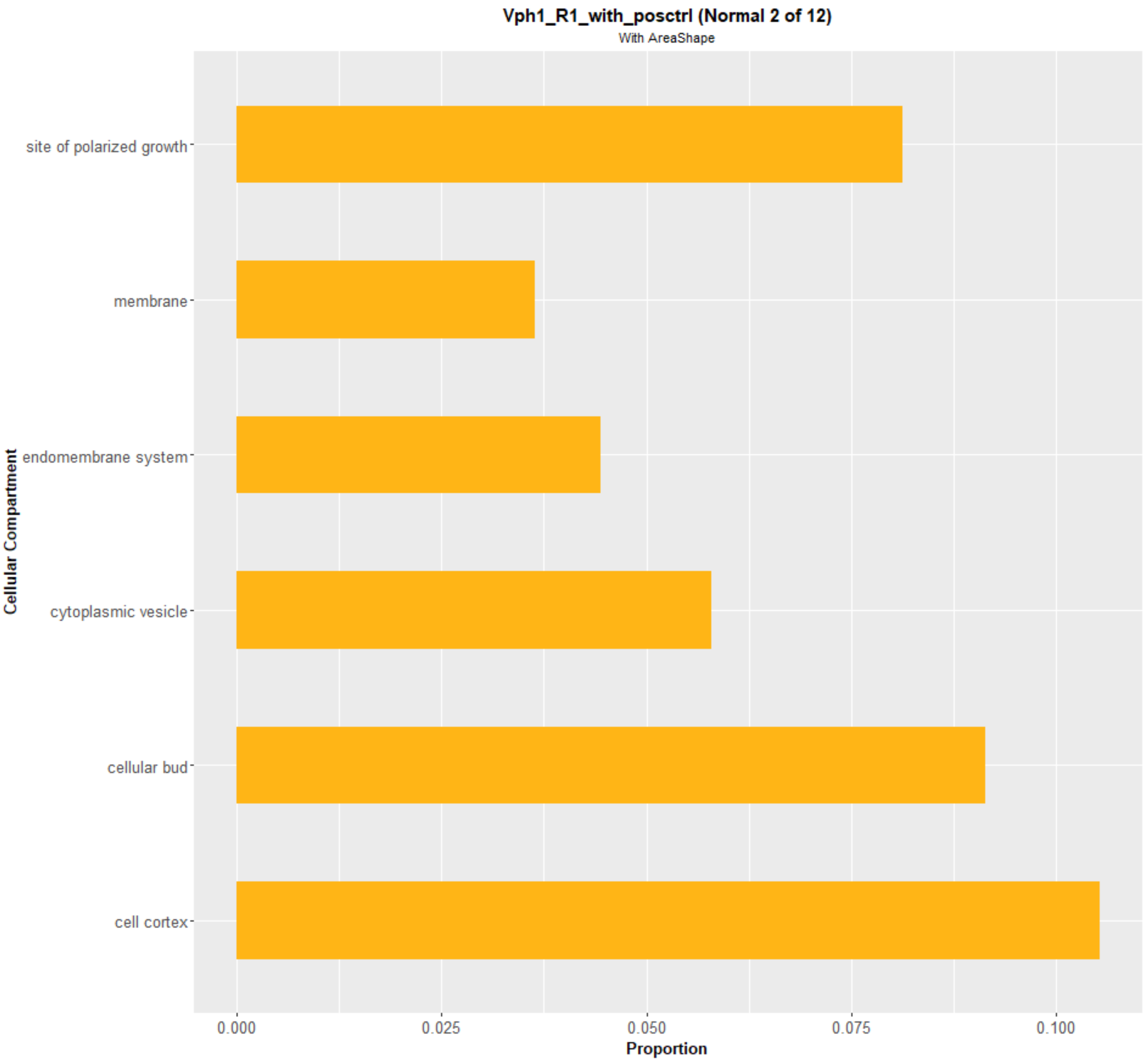
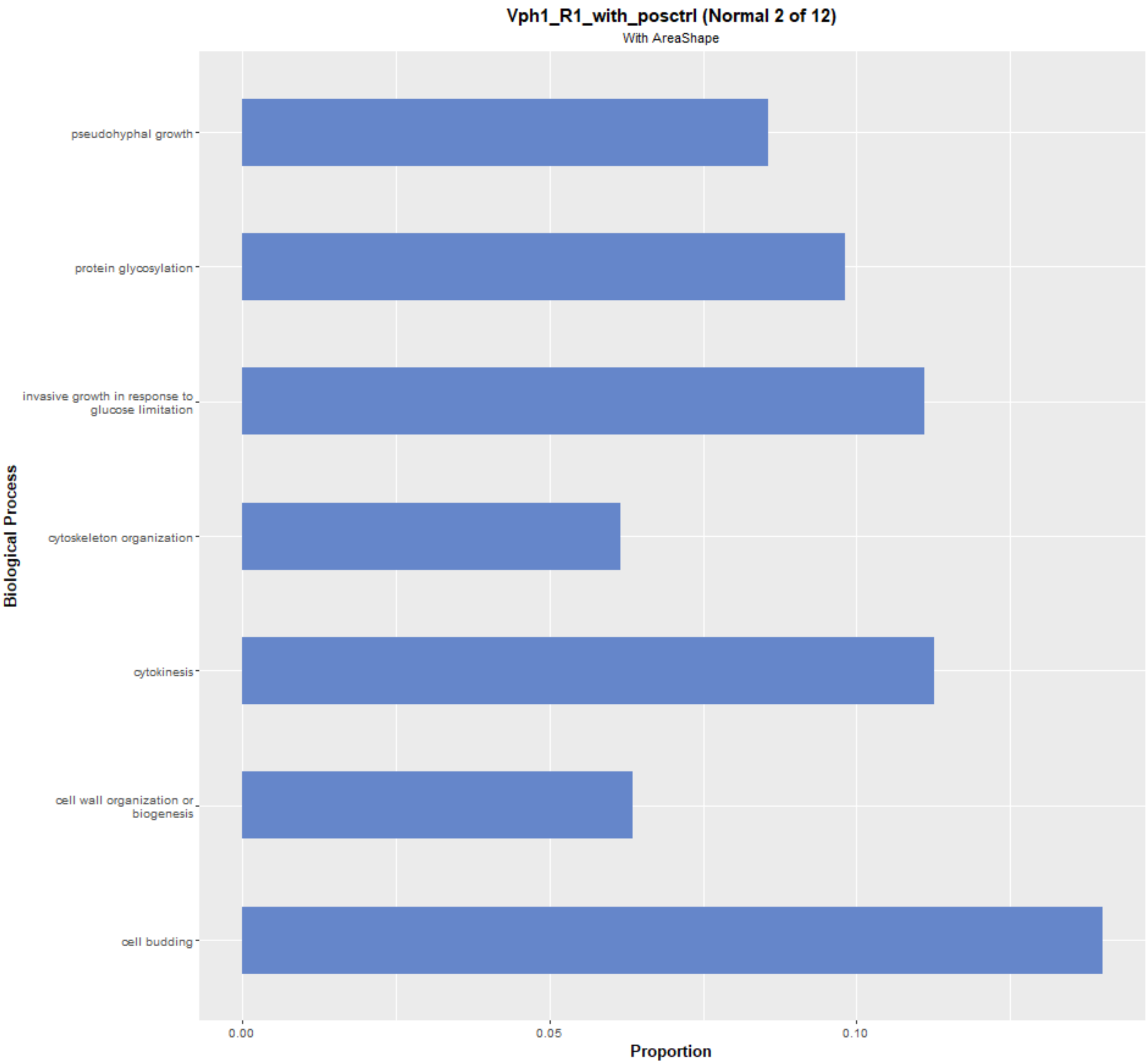
Vph1_R1_with_posctrl (Normal 0 of 12)

