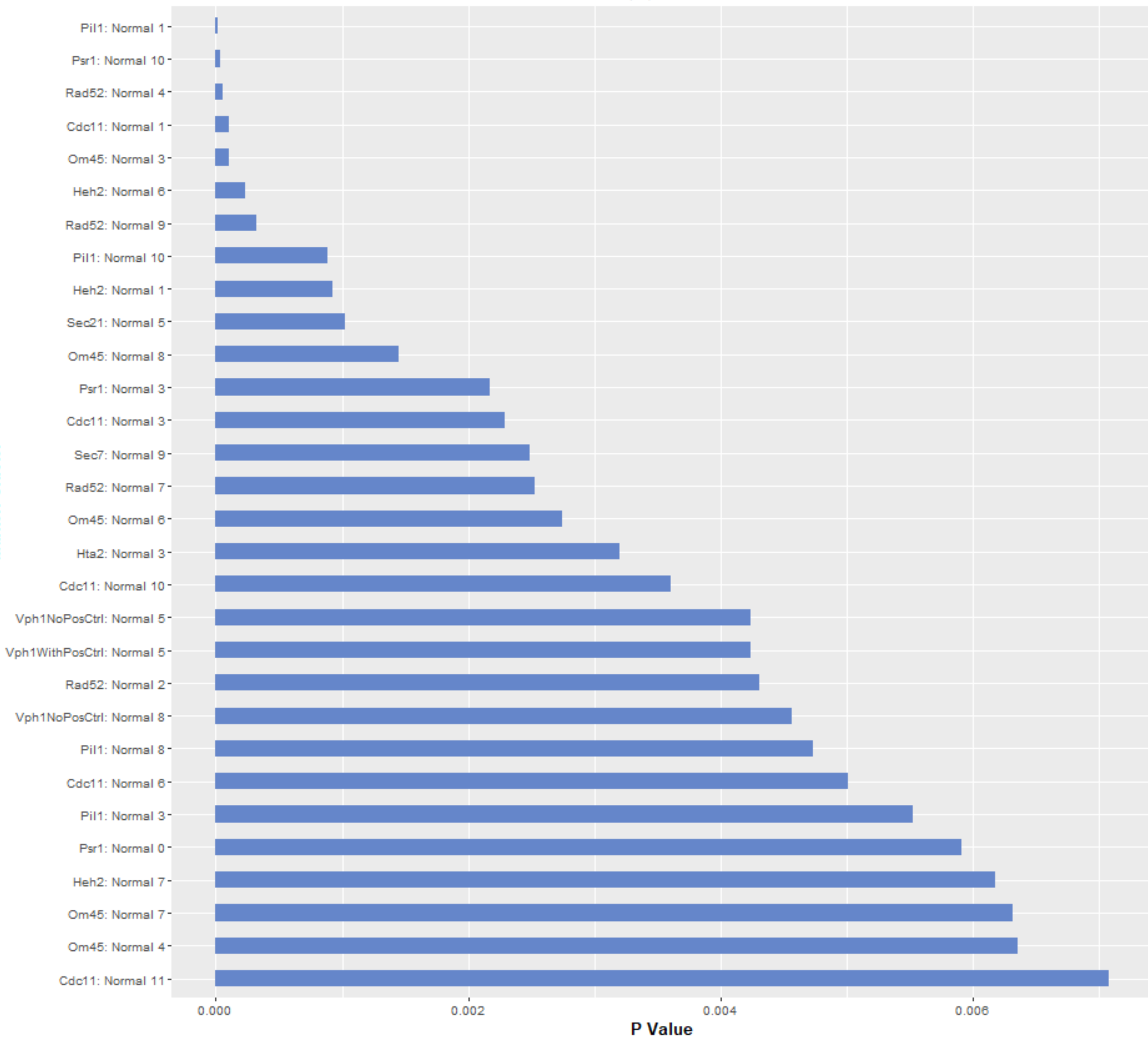


cell budding

With AreaShape | Unordered Gene Set

Marker: Cluster



cell morphogenesis

With AreaShape | Unordered Gene Set

Marker: Cluster

Om45: Normal 3

0.000000

0.000025

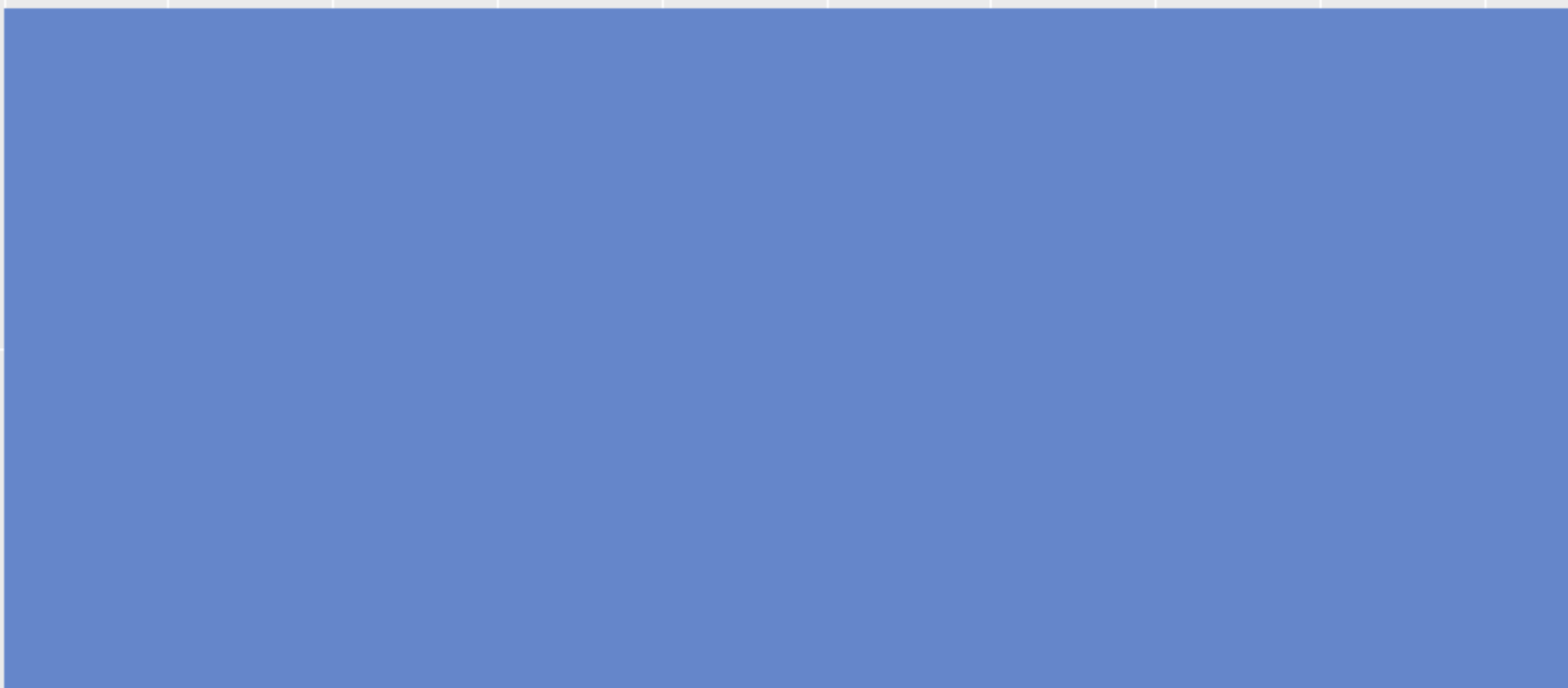
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0.000075

