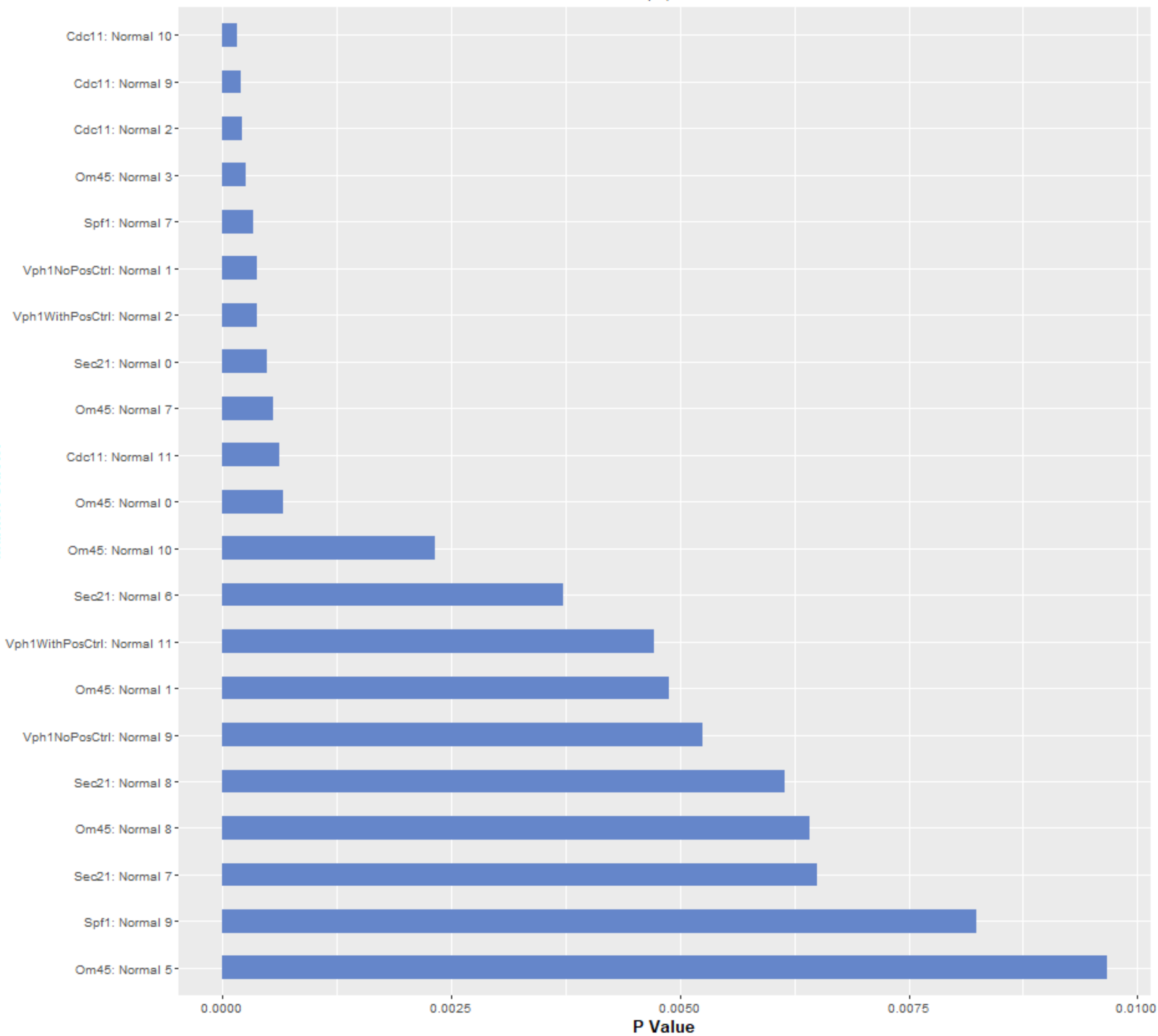


# cell budding

Without AreaShape | Ordered Gene Set

Marker: Cluster



# cell morphogenesis

Without AreaShape | Ordered Gene Set

Marker: Cluster

Cdc11: Normal 10

Om45: Normal 3

0.000

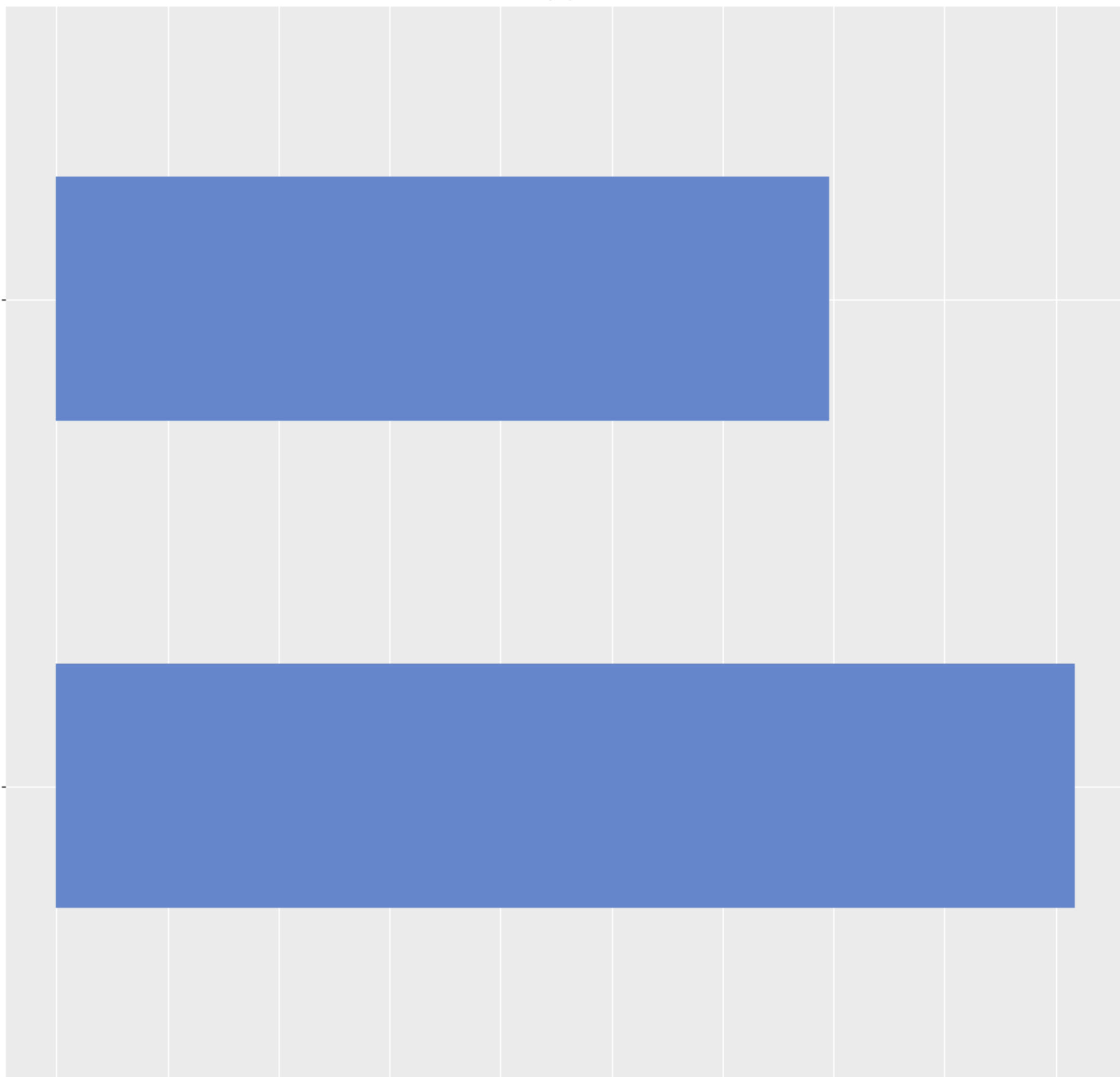
0.001

0.002

0.003

0.004

P Value



# cell wall organization or biogenesis

Without AreaShape | Ordered Gene Set

Marker: Cluster

Cdc11: Normal 2

Vph1WithPosCtrl: Normal 0

Vph1NoPosCtrl: Normal 2

Vph1WithPosCtrl: Normal 6

Vph1NoPosCtrl: Normal 8

0.000

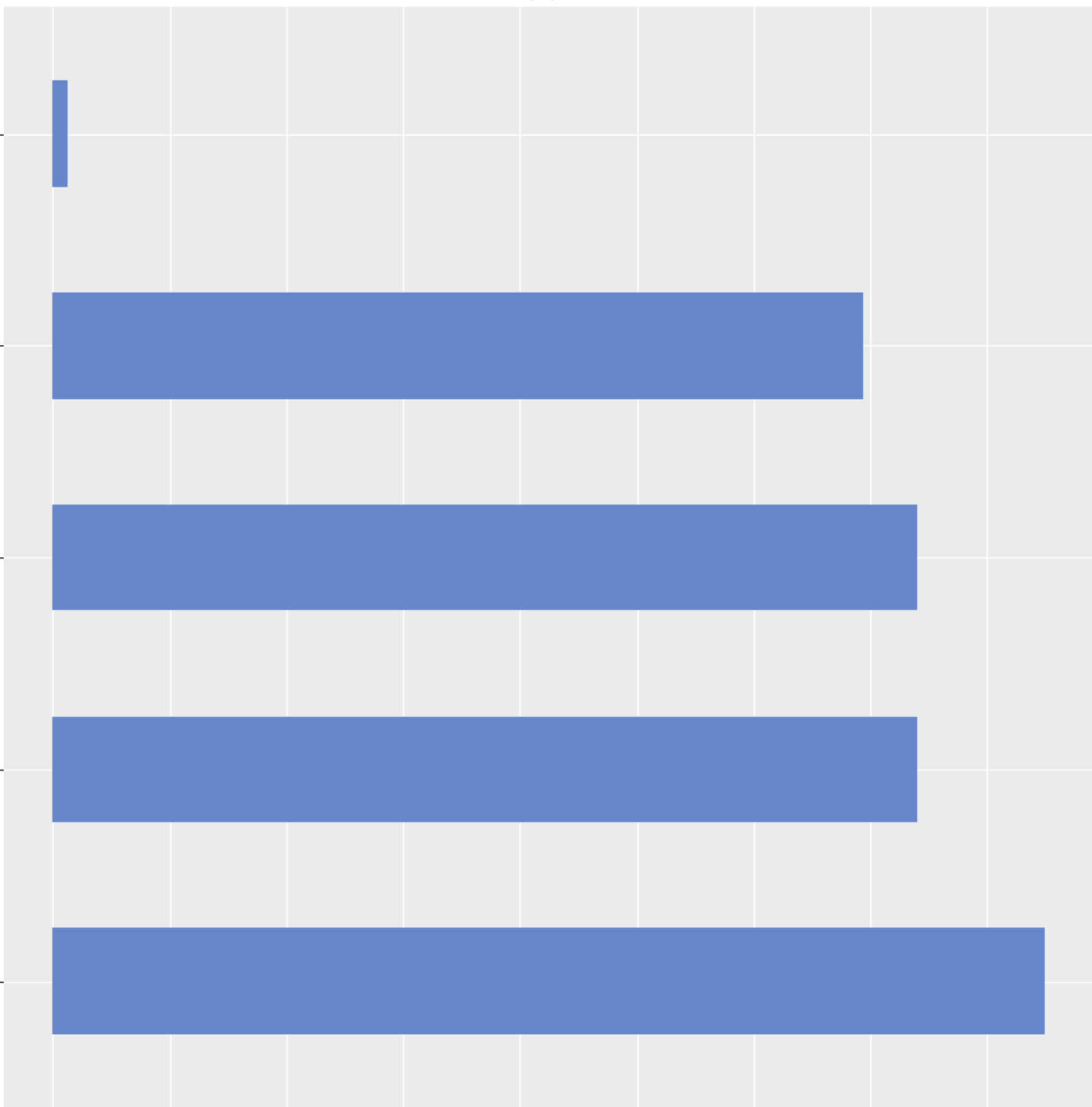
0.002

0.004

0.006

0.008

P Value



# cellular amino acid metabolic process

Without AreaShape | Ordered Gene Set

Marker: Cluster

Vph1NoPosCtrl: Normal 9

Vph1WithPosCtrl: Normal 11

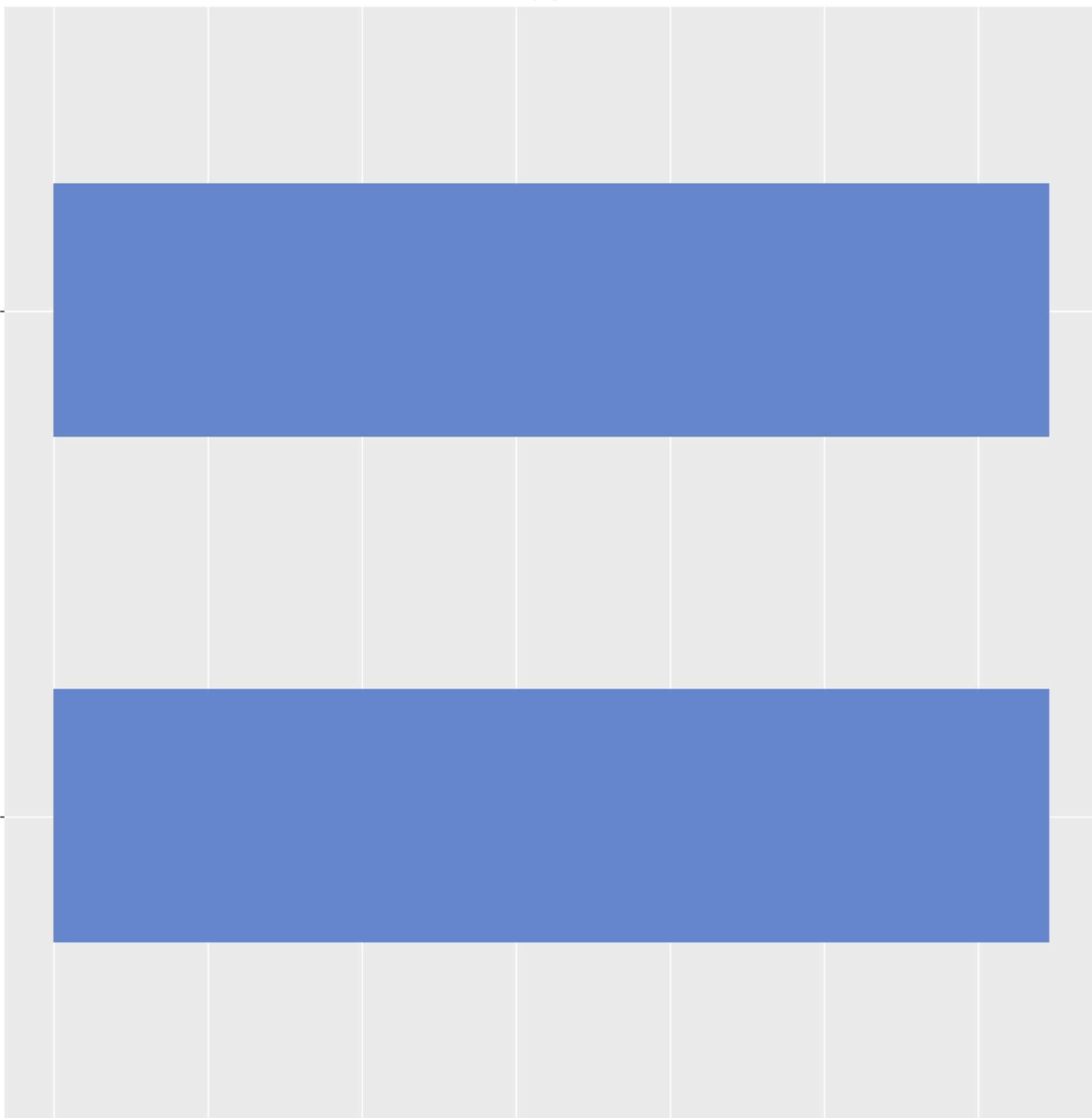
0.000

0.002

0.004

0.006

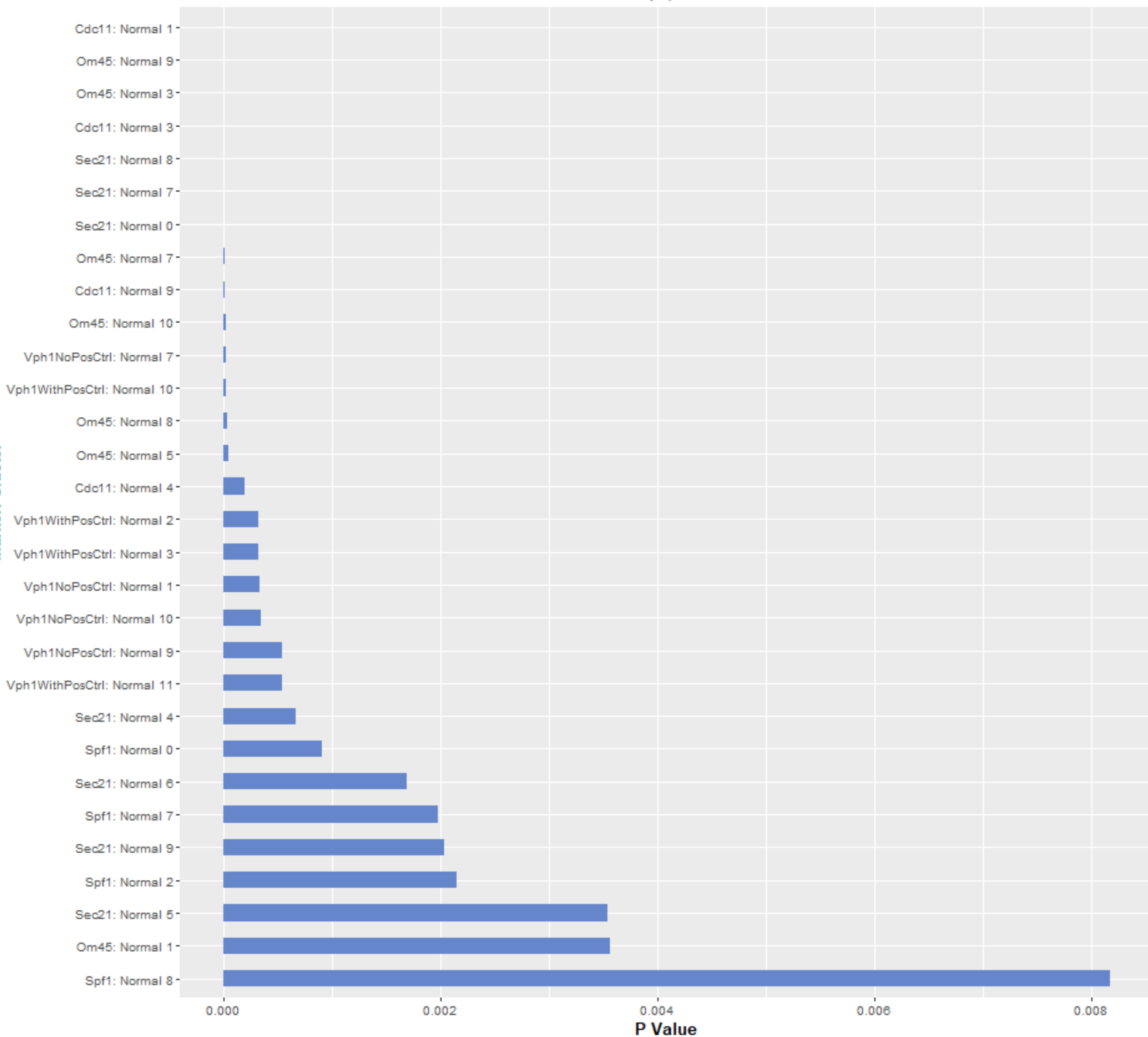
P Value



# cellular response to DNA damage stimulus

Without AreaShape | Ordered Gene Set

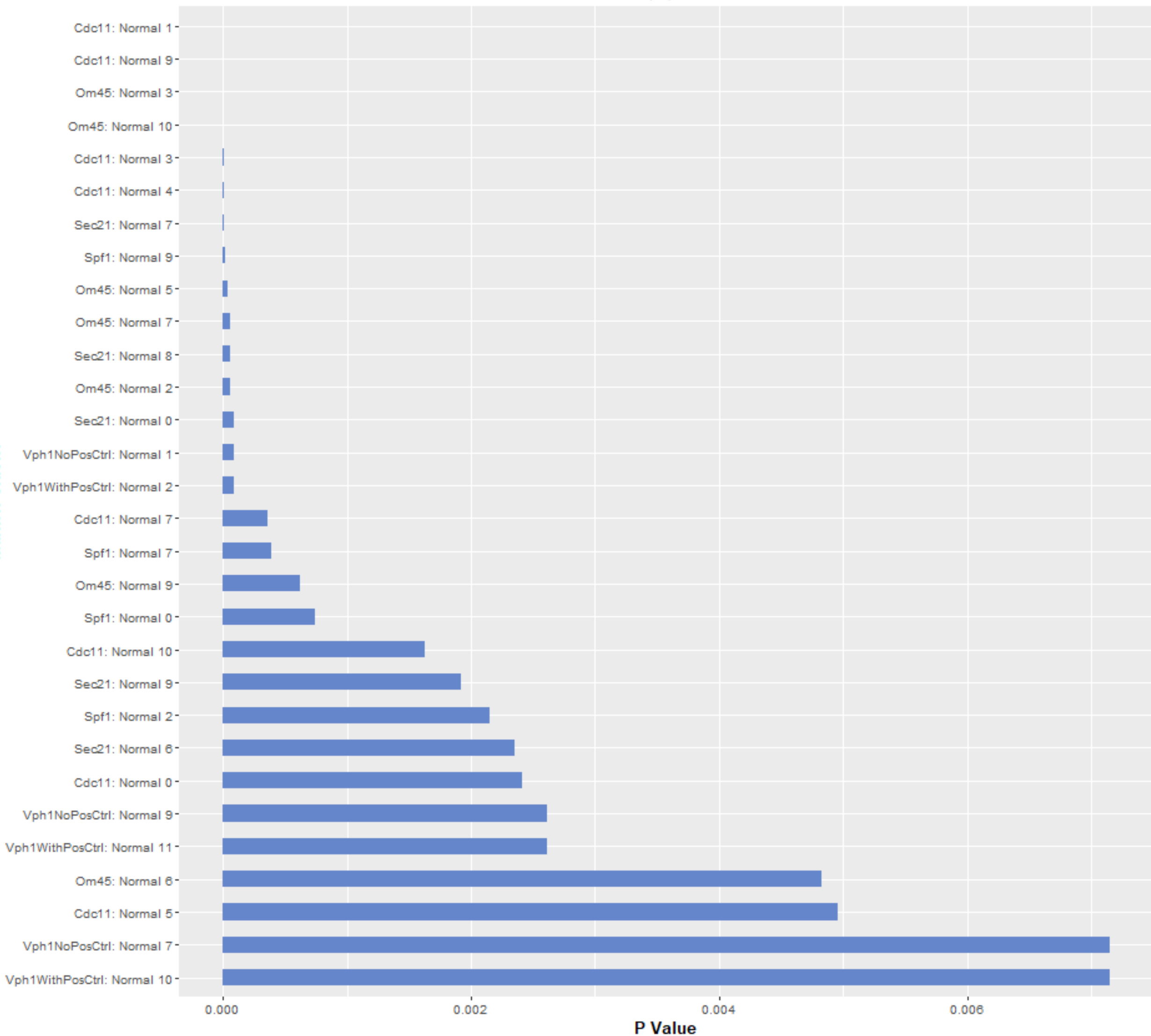
Marker: Cluster



# chromatin organization

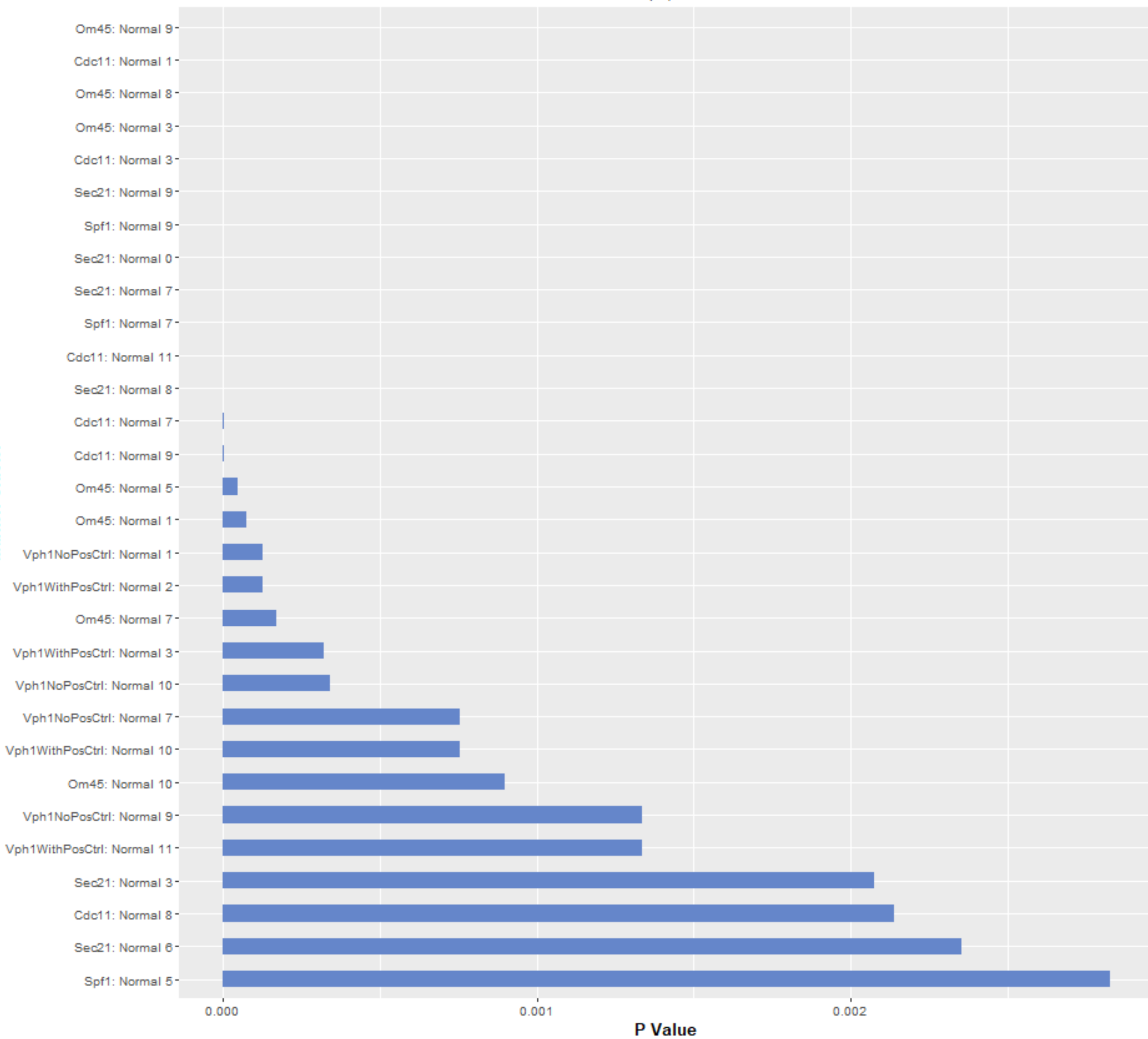
Without AreaShape | Ordered Gene Set

Marker: Cluster



**chromosome segregation**  
Without AreaShape | Ordered Gene Set

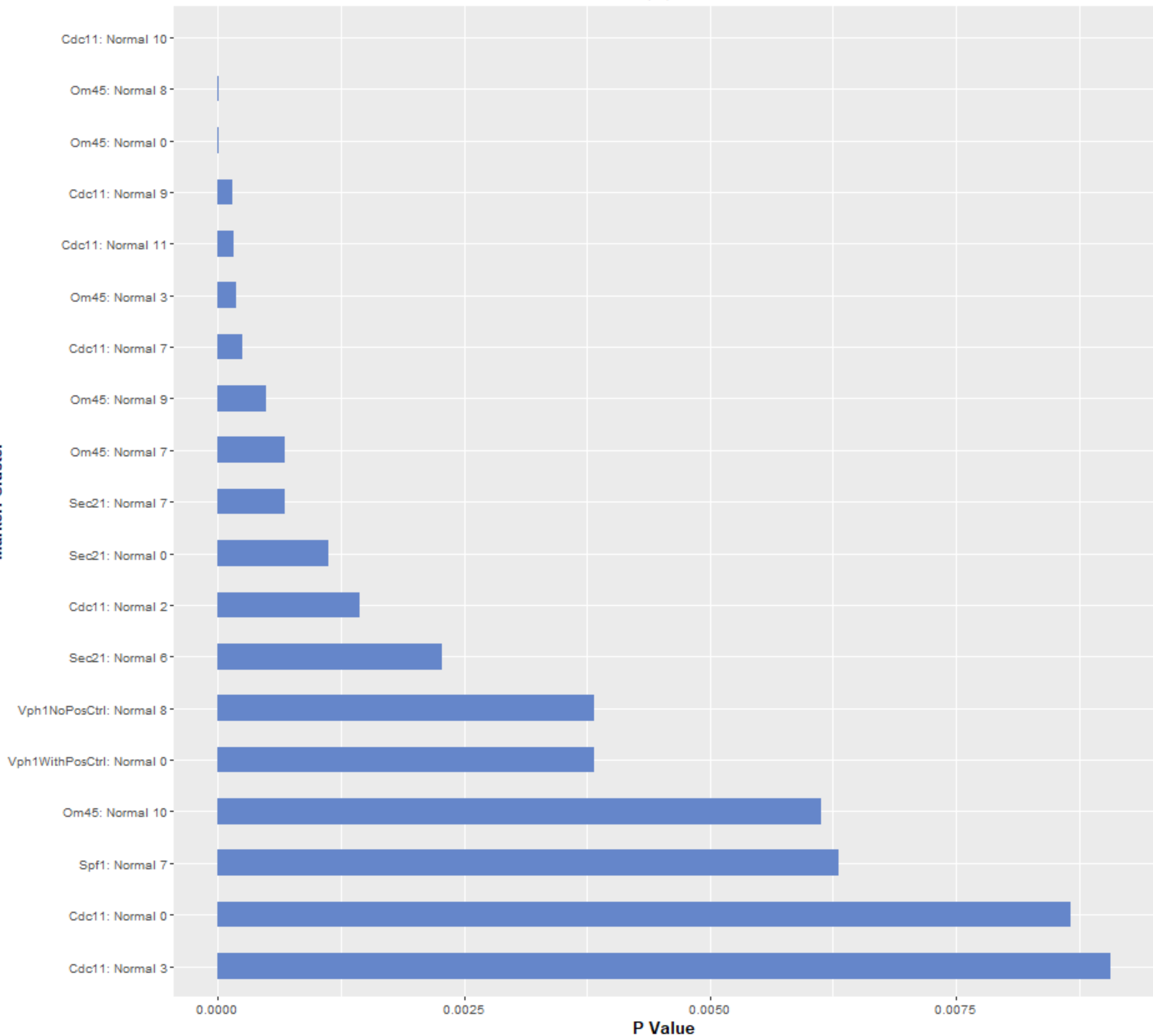
**Marker: Cluster**



# cytokinesis

Without AreaShape | Ordered Gene Set

Marker: Cluster





# cytoplasmic translation

Without AreaShape | Ordered Gene Set

Marker: Cluster

Vph1NoPosCtrl: Normal 9

Vph1WithPosCtrl: Normal 11

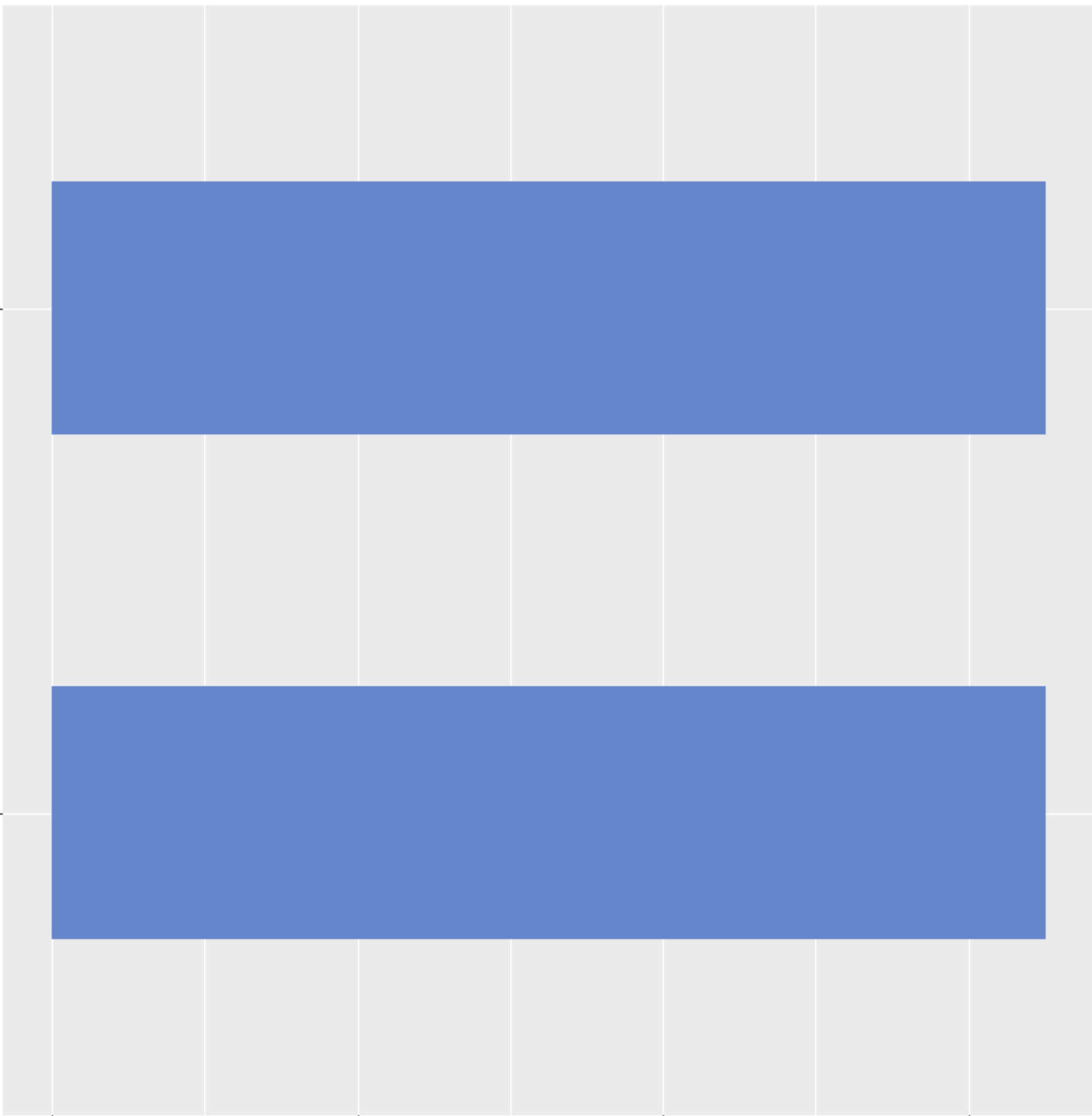