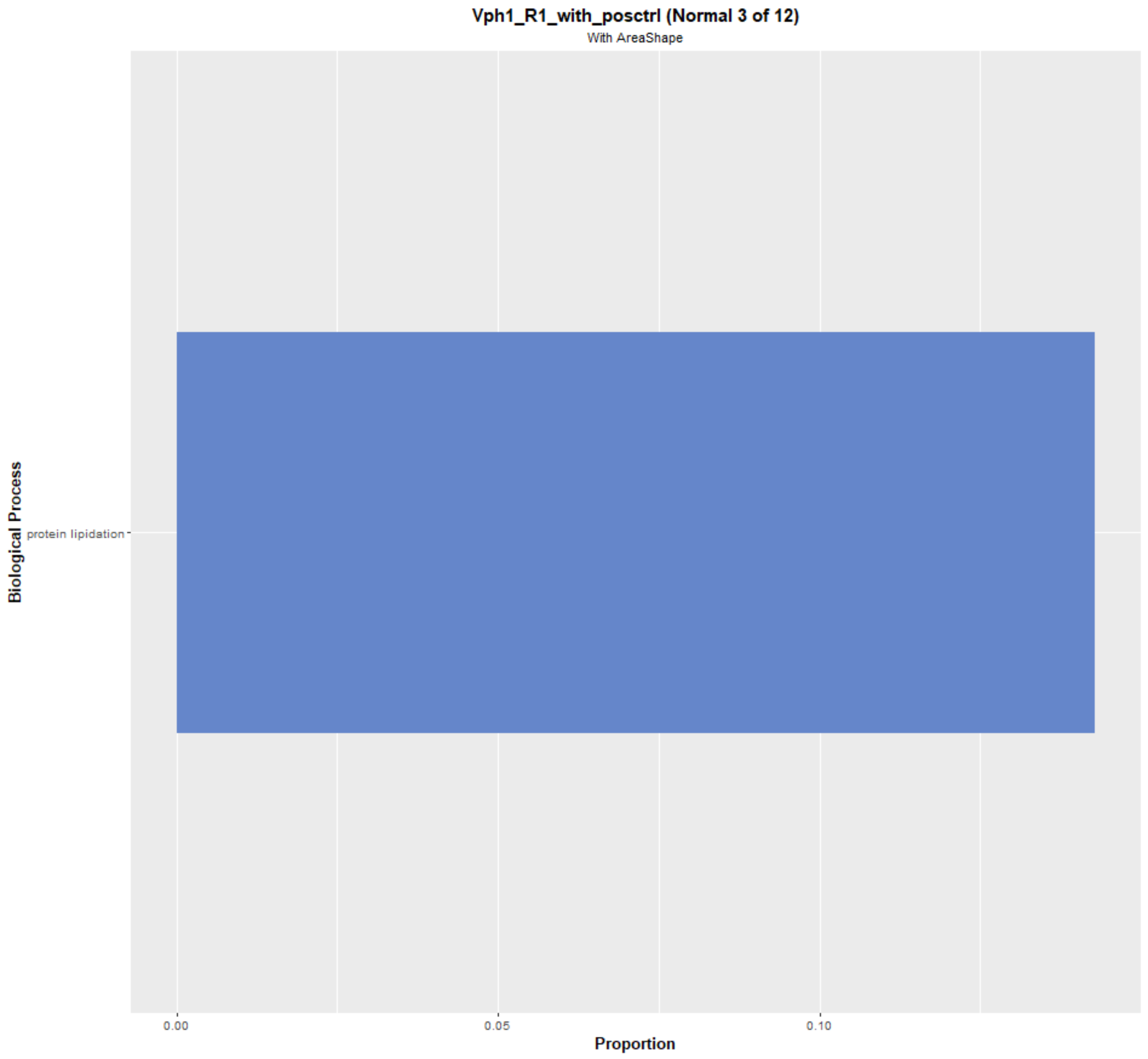
With AreaShape

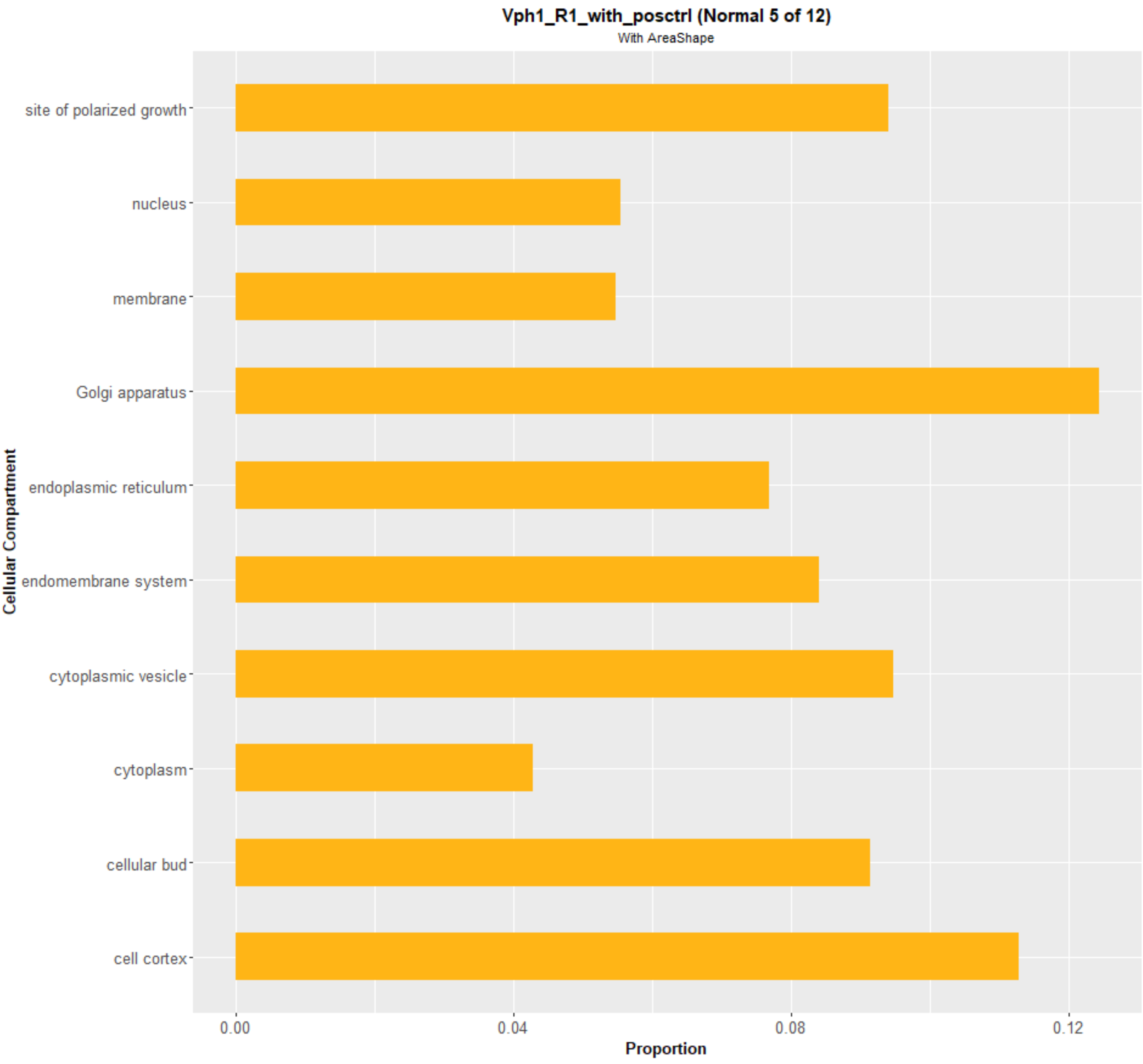


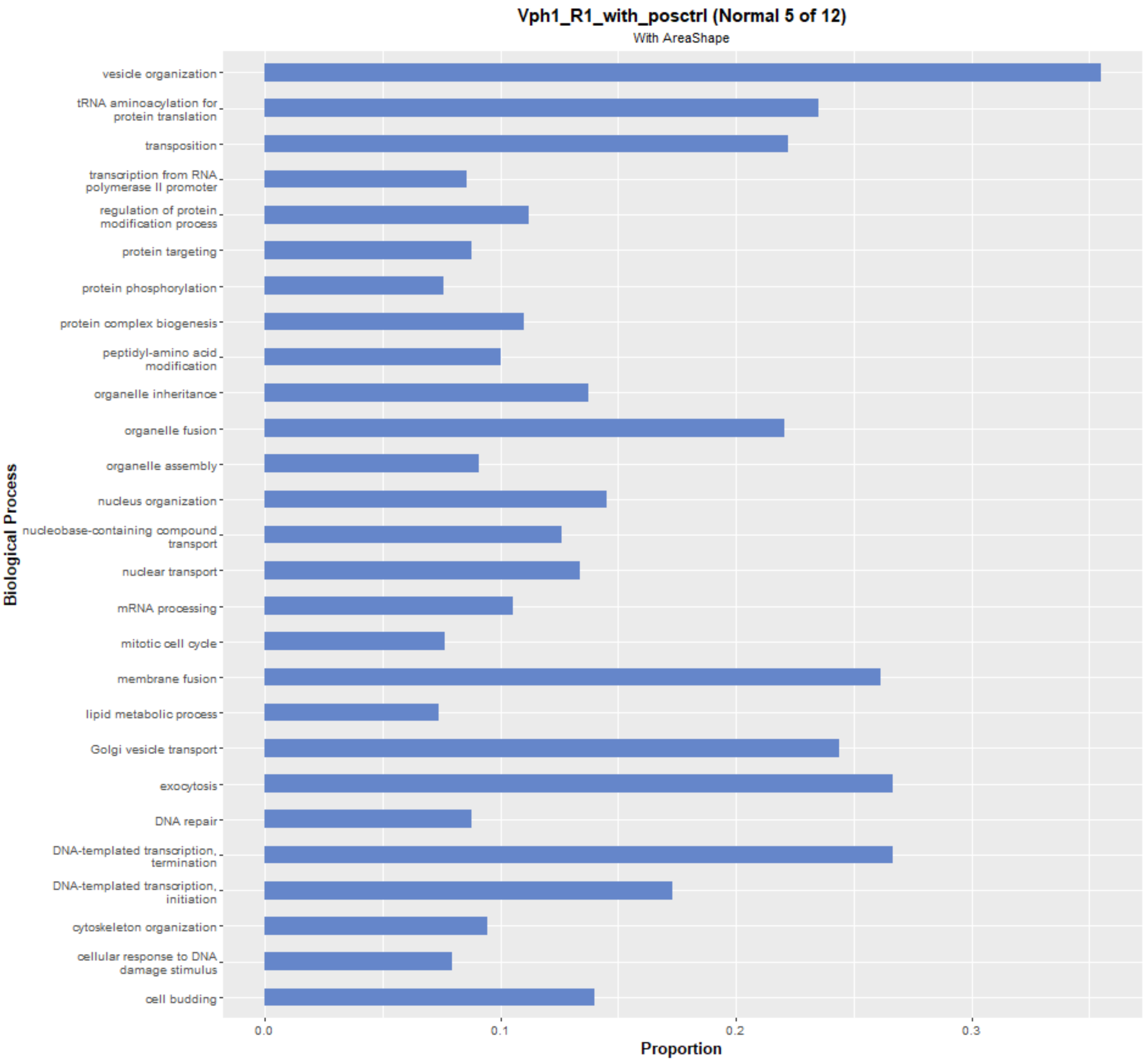

Vph1_R1_with_posctrl (Normal 0 of 12)
With AreaShape
vesicle organization
vacuole organization
tRNA aminoacylation for protein translation
transcription from RNA polymerase II promoter
regulation of transport
regulation of protein modification process
protein targeting
protein complex biogenesis
peptidyl-amino acid modification
organelle inheritance
organelle fusion
nucleus organization
nucleobase-containing compound transport
nuclear transport
mRNA processing
mitotic cell cycle
membrane fusion
lipid metabolic process
Golgi vesicle transport
exocytosis
DNA repair
DNA-templated transcription, termination
DNA-templated transcription, initiation
DNA-templated transcription, elongation
cytoskeleton organization
cellular response to DNA damage stimulus
cell budding
Biological Process
0.0
0.1
0.2
0.3
Proportion

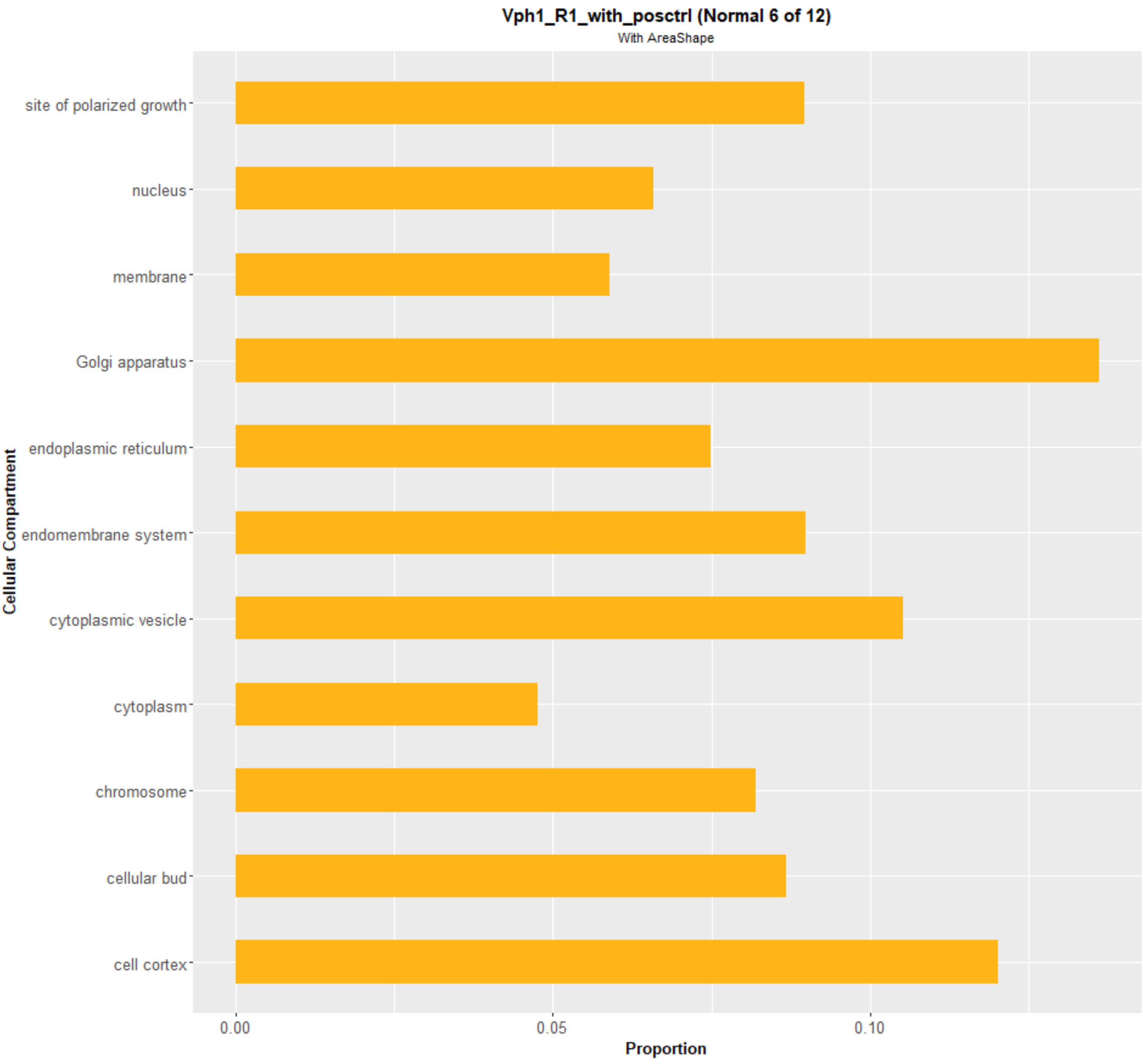

Vph1_R1_with_posctrl (Normal 2 of 12)
With AreaShape
site of polarized growth
membrane
endomembrane system
cytoplasmic vesicle
cellular bud
cell cortex
Cellular Compartment
0.000
0.025
0.050
0.075
0.100
Proportion


Vph1_R1_with_posctrl (Normal 2 of 12)
With AreaShape
pseudohyphal growth
protein glycosylation
invasive growth in response to glucose limitation
cytoskeleton organization
cytokinesis
cell wall organization or biogenesis
cell budding
Biological Process
0.00
0.05
0.10
Proportion


Vph1_R1_with_posctrl (Normal 3 of 12)
With AreaShape
Biological Process
protein lipidation
0.00
0.05
0.10
Proportion


Vph1_R1_with_posctrl (Normal 5 of 12)
With AreaShape
site of polarized growth
nucleus
membrane
Golgi apparatus
endoplasmic reticulum
endomembrane system
cytoplasmic vesicle
cytoplasm
cellular bud
cell cortex
Cellular Compartment
0.00
0.04
0.08
0.12
Proportion


Vph1_R1_with_posctrl (Normal 5 of 12)
With AreaShape
vesicle organization
tRNA aminoacylation for protein translation
transposition
transcription from RNA polymerase II promoter
regulation of protein modification process
protein targeting
protein phosphorylation
protein complex biogenesis
peptidyl-amino acid modification
organelle inheritance
organelle fusion
organelle assembly
nucleus organization
nucleobase-containing compound transport
nuclear transport
mRNA processing
mitotic cell cycle
membrane fusion
lipid metabolic process
Golgi vesicle transport
exocytosis
DNA repair
DNA-templated transcription, termination
DNA-templated transcription, initiation
cytoskeleton organization
cellular response to DNA damage stimulus
cell budding
Biological Process
0.0
0.1
0.2
0.3
Proportion


Vph1_R1_with_posctrl (Normal 6 of 12)
With AreaShape
site of polarized growth
nucleus
membrane
Golgi apparatus
endoplasmic reticulum
endomembrane system
cytoplasmic vesicle
cytoplasm
chromosome
cellular bud
cell cortex
Cellular Compartment
0.00
0.05
0.10
Proportion

Vph1_R1_with_posctrl (Normal 6 of 12)