0.000100

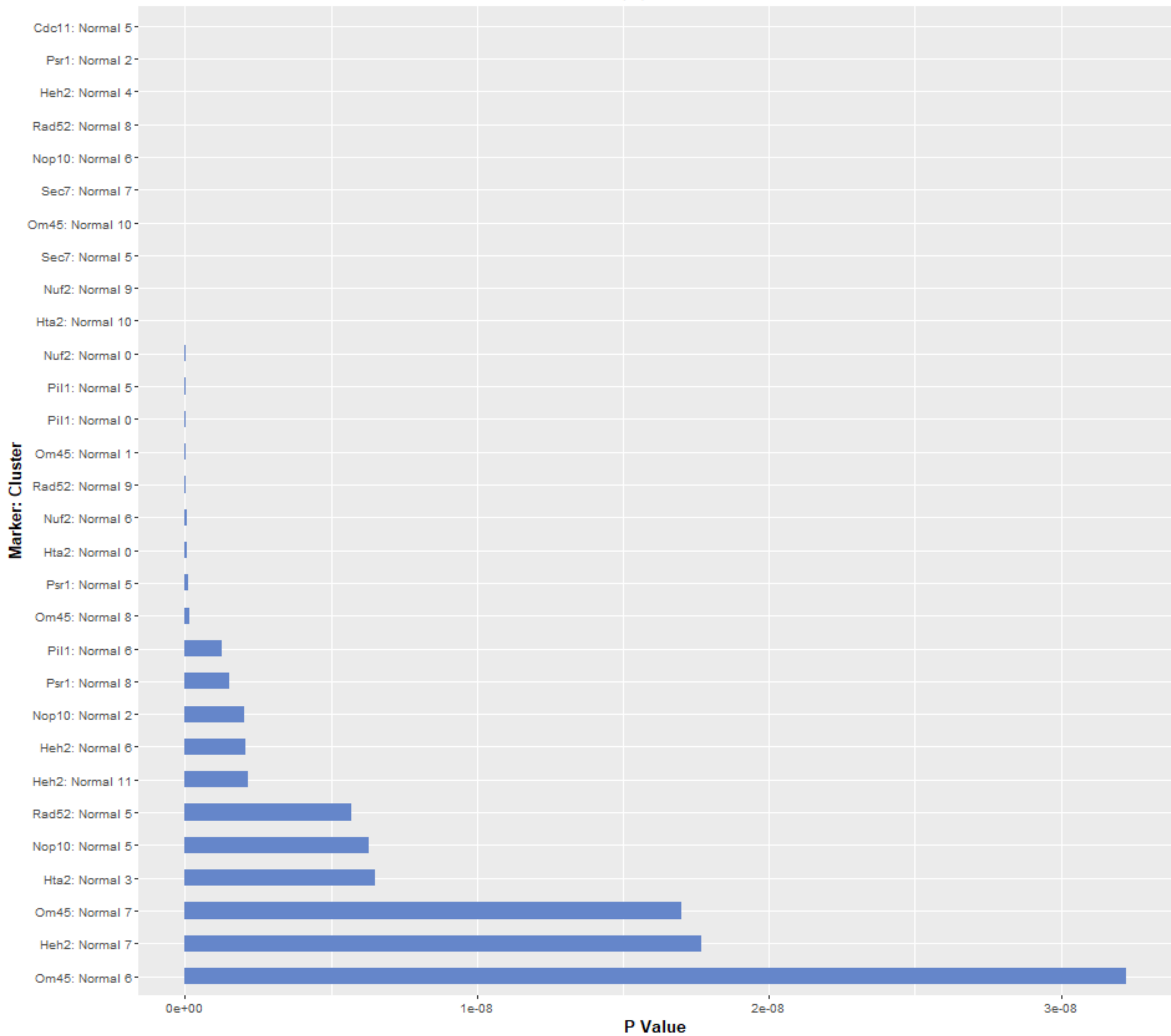
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P Value



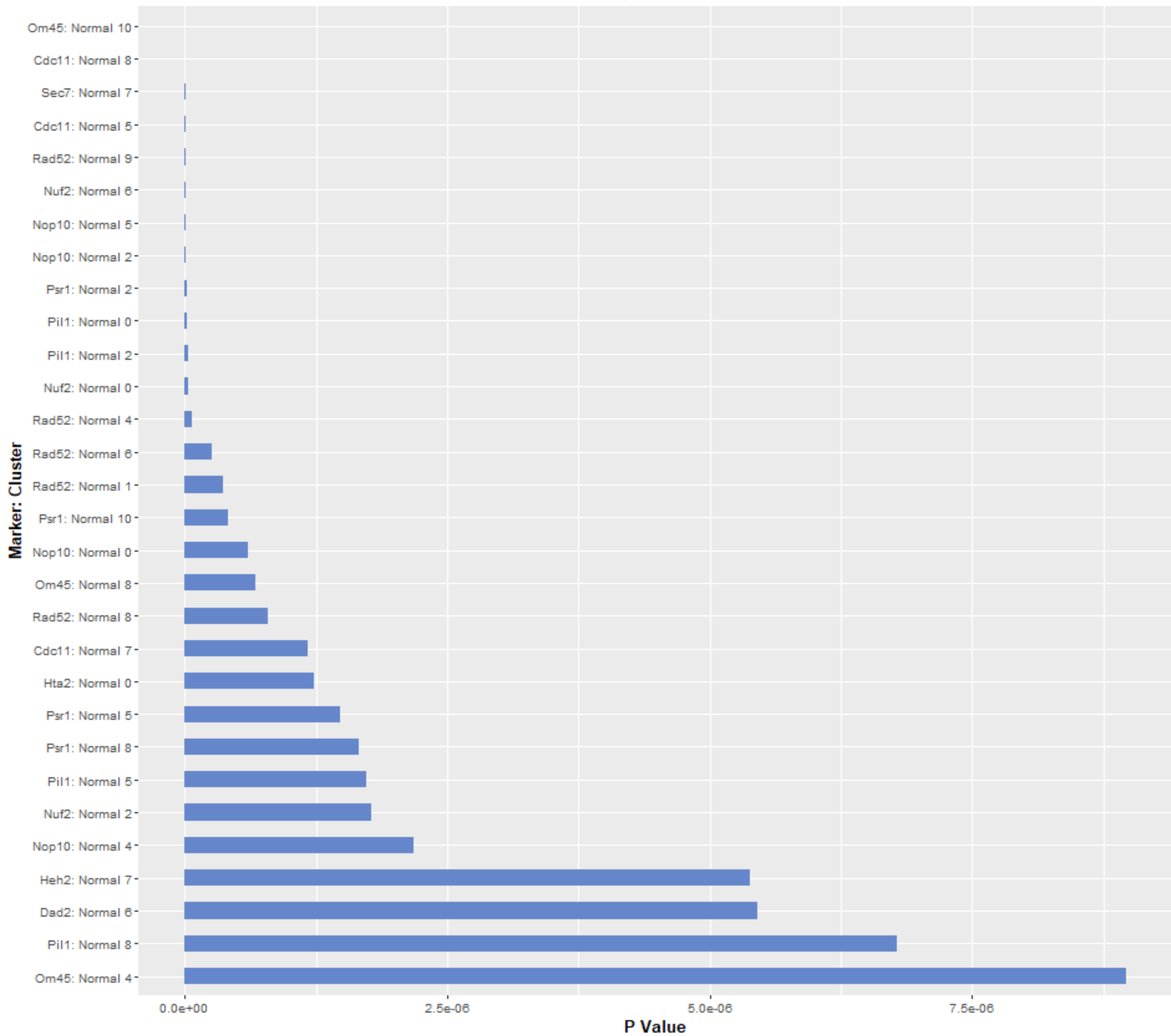
cellular response to DNA damage stimulus