0.000

0.002

0.004

0.006

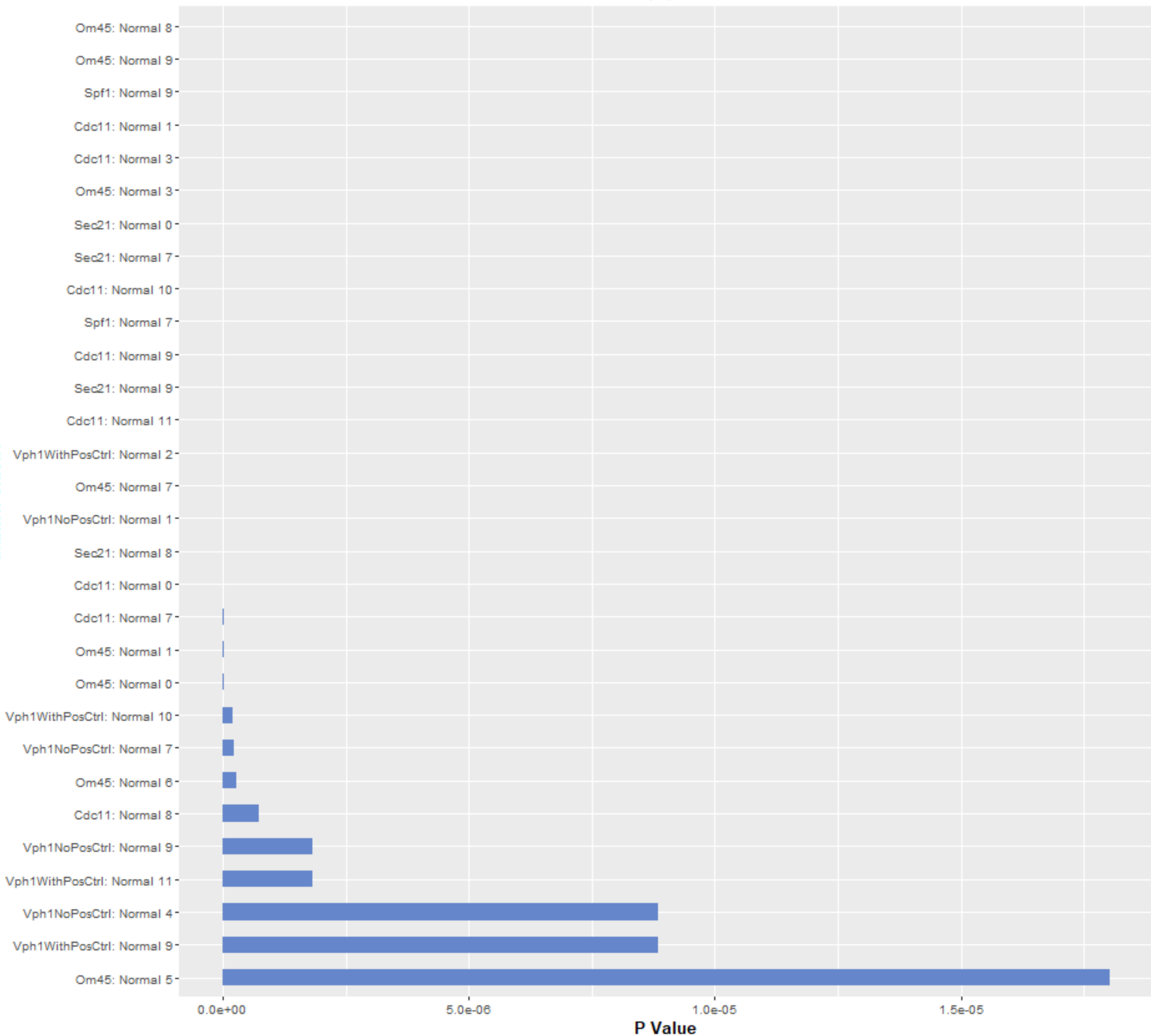
P Value



# cytoskeleton organization

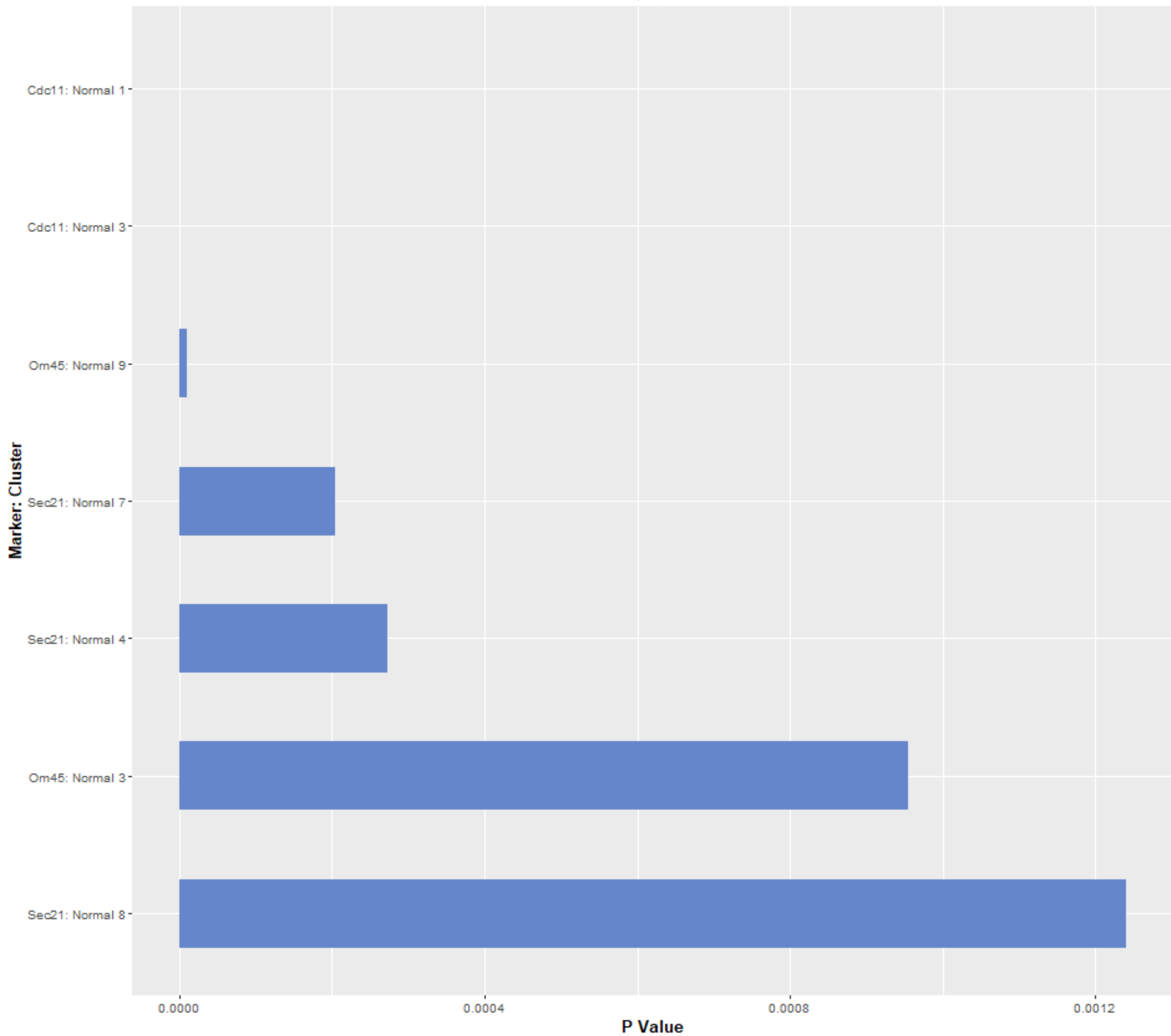
Without AreaShape | Ordered Gene Set

Marker: Cluster



# DNA recombination

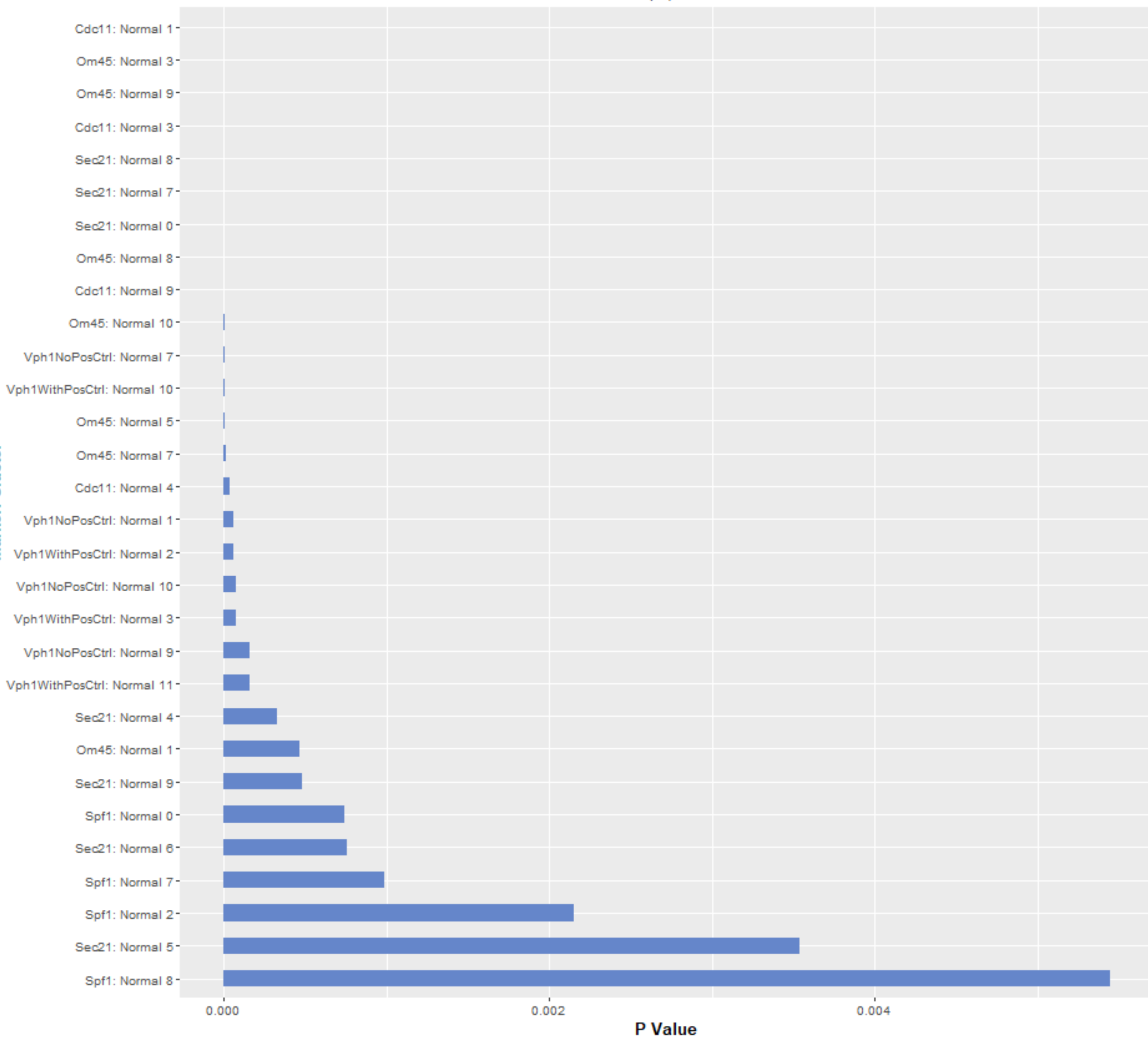
Without AreaShape | Ordered Gene Set



# DNA repair

Without AreaShape | Ordered Gene Set

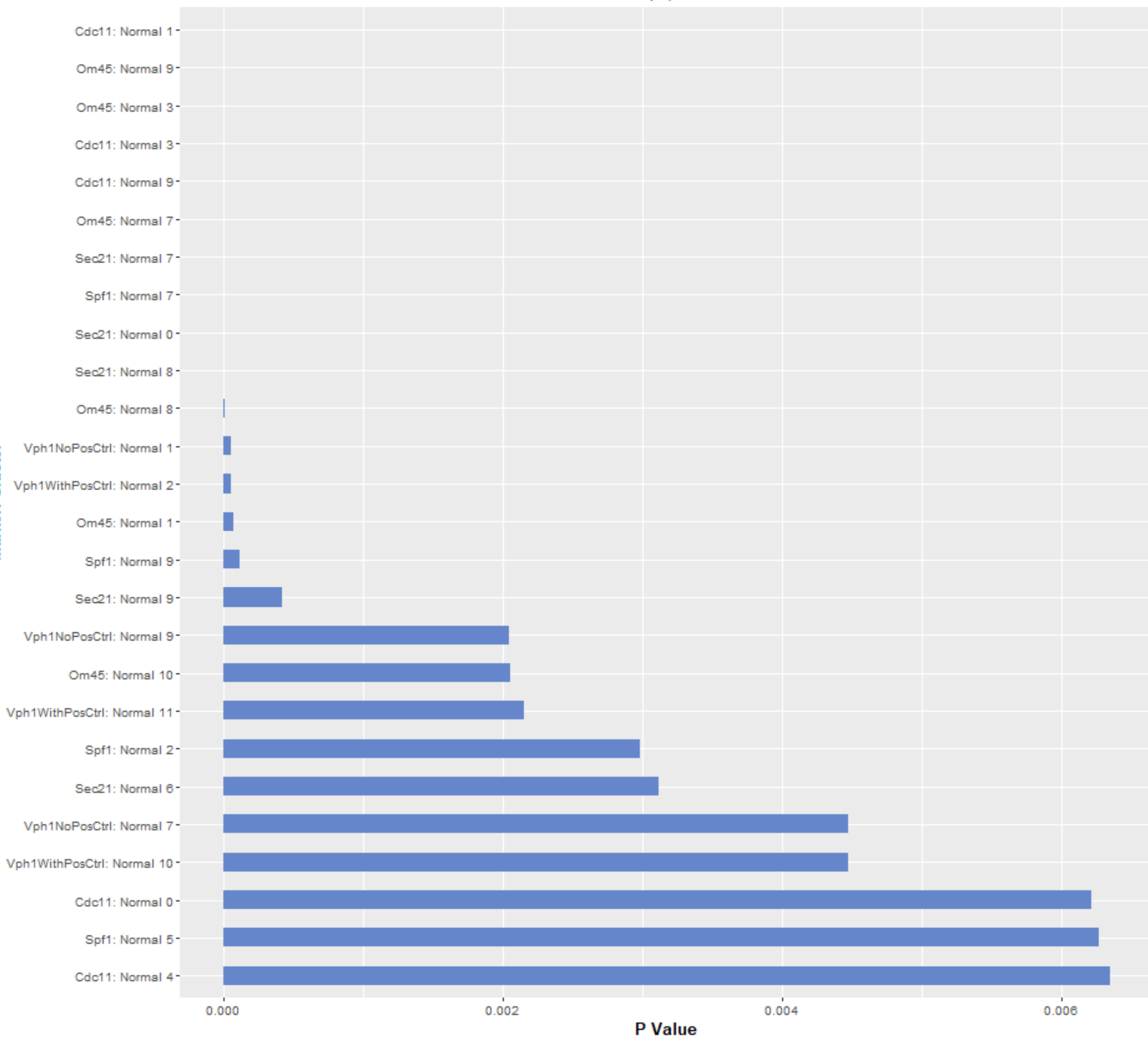
Marker: Cluster



# DNA replication

Without AreaShape | Ordered Gene Set

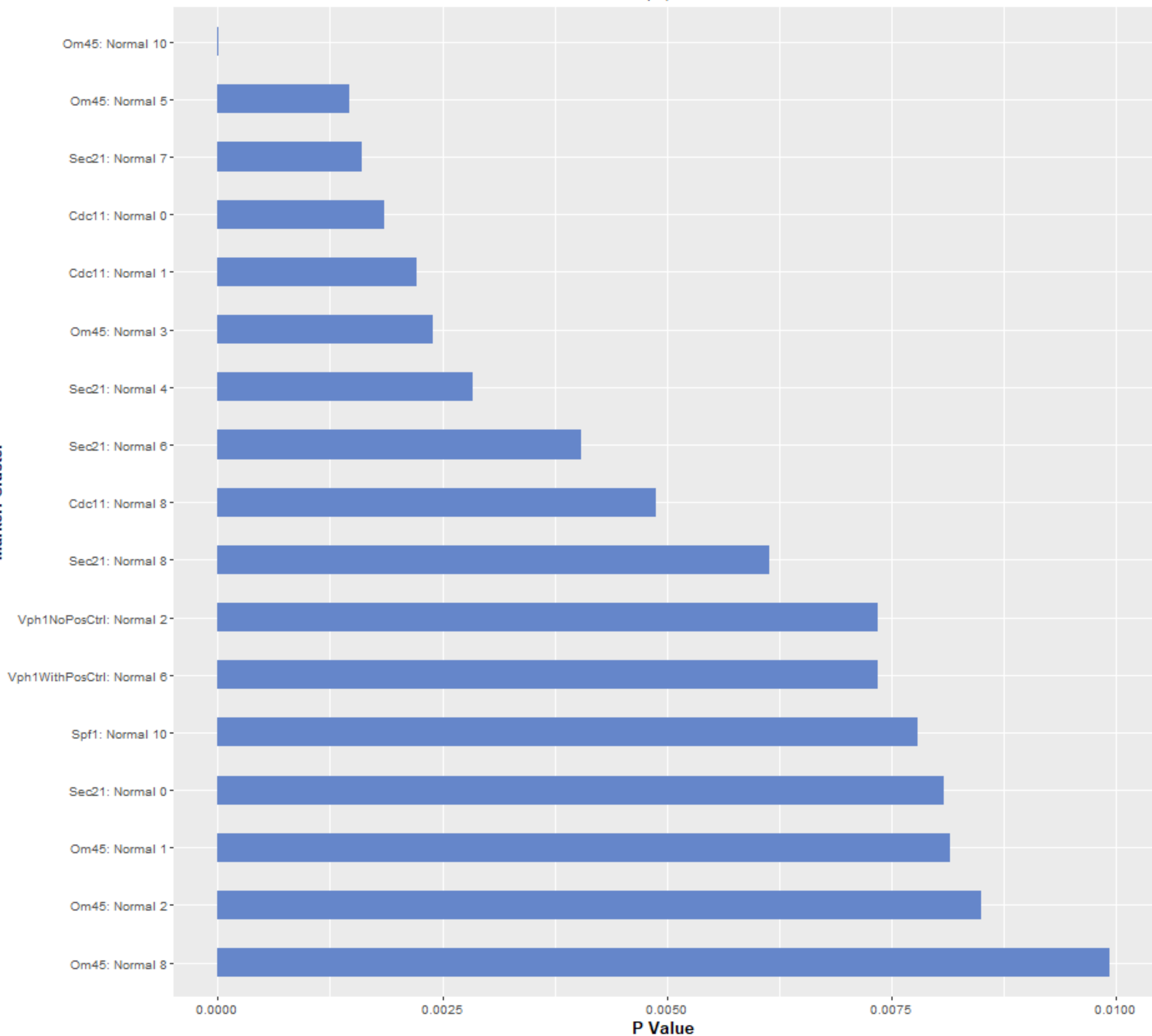
Marker: Cluster



# DNA-templated transcription, elongation

Without AreaShape | Ordered Gene Set

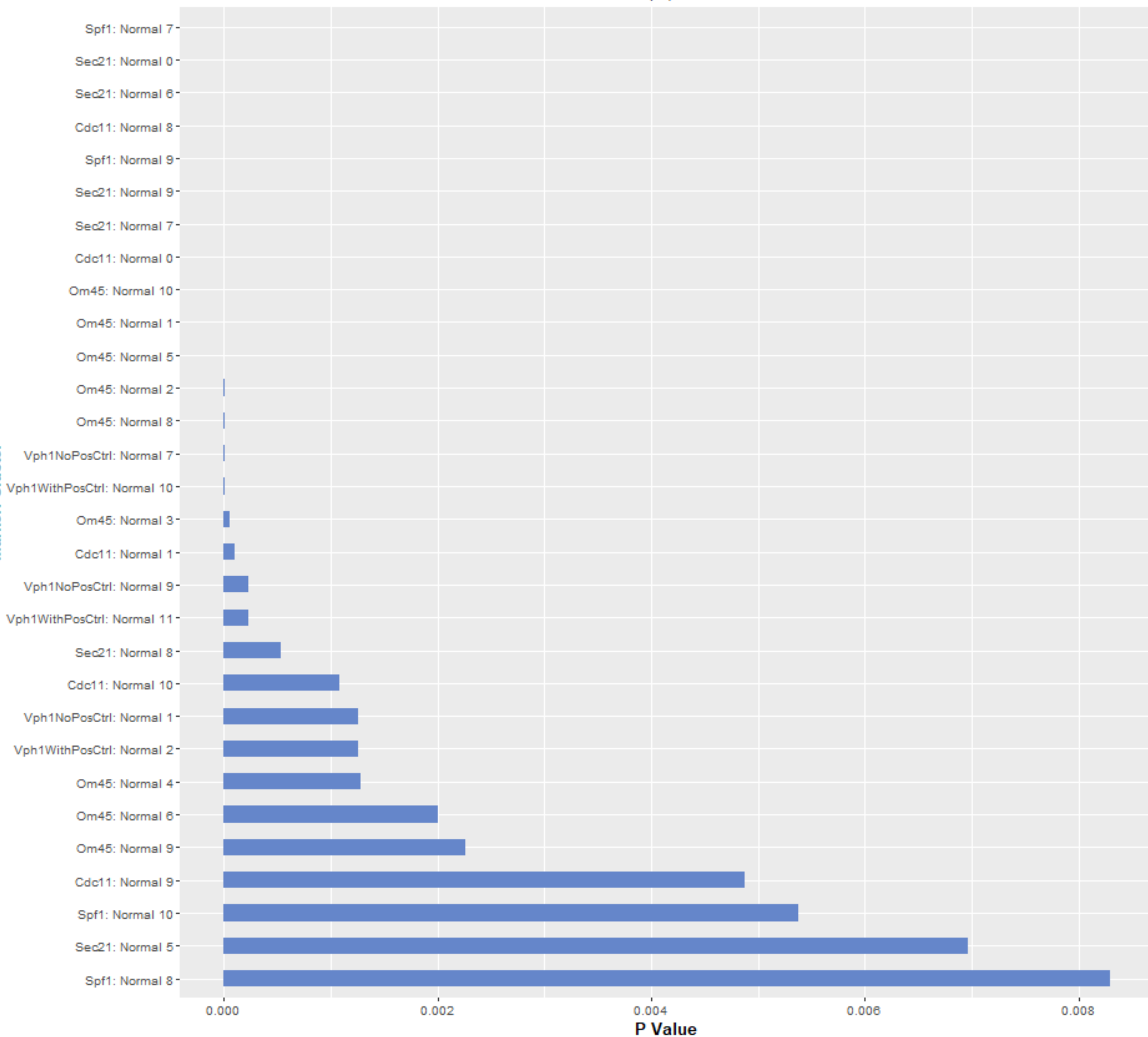
Marker: Cluster



# DNA-templated transcription, initiation

Without AreaShape | Ordered Gene Set

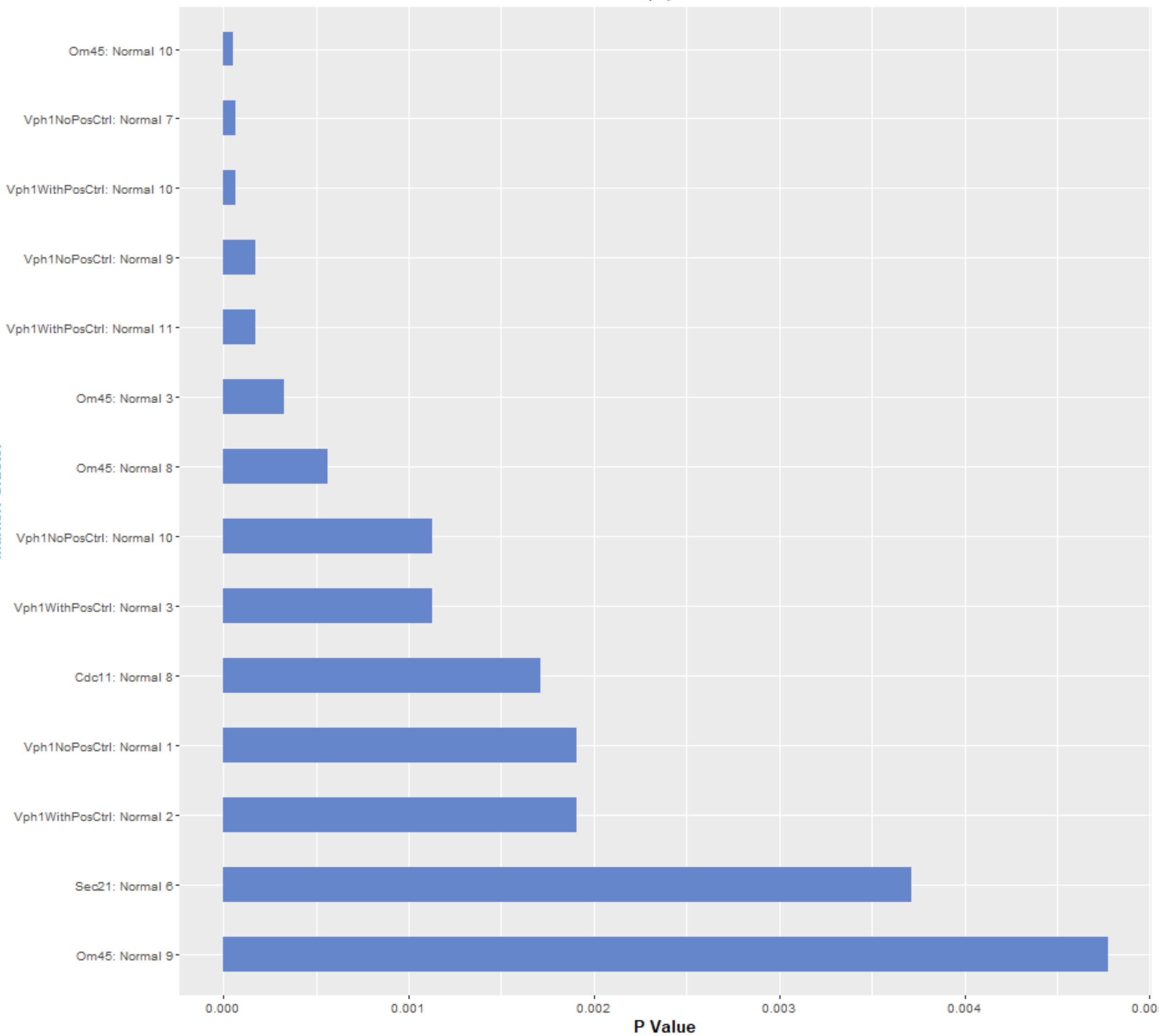
Marker: Cluster



# DNA-templated transcription, termination

Without AreaShape | Ordered Gene Set

Marker: Cluster

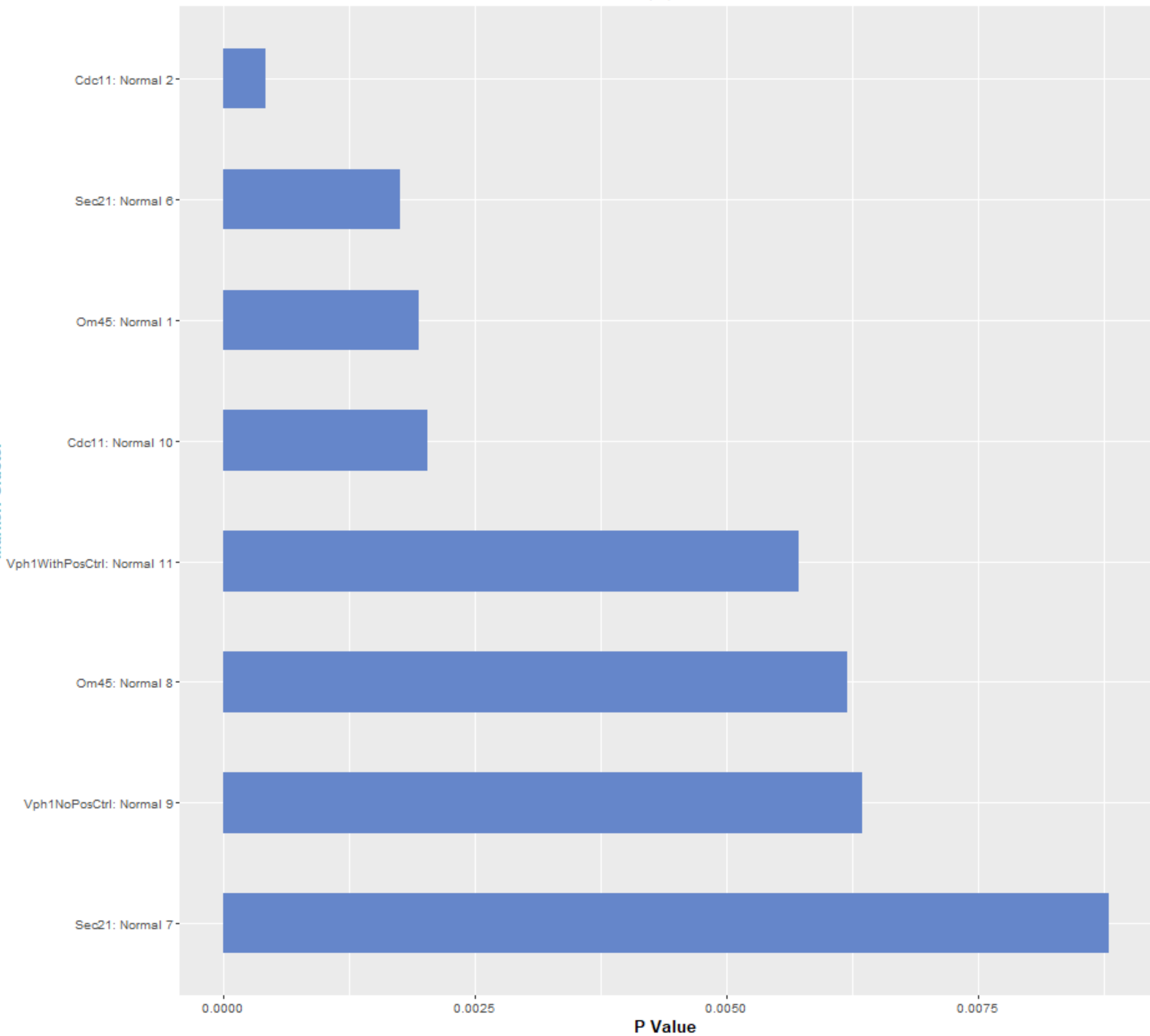




# endocytosis

Without AreaShape | Ordered Gene Set

Marker: Cluster



# endosomal transport

Without AreaShape | Ordered Gene Set

Marker: Cluster

Vph1WithPosCtrl: Normal 8

Vph1NoPosCtrl: Normal 5

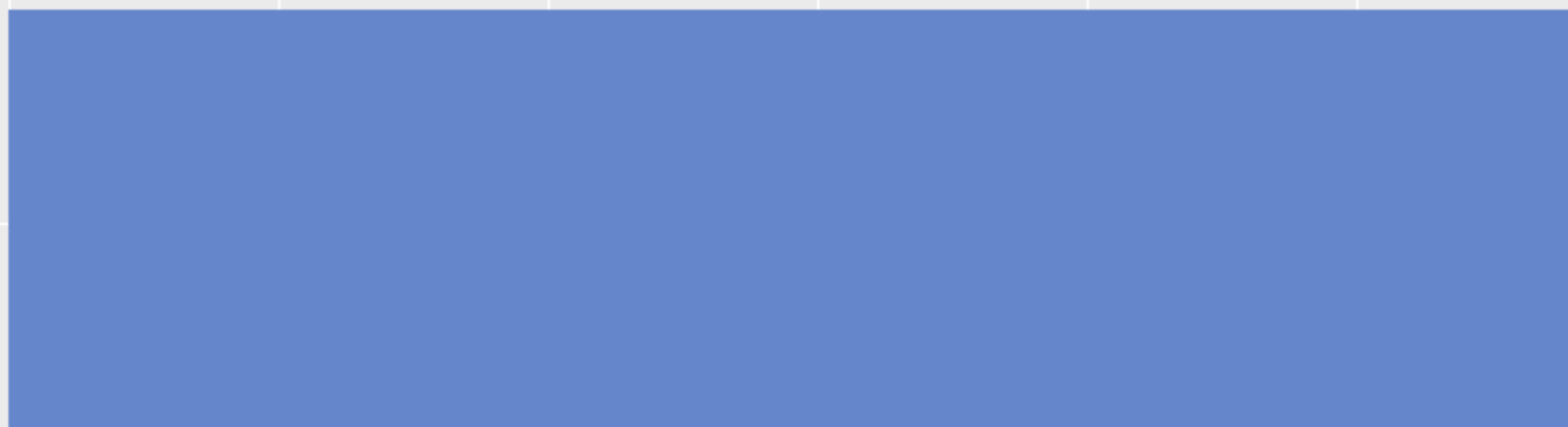
0.0000

0.0004

P Value

0.0008

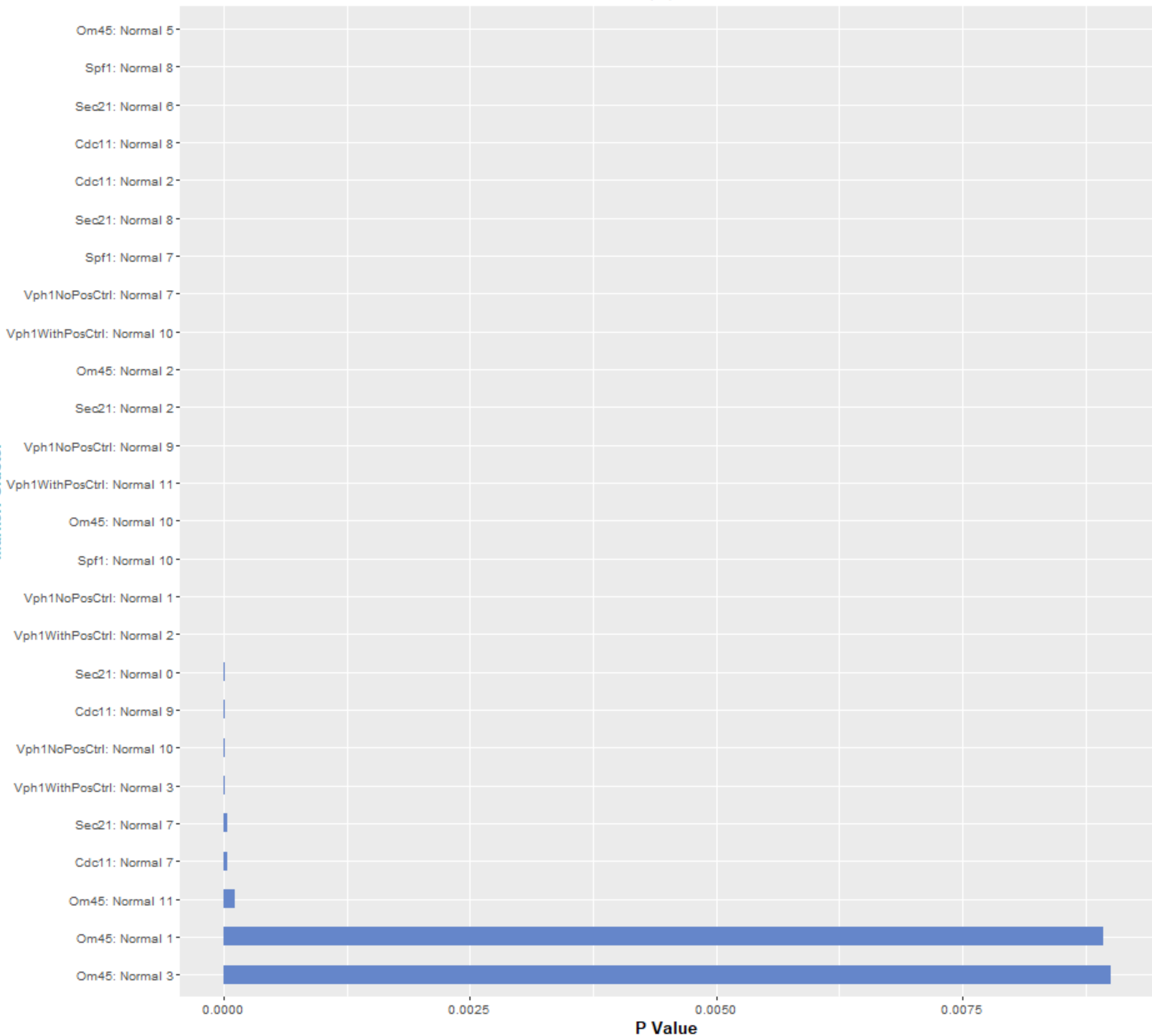
0.0012



# exocytosis

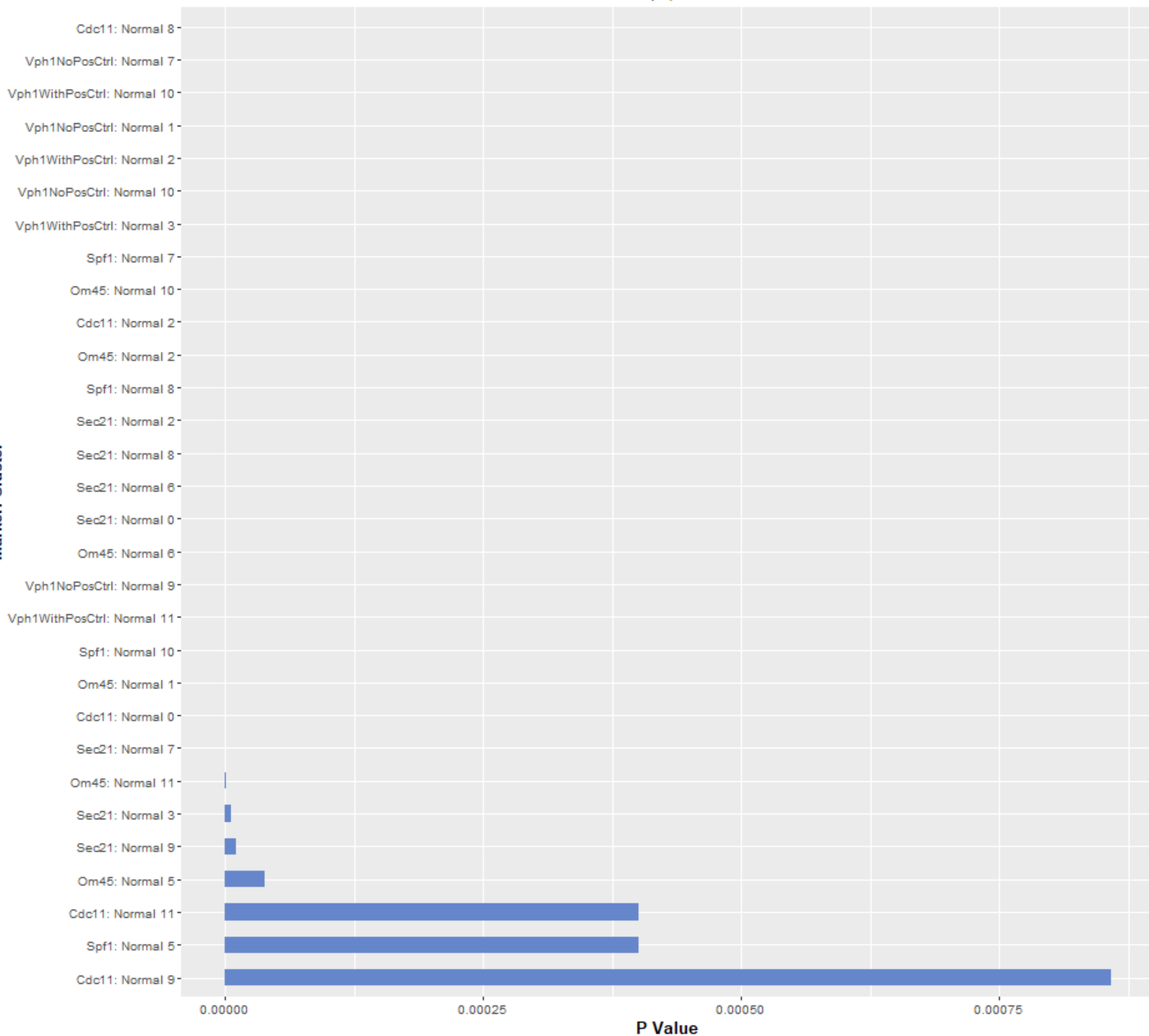
Without AreaShape | Ordered Gene Set

Marker: Cluster



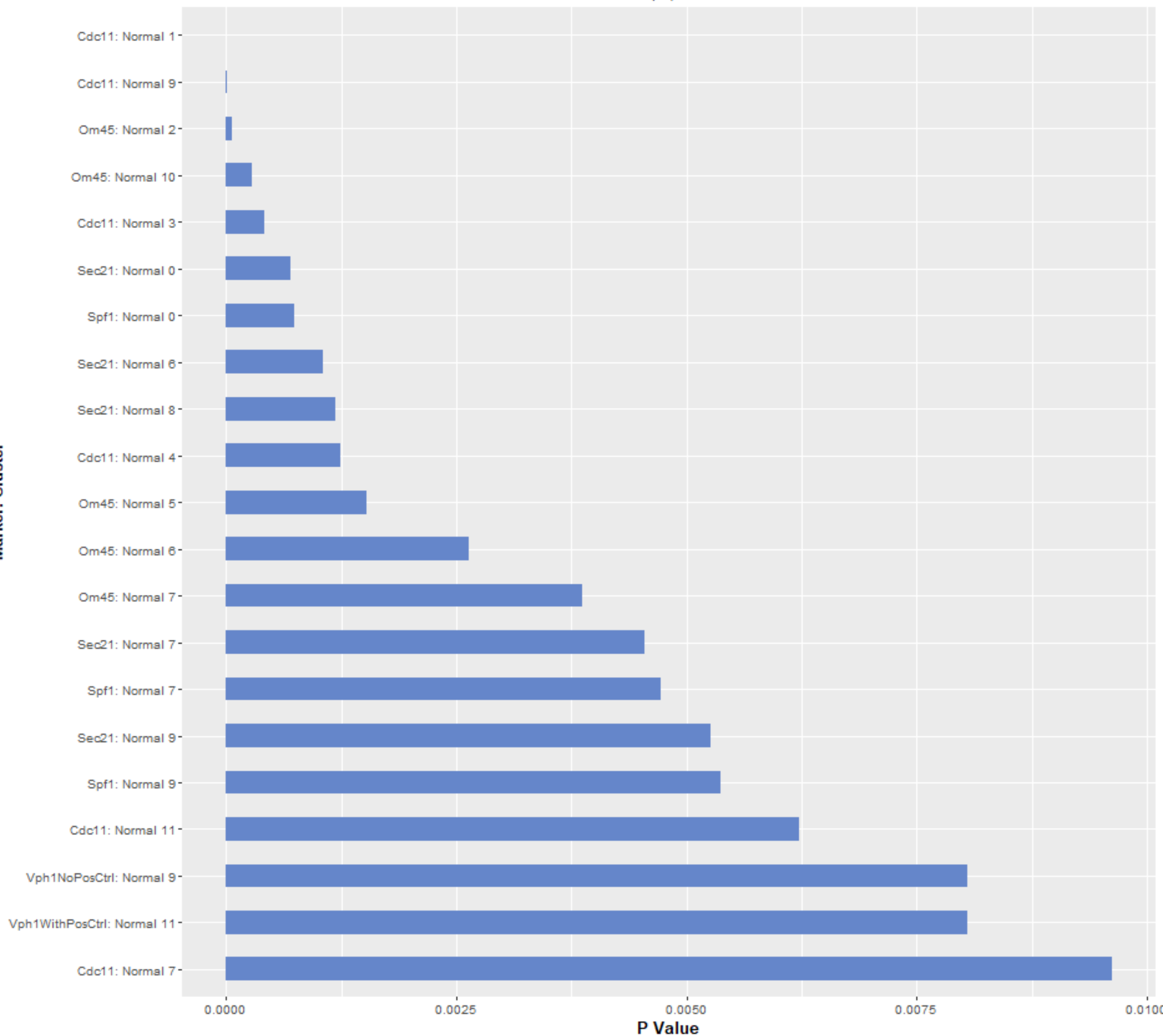
**Golgi vesicle transport**  
Without AreaShape | Ordered Gene Set