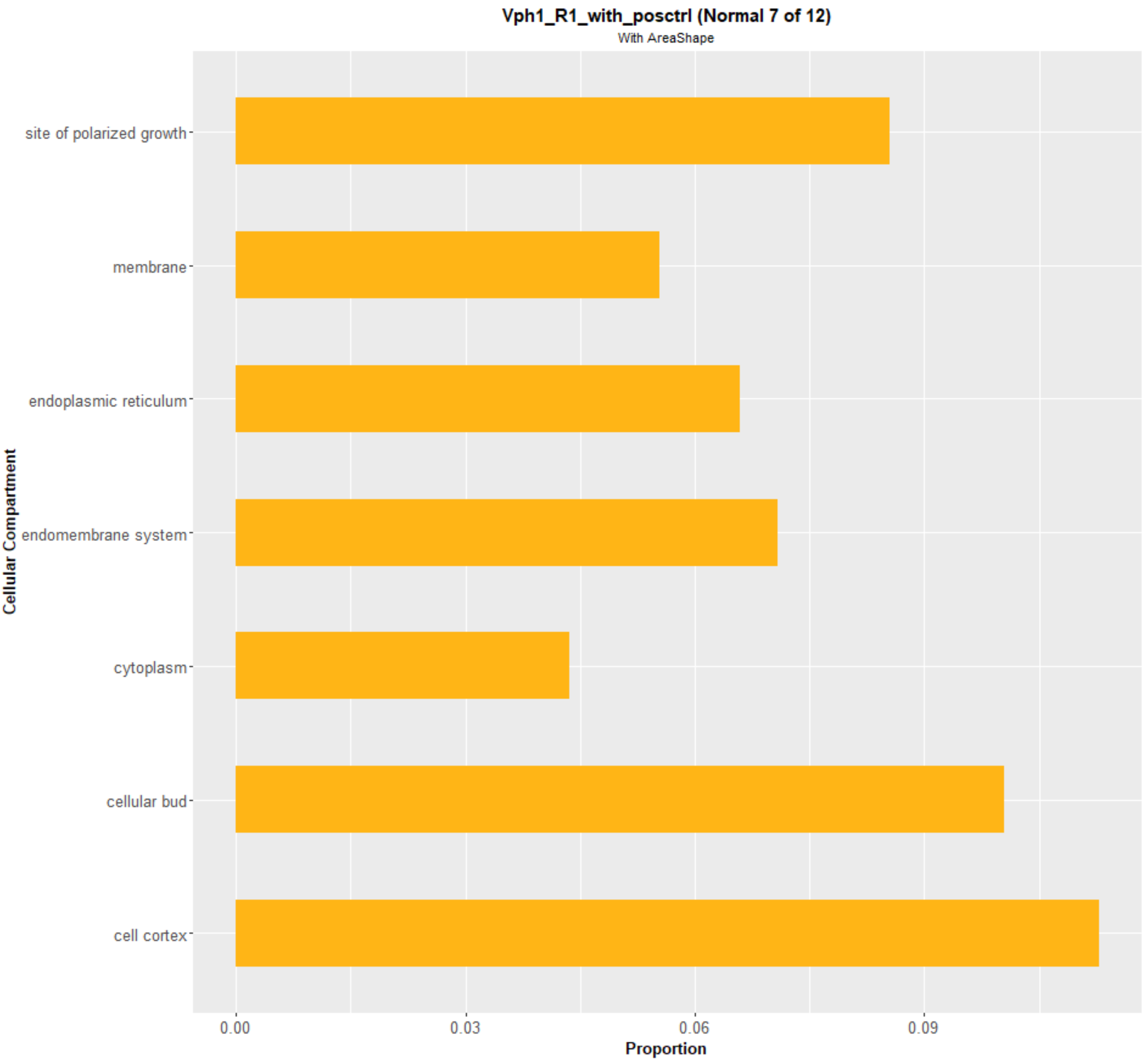
With AreaShape

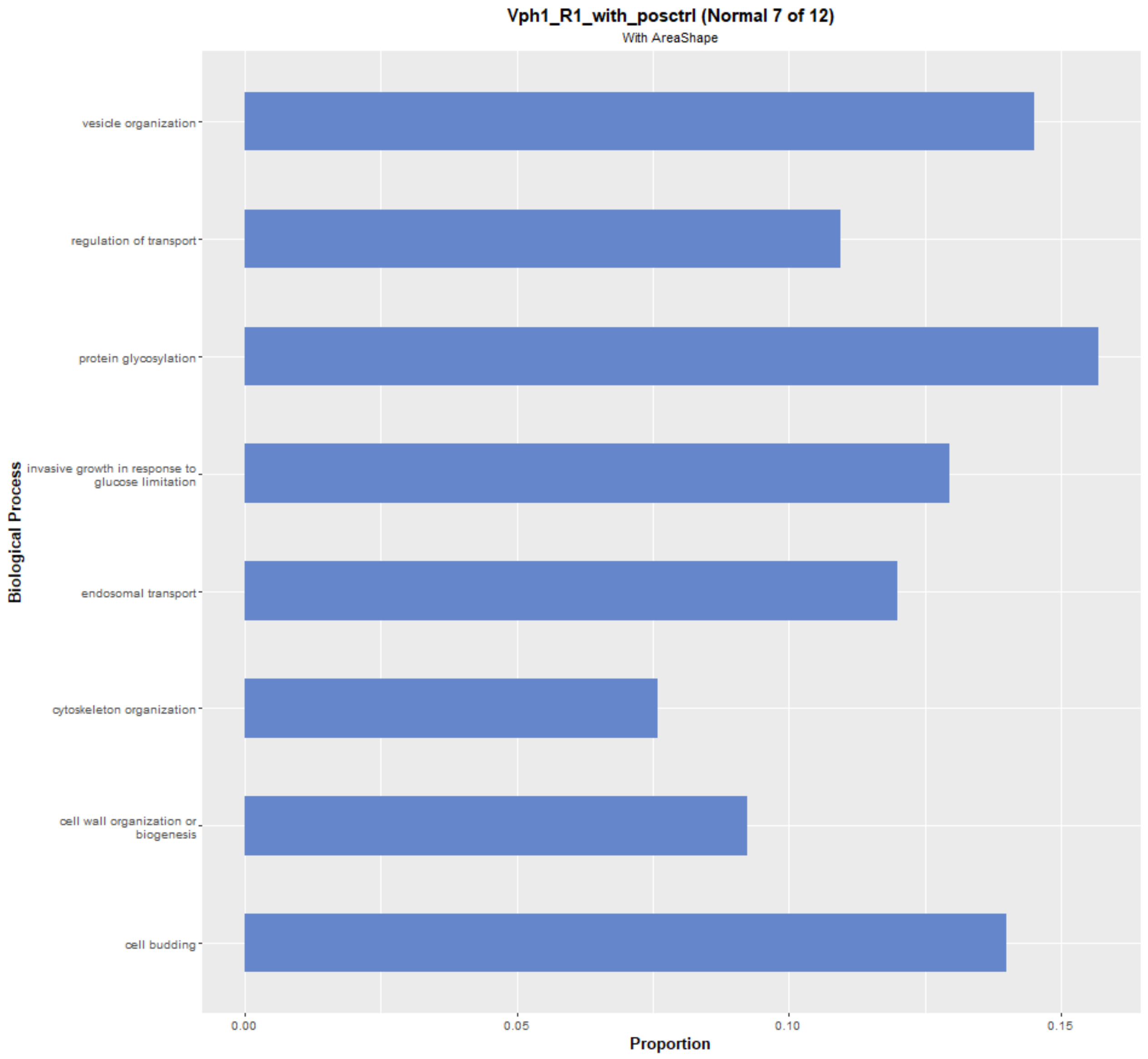
vesicle organization
tRNA aminoacylation for protein translation
transposition
translational initiation
transcription from RNA polymerase II promoter
regulation of protein modification process
regulation of organelle organization
protein targeting
protein complex biogenesis
peptidyl-amino acid modification
organelle inheritance
organelle fusion
nucleus organization
nucleobase-containing compound transport
nuclear transport
mRNA processing
mitotic cell cycle
membrane fusion
lipid metabolic process
Golgi vesicle transport
exocytosis
DNA repair
DNA-templated transcription, termination
DNA-templated transcription, initiation
cytoskeleton organization
cellular response to DNA damage stimulus

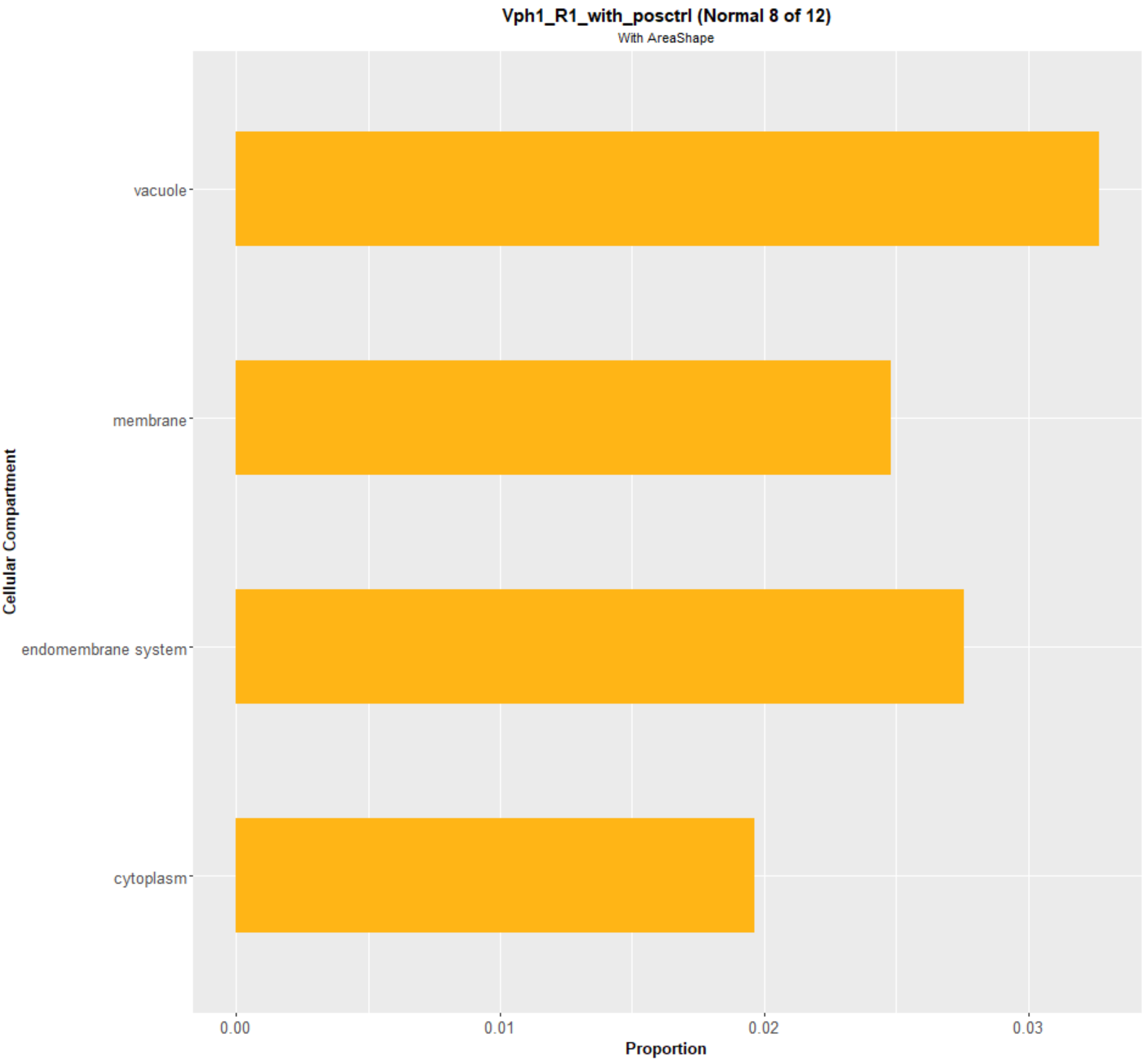
Biological Process

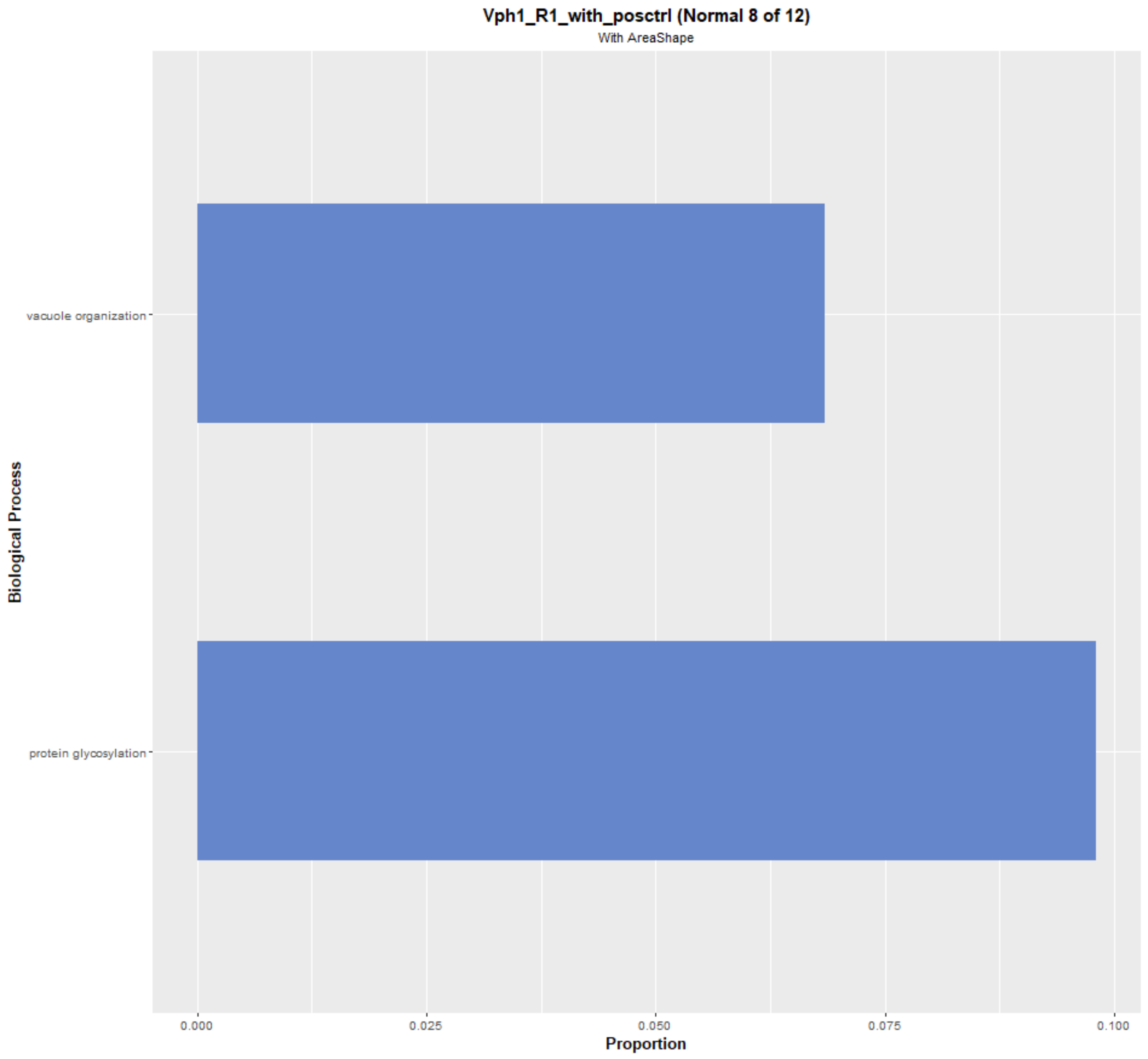
0.0
0.1
0.2
0.3

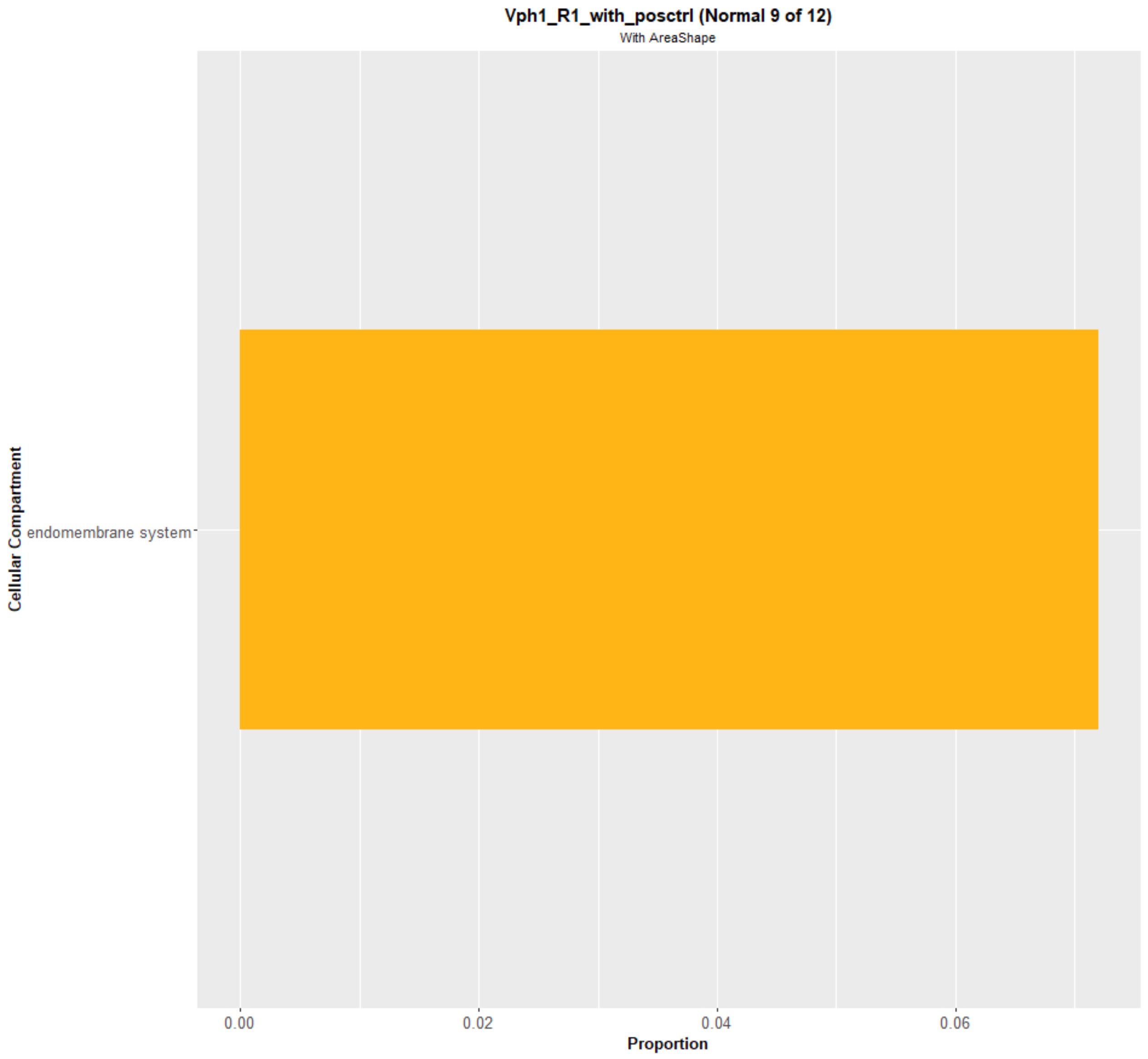
Proportion


Vph1_R1_with_posctrl (Normal 7 of 12)
With AreaShape
site of polarized growth
membrane
endoplasmic reticulum
endomembrane system
cytoplasm
cellular bud
cell cortex
Cellular Compartment
0.00
0.03
0.06
0.09
Proportion