With AreaShape | Unordered Gene Set



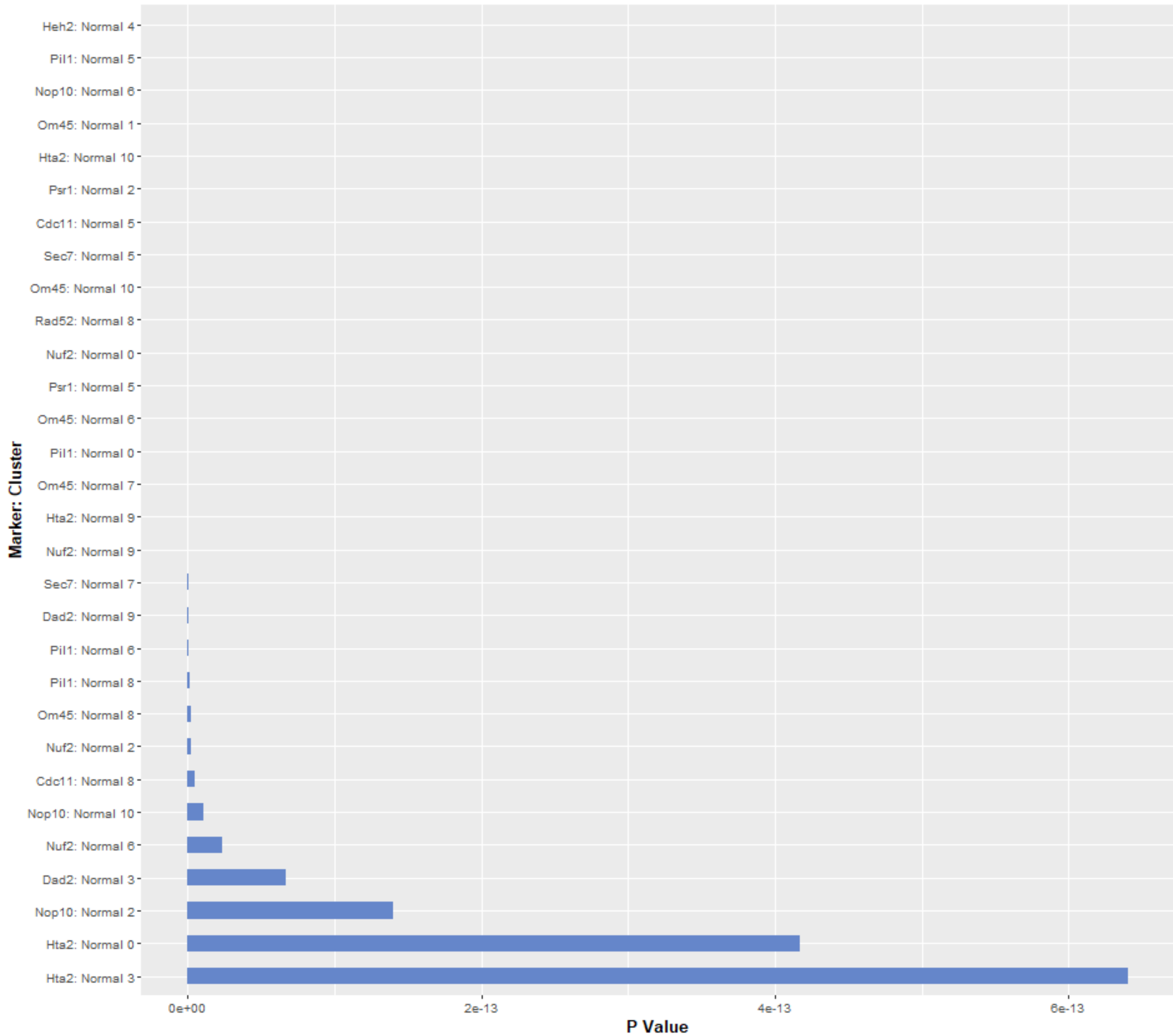
chromatin organization

With AreaShape | Unordered Gene Set



chromosome segregation

With AreaShape | Unordered Gene Set