Marker: Cluster



histone modification  
Without AreaShape | Ordered Gene Set

Marker: Cluster



# invasive growth in response to glucose limitation

Without AreaShape | Ordered Gene Set

Marker: Cluster

Om45: Normal 0

Cdc11: Normal 9

Cdc11: Normal 5

Spf1: Normal 2

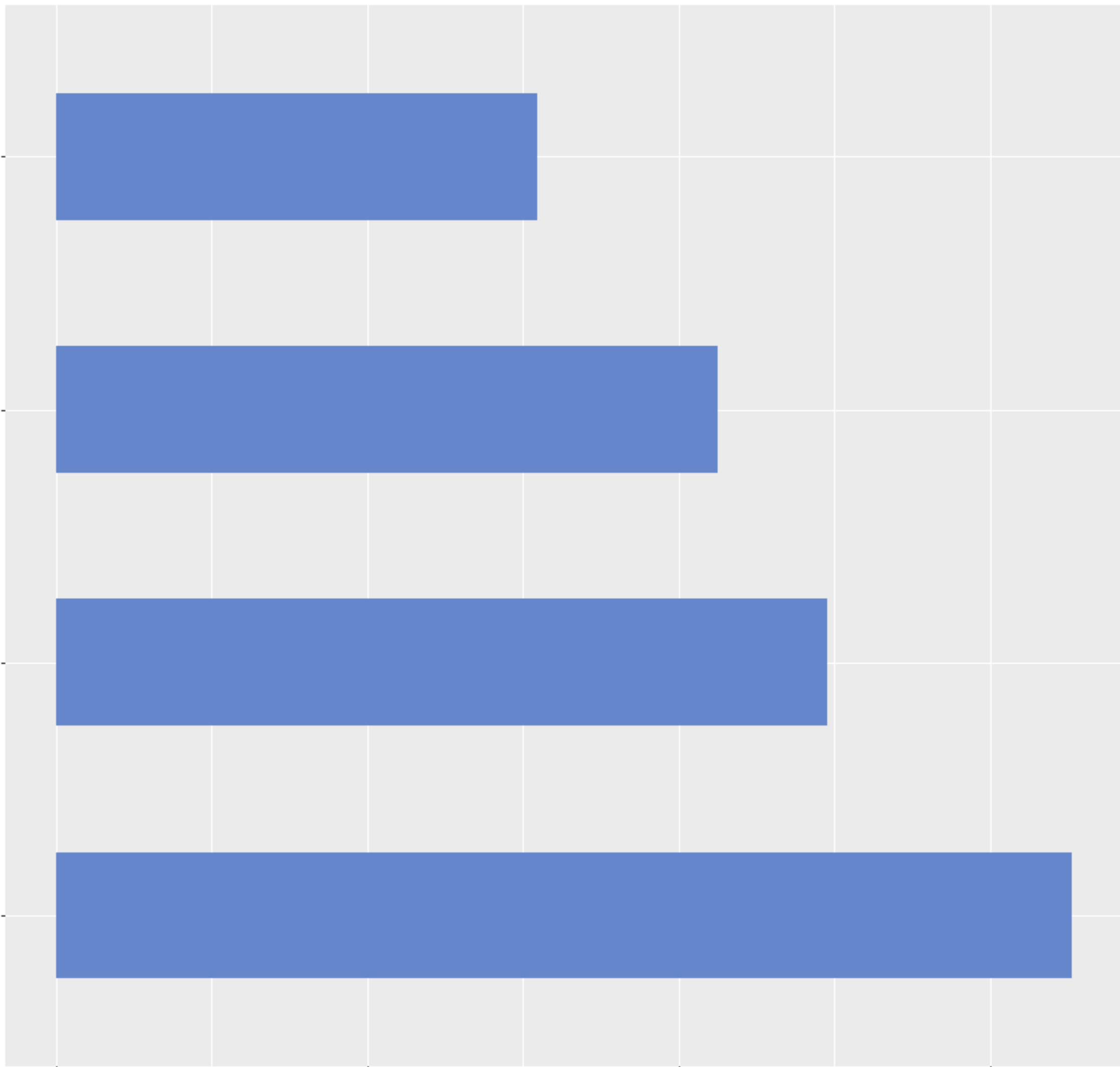
0.000

0.002

0.004

0.006

P Value

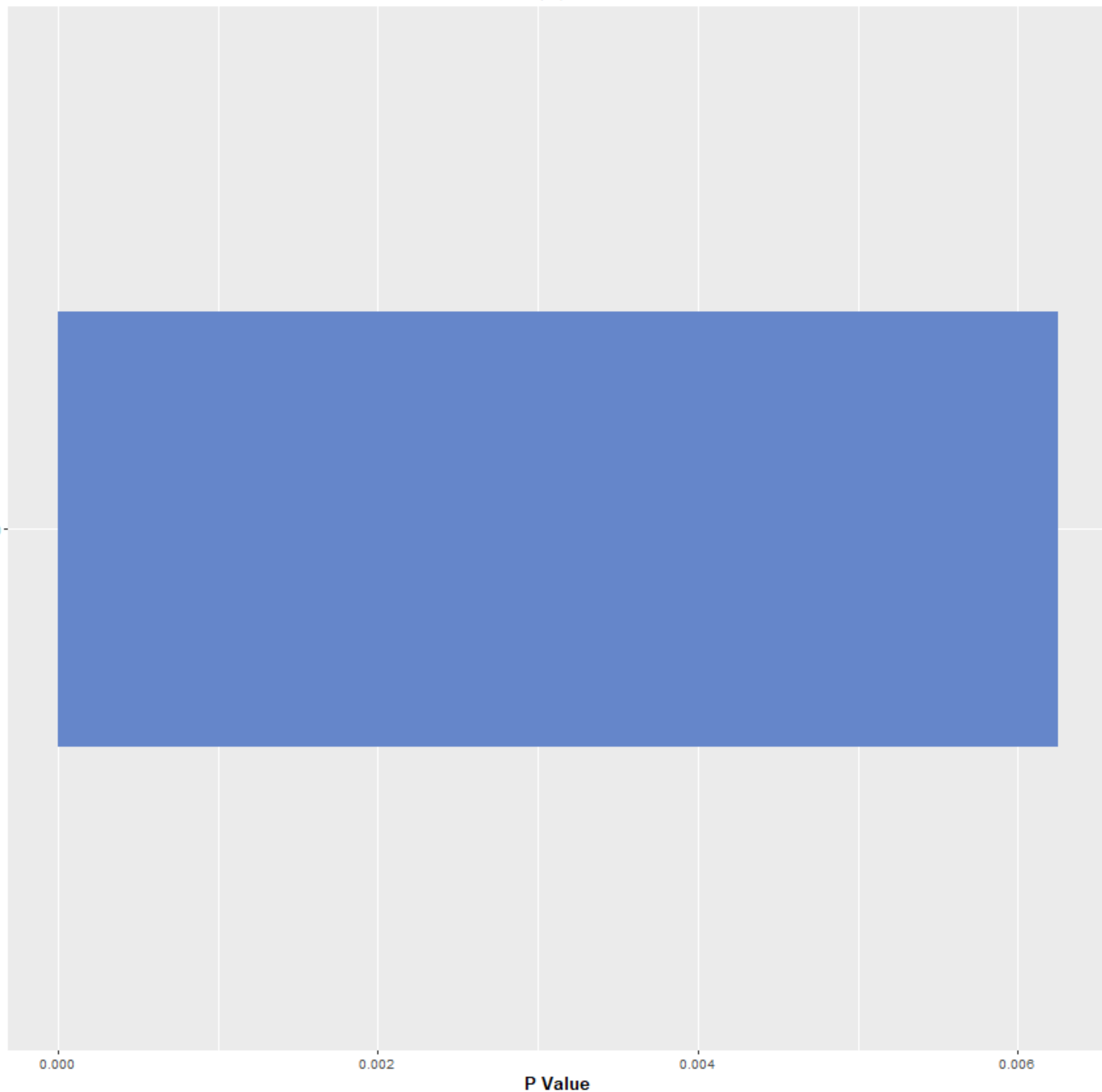


# ion transport

Without AreaShape | Ordered Gene Set

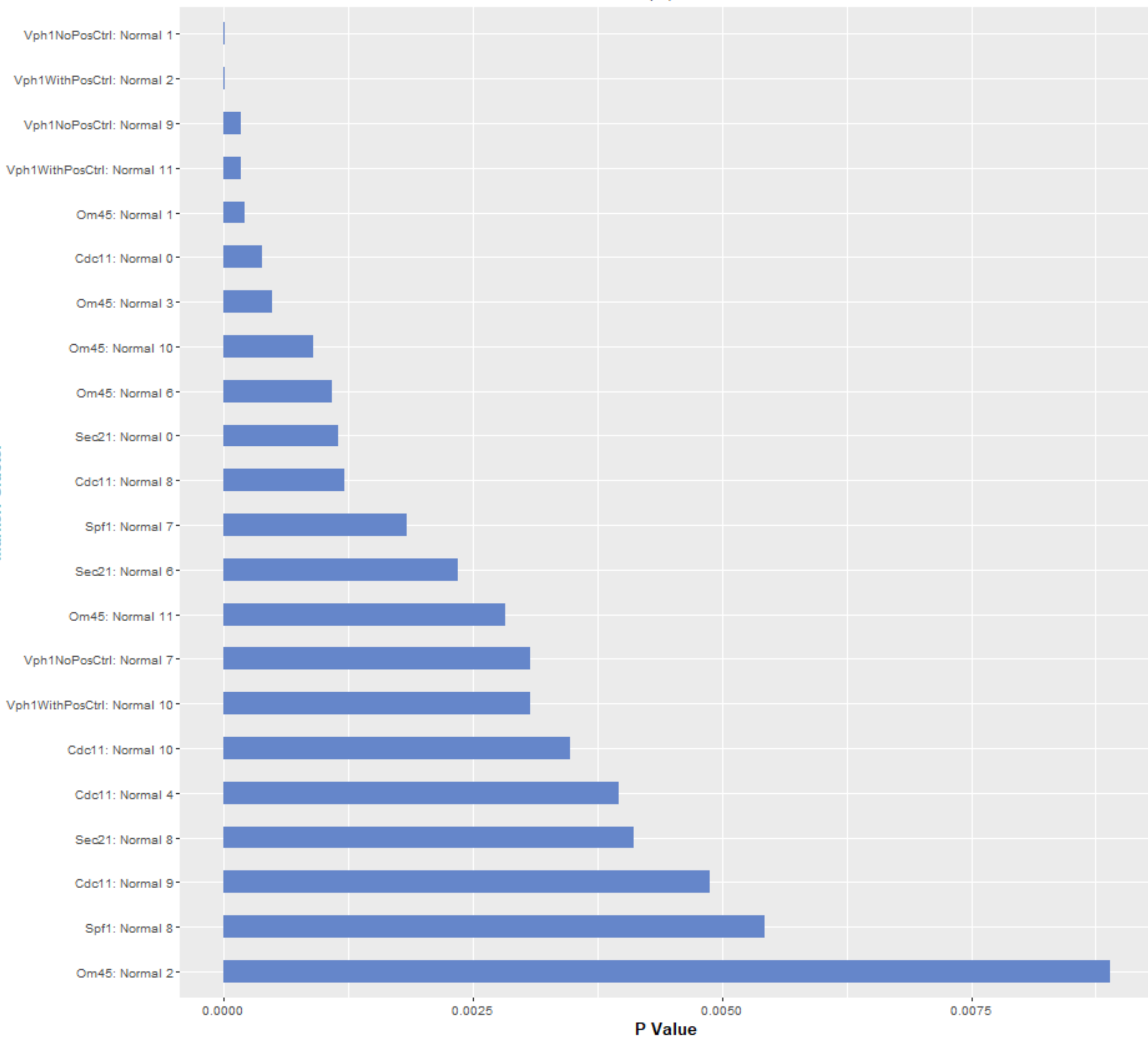
Marker: Cluster

Om45: Normal 0



**lipid metabolic process**  
Without AreaShape | Ordered Gene Set

Marker: Cluster





# meiotic cell cycle

Without AreaShape | Ordered Gene Set

Marker: Cluster

Cdc11: Normal 1

Cdc11: Normal 7

Sec21: Normal 7

Cdc11: Normal 11

Om45: Normal 5

0.0000

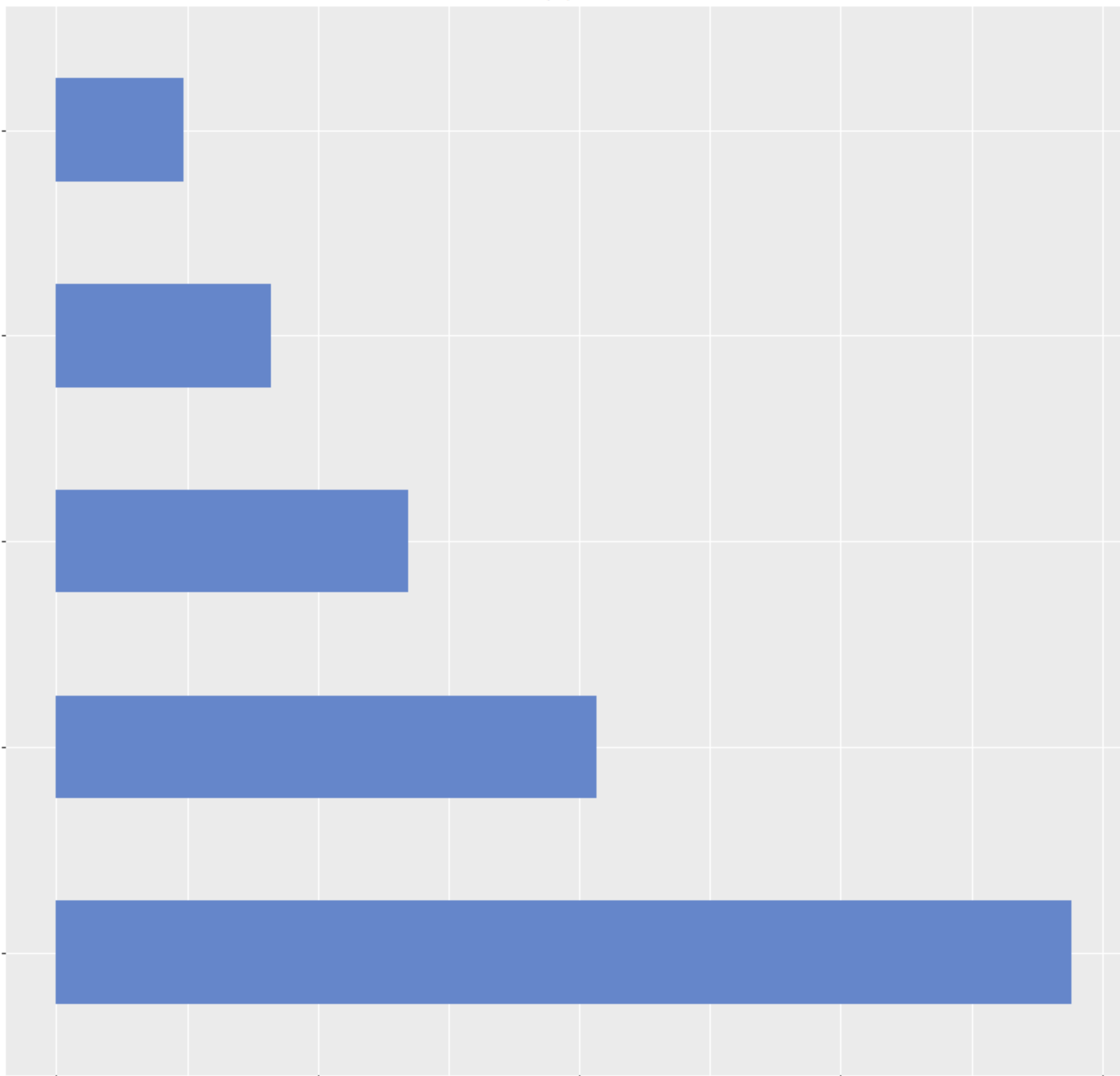
0.0025

0.0050

0.0075

0.0100

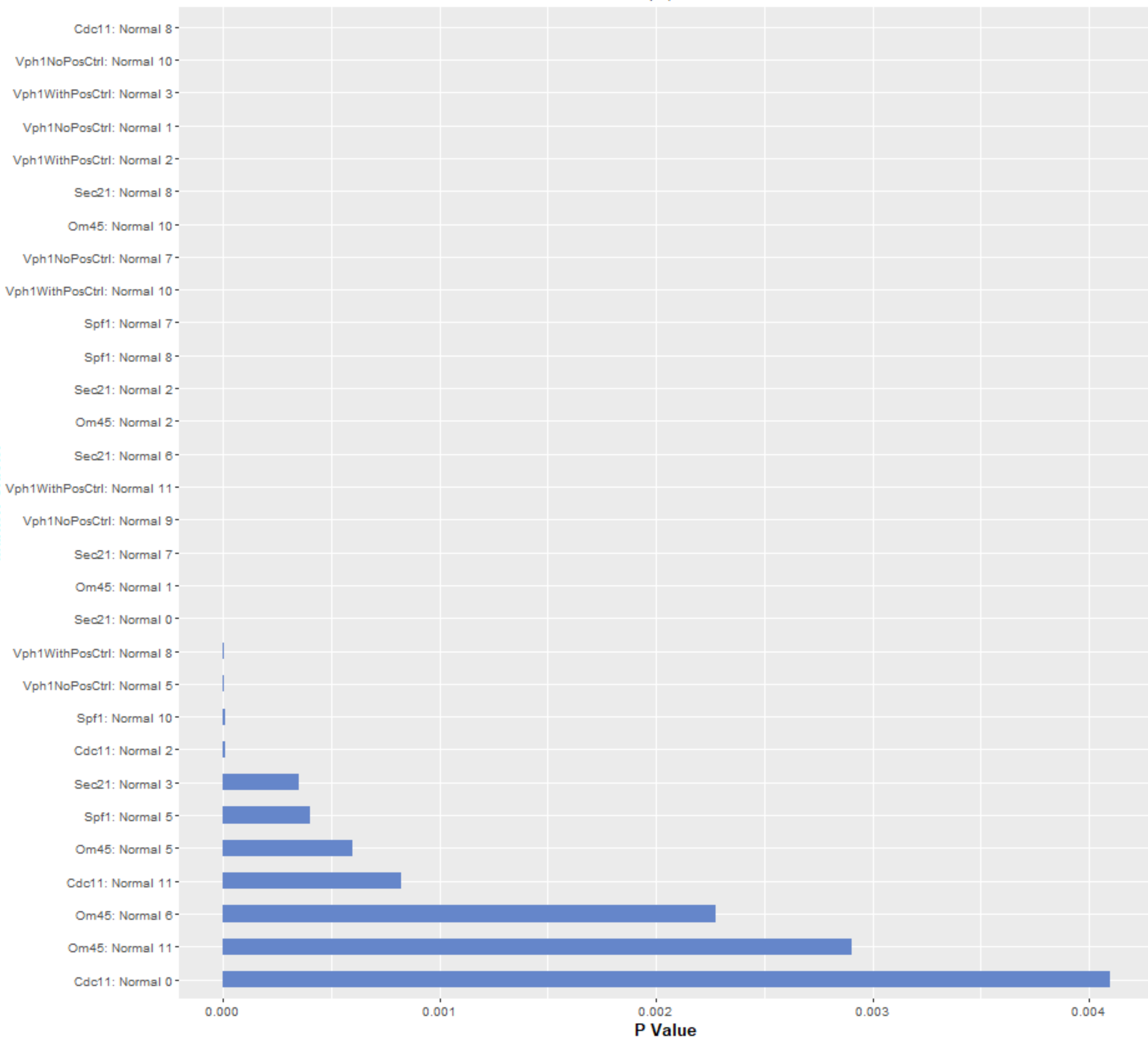
P Value



# membrane fusion

Without AreaShape | Ordered Gene Set

Marker: Cluster



# mitochondrion organization

Without AreaShape | Ordered Gene Set

Marker: Cluster

Om45: Normal 0

0.000

0.002

0.004

0.006

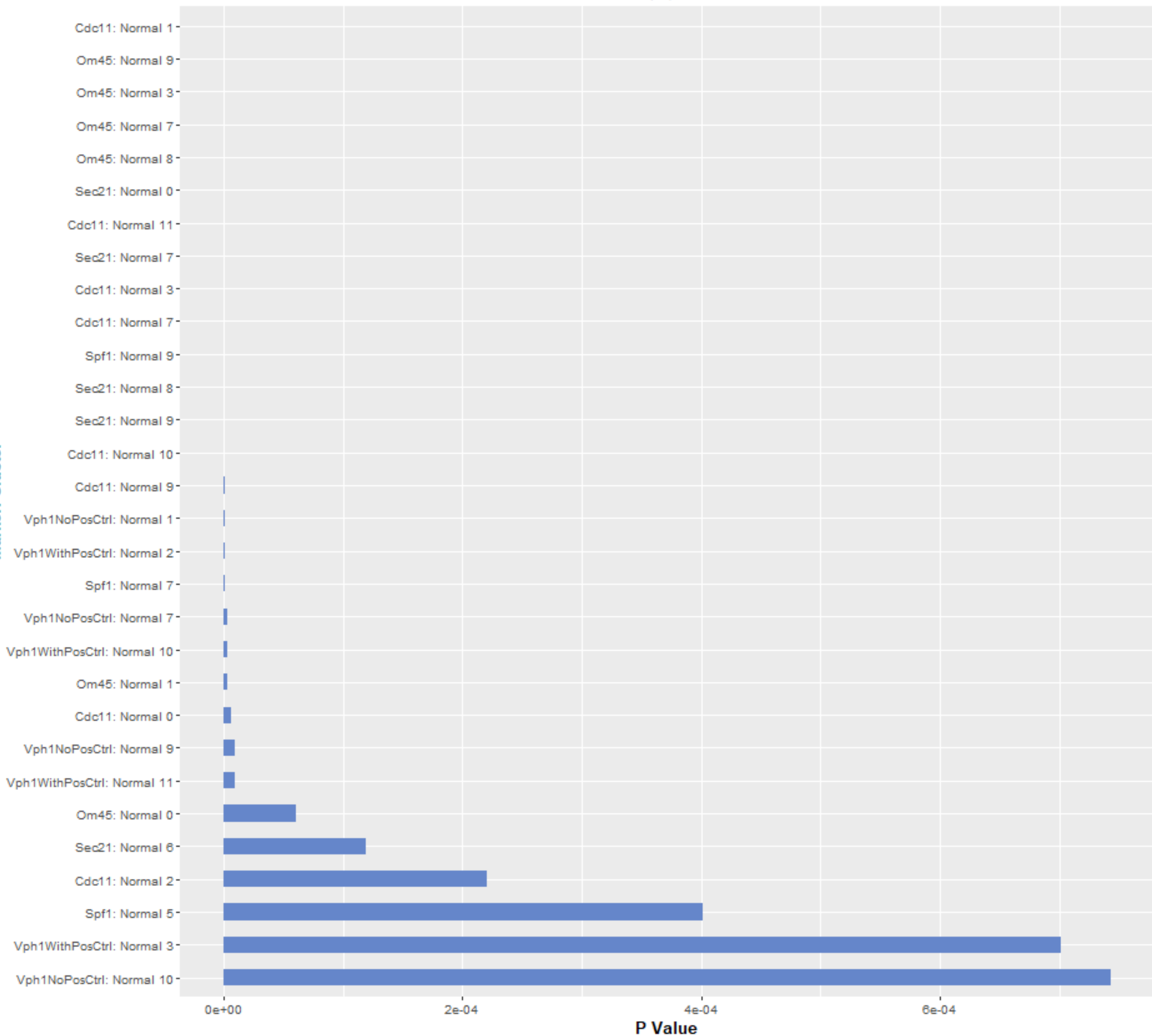
0.008

P Value

# mitotic cell cycle

Without AreaShape | Ordered Gene Set

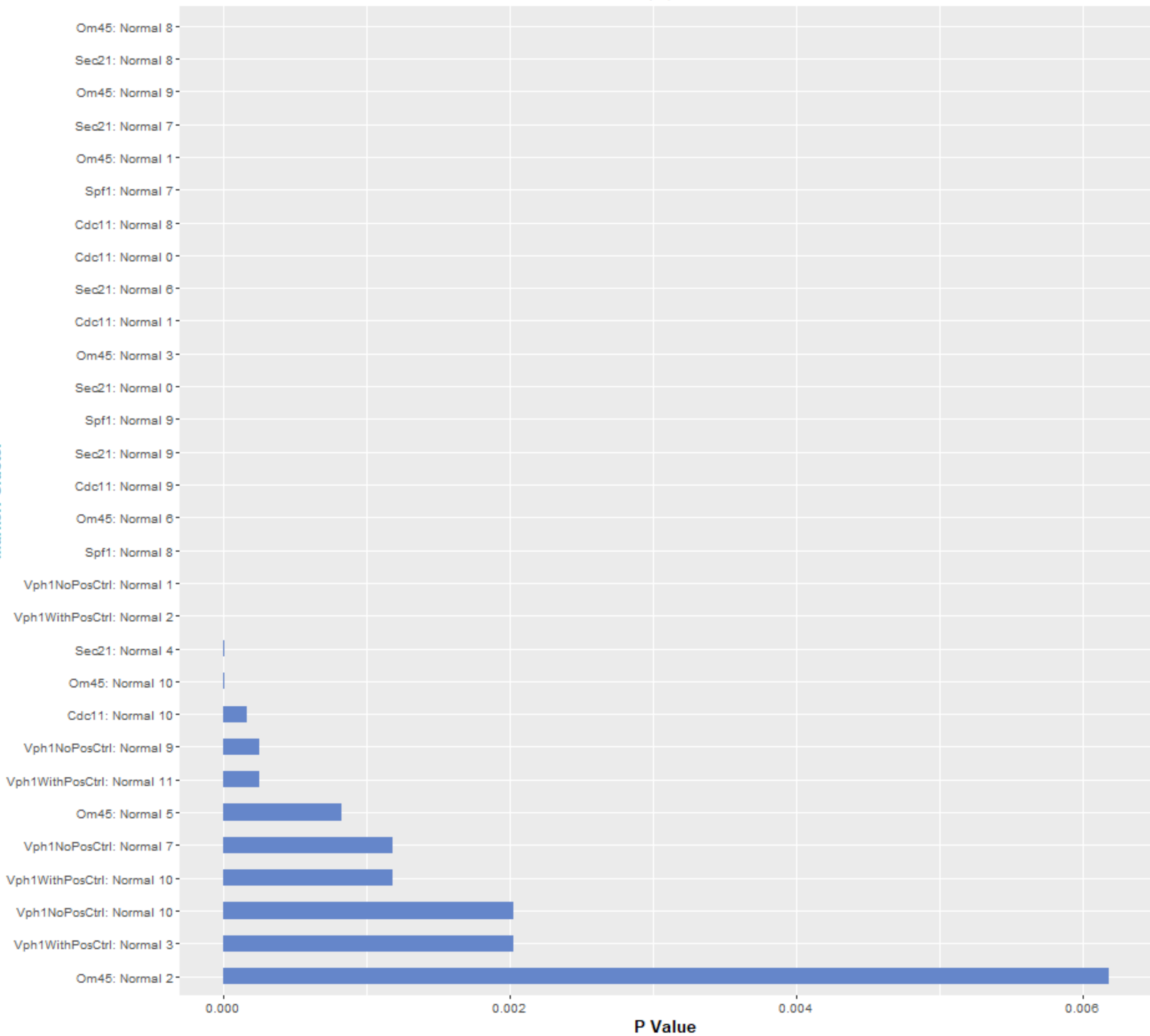
Marker: Cluster



# mRNA processing

Without AreaShape | Ordered Gene Set

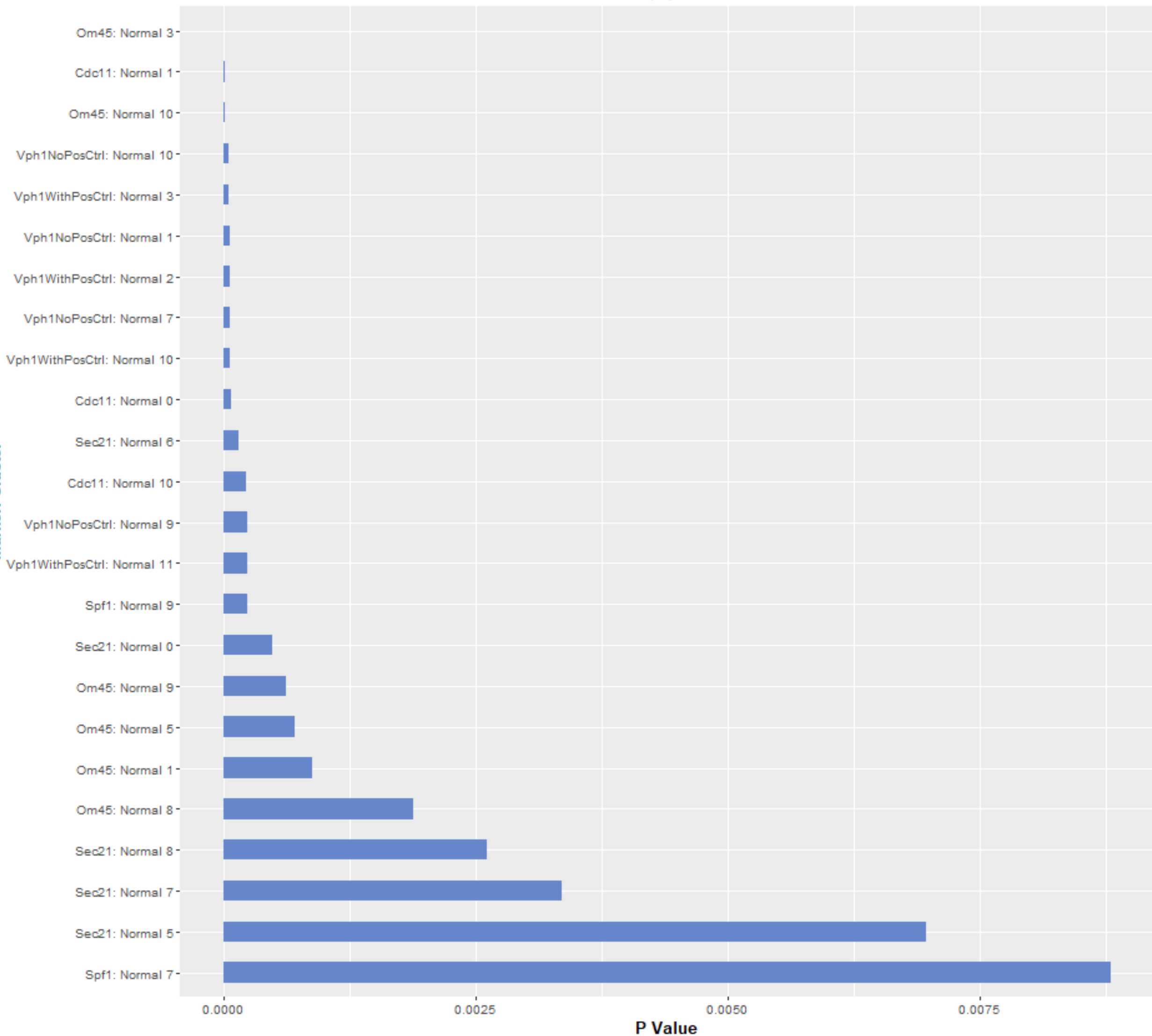
Marker: Cluster



# nuclear transport

Without AreaShape | Ordered Gene Set

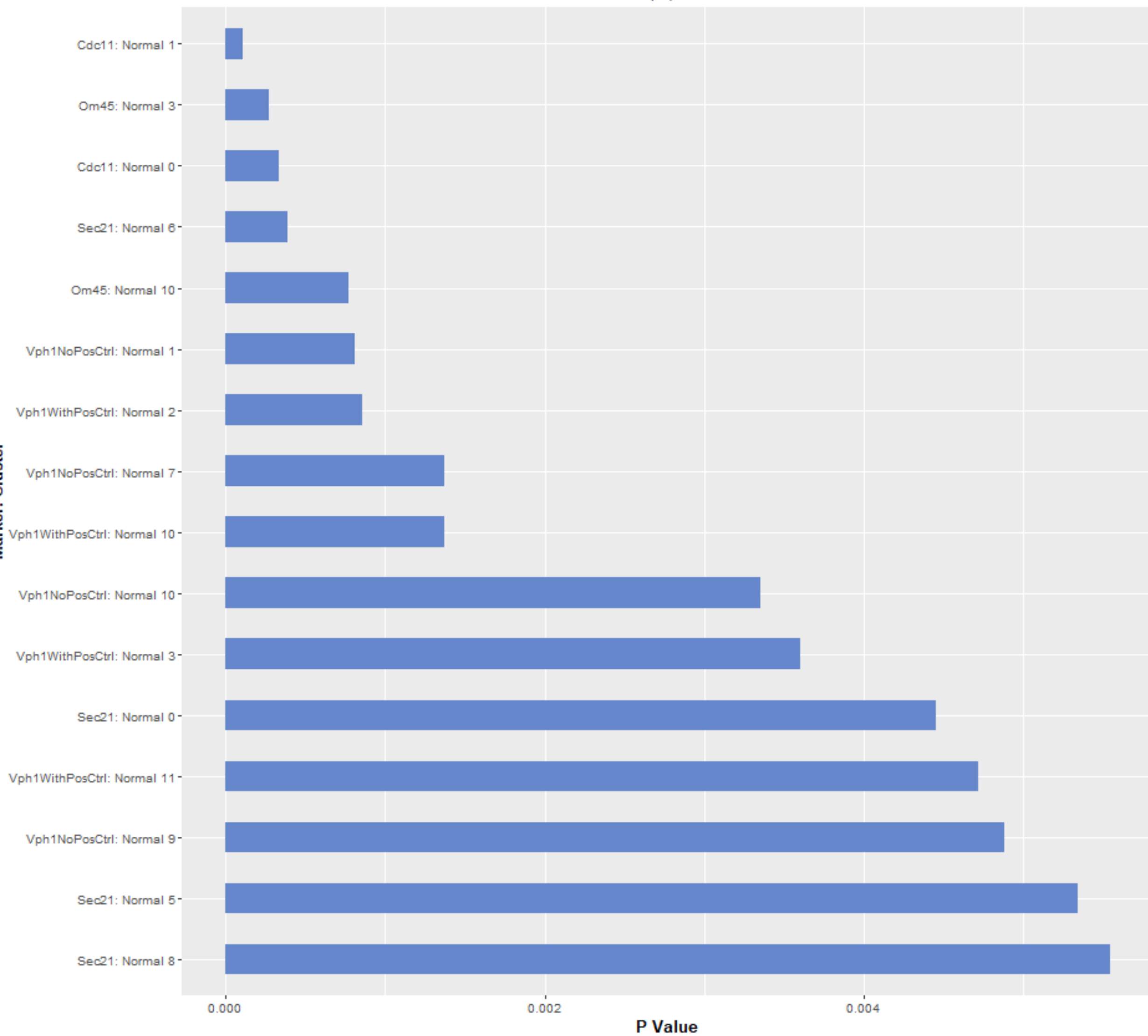
Marker: Cluster



# nucleobase-containing compound transport

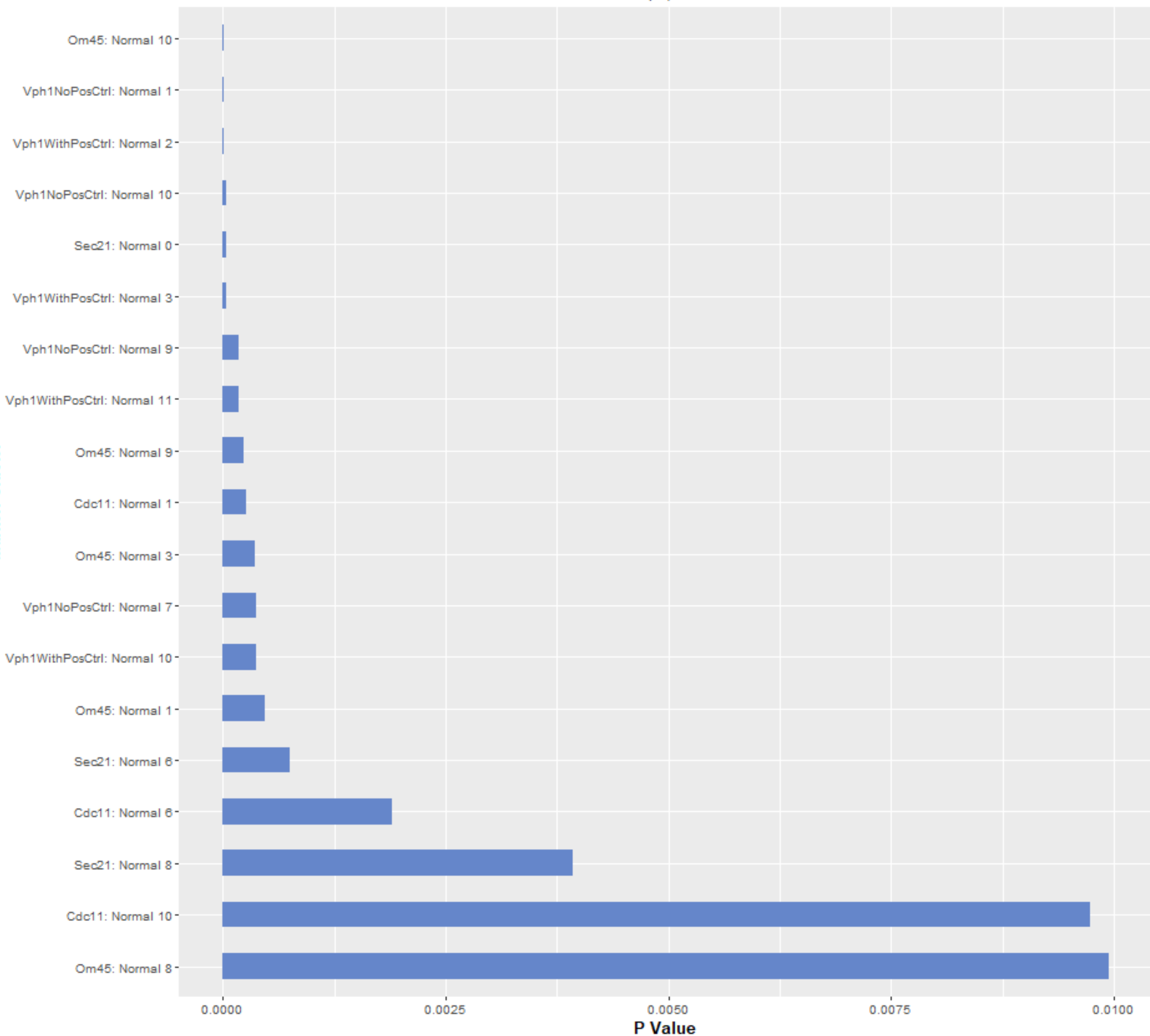
Without AreaShape | Ordered Gene Set

Marker: Cluster



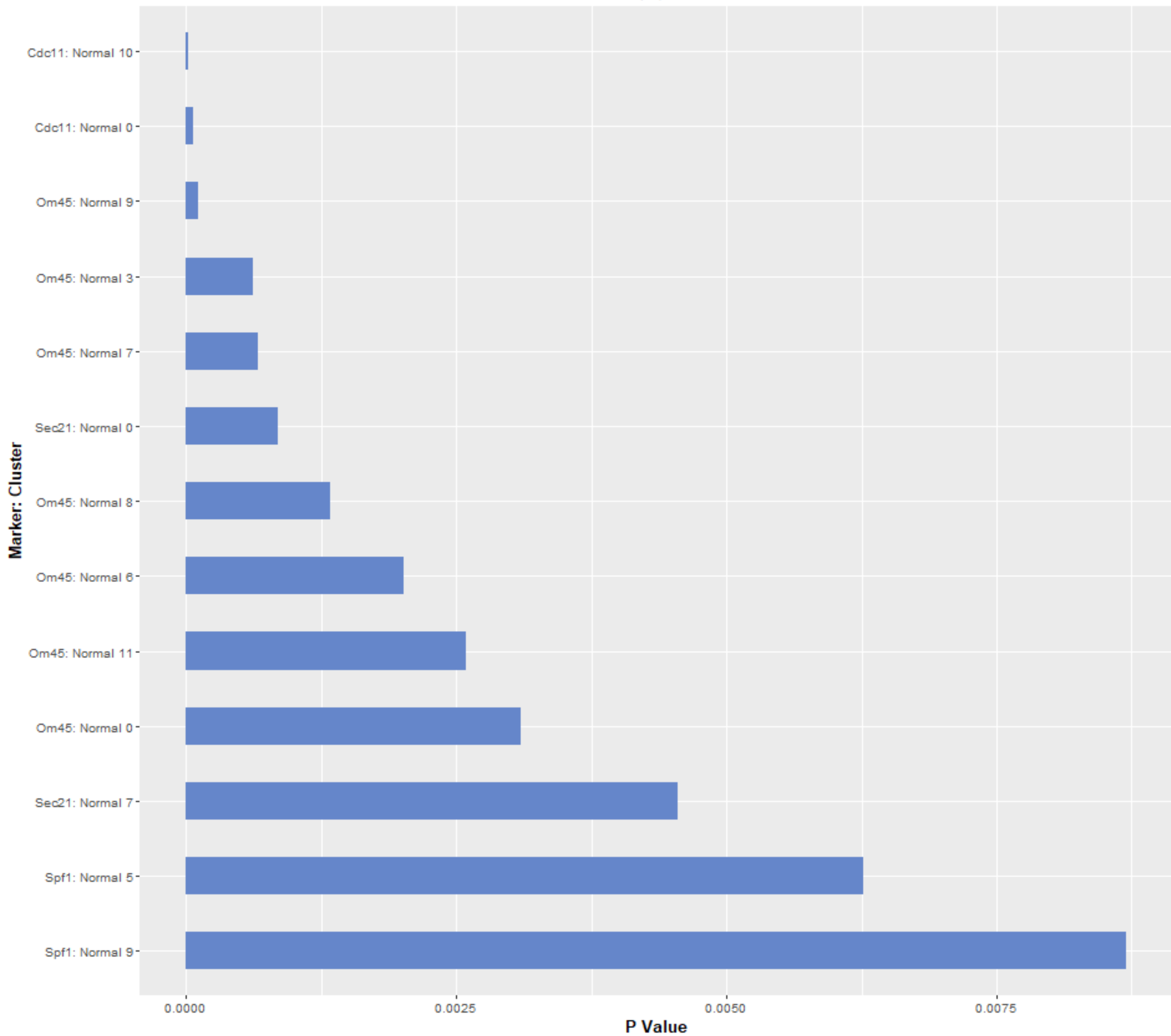
**nucleus organization**  
Without AreaShape | Ordered Gene Set