Vph1_R1_with_posctrl (Normal 7 of 12)
With AreaShape
vesicle organization
regulation of transport
protein glycosylation
invasive growth in response to glucose limitation
endosomal transport
cytoskeleton organization
cell wall organization or biogenesis
cell budding
Biological Process
0.00
0.05
0.10
0.15
Proportion


Vph1_R1_with_posctrl (Normal 8 of 12)
With AreaShape
vacuole
membrane
endomembrane system
cytoplasm
Cellular Compartment
0.00
0.01
0.02
0.03
Proportion


Vph1_R1_with_posctrl (Normal 8 of 12)
With AreaShape
vacuole organization
protein glycosylation
Biological Process
0.000
0.025
0.050
0.075
0.100
Proportion


Vph1_R1_with_posctrl (Normal 9 of 12)
With AreaShape
Cellular Compartment
endomembrane system
0.00
0.02
0.04
0.06
Proportion

Vph1_R1_with_posctrl (Normal 9 of 12)

With AreaShape

Biological Process

protein lipidation

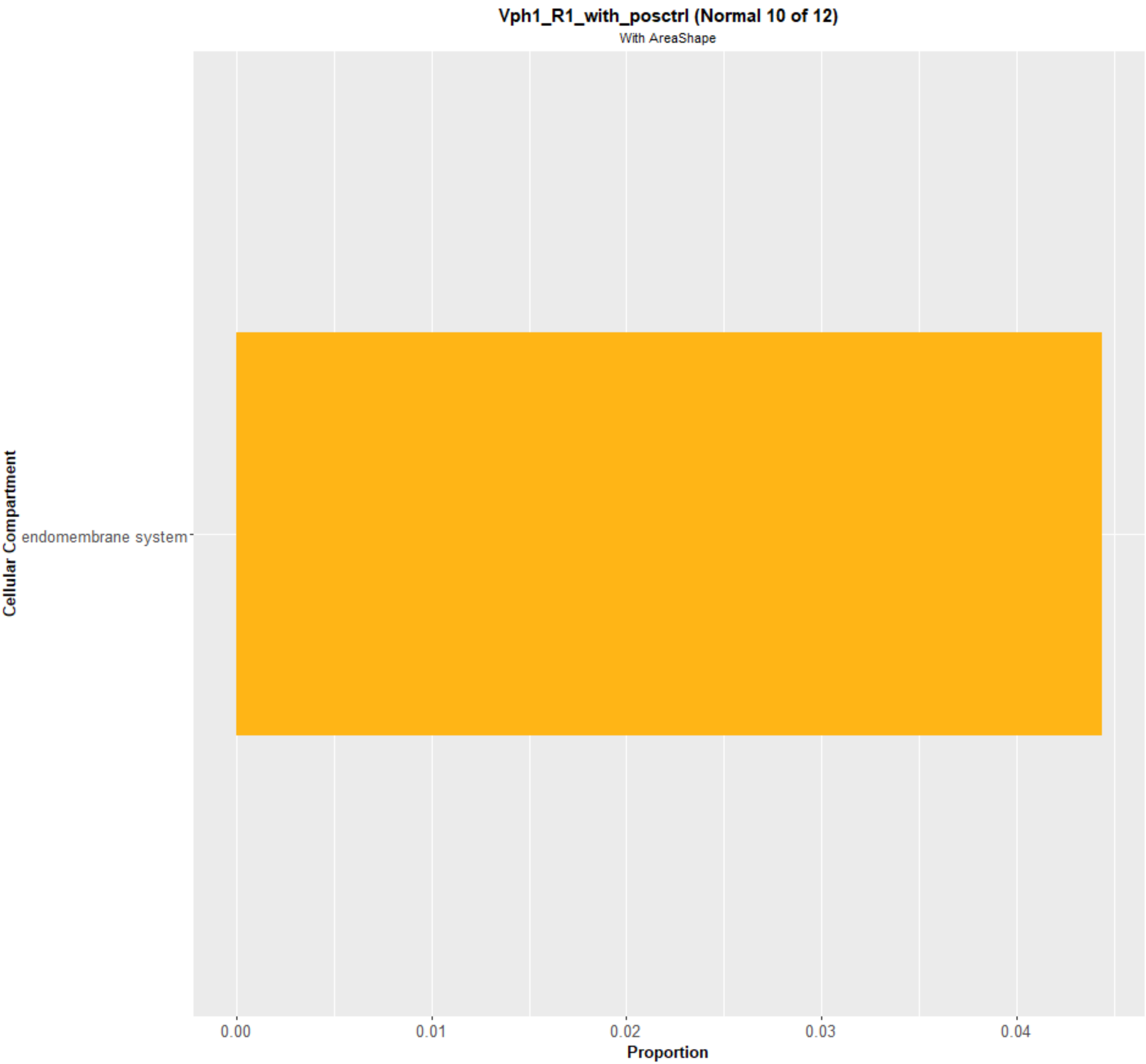
0.00

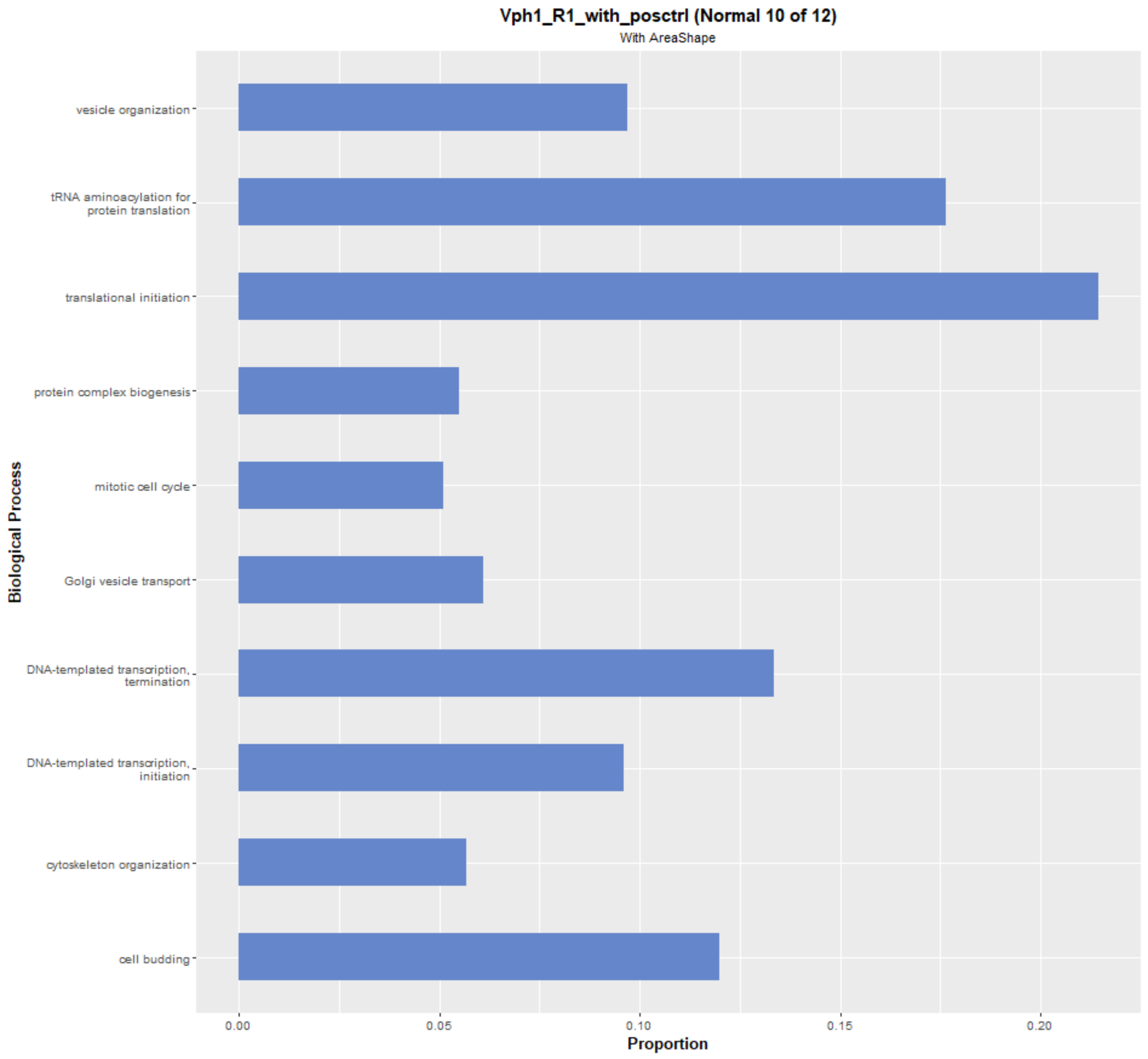
0.05

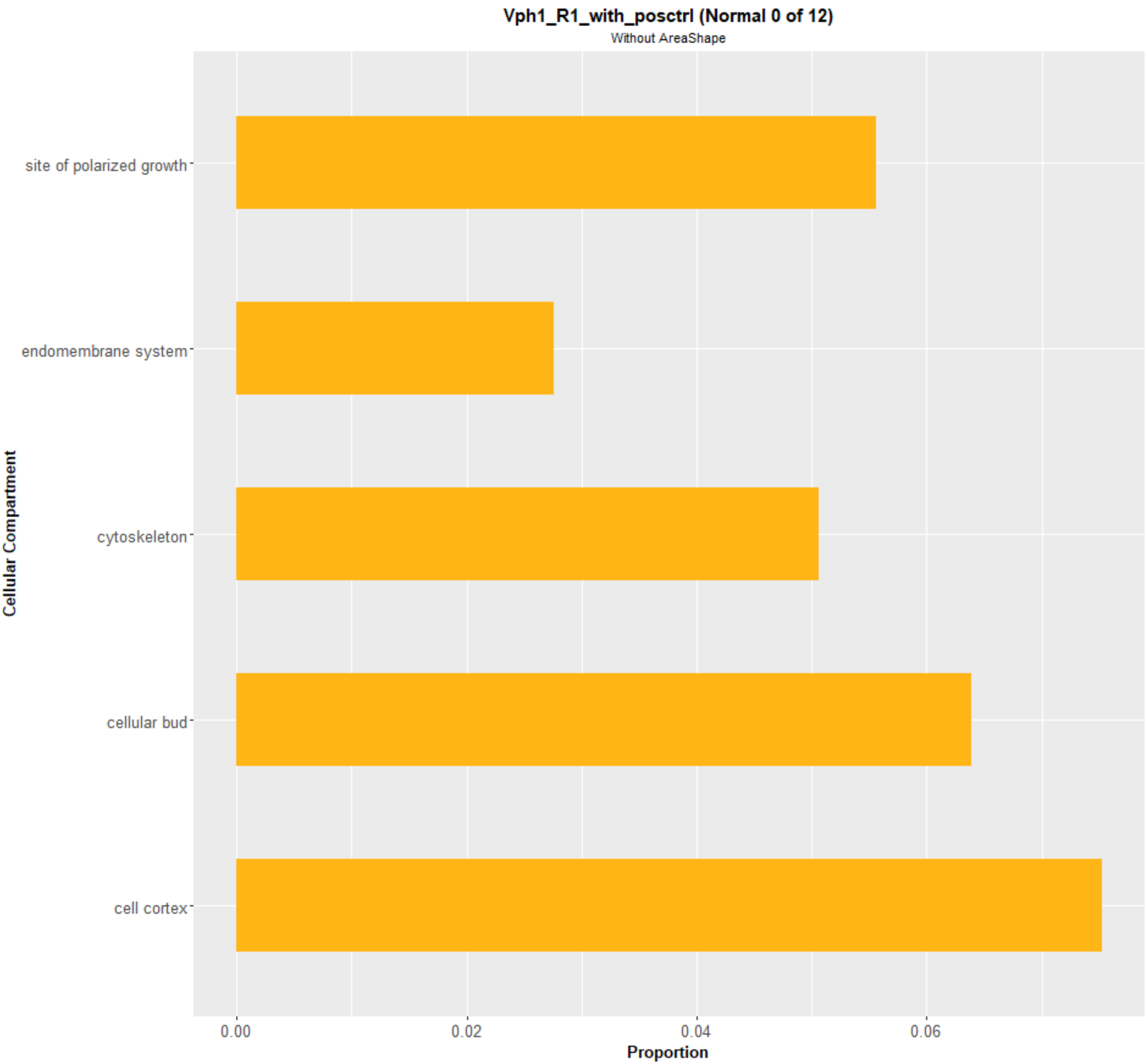
0.10

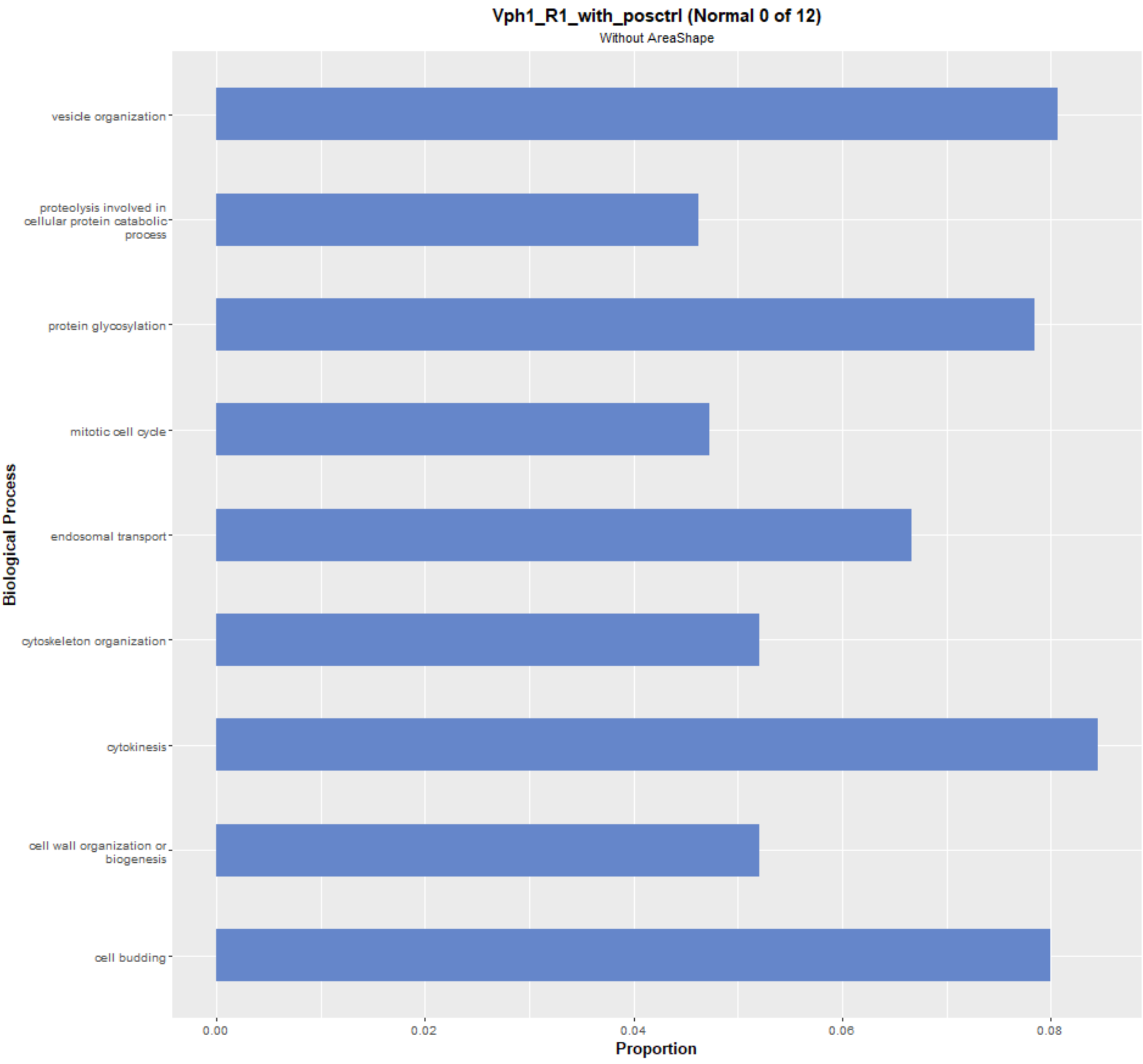
0.15

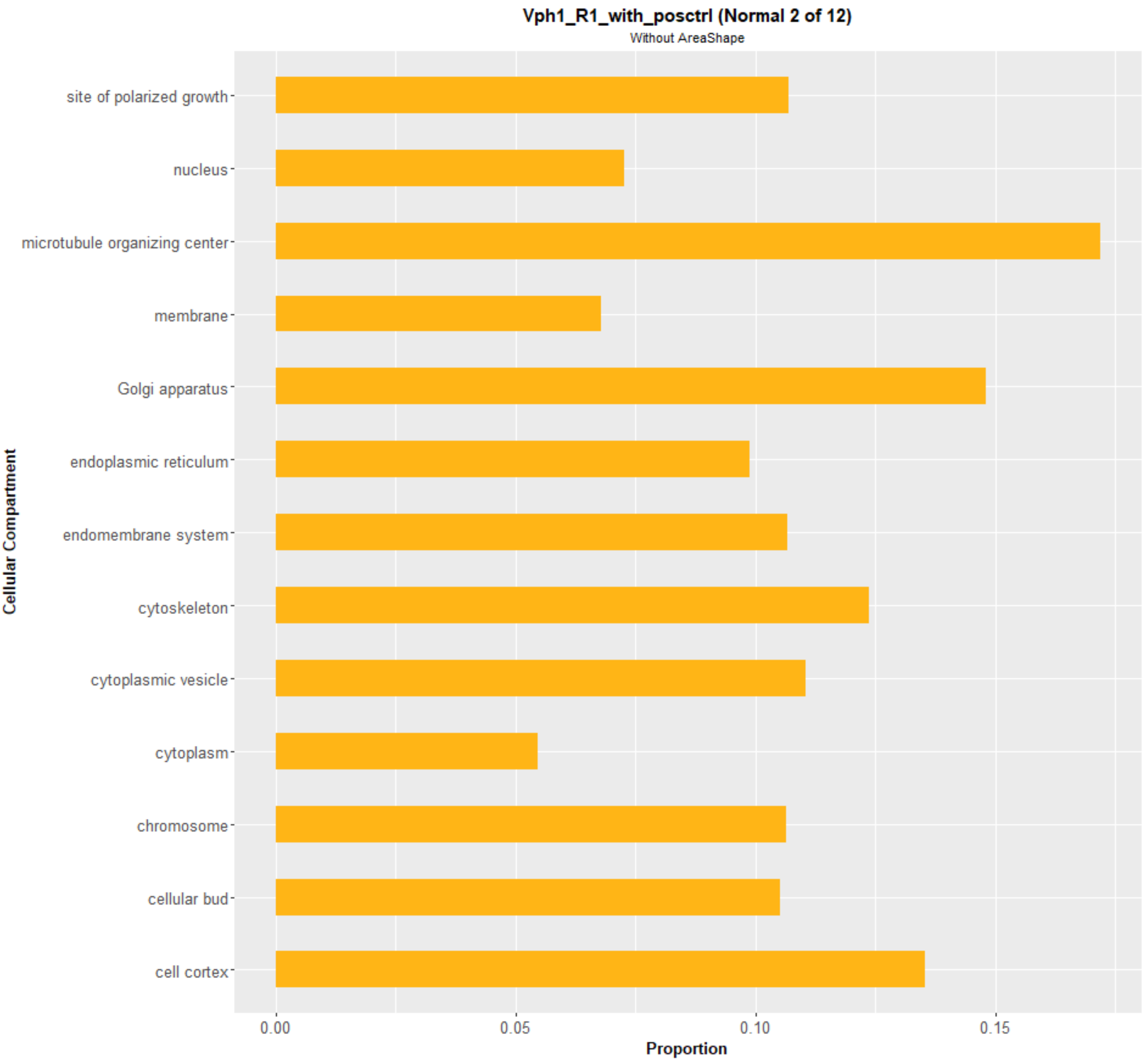
Proportion

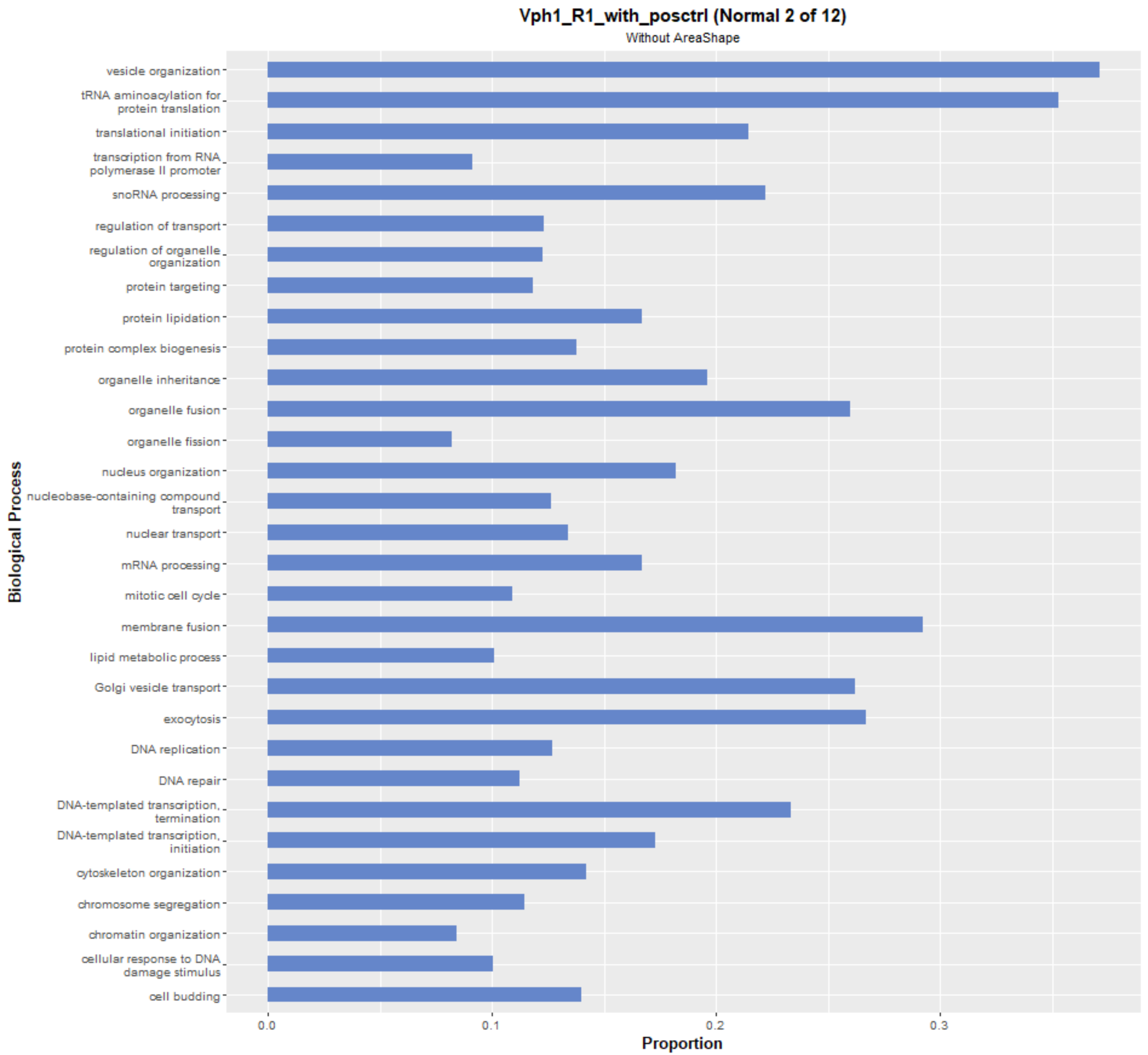

Vph1_R1_with_posctrl (Normal 10 of 12)
With AreaShape
Cellular Compartment
endomembrane system
0.00
0.01
0.02
0.03
0.04
Proportion

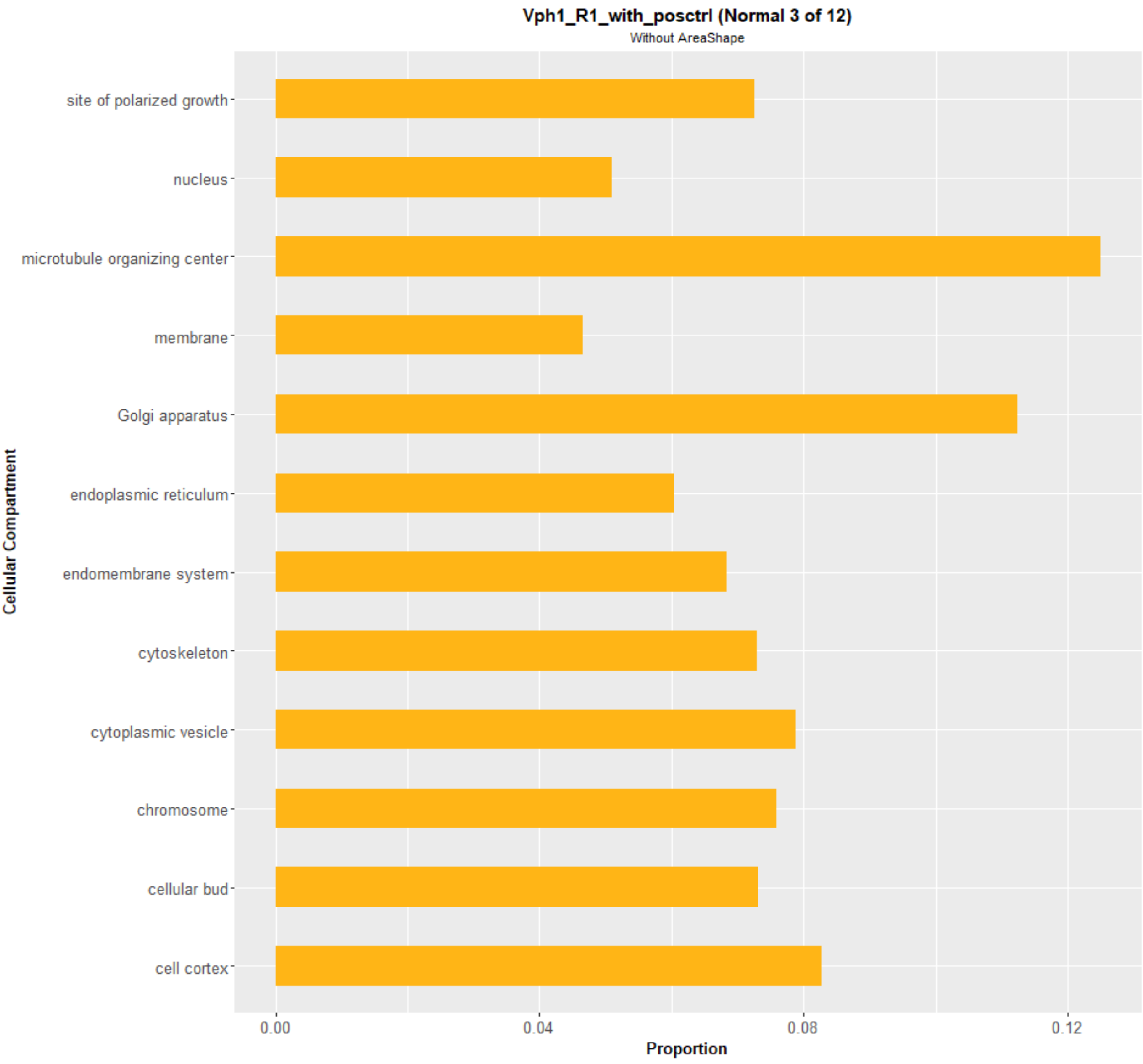

Vph1_R1_with_posctrl (Normal 10 of 12)
With AreaShape
vesicle organization
tRNA aminoacylation for protein translation
translational initiation
protein complex biogenesis
mitotic cell cycle
Golgi vesicle transport
DNA-templated transcription, termination
DNA-templated transcription, initiation
cytoskeleton organization
cell budding
Biological Process
0.00
0.05
0.10
0.15
0.20
Proportion