conjugation

With AreaShape | Unordered Gene Set

Marker: Cluster

Sec7: Normal 10

0.00000

0.00005

0.00010

0.00015

0.00020

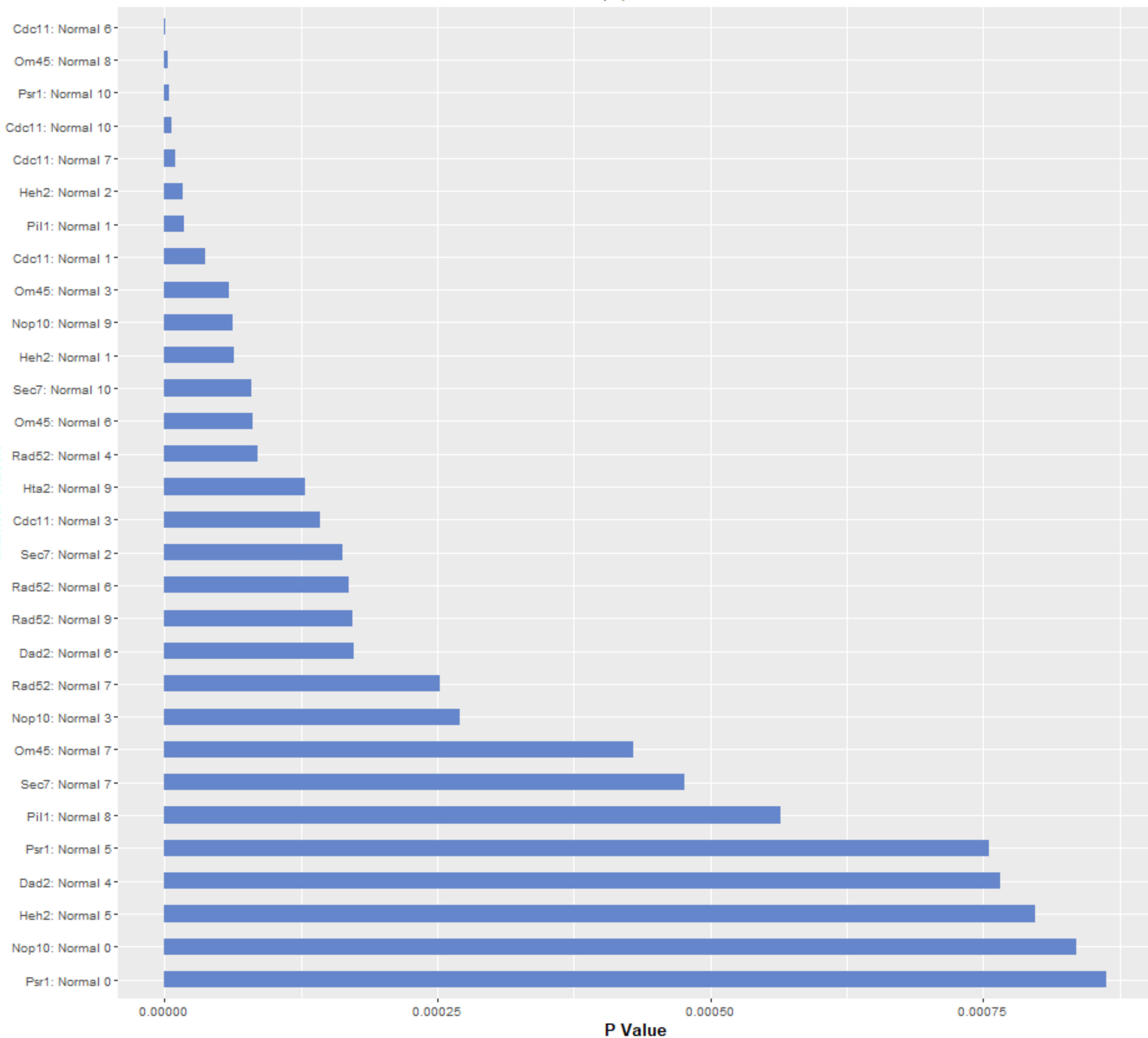
P Value