**Marker: Cluster**





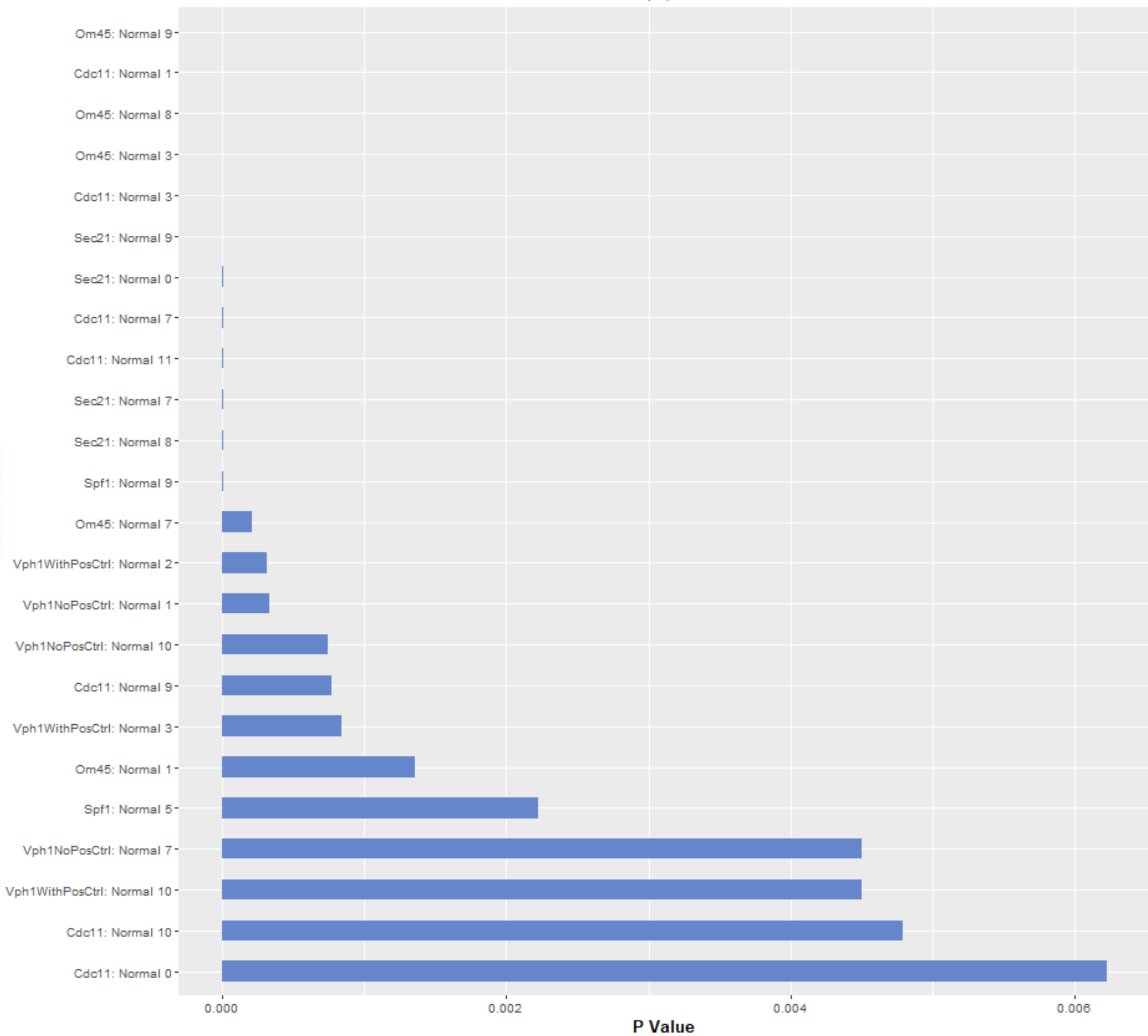
**organelle assembly**  
Without AreaShape | Ordered Gene Set