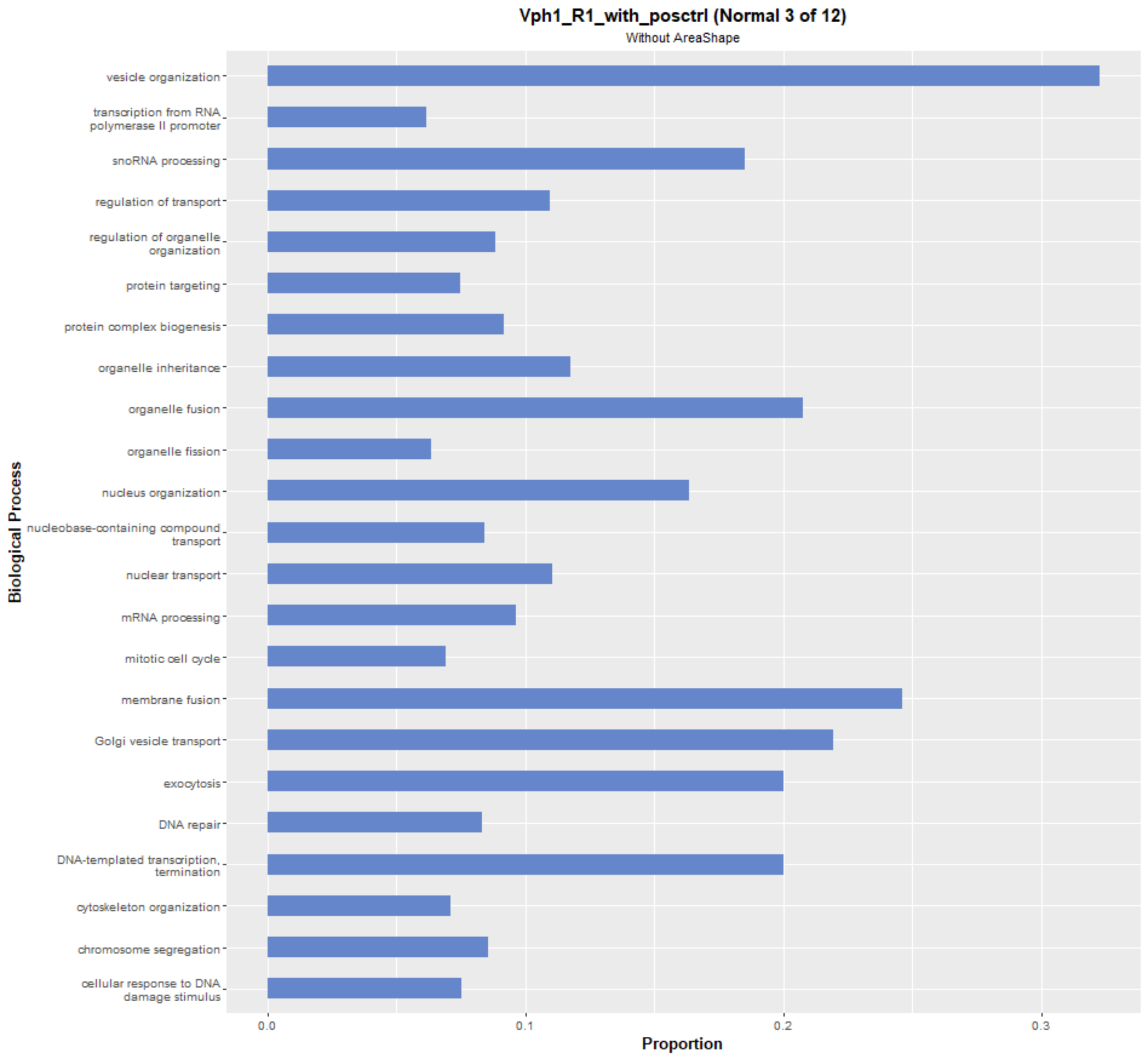

Vph1_R1_with_posctrl (Normal 0 of 12)
Without AreaShape
Cellular Compartment
site of polarized growth
endomembrane system
cytoskeleton
cellular bud
cell cortex
0.00
0.02
0.04
0.06
Proportion

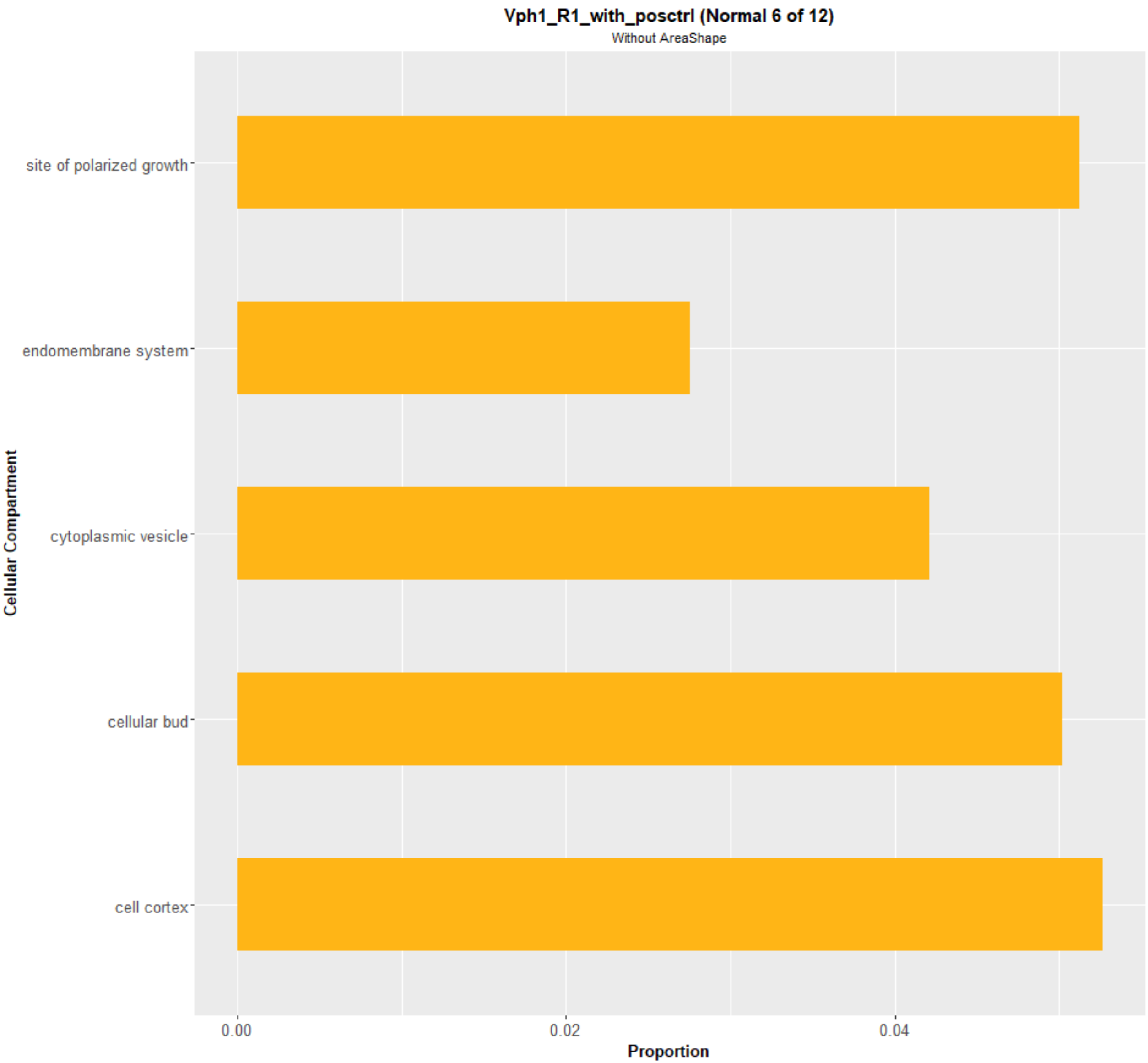

Vph1_R1_with_posctrl (Normal 0 of 12)
Without AreaShape
vesicle organization
proteolysis involved in cellular protein catabolic process
protein glycosylation
mitotic cell cycle
endosomal transport
cytoskeleton organization
cytokinesis
cell wall organization or biogenesis
cell budding
Biological Process
0.00
0.02
0.04
0.06
0.08
Proportion


Vph1_R1_with_posctrl (Normal 2 of 12)
Without AreaShape
site of polarized growth
nucleus
microtubule organizing center
membrane
Golgi apparatus
endoplasmic reticulum
endomembrane system
cytoskeleton
cytoplasmic vesicle
cytoplasm
chromosome
cellular bud
cell cortex
Cellular Compartment
0.00
0.05
0.10
0.15
Proportion


Vph1_R1_with_posctrl (Normal 2 of 12)
Without AreaShape
Biological Process
vesicle organization
tRNA aminoacylation for protein translation
translational initiation
transcription from RNA polymerase II promoter
snoRNA processing
regulation of transport
regulation of organelle organization
protein targeting
protein lipidation
protein complex biogenesis
organelle inheritance
organelle fusion
organelle fission
nucleus organization
nucleobase-containing compound transport
nuclear transport
mRNA processing
mitotic cell cycle
membrane fusion
lipid metabolic process
Golgi vesicle transport
exocytosis
DNA replication
DNA repair
DNA-templated transcription, termination
DNA-templated transcription, initiation
cytoskeleton organization
chromosome segregation
chromatin organization
cellular response to DNA damage stimulus
cell budding
0.0
0.1
0.2
0.3
Proportion


Vph1_R1_with_posctrl (Normal 3 of 12)
Without AreaShape
site of polarized growth
nucleus
microtubule organizing center
membrane
Golgi apparatus
endoplasmic reticulum
endomembrane system
cytoskeleton
cytoplasmic vesicle
chromosome
cellular bud
cell cortex
Cellular Compartment
0.00
0.04
0.08
0.12
Proportion


Vph1_R1_with_posctrl (Normal 3 of 12)
Without AreaShape
vesicle organization
transcription from RNA polymerase II promoter
snoRNA processing
regulation of transport
regulation of organelle organization
protein targeting
protein complex biogenesis
organelle inheritance
organelle fusion
organelle fission
nucleus organization
nucleobase-containing compound transport
nuclear transport
mRNA processing
mitotic cell cycle
membrane fusion
Golgi vesicle transport
exocytosis
DNA repair
DNA-templated transcription, termination
cytoskeleton organization
chromosome segregation
cellular response to DNA damage stimulus
Biological Process
0.0
0.1
0.2
0.3
Proportion


Vph1_R1_with_posctrl (Normal 6 of 12)
Without AreaShape
site of polarized growth
endomembrane system
cytoplasmic vesicle
cellular bud
cell cortex
Cellular Compartment
0.00
0.02
0.04
Proportion

Vph1_R1_with_posctrl (Normal 6 of 12)