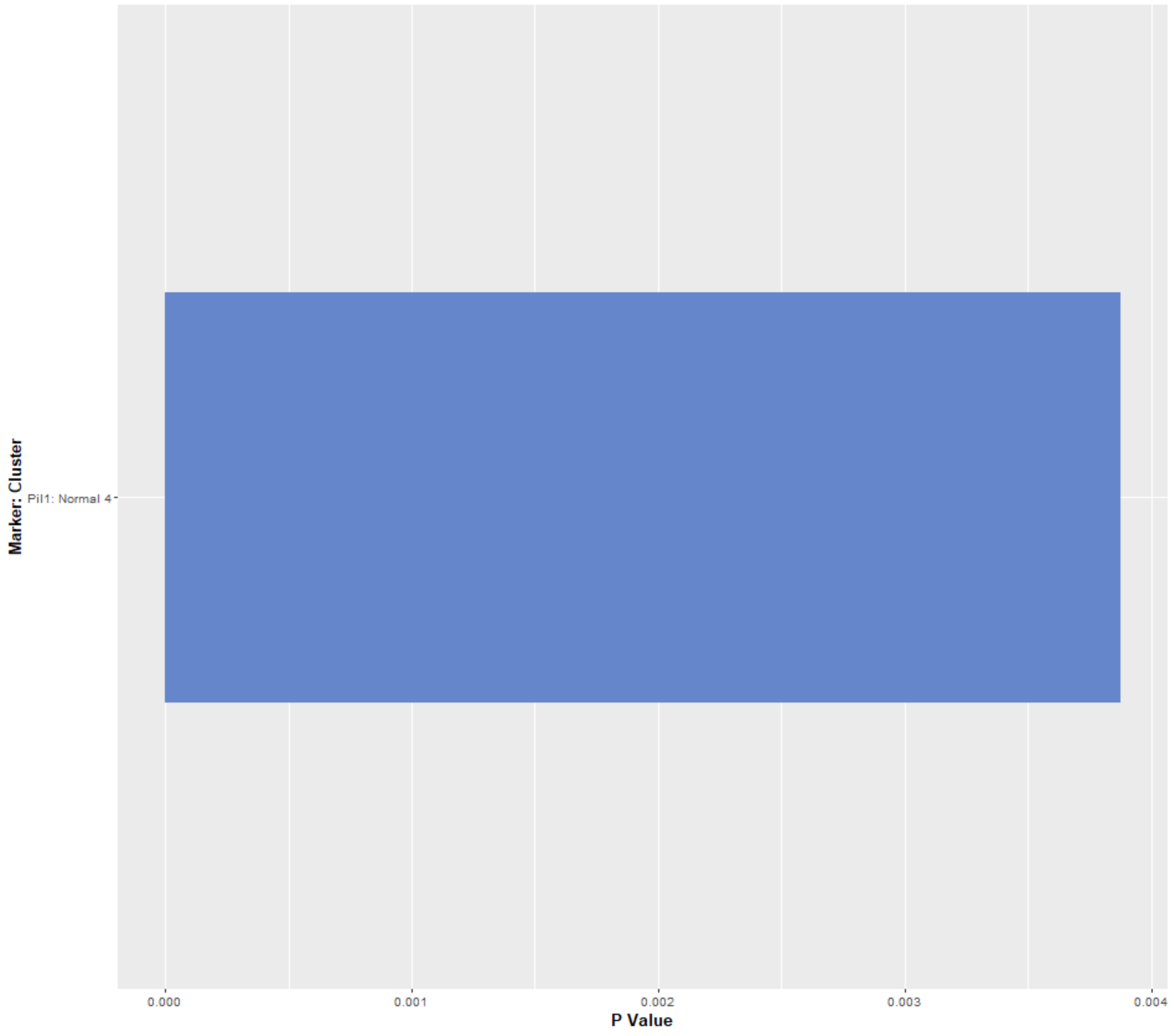
cytokinesis

With AreaShape | Unordered Gene Set

Marker: Cluster



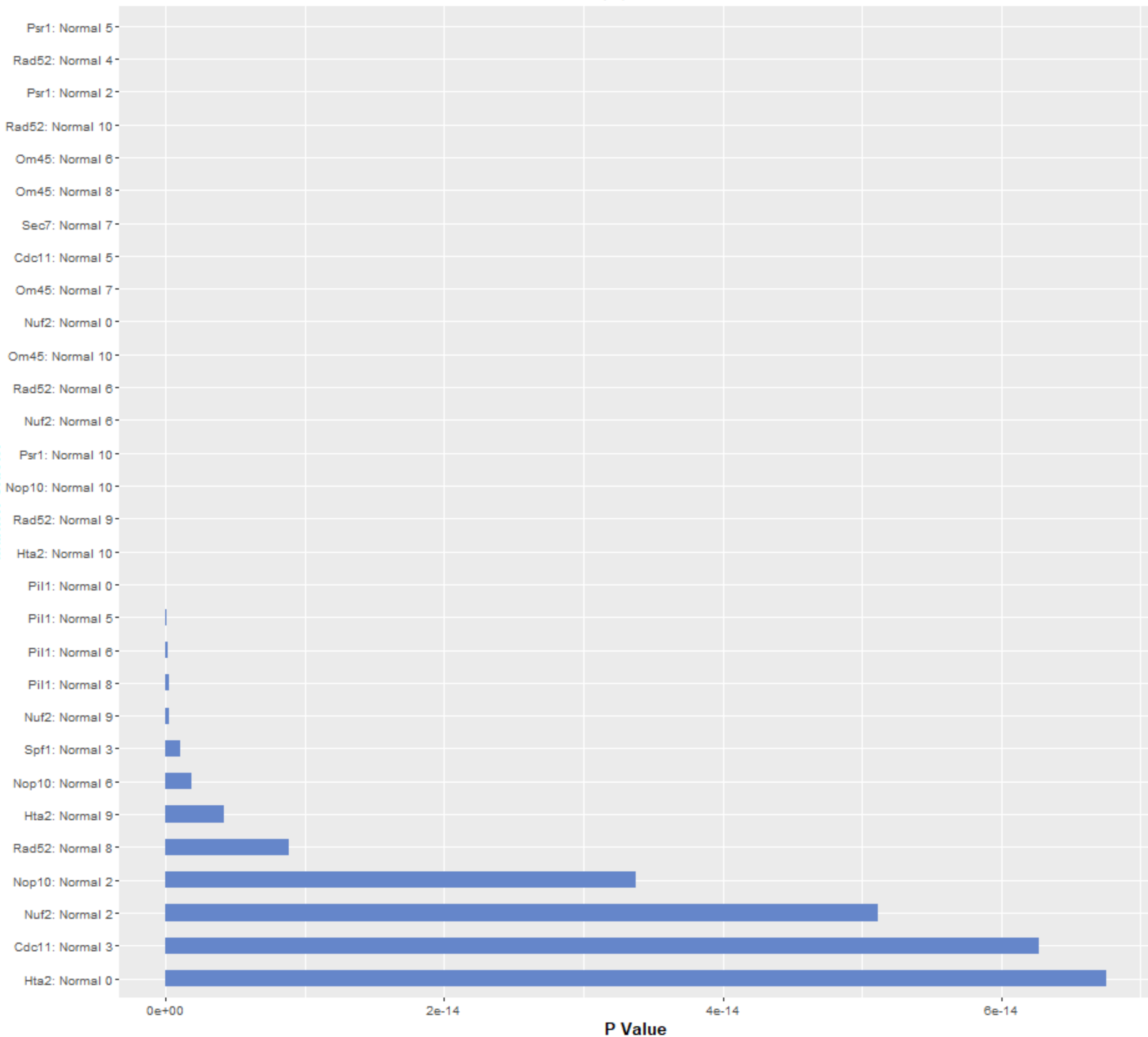
cytoplasmic translation
With AreaShape | Unordered Gene Set