# organelle fission

Without AreaShape | Ordered Gene Set

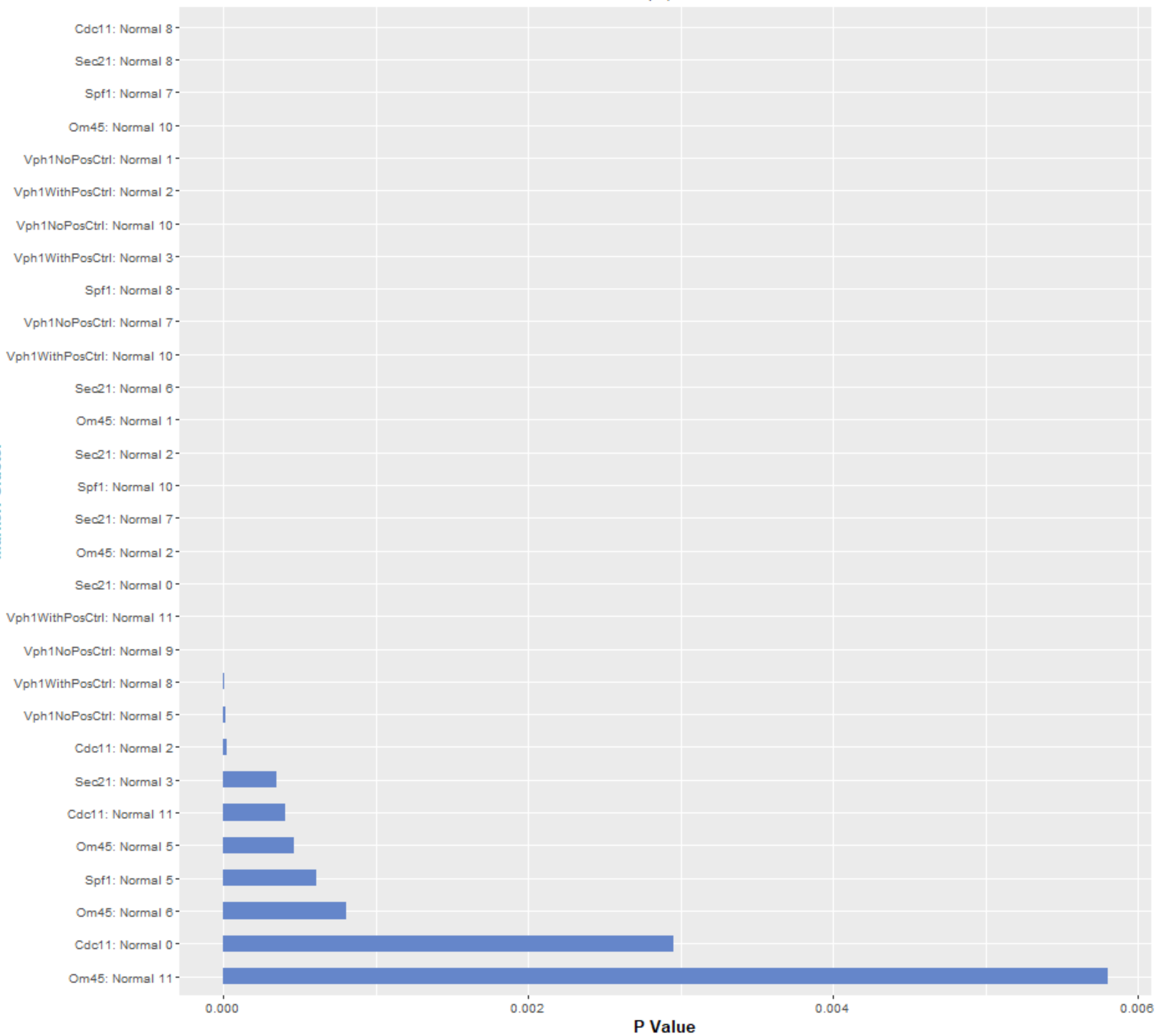
Marker: Cluster



# organelle fusion

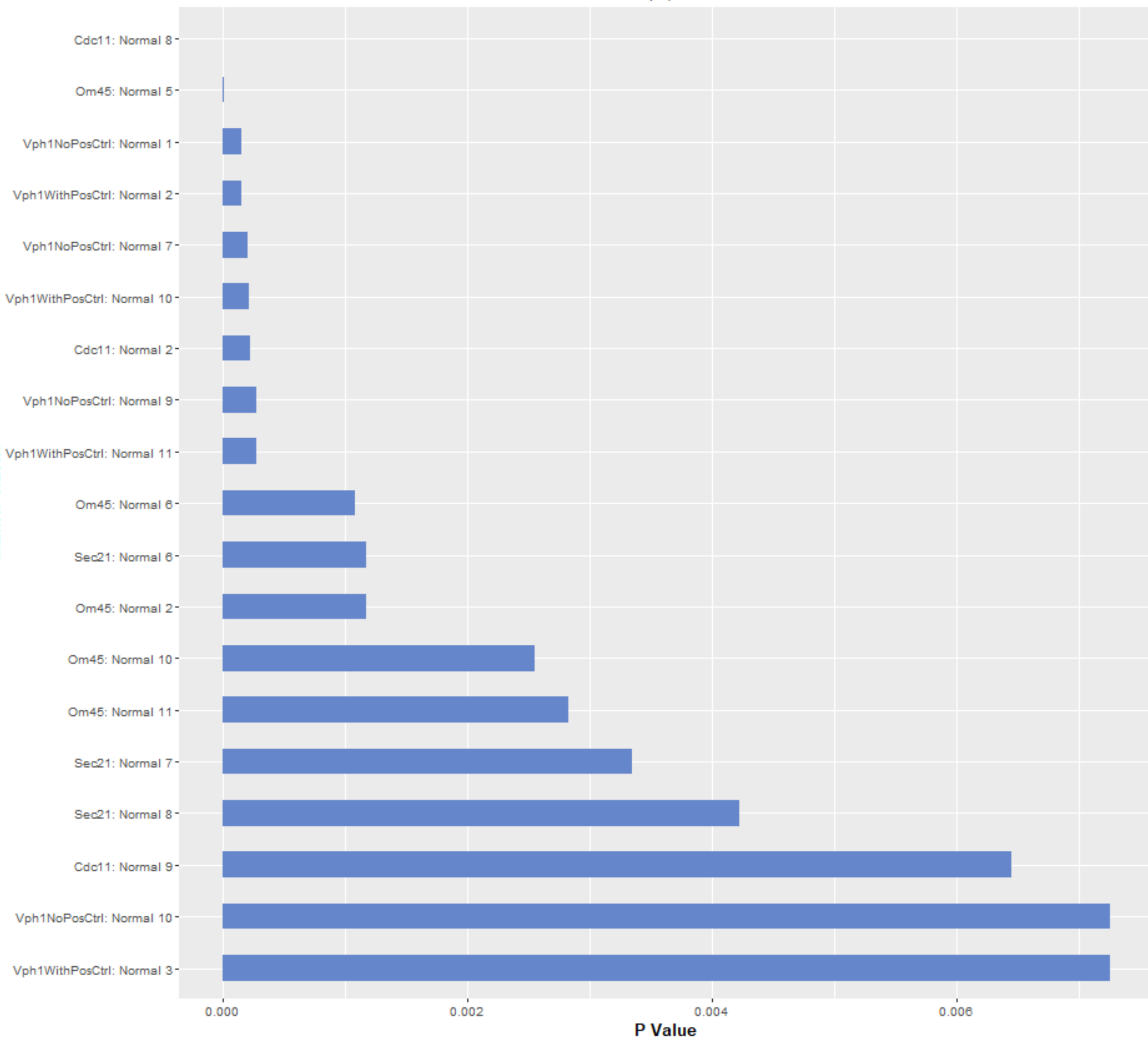
Without AreaShape | Ordered Gene Set

Marker: Cluster



**organelle inheritance**  
Without AreaShape | Ordered Gene Set

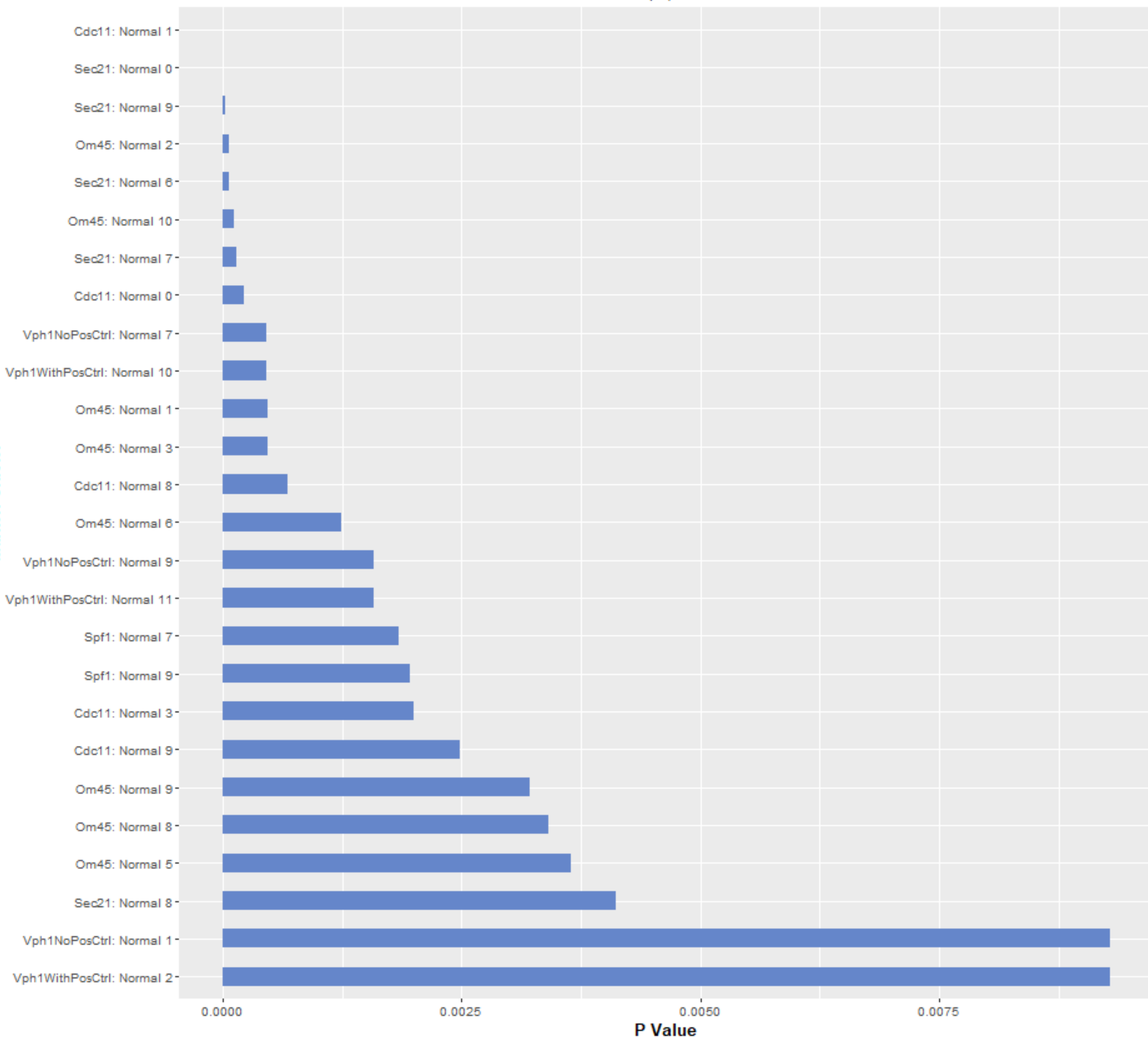
**Marker: Cluster**



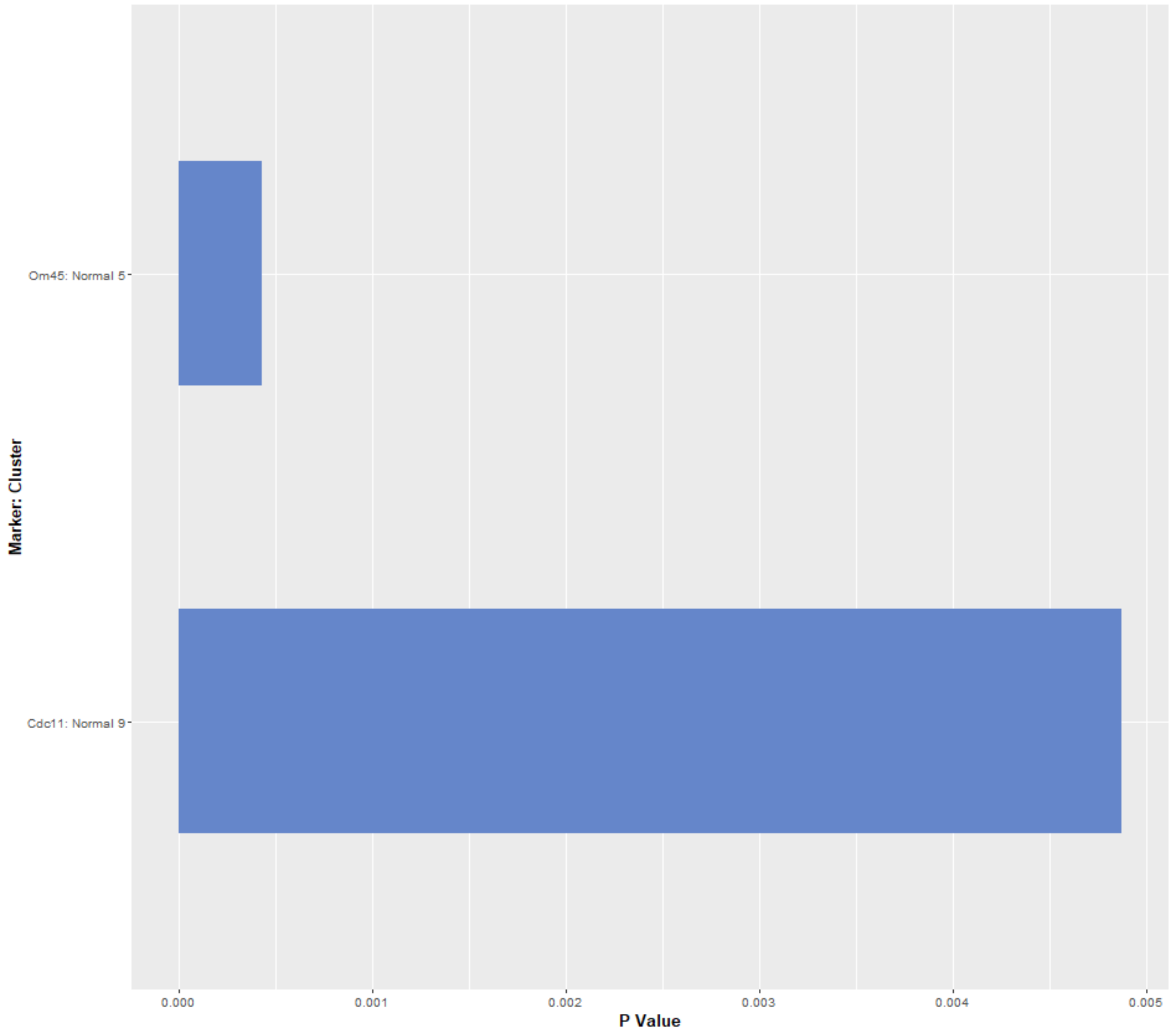
# peptidyl-amino acid modification

Without AreaShape | Ordered Gene Set

Marker: Cluster

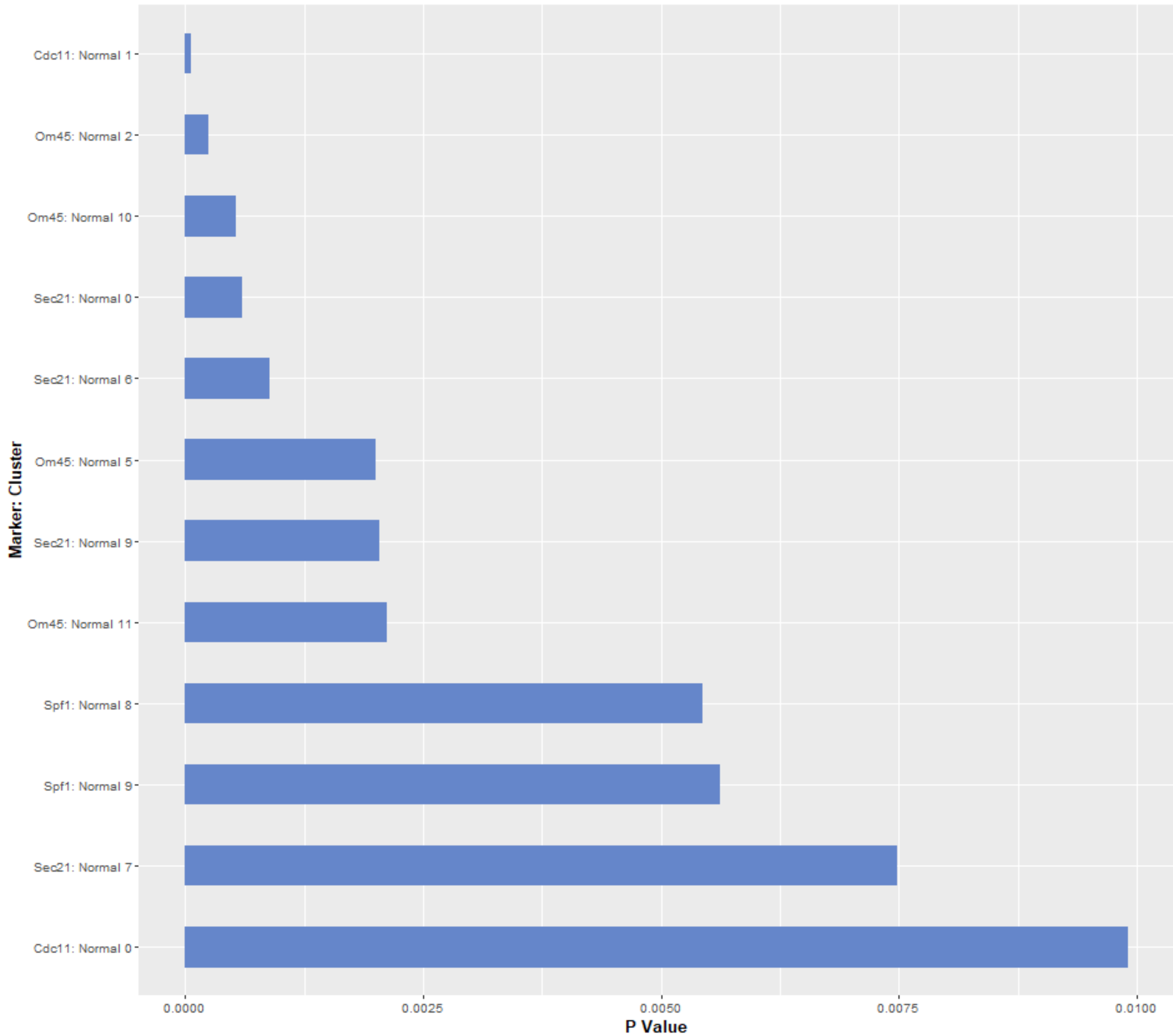


**peroxisome organization**  
Without AreaShape | Ordered Gene Set



# protein acylation

Without AreaShape | Ordered Gene Set



# protein alkylation

Without AreaShape | Ordered Gene Set

Marker: Cluster

Vph1NoPosCtrl: Normal 2

Vph1WithPosCtrl: Normal 6

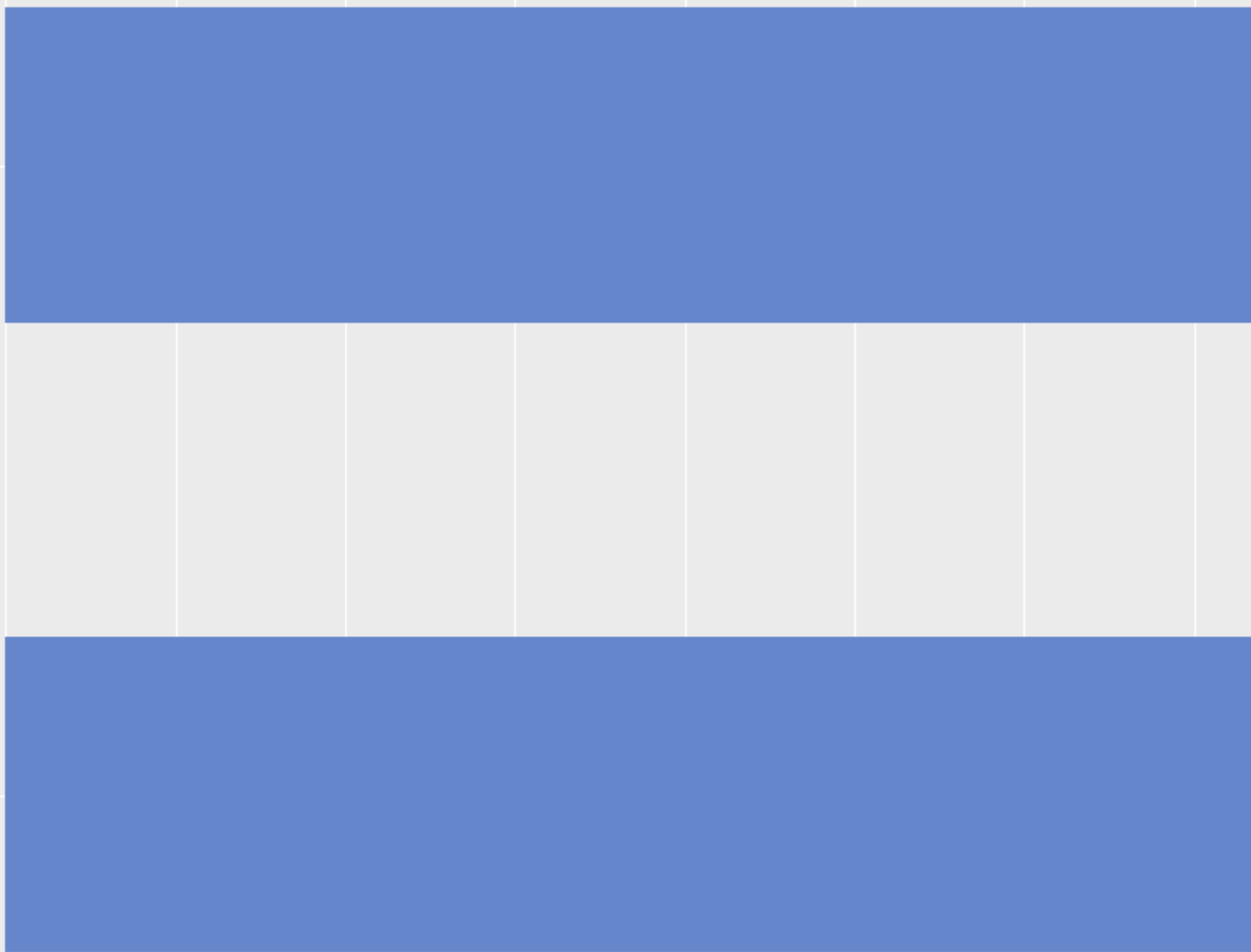
0.000

0.002

0.004

0.006

P Value

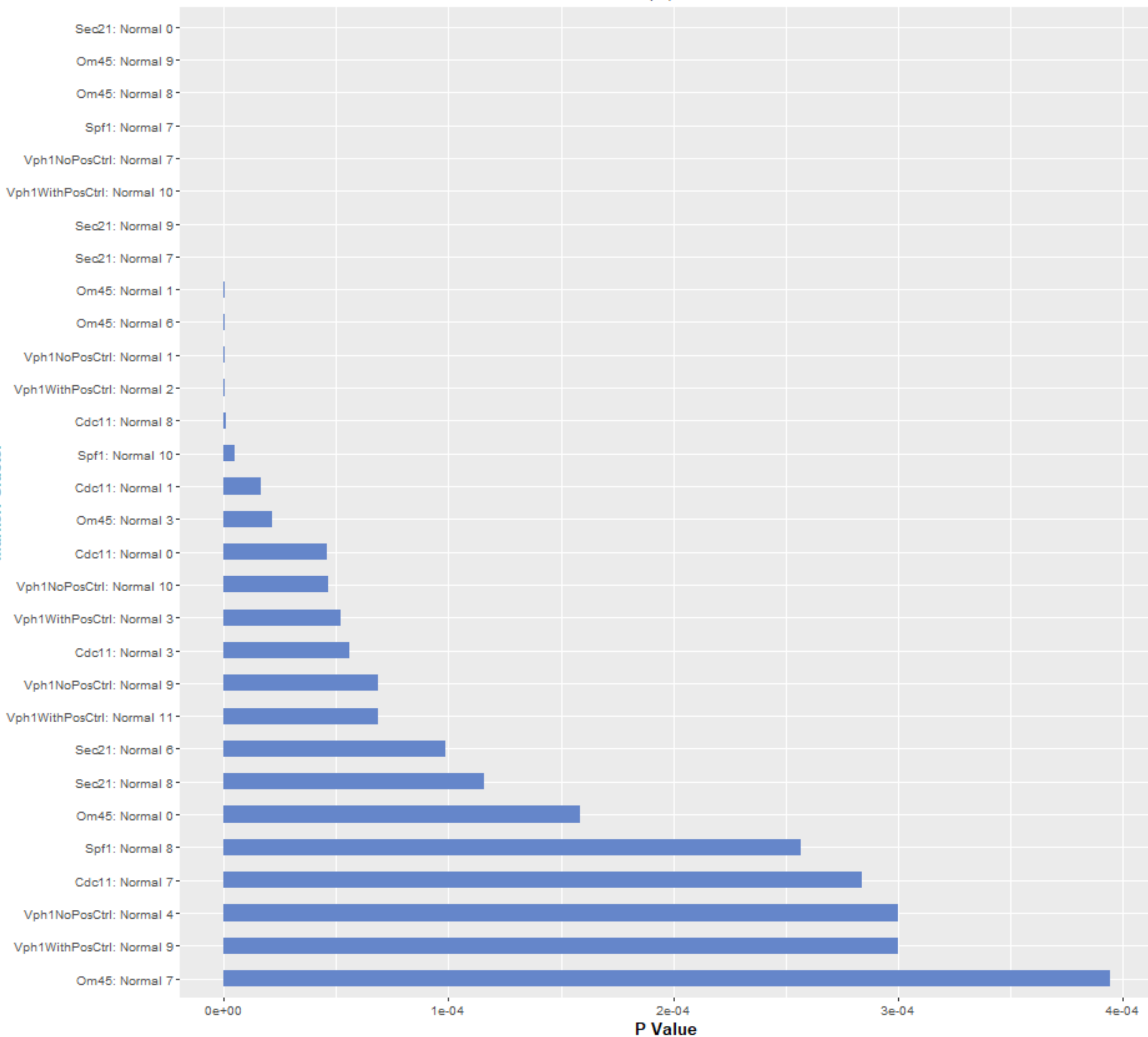




# protein complex biogenesis

Without AreaShape | Ordered Gene Set

Marker: Cluster



# protein dephosphorylation

Without AreaShape | Ordered Gene Set