Without AreaShape

vesicle organization

transcription from RNA polymerase II promoter

mitotic cell cycle

membrane fusion

Golgi vesicle transport

DNA repair

chromatin organization

cellular response to DNA damage stimulus

cell wall organization or biogenesis

Biological Process

0.000

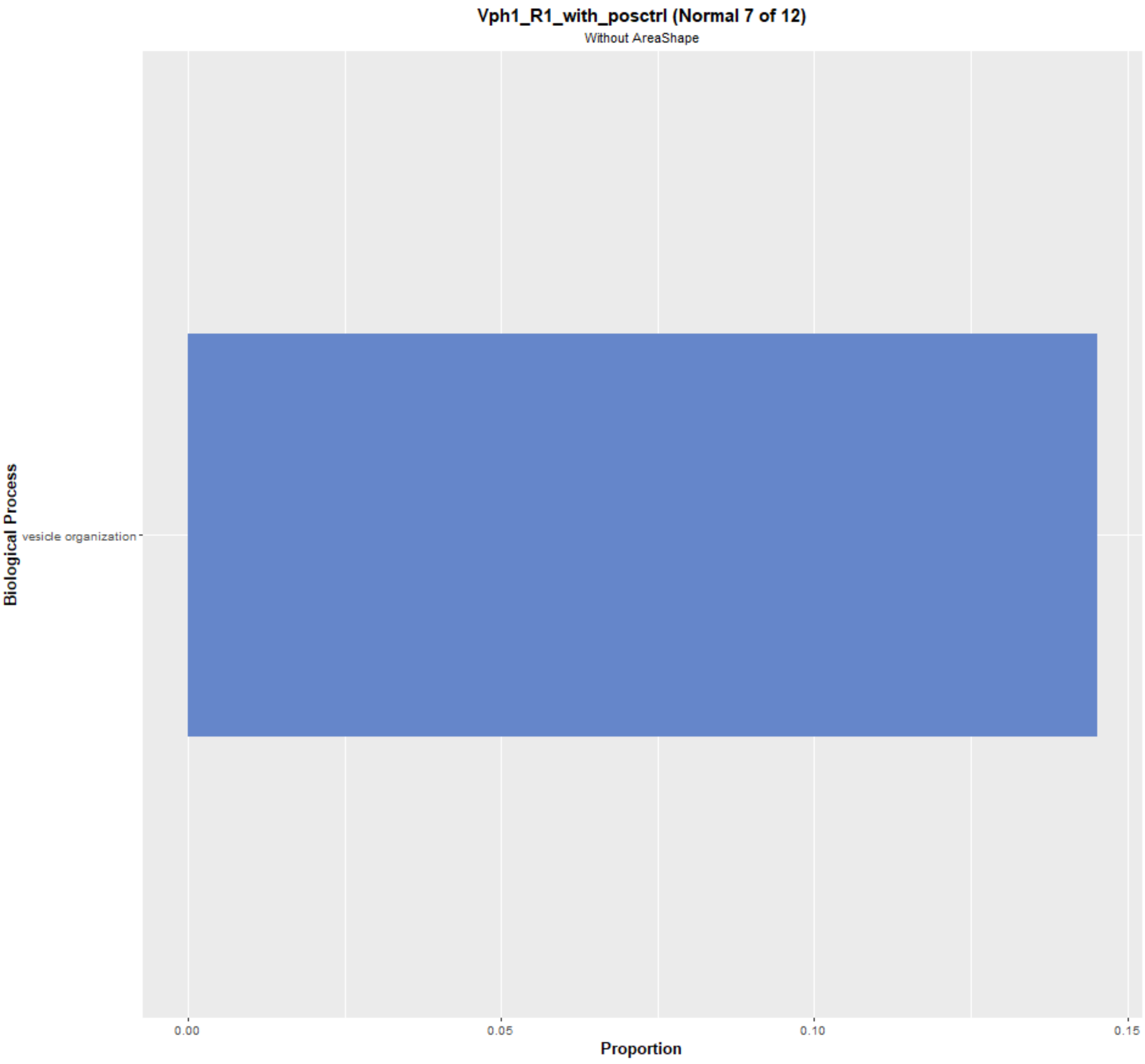
0.025

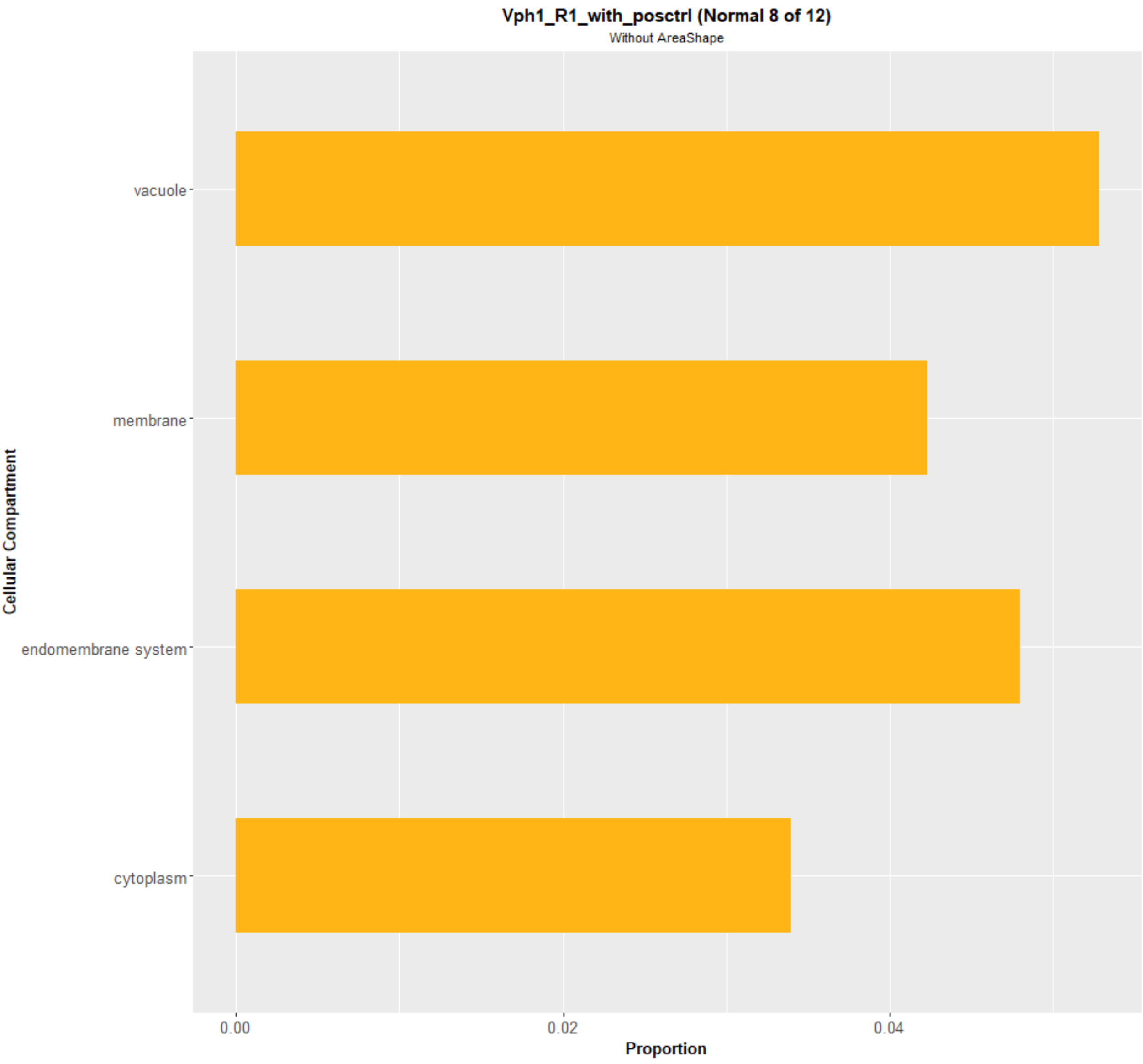
0.050

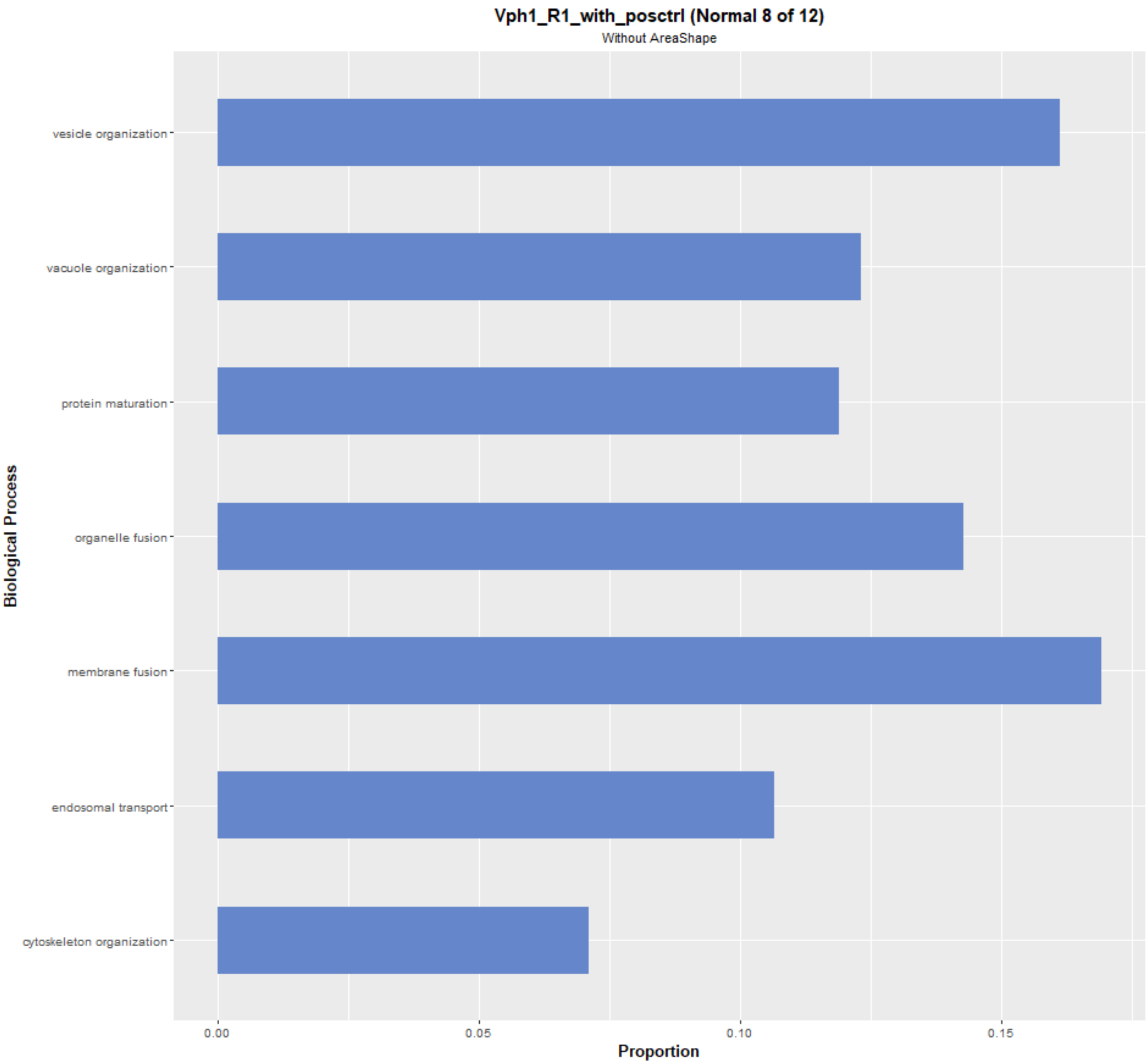
0.075

0.100

Proportion


Vph1_R1_with_posctrl (Normal 7 of 12)
Without AreaShape
Biological Process
vesicle organization
0.00
0.05
0.10
0.15
Proportion


Vph1_R1_with_posctrl (Normal 8 of 12)
Without AreaShape
vacuole
membrane
endomembrane system
cytoplasm
Cellular Compartment
0.00
0.02
0.04
Proportion


Vph1_R1_with_posctrl (Normal 8 of 12)
Without AreaShape
vesicle organization
vacuole organization
protein maturation
organelle fusion
membrane fusion
endosomal transport
cytoskeleton organization
Biological Process
0.00
0.05
0.10
0.15
Proportion

Vph1_R1_with_posctrl (Normal 9 of 12)