cytoskeleton organization

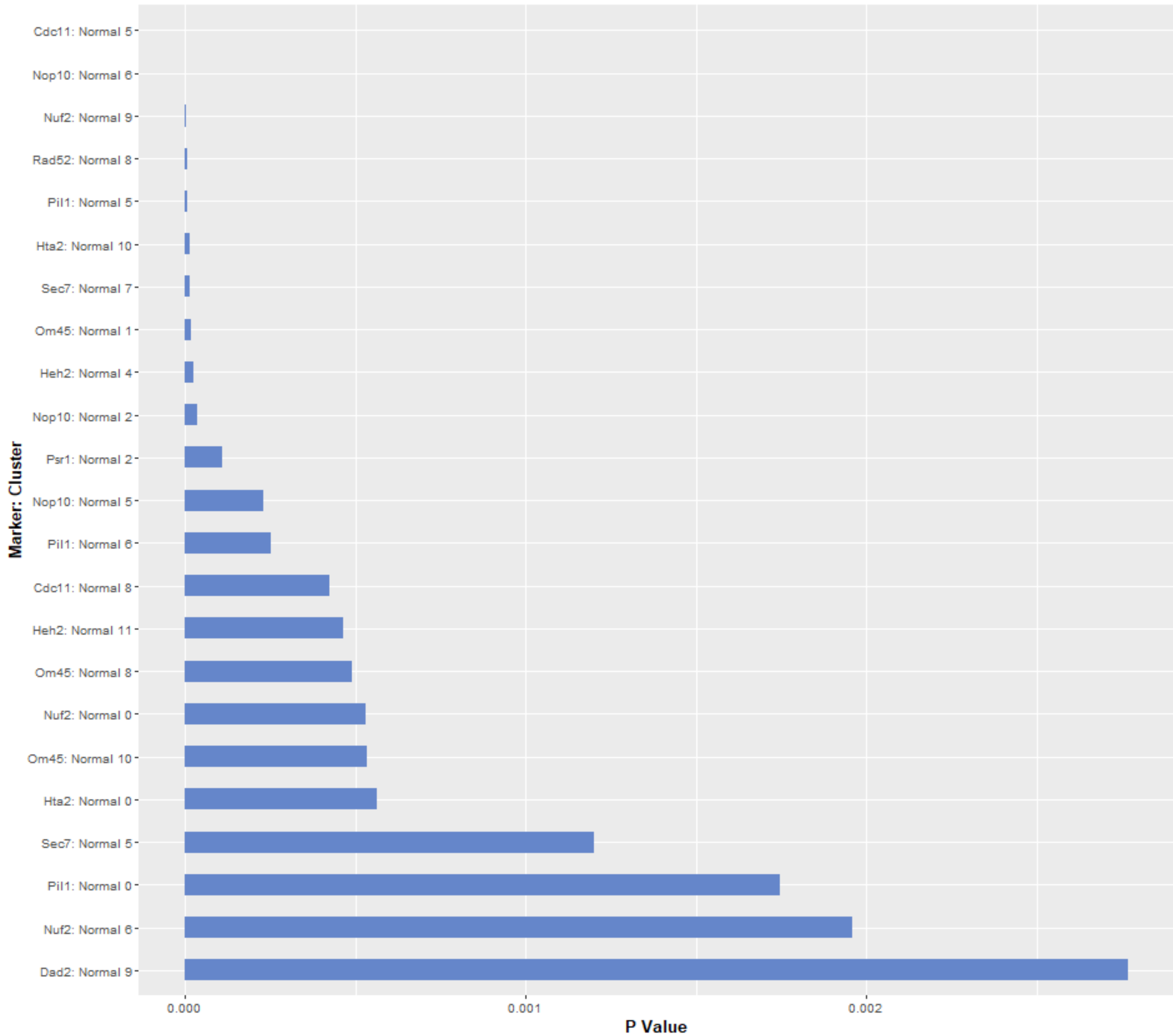
With AreaShape | Unordered Gene Set

Marker: Cluster



DNA recombination

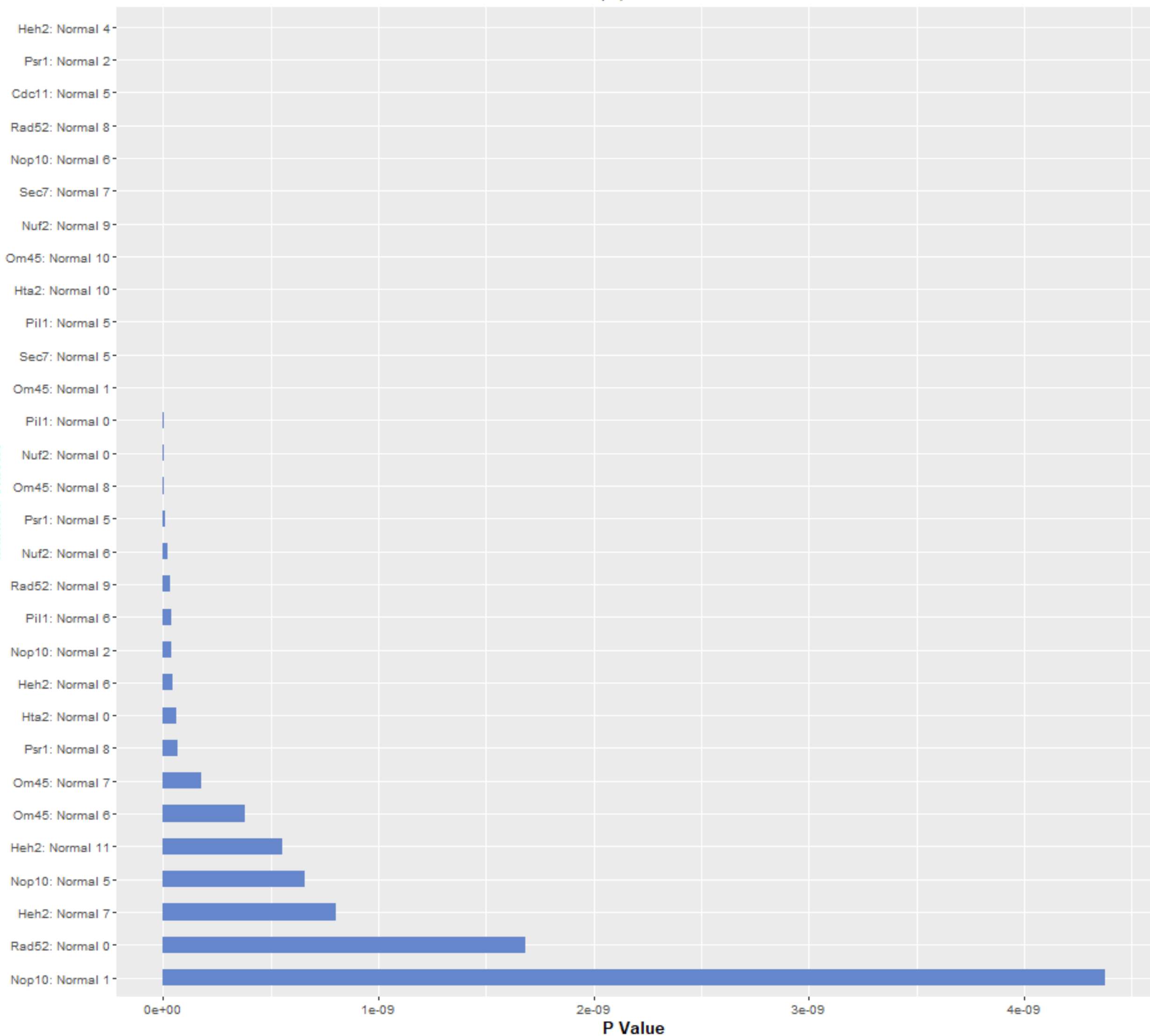
With AreaShape | Unordered Gene Set