Marker: Cluster

Cdc11: Normal 11

0e+00

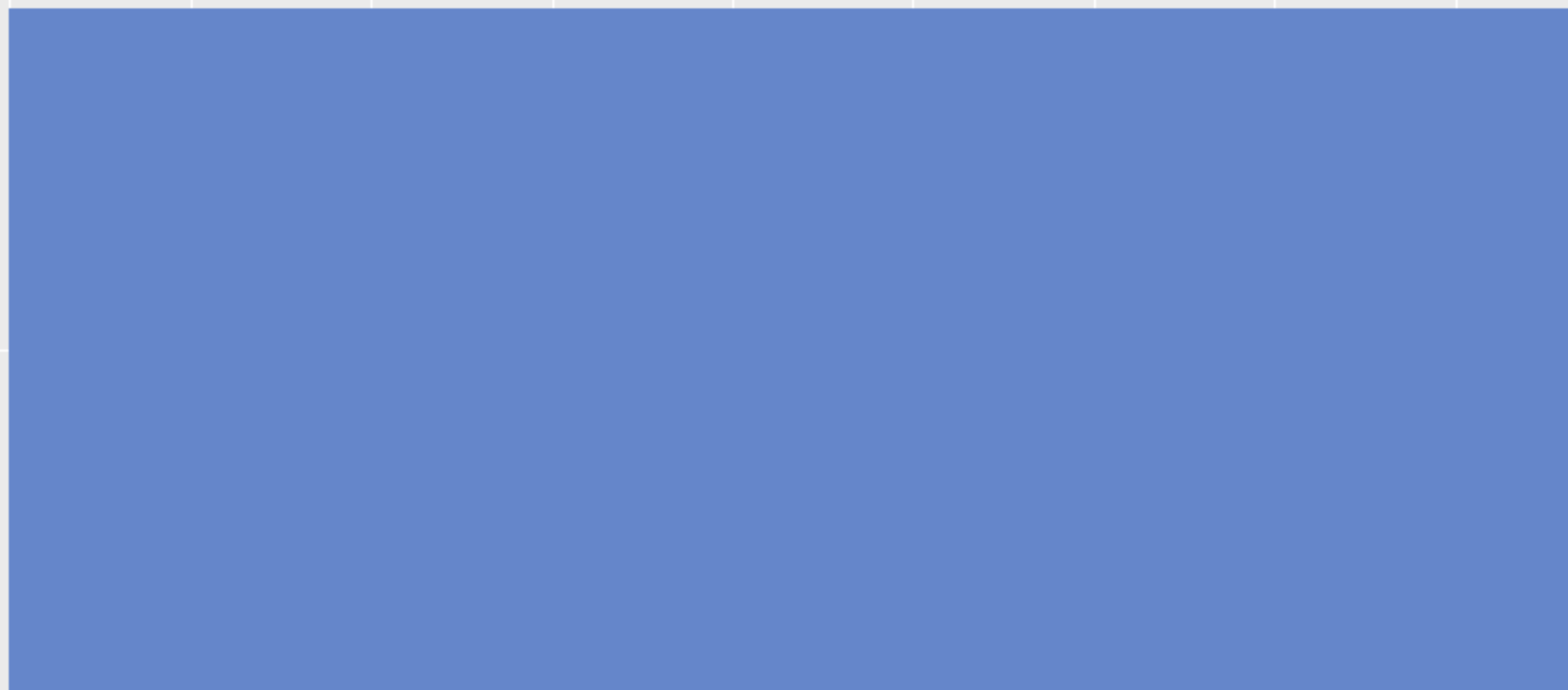
1e-04

2e-04

3e-04

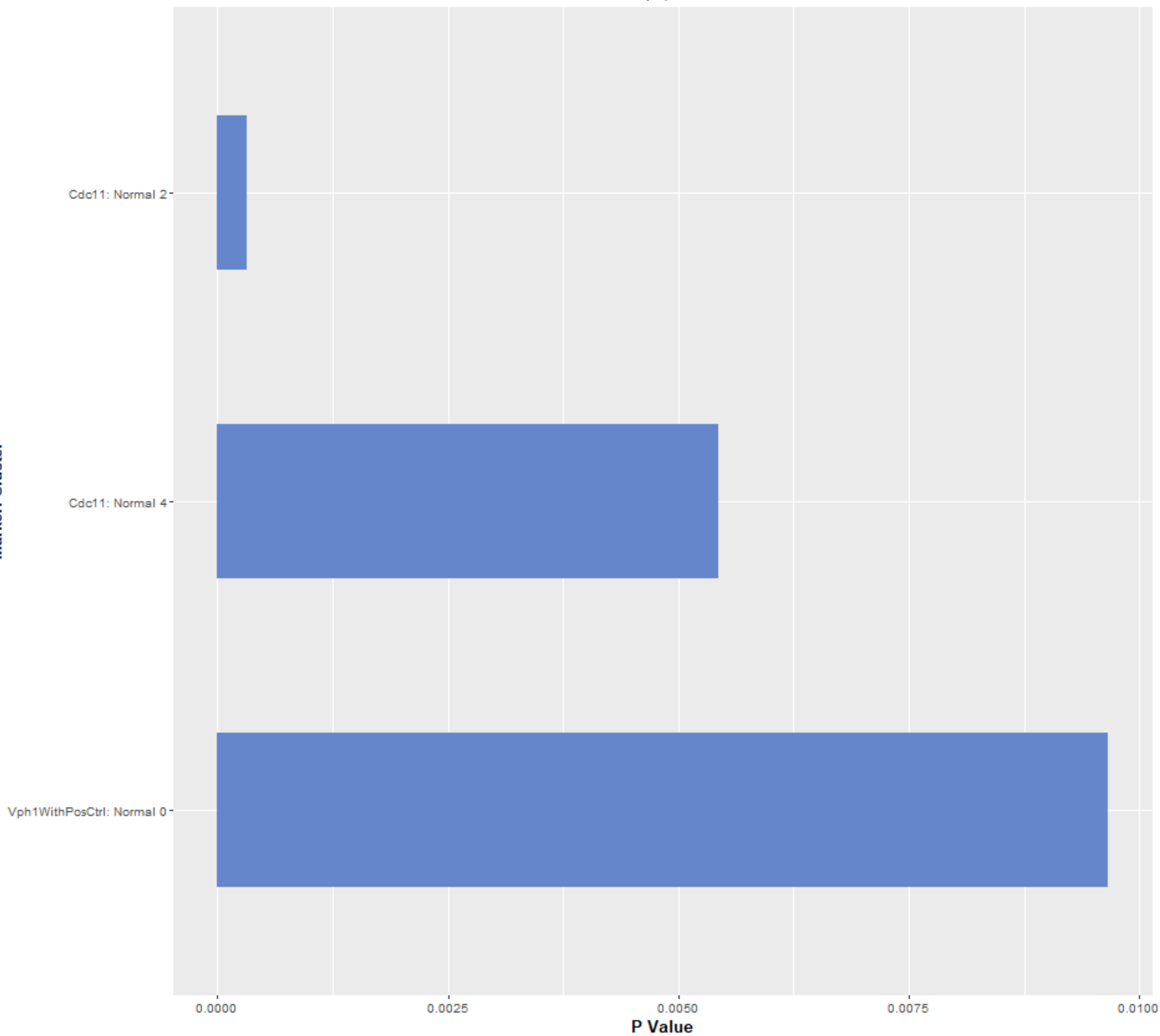
4e-04

P Value



**protein glycosylation**  
Without AreaShape | Ordered Gene Set

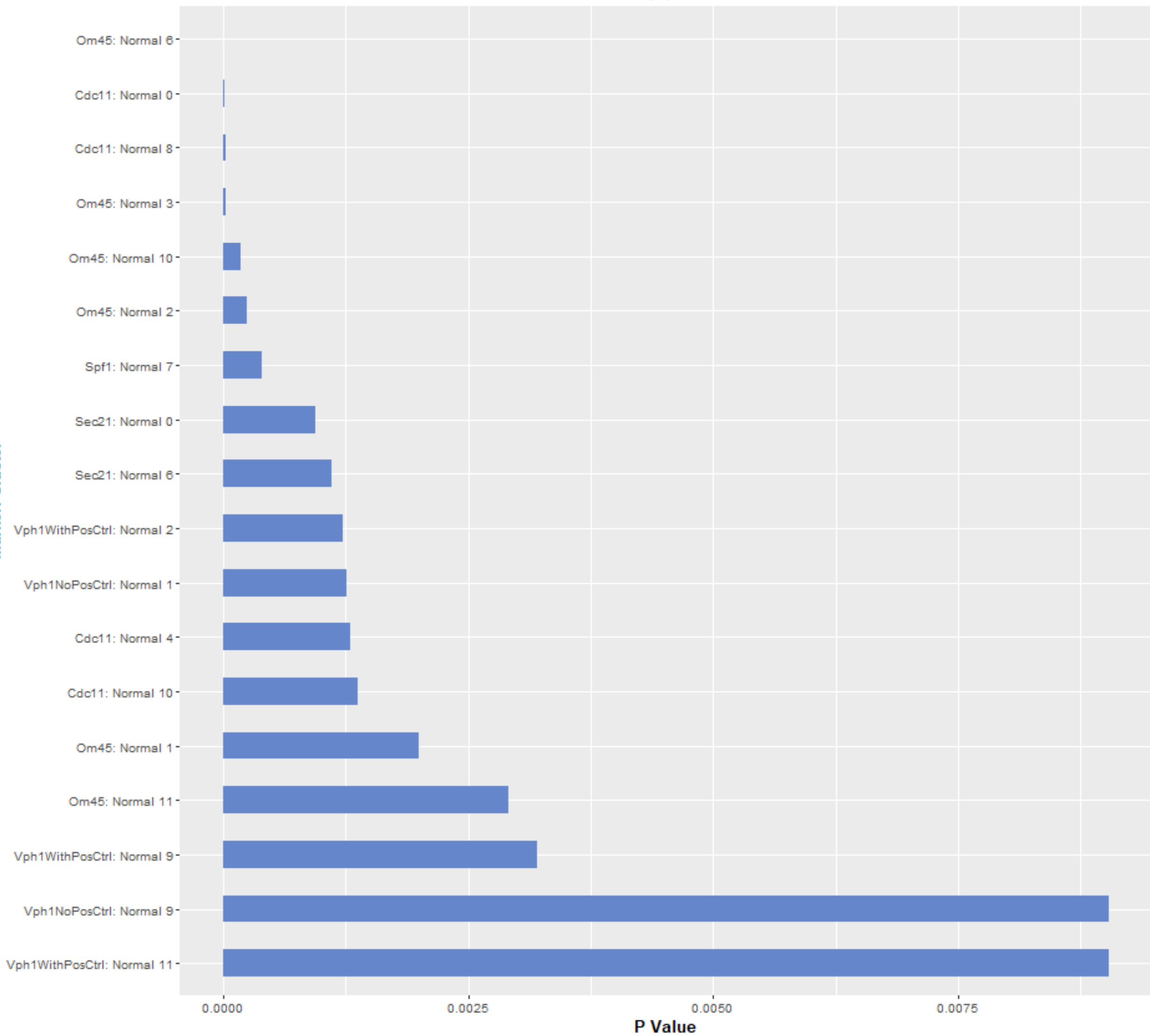
Marker: Cluster



# protein lipidation

Without AreaShape | Ordered Gene Set

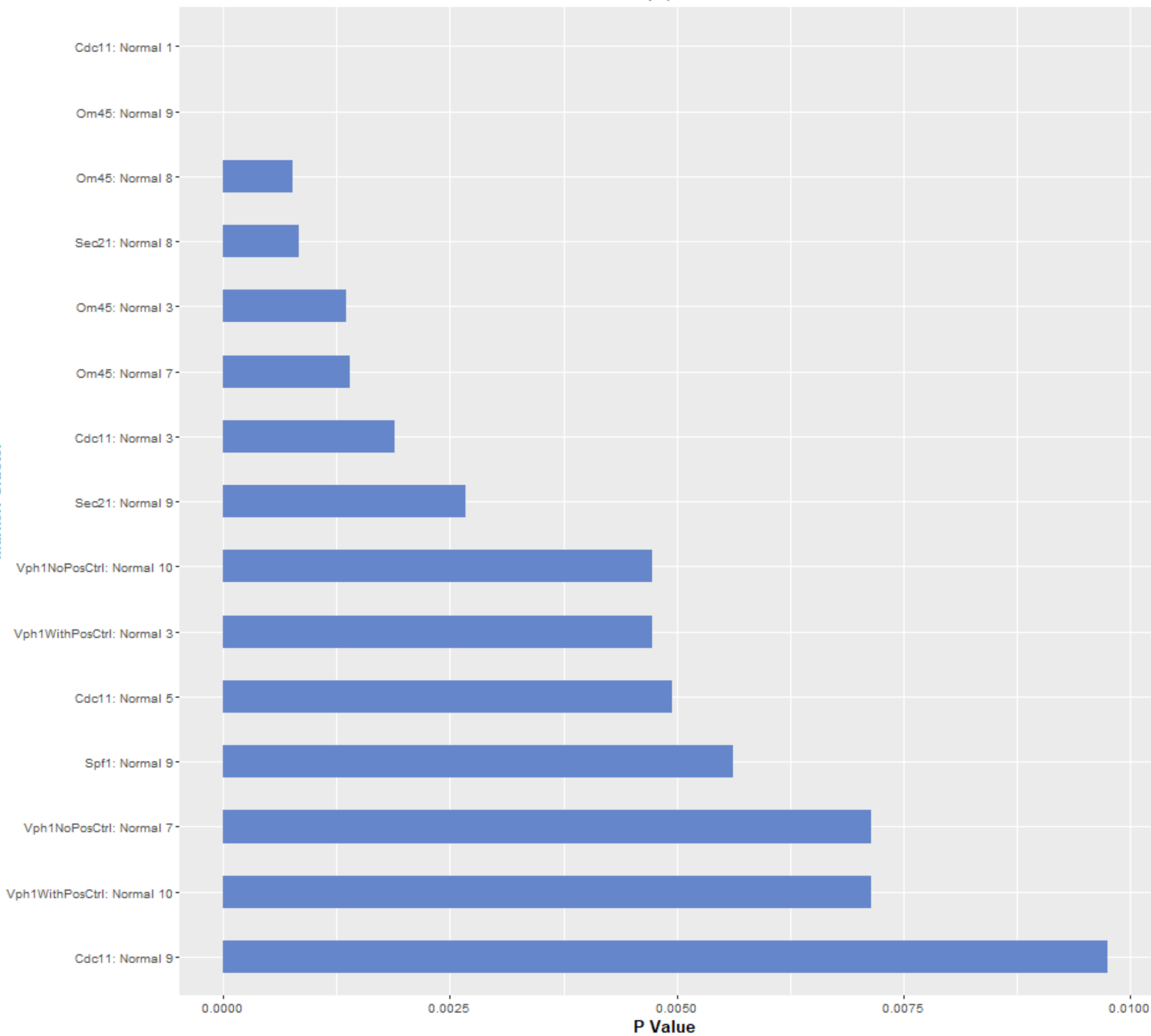
Marker: Cluster



# protein modification by small protein conjugation or removal

Without AreaShape | Ordered Gene Set

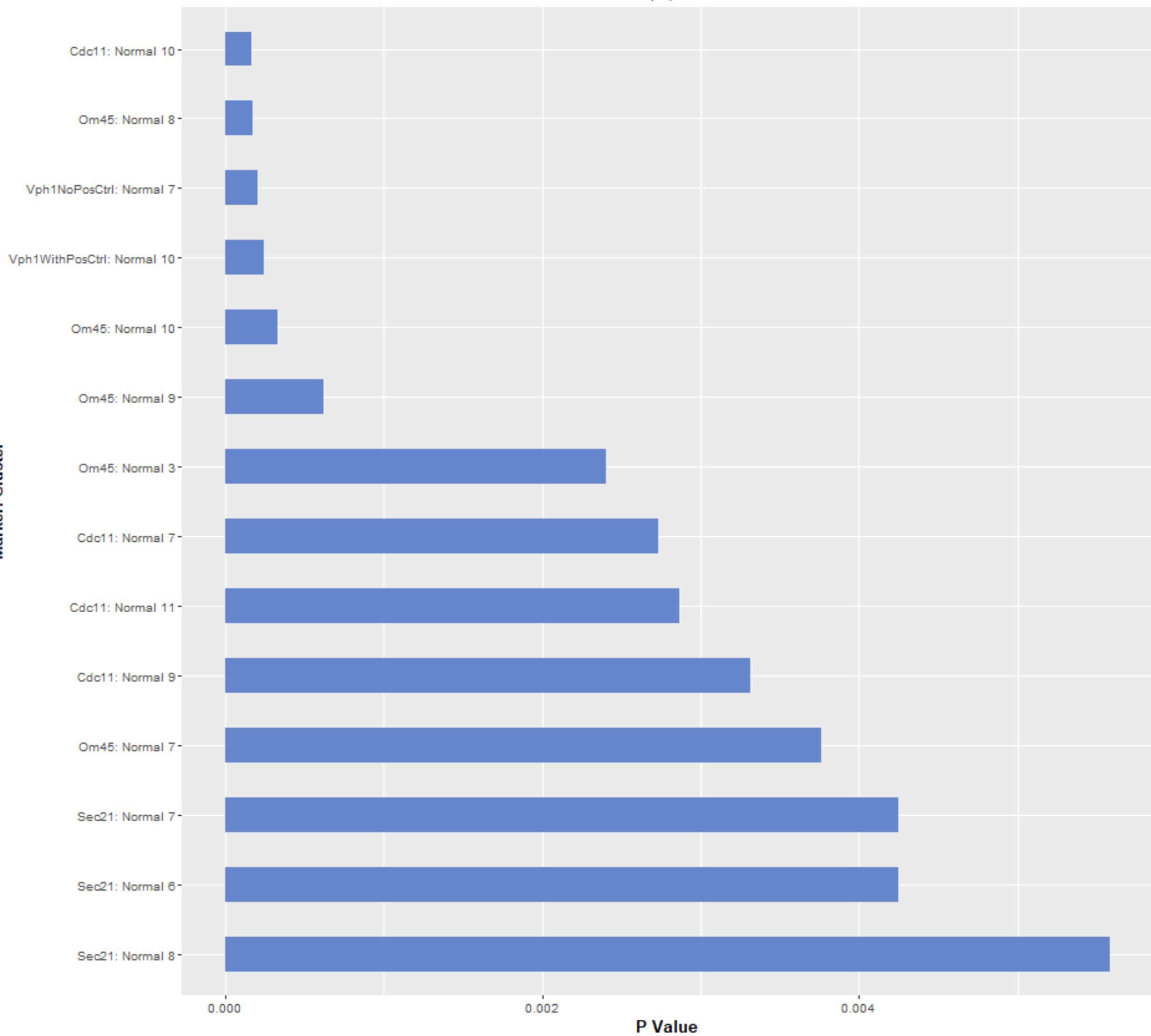
Marker: Cluster



# protein phosphorylation

Without AreaShape | Ordered Gene Set

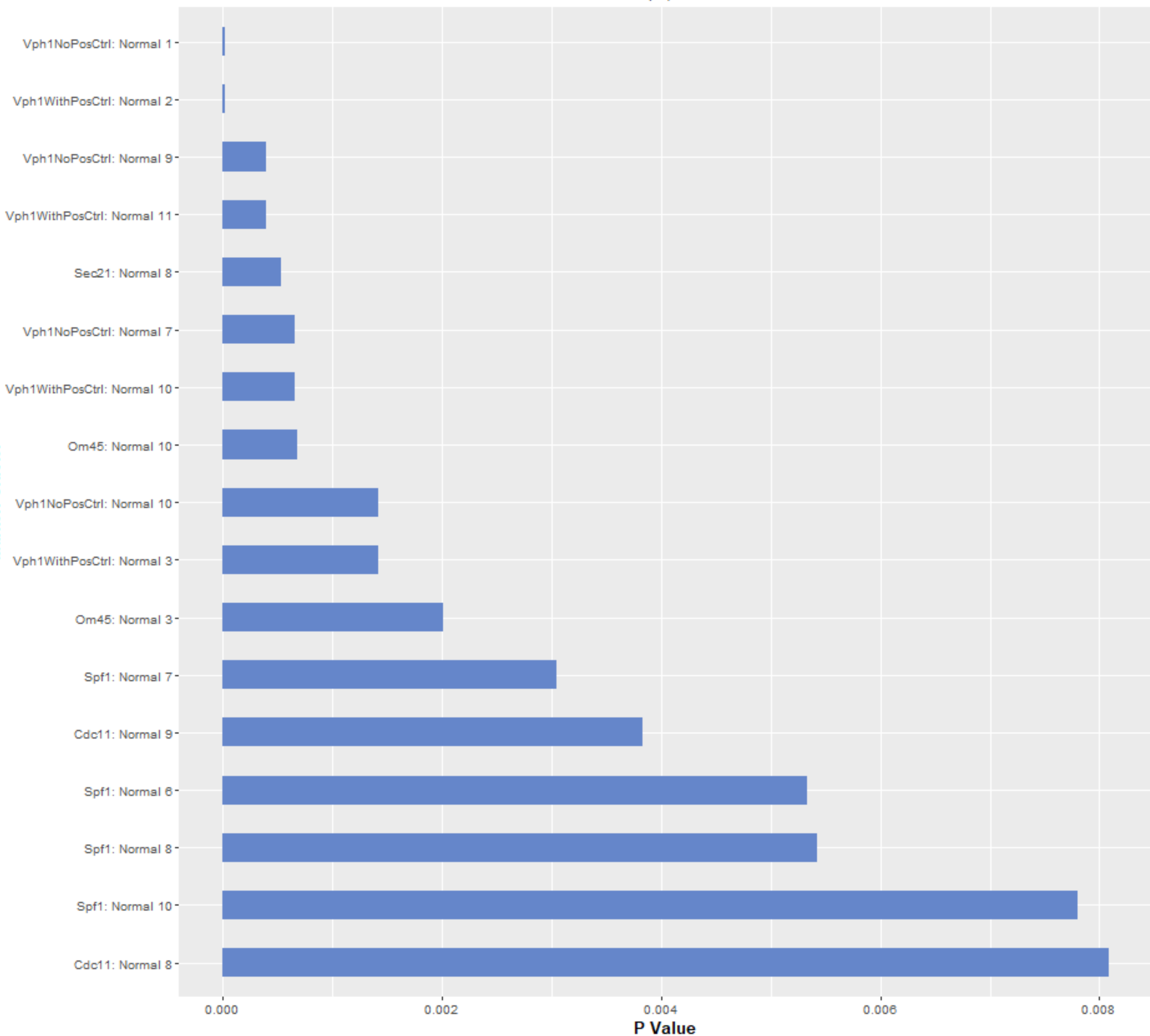
Marker: Cluster



# protein targeting

Without AreaShape | Ordered Gene Set

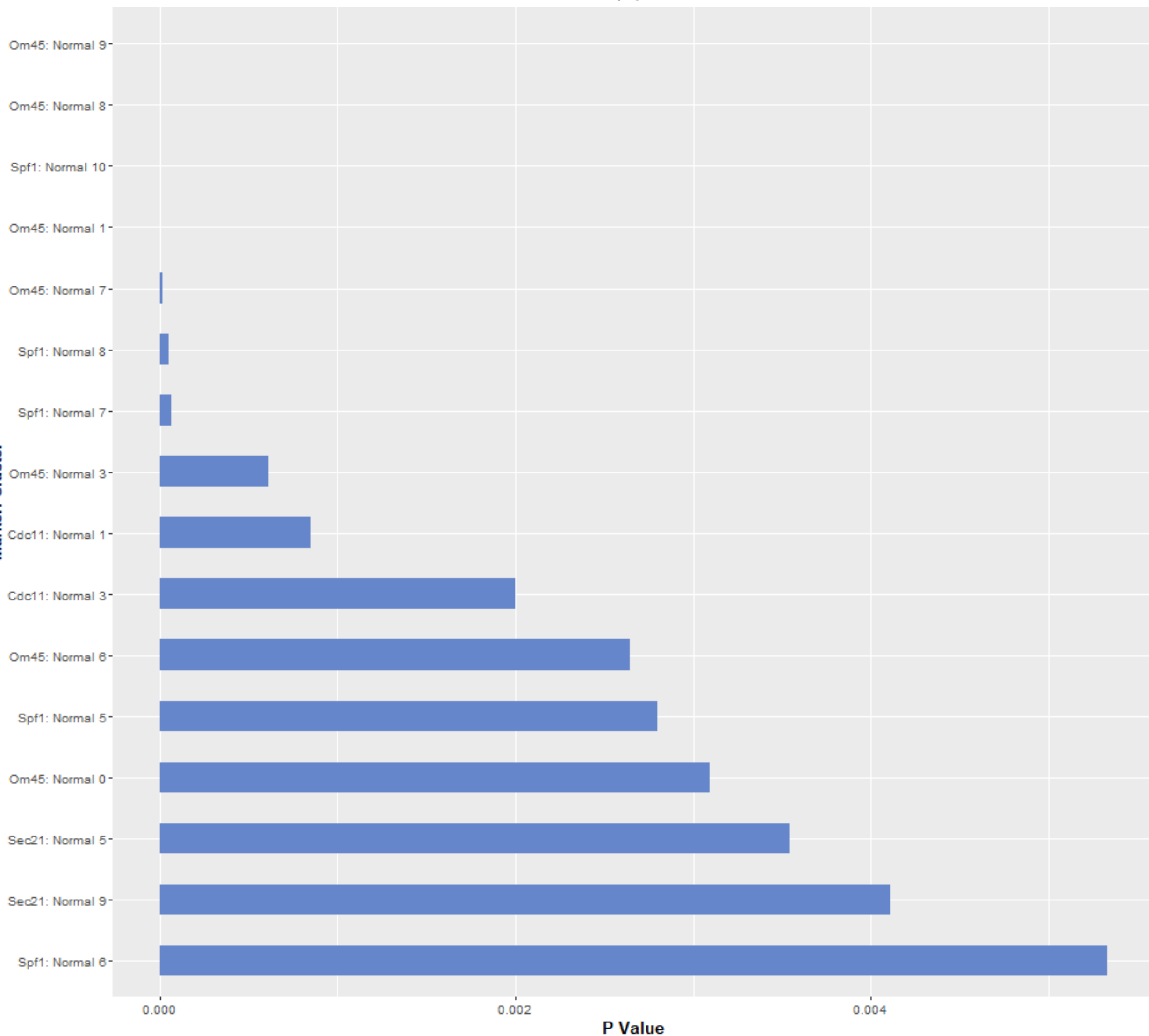
Marker: Cluster



# proteolysis involved in cellular protein catabolic process

Without AreaShape | Ordered Gene Set

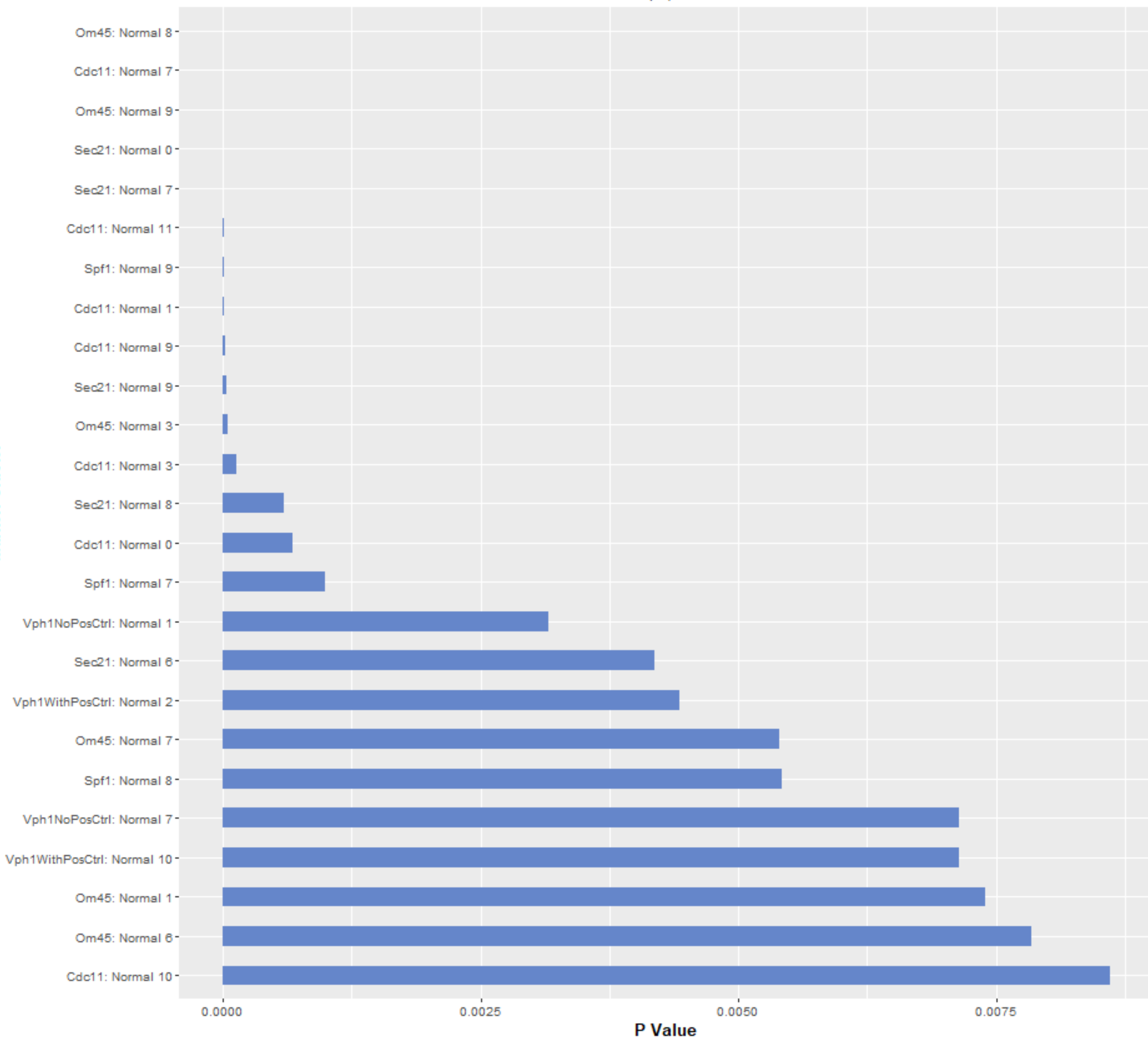
Marker: Cluster





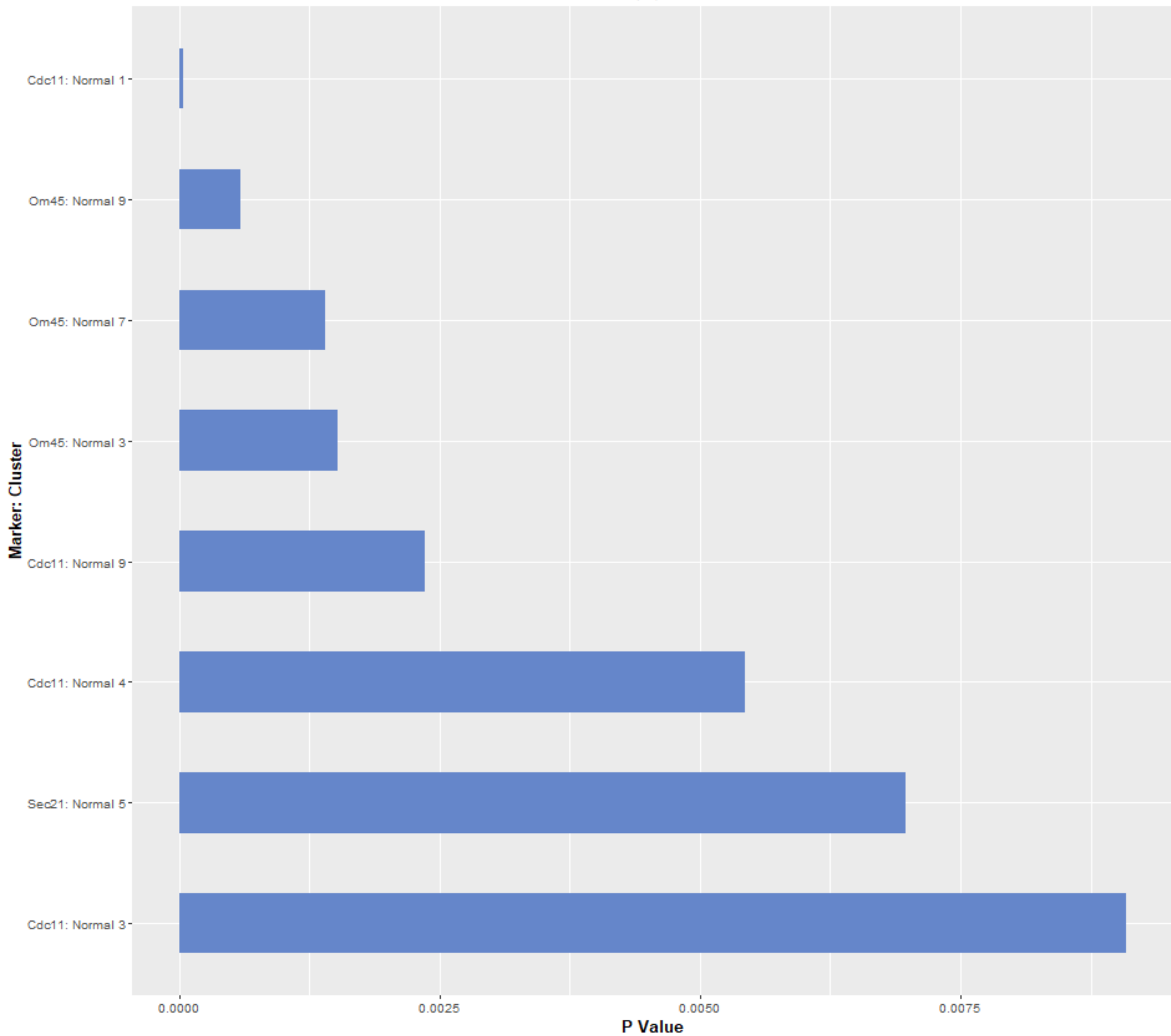
**regulation of cell cycle**  
Without AreaShape | Ordered Gene Set