Without AreaShape

Cellular Compartment

cytoskeleton

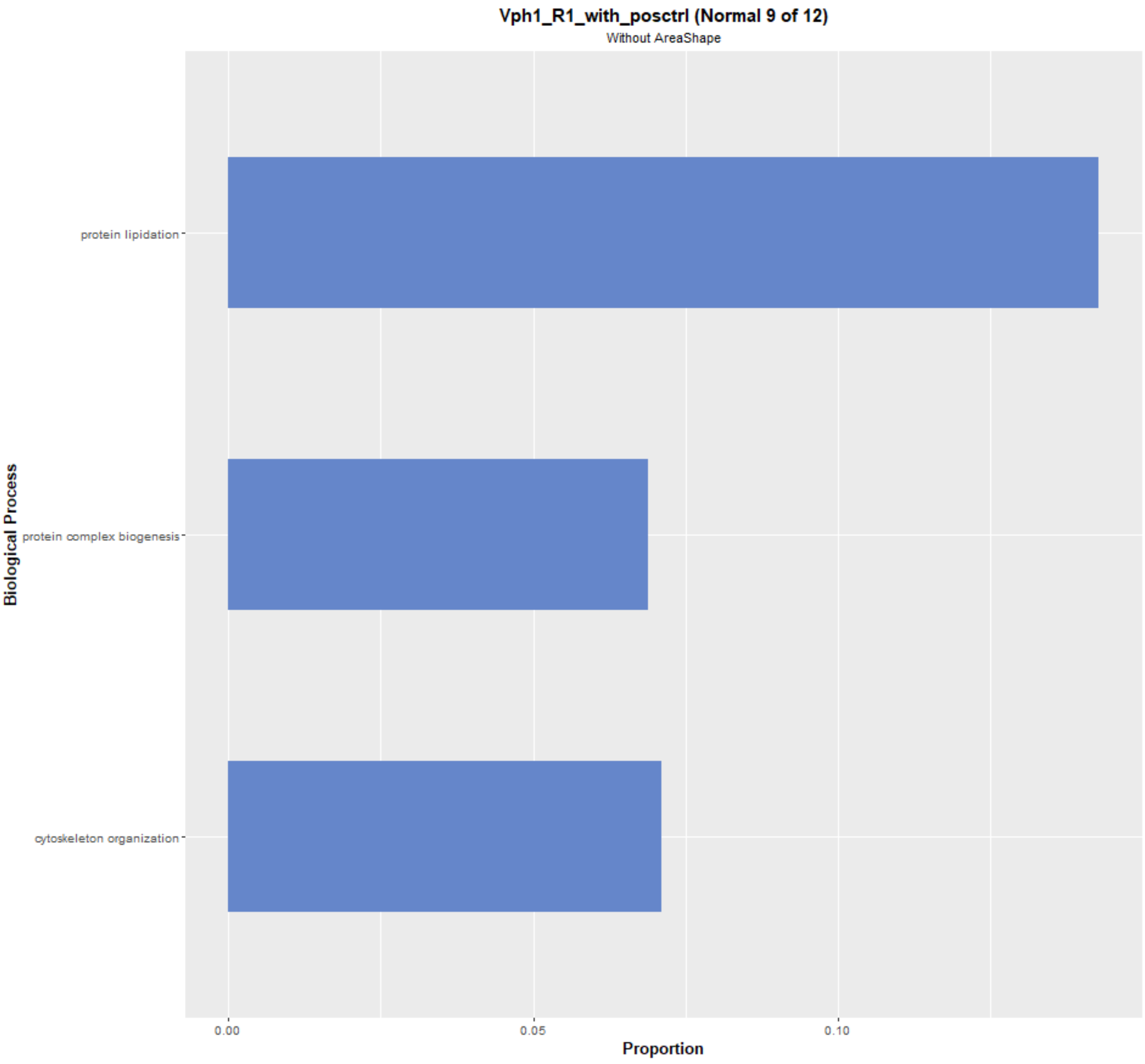
0.00

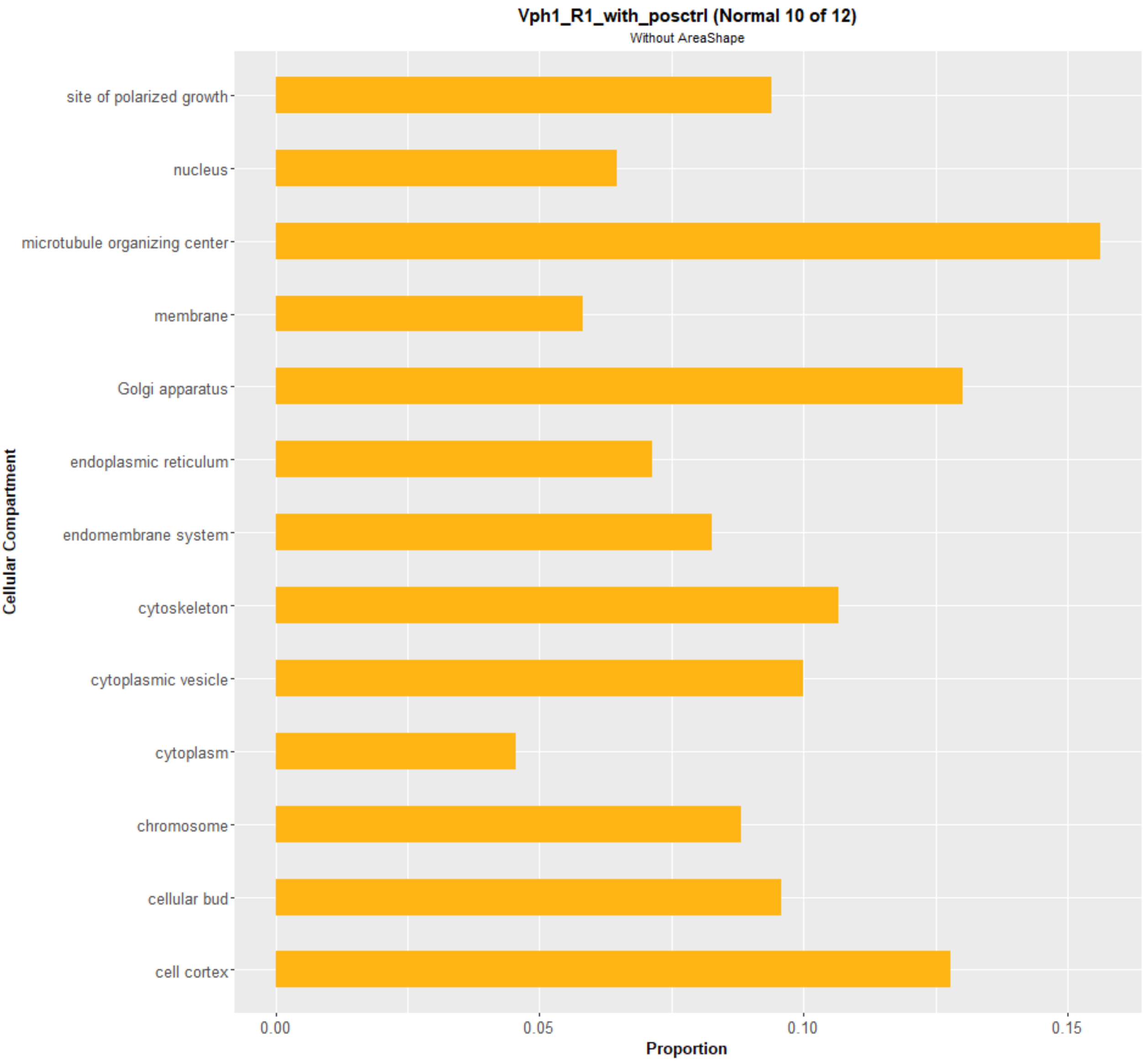
0.02

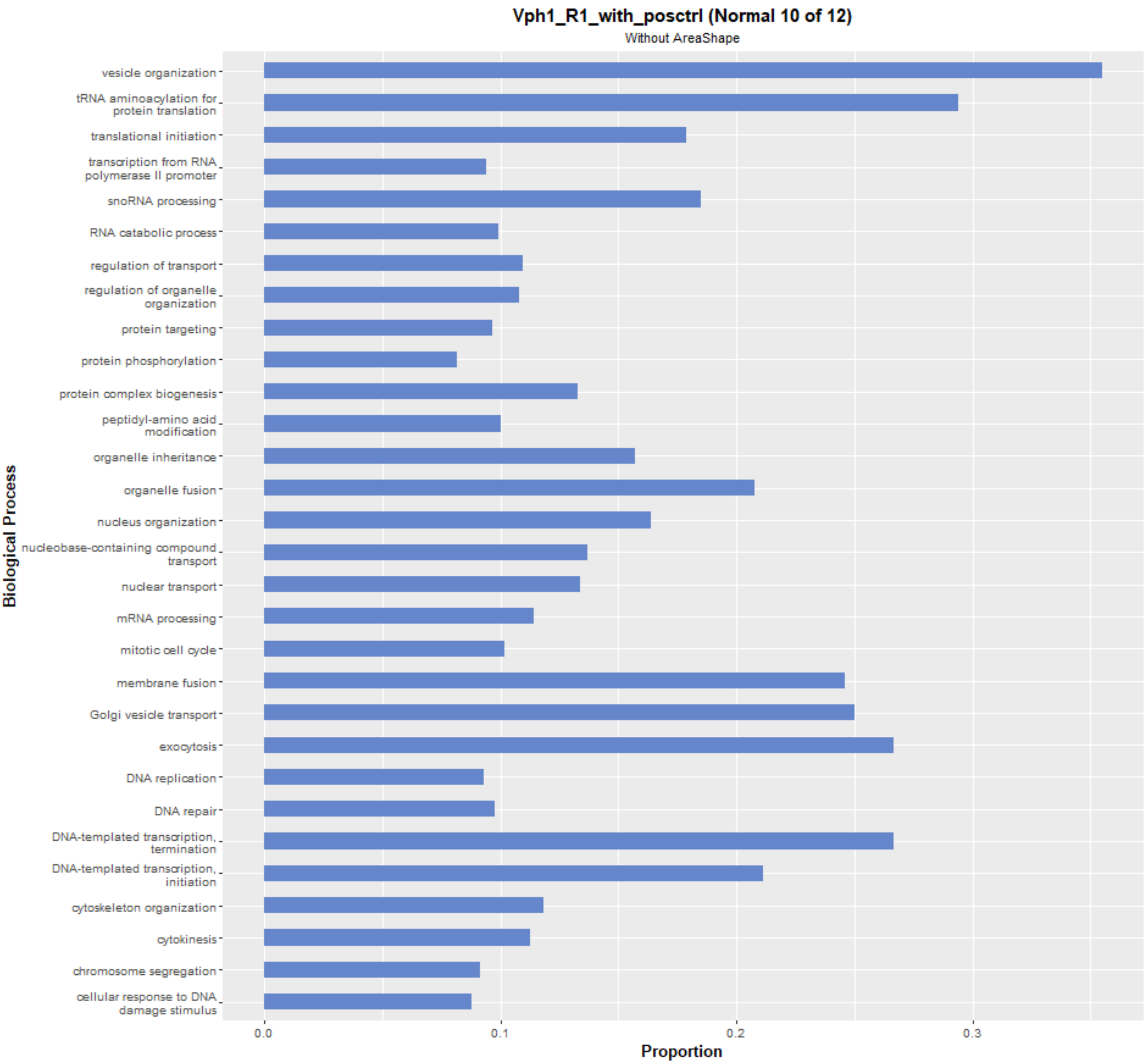
0.04

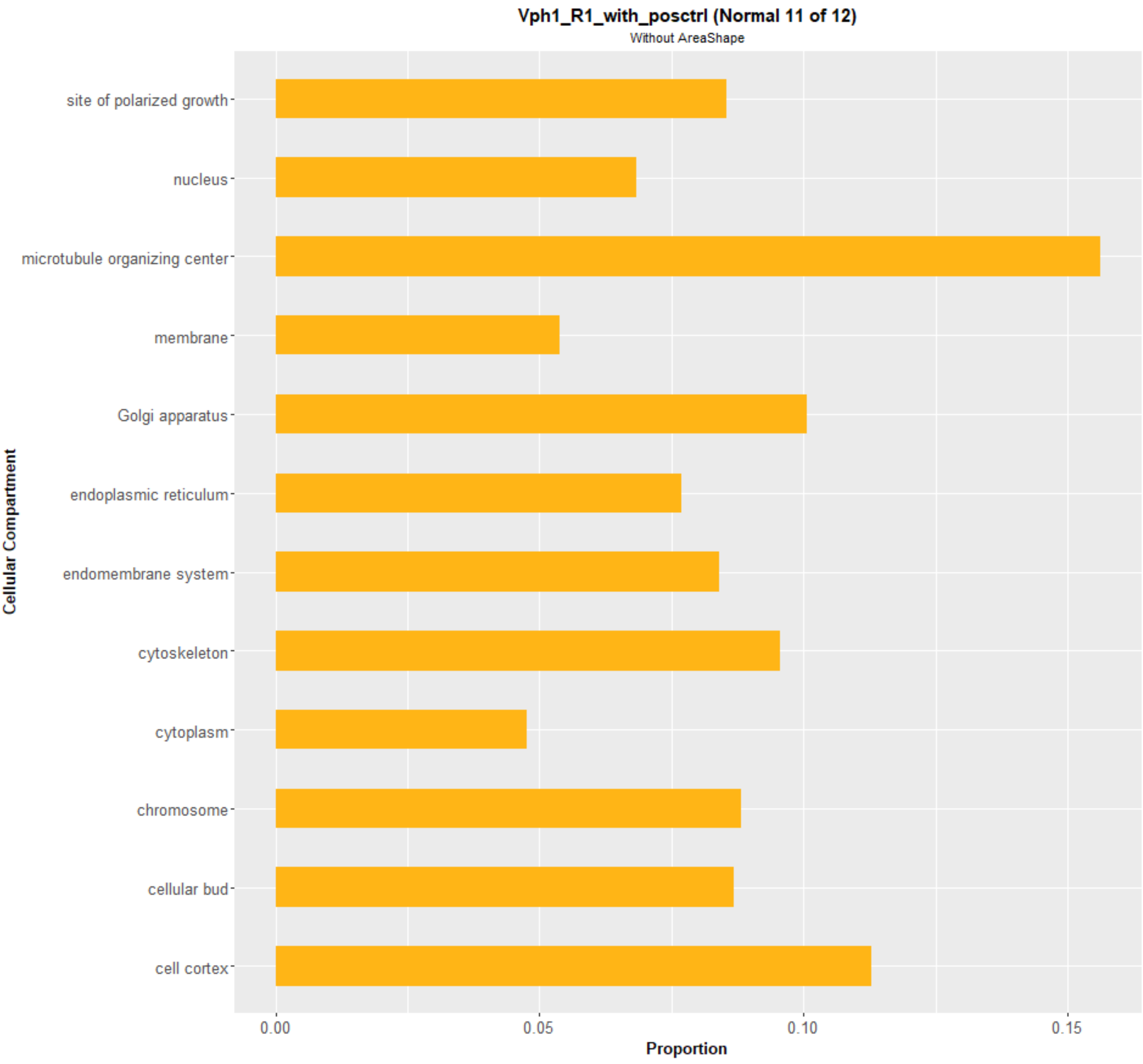
0.06

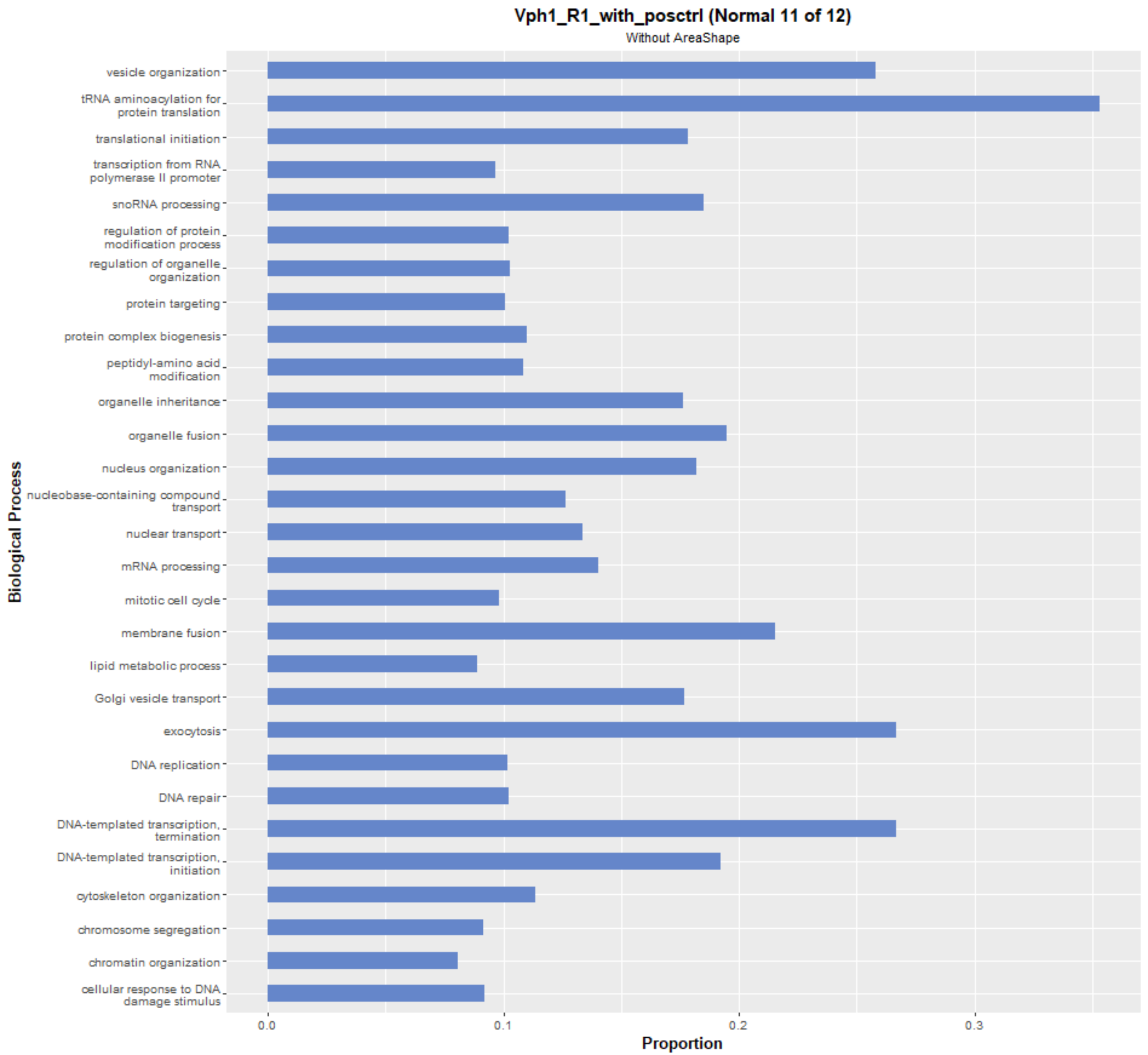
Proportion


Vph1_R1_with_posctrl (Normal 9 of 12)
Without AreaShape
protein lipidation
protein complex biogenesis
cytoskeleton organization
Biological Process
0.00
0.05
0.10
Proportion


Vph1_R1_with_posctrl (Normal 10 of 12)
Without AreaShape
site of polarized growth
nucleus
microtubule organizing center
membrane
Golgi apparatus
endoplasmic reticulum
endomembrane system
cytoskeleton
cytoplasmic vesicle
cytoplasm
chromosome
cellular bud
cell cortex
Cellular Compartment
0.00
0.05
0.10
0.15
Proportion


Vph1_R1_with_posctrl (Normal 10 of 12)
Without AreaShape
vesicle organization
tRNA aminoacylation for protein translation
translational initiation
transcription from RNA polymerase II promoter
snoRNA processing
RNA catabolic process
regulation of transport
regulation of organelle organization
protein targeting
protein phosphorylation
protein complex biogenesis
peptidyl-amino acid modification
organelle inheritance
organelle fusion
nucleus organization
nucleobase-containing compound transport
nuclear transport
mRNA processing
mitotic cell cycle
membrane fusion
Golgi vesicle transport
exocytosis
DNA replication
DNA repair
DNA-templated transcription, termination
DNA-templated transcription, initiation
cytoskeleton organization
cytokinesis
chromosome segregation
cellular response to DNA damage stimulus
Biological Process
0.0
0.1
0.2
0.3
Proportion


Vph1_R1_with_posctrl (Normal 11 of 12)
Without AreaShape
site of polarized growth
nucleus
microtubule organizing center
membrane
Golgi apparatus
endoplasmic reticulum
endomembrane system
cytoskeleton
cytoplasm
chromosome
cellular bud
cell cortex
Cellular Compartment
0.00
0.05
0.10
0.15
Proportion


Vph1_R1_with_posctrl (Normal 11 of 12)
Without AreaShape
Biological Process
vesicle organization
tRNA aminoacylation for protein translation
translational initiation
transcription from RNA polymerase II promoter
snoRNA processing
regulation of protein modification process
regulation of organelle organization
protein targeting
protein complex biogenesis
peptidyl-amino acid modification
organelle inheritance
organelle fusion
nucleus organization
nucleobase-containing compound transport
nuclear transport
mRNA processing
mitotic cell cycle
membrane fusion
lipid metabolic process
Golgi vesicle transport
exocytosis
DNA replication
DNA repair
DNA-templated transcription, termination
DNA-templated transcription, initiation
cytoskeleton organization
chromosome segregation
chromatin organization
cellular response to DNA damage stimulus
0.0
0.1
0.2
0.3
Proportion