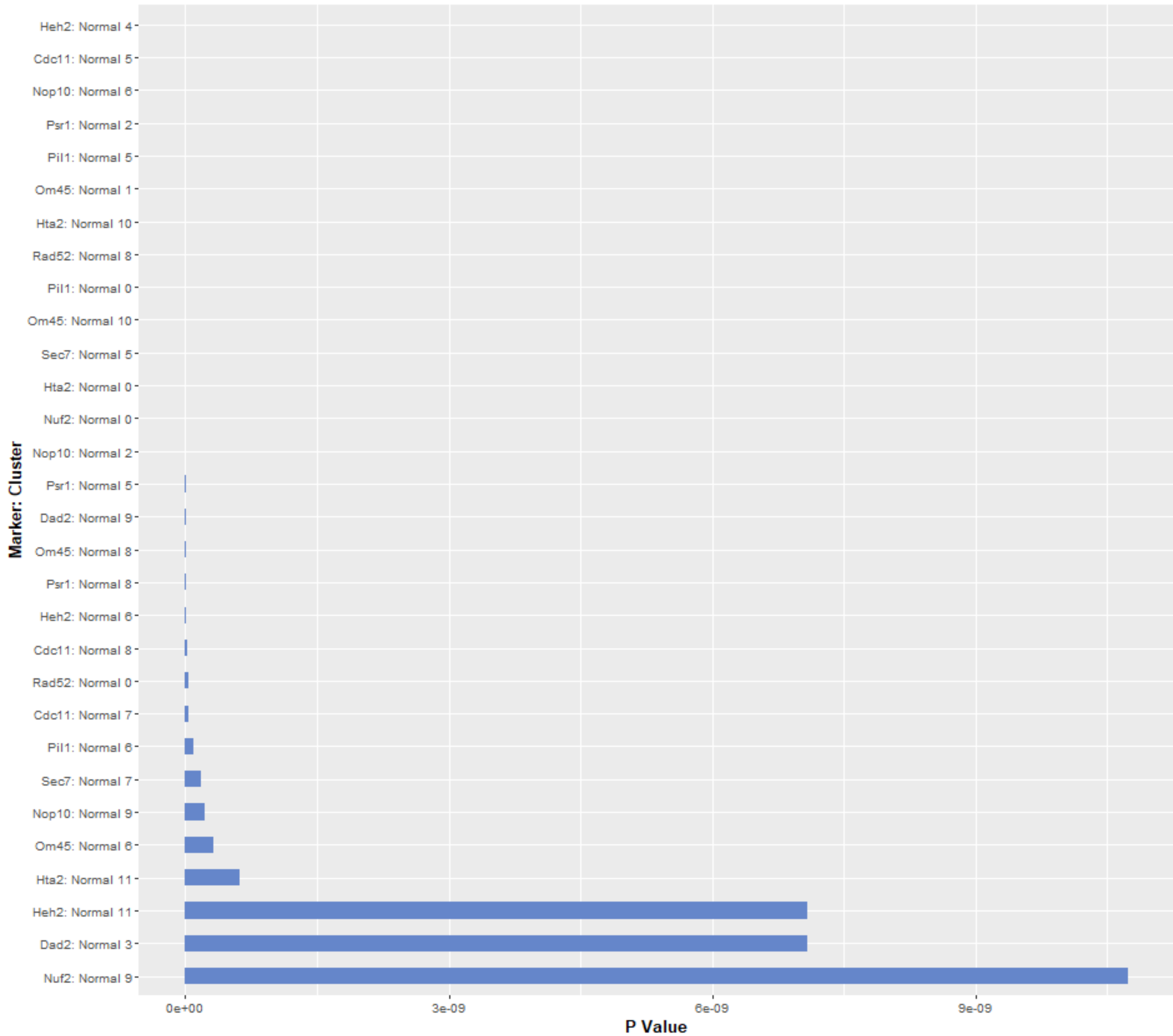
DNA repair

With AreaShape | Unordered Gene Set

Marker: Cluster



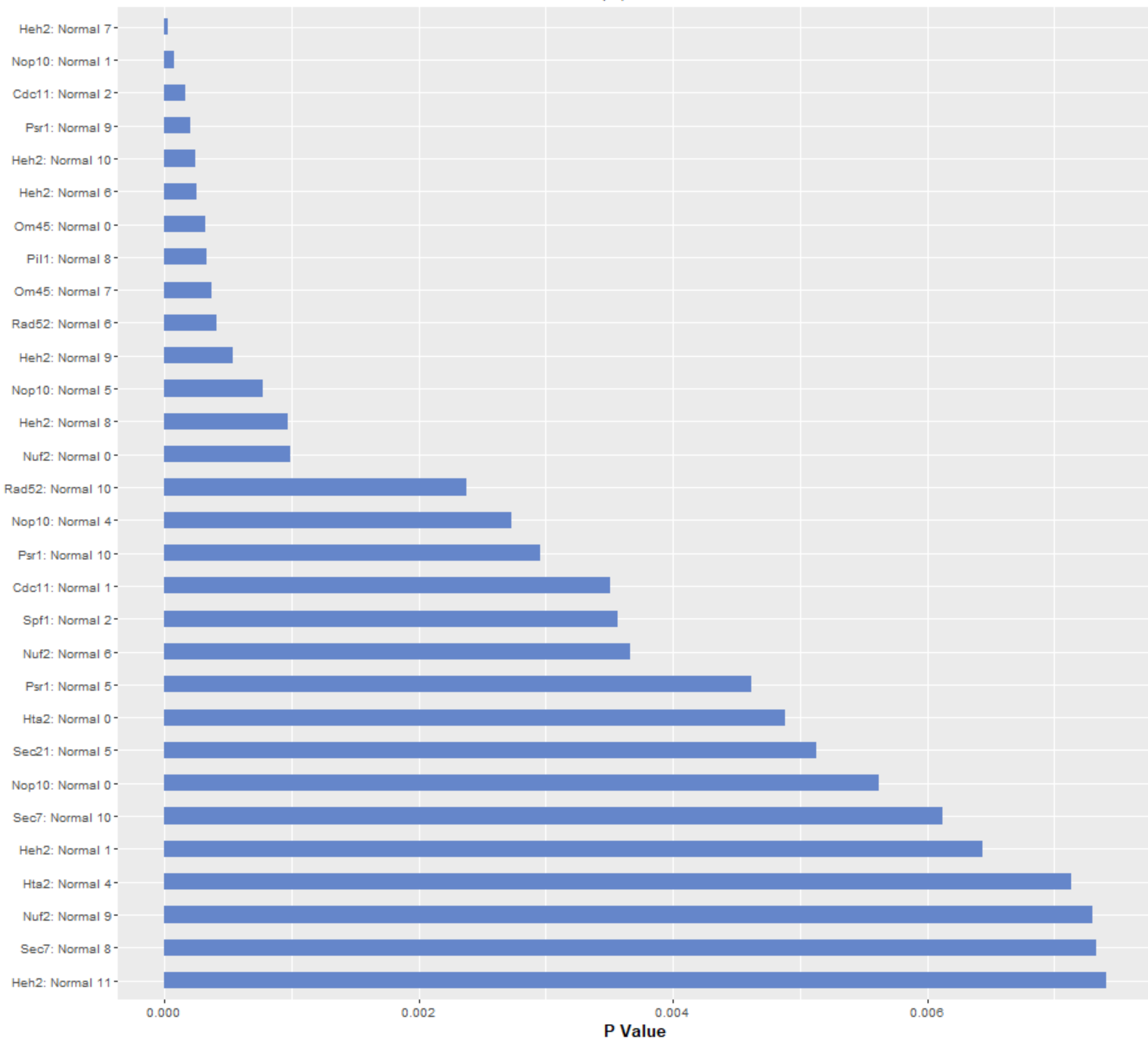
DNA replication
With AreaShape | Unordered Gene Set