Marker: Cluster



# regulation of DNA metabolic process

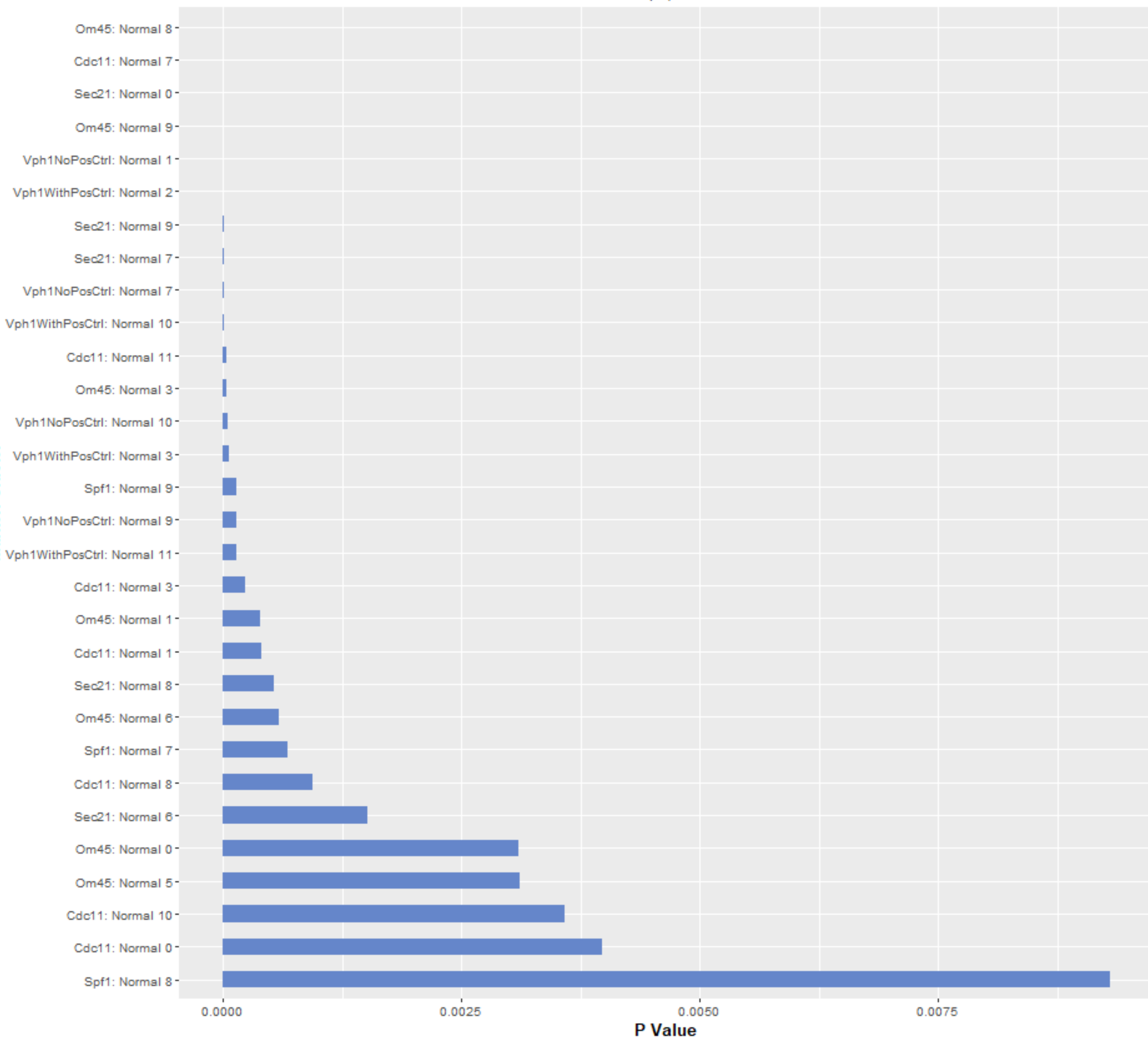
Without AreaShape | Ordered Gene Set



# regulation of organelle organization

Without AreaShape | Ordered Gene Set

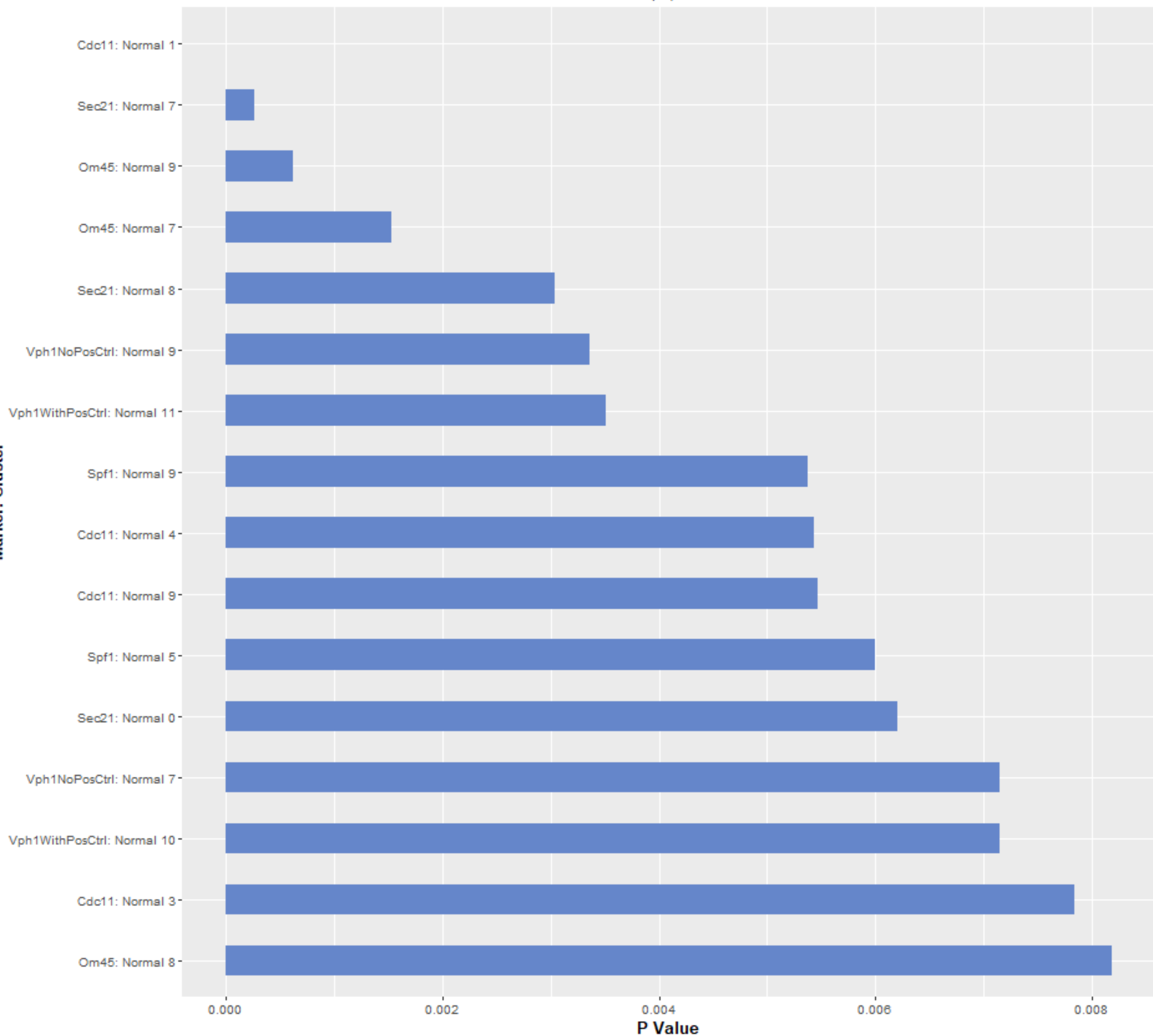
Marker: Cluster



# regulation of protein modification process

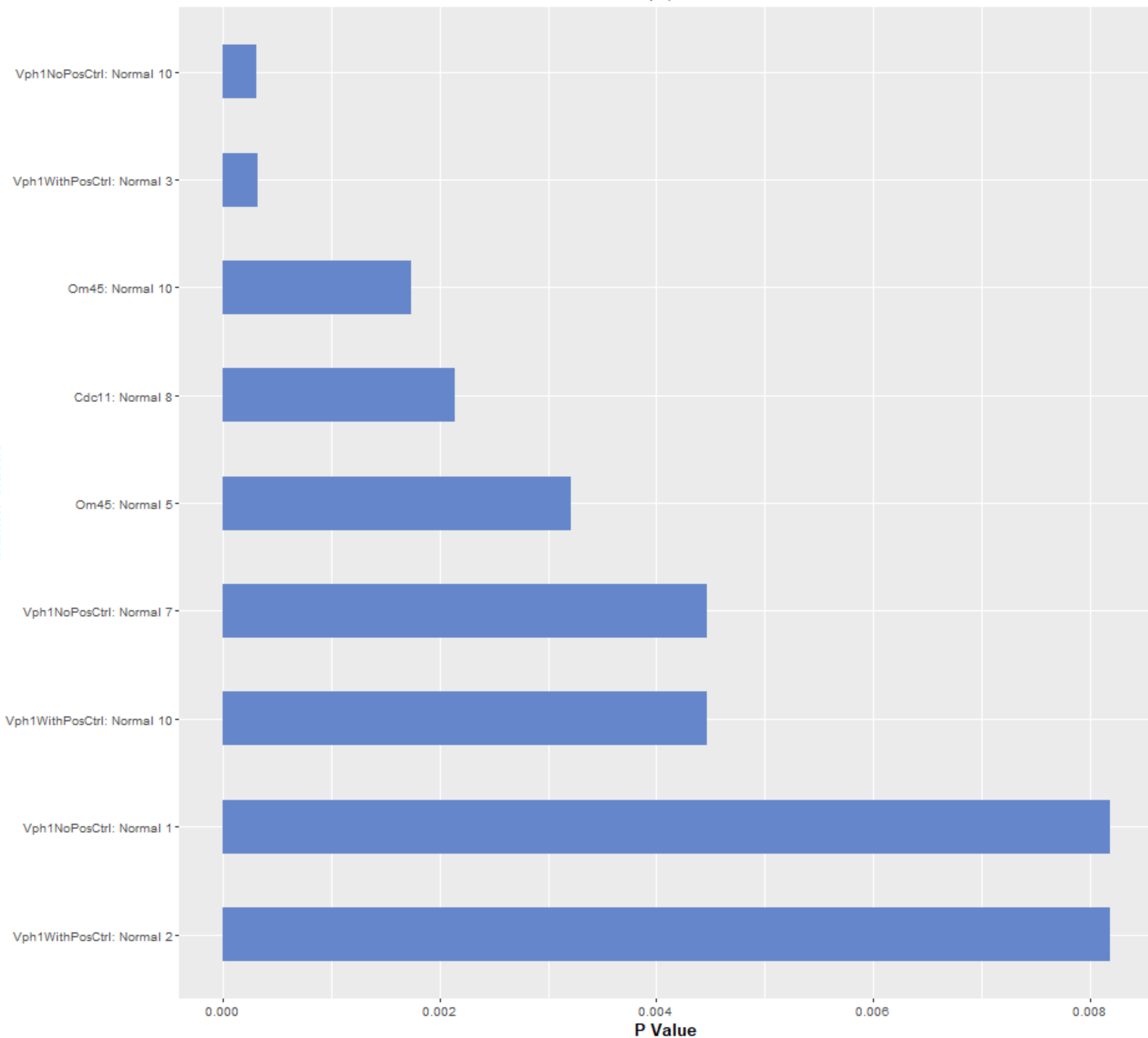
Without AreaShape | Ordered Gene Set

Marker: Cluster



**regulation of transport**  
Without AreaShape | Ordered Gene Set

**Marker: Cluster**



# response to heat

Without AreaShape | Ordered Gene Set

Marker: Cluster

Cdc11: Normal 5

0.000

0.001

0.002

0.003

0.004

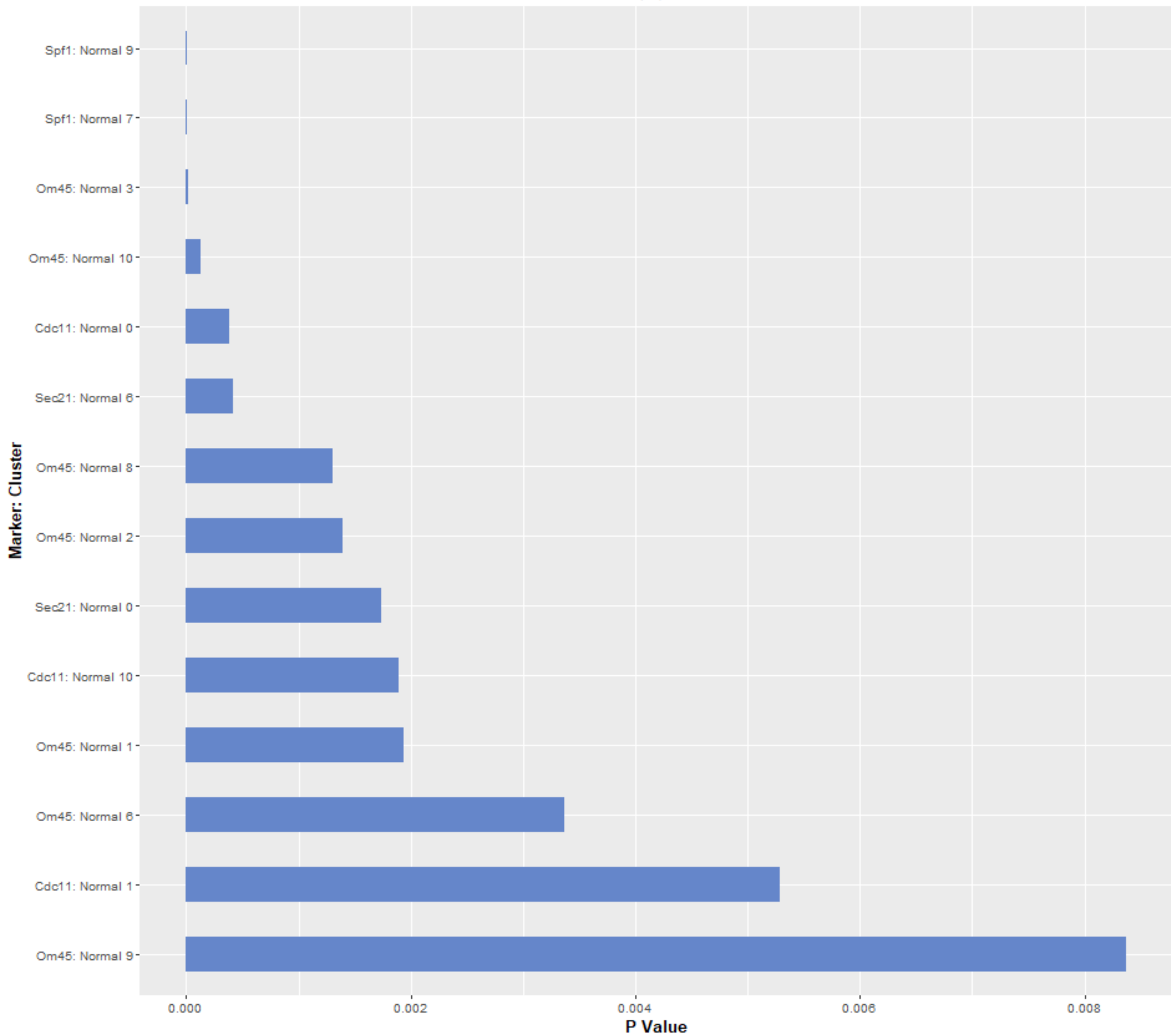
0.005

P Value



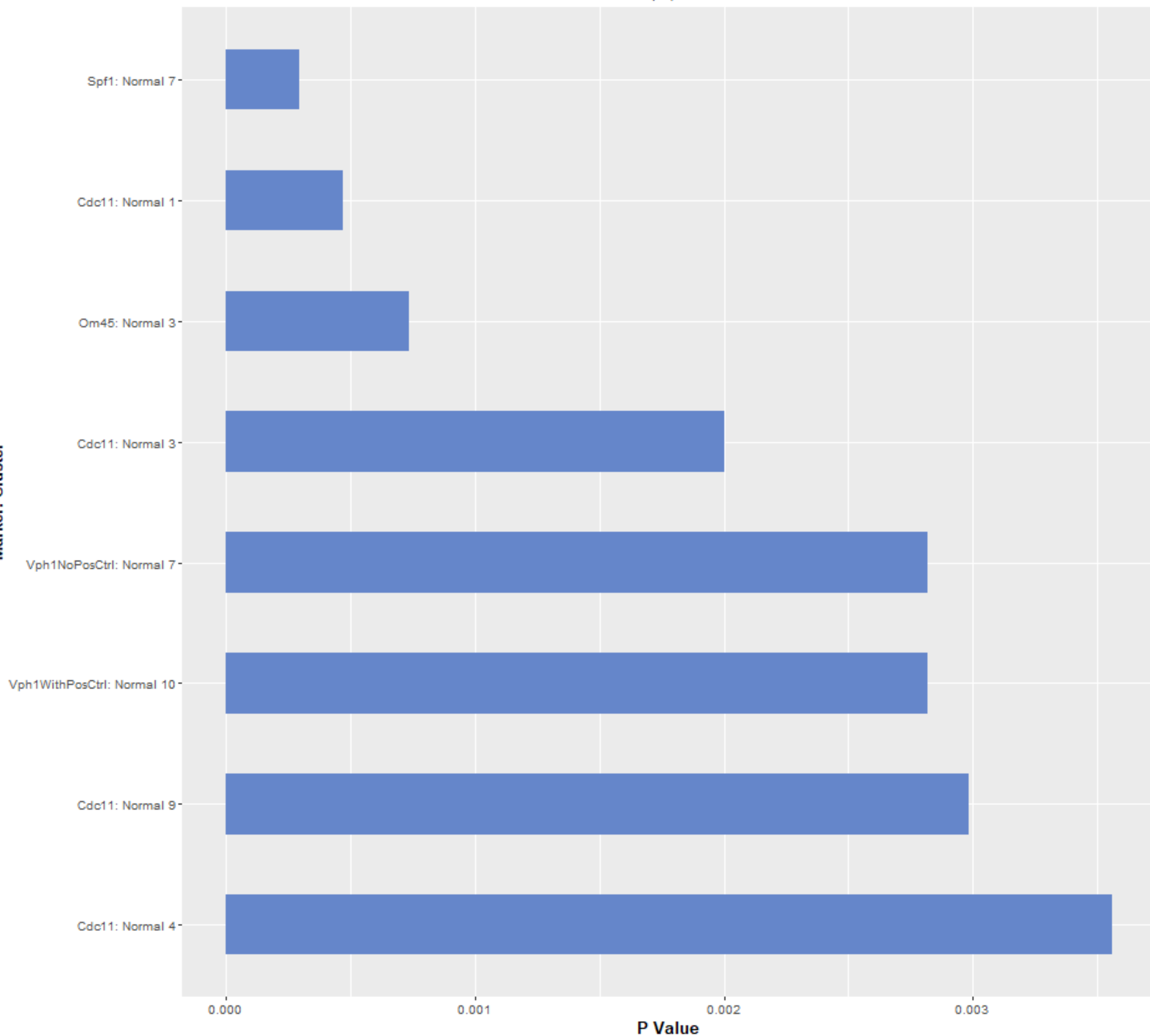
# ribosomal subunit export from nucleus

Without AreaShape | Ordered Gene Set



**RNA catabolic process**  
Without AreaShape | Ordered Gene Set

**Marker: Cluster**





# RNA modification

Without AreaShape | Ordered Gene Set

Marker: Cluster

Cdc11: Normal 4

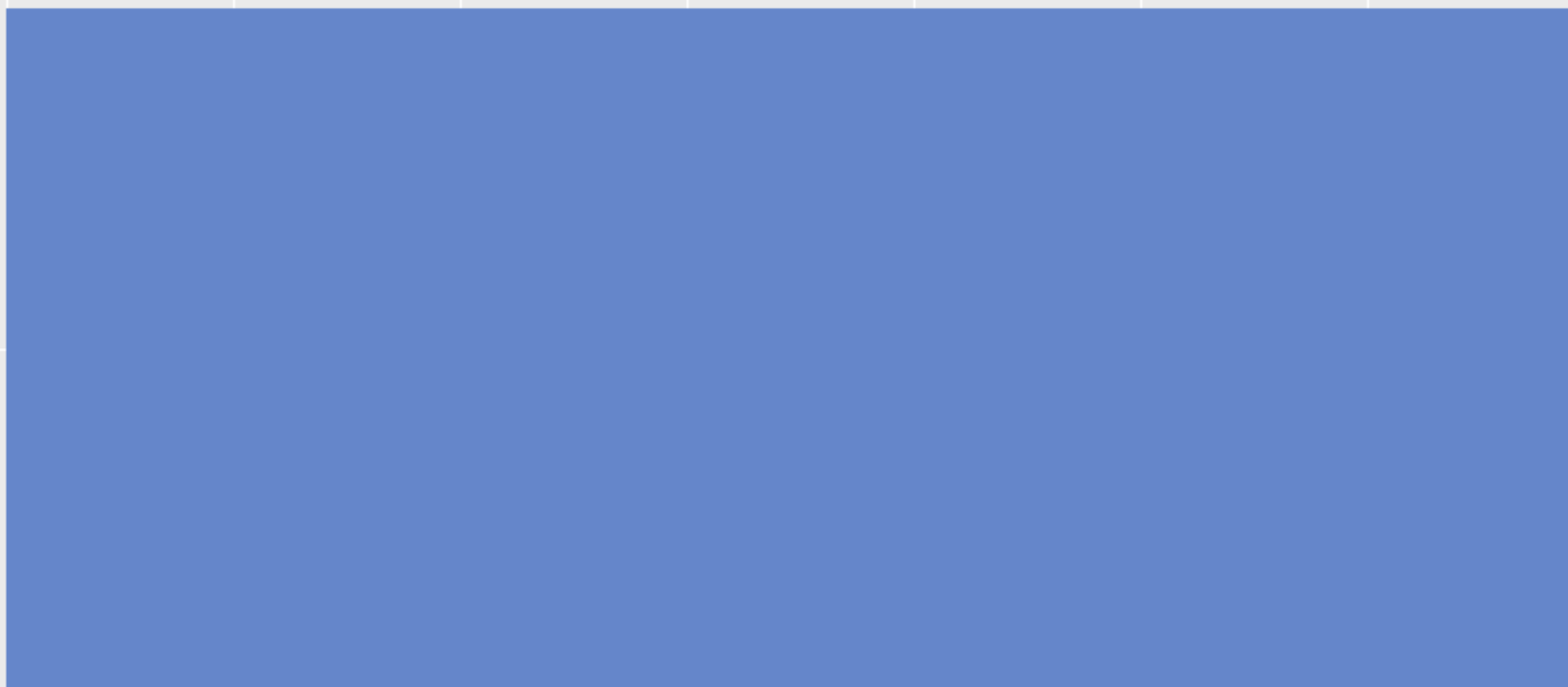
0.0000

0.0025

0.0050

0.0075

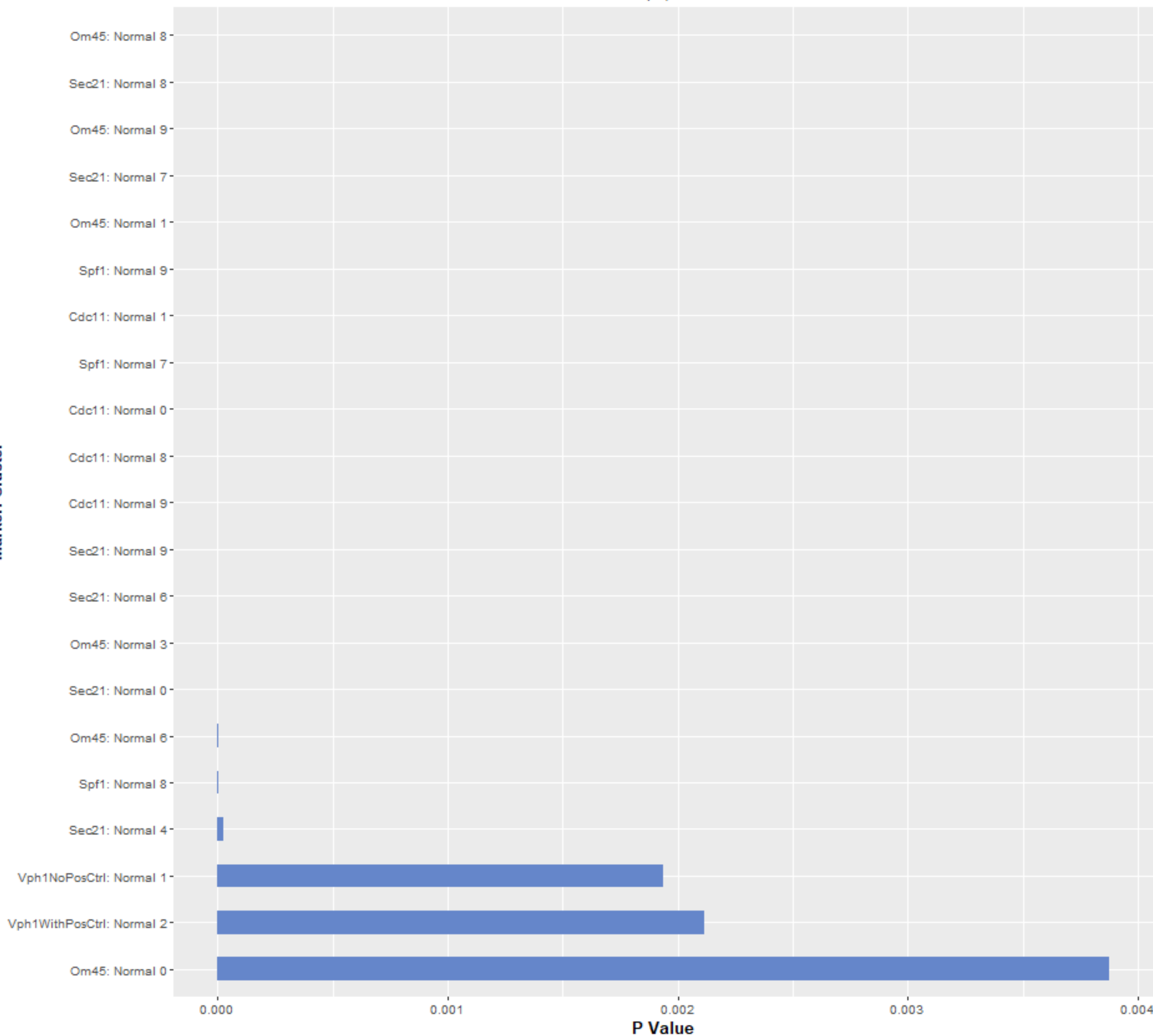
P Value



# RNA splicing

Without AreaShape | Ordered Gene Set

Marker: Cluster



# signaling

Without AreaShape | Ordered Gene Set

Marker: Cluster

Cdc11: Normal 9

Spf1: Normal 7

0.000

0.001

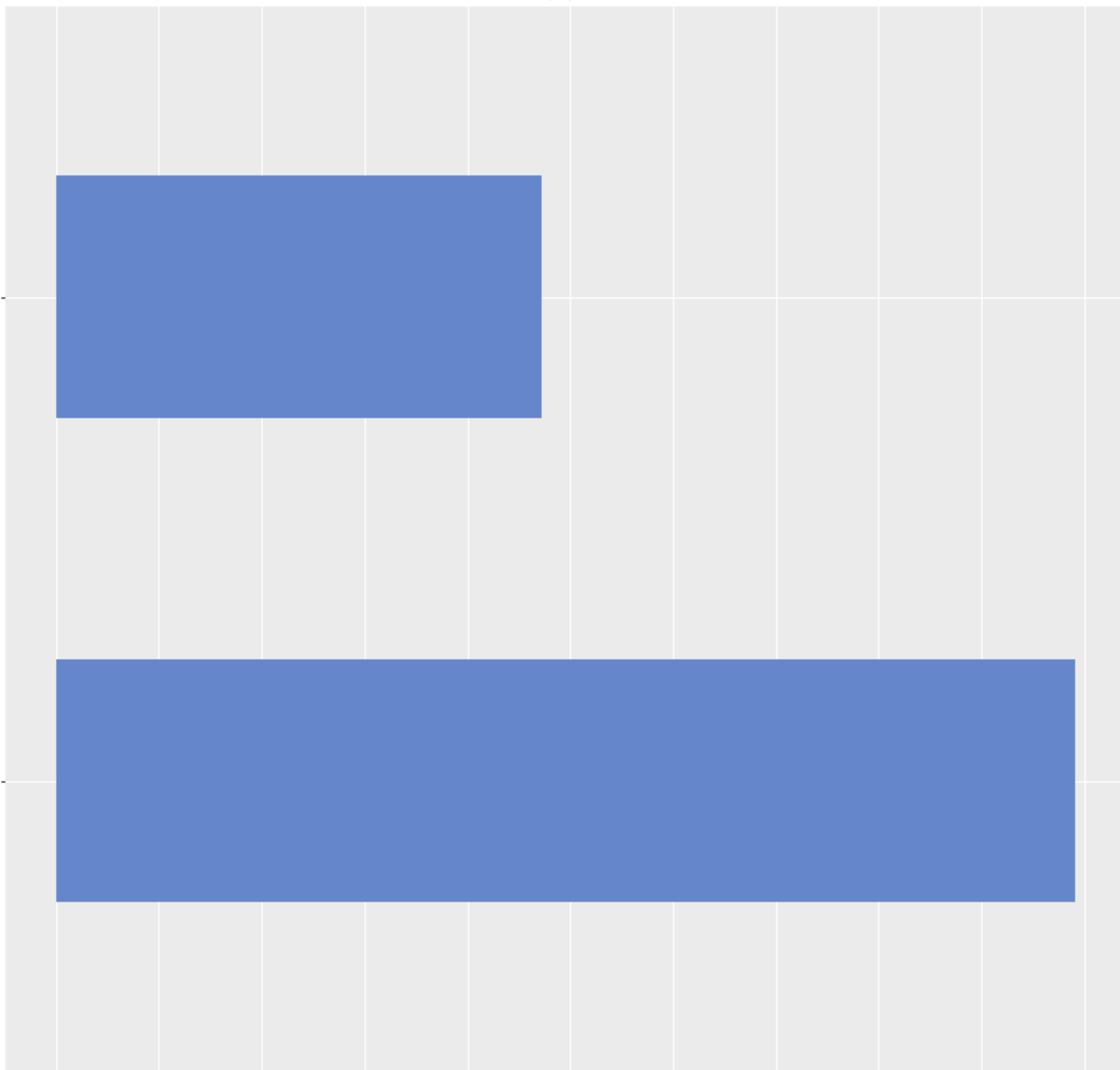
0.002

0.003

0.004

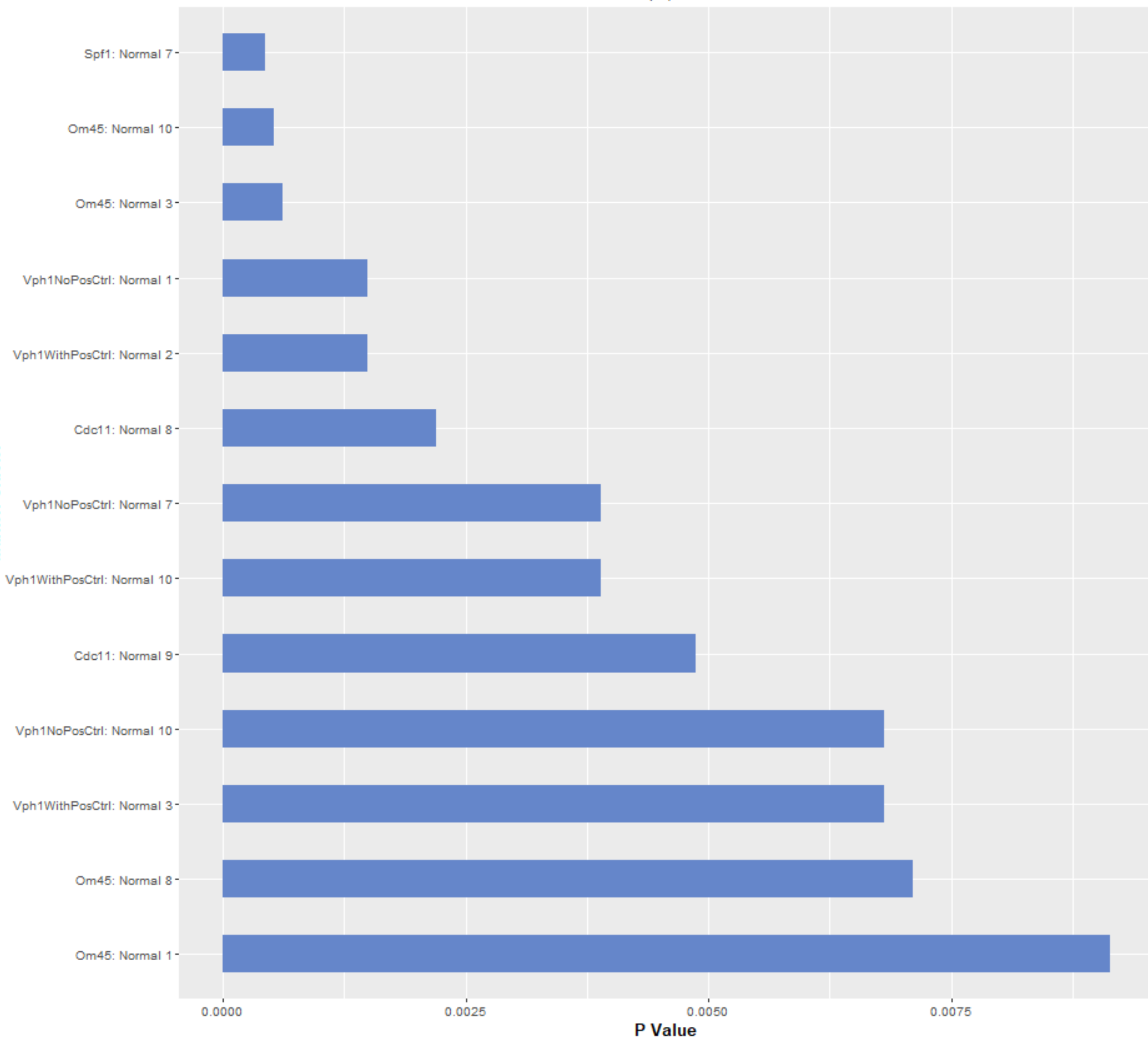
0.005

P Value



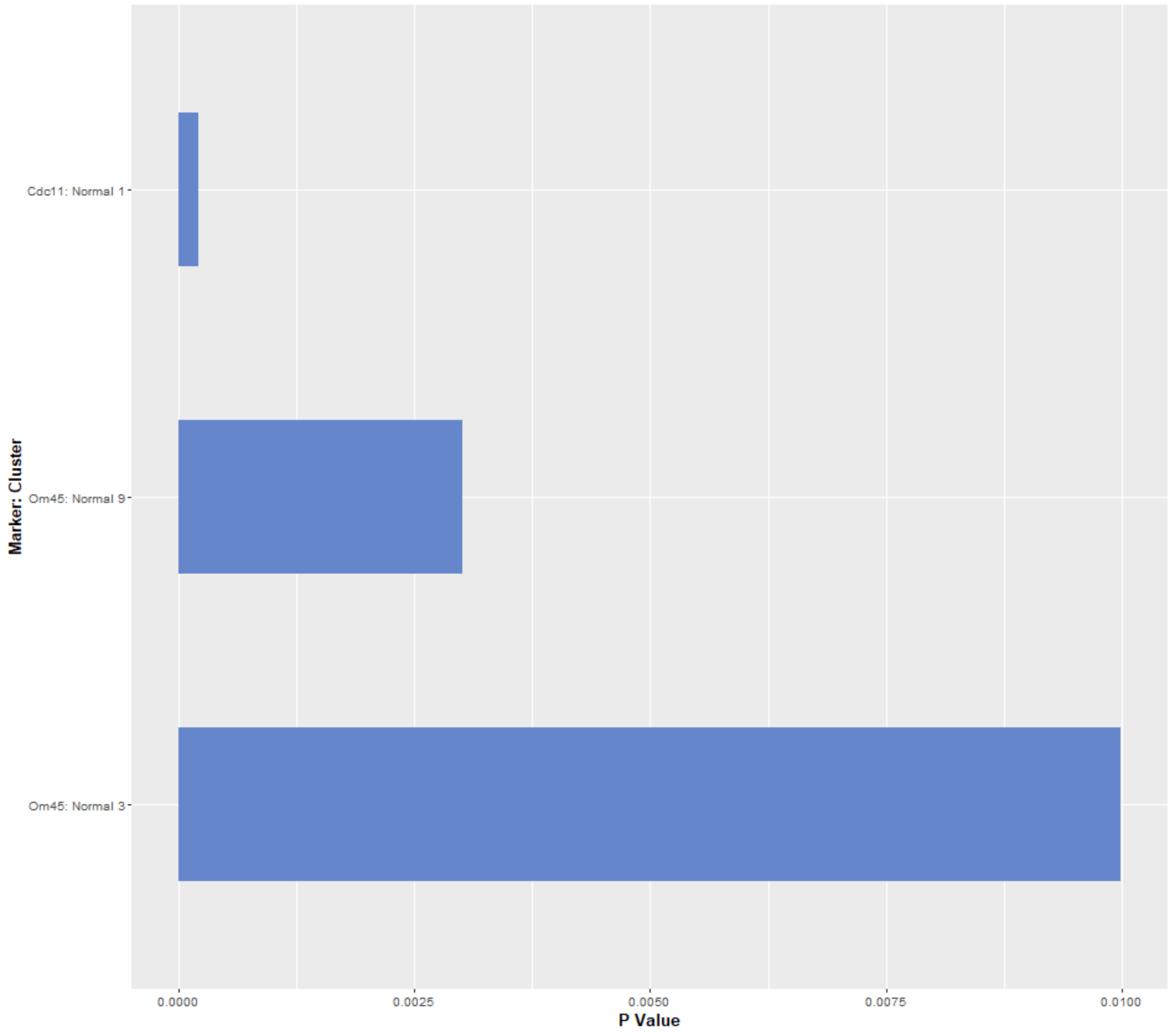
**snoRNA processing**  
Without AreaShape | Ordered Gene Set

**Marker: Cluster**



# telomere organization

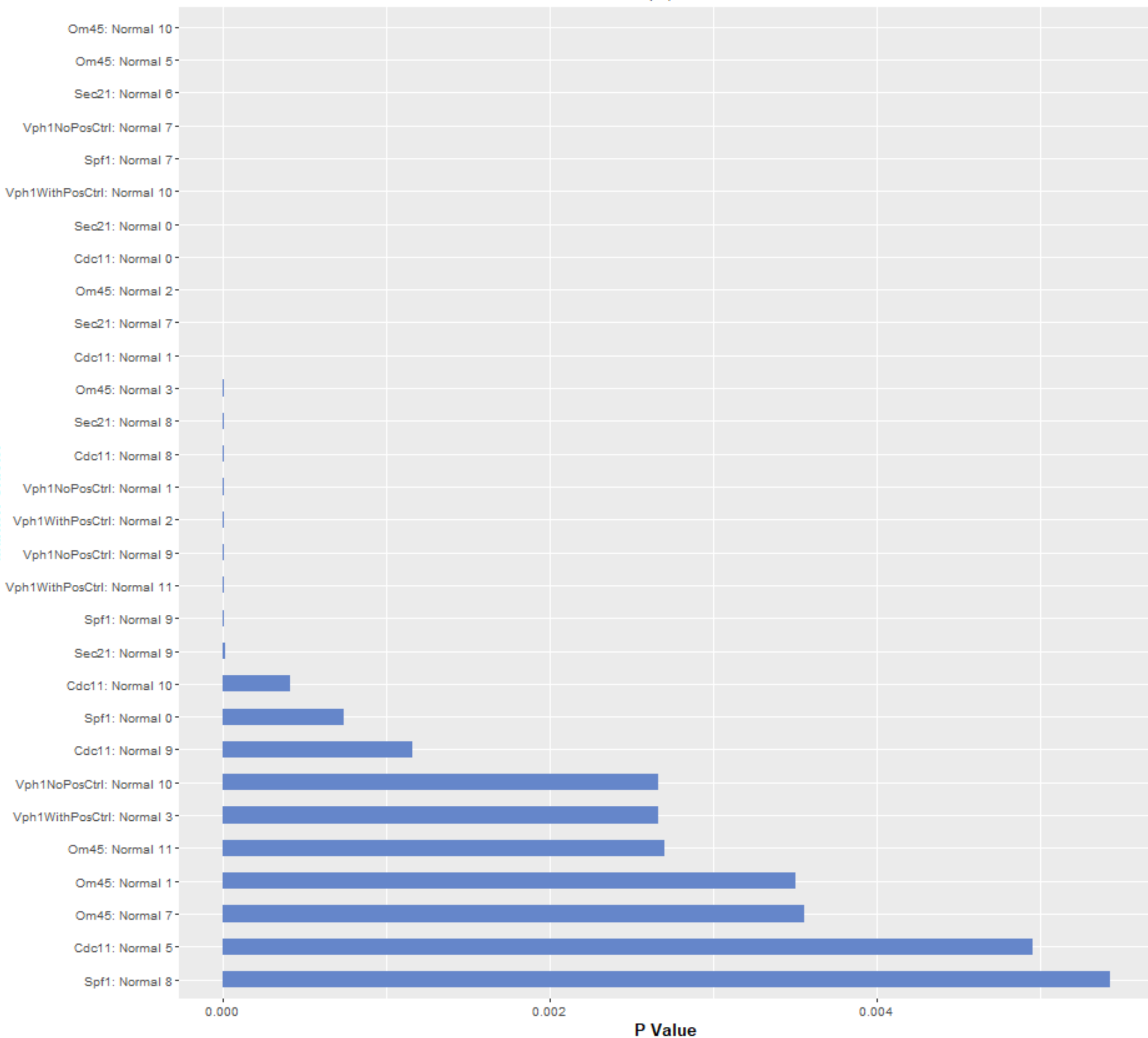
Without AreaShape | Ordered Gene Set



# transcription from RNA polymerase II promoter

Without AreaShape | Ordered Gene Set

Marker: Cluster



# translational initiation

Without AreaShape | Ordered Gene Set

Marker: Cluster

Vph1NoPosCtrl: Normal 9

Vph1WithPosCtrl: Normal 11

Om45: Normal 10

Om45: Normal 3

0.000

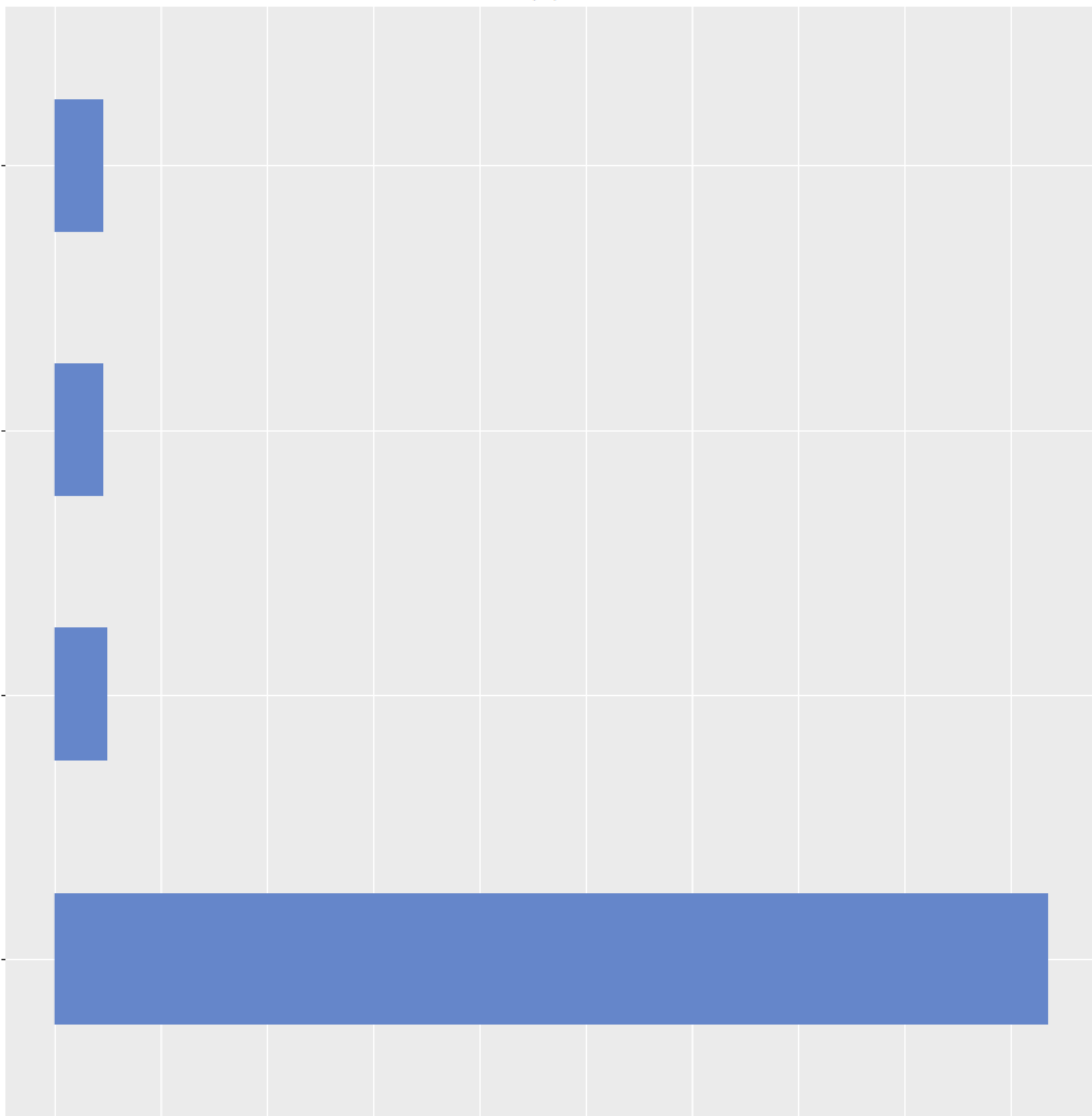
0.001

0.002

0.003

0.004

P Value



# transposition

Without AreaShape | Ordered Gene Set

Marker: Cluster

Cdc11: Normal 1

Vph1NoPosCtrl: Normal 7

Vph1WithPosCtrl: Normal 10

Spf1: Normal 1

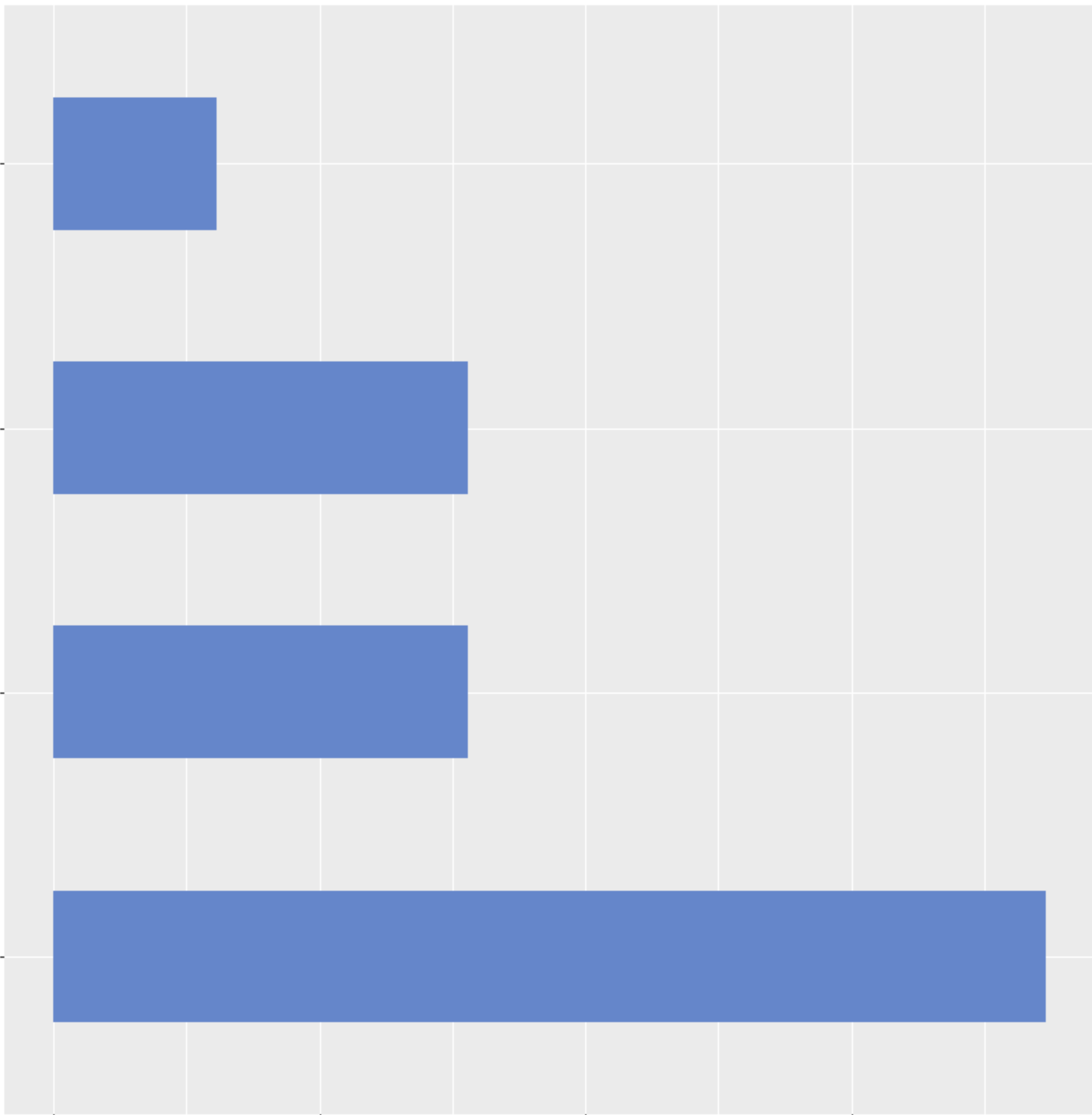
0.0000

0.0025

0.0050

0.0075

P Value

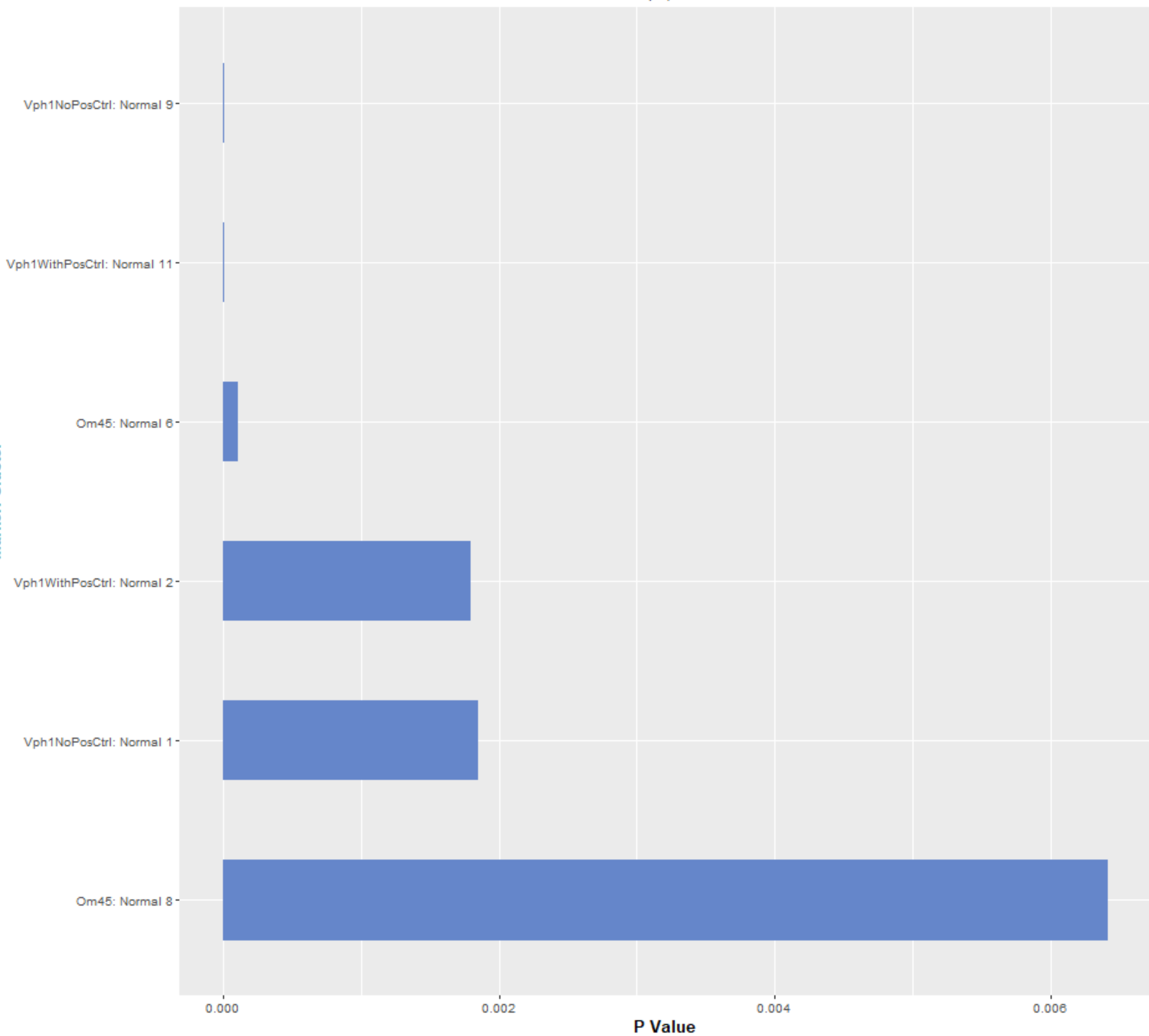




# tRNA aminoacylation for protein translation

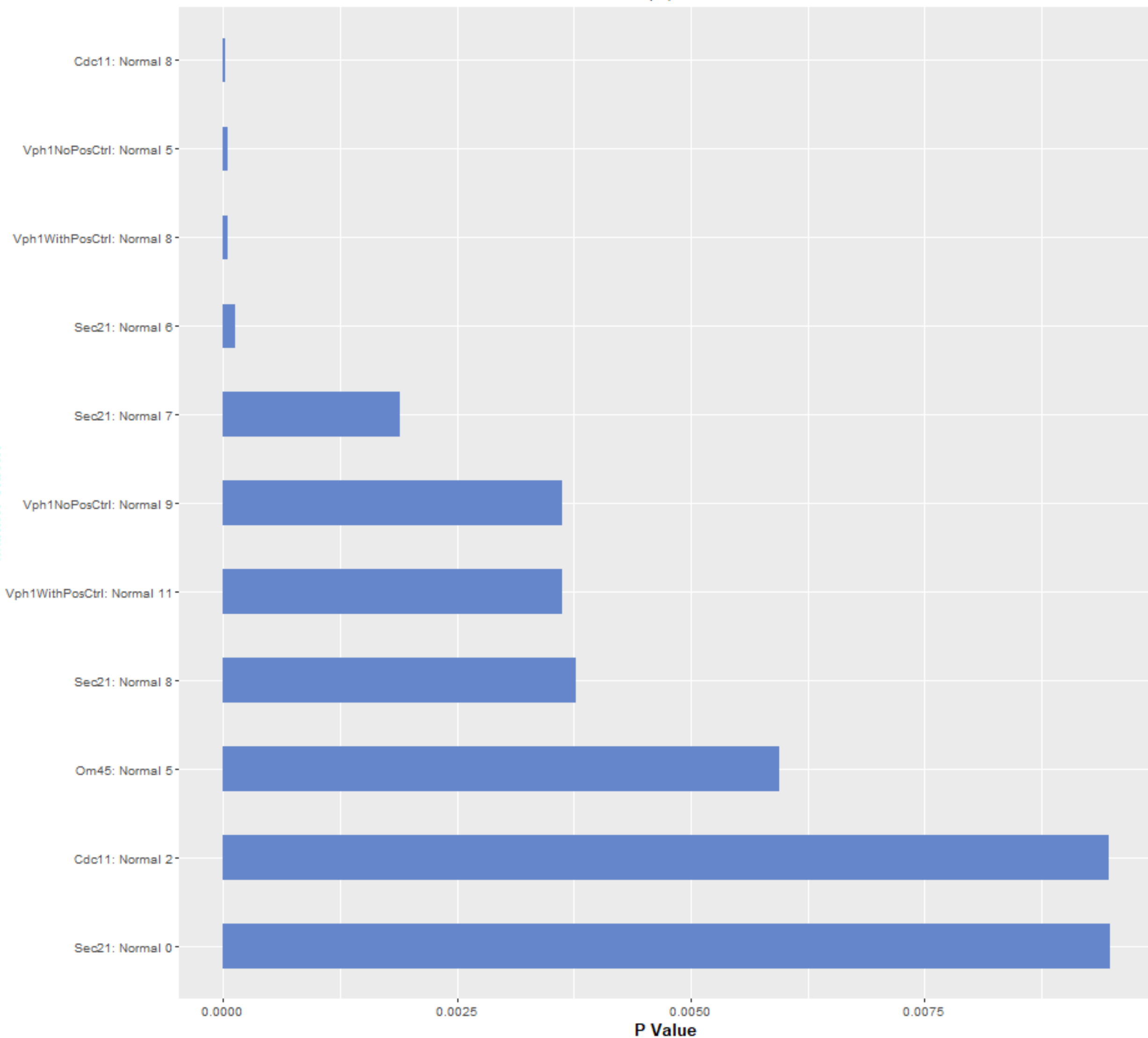
Without AreaShape | Ordered Gene Set

Marker: Cluster



**vacuole organization**  
Without AreaShape | Ordered Gene Set

**Marker: Cluster**



**vesicle organization**  
Without AreaShape | Ordered Gene Set

**Marker: Cluster**