DNA-templated transcription, elongation

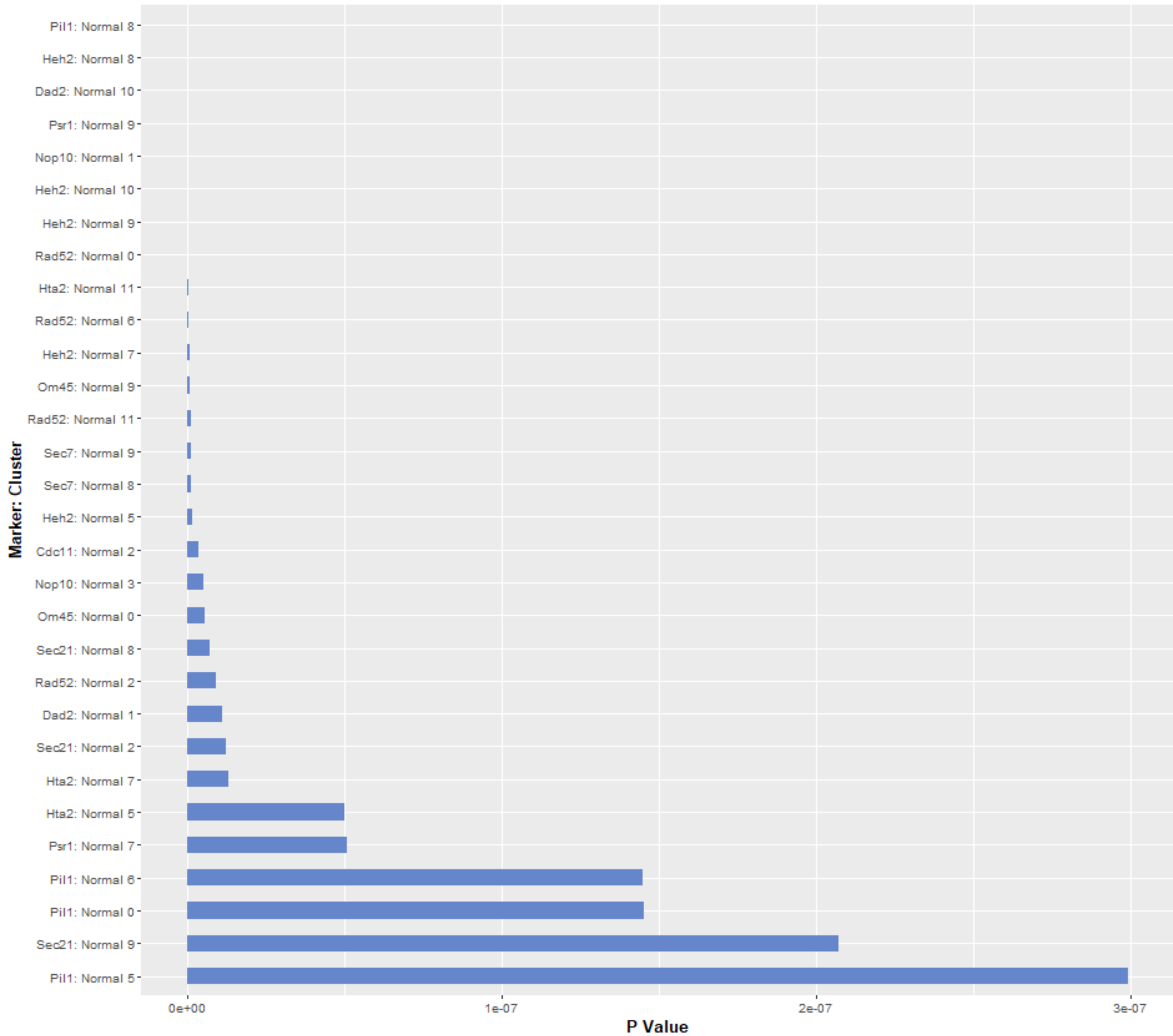
With AreaShape | Unordered Gene Set

Marker: Cluster



DNA-templated transcription, initiation

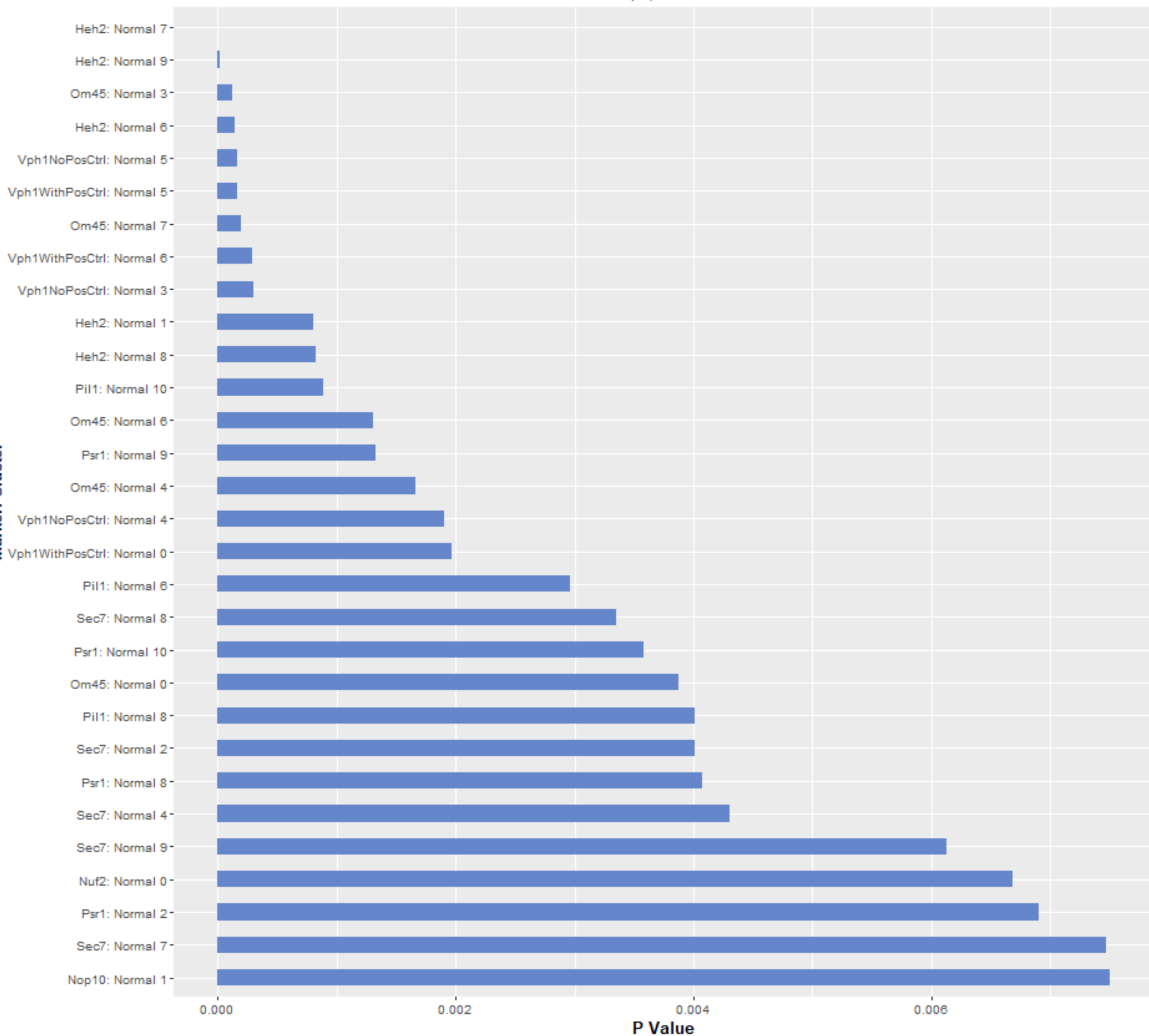
With AreaShape | Unordered Gene Set



DNA-templated transcription, termination

With AreaShape | Unordered Gene Set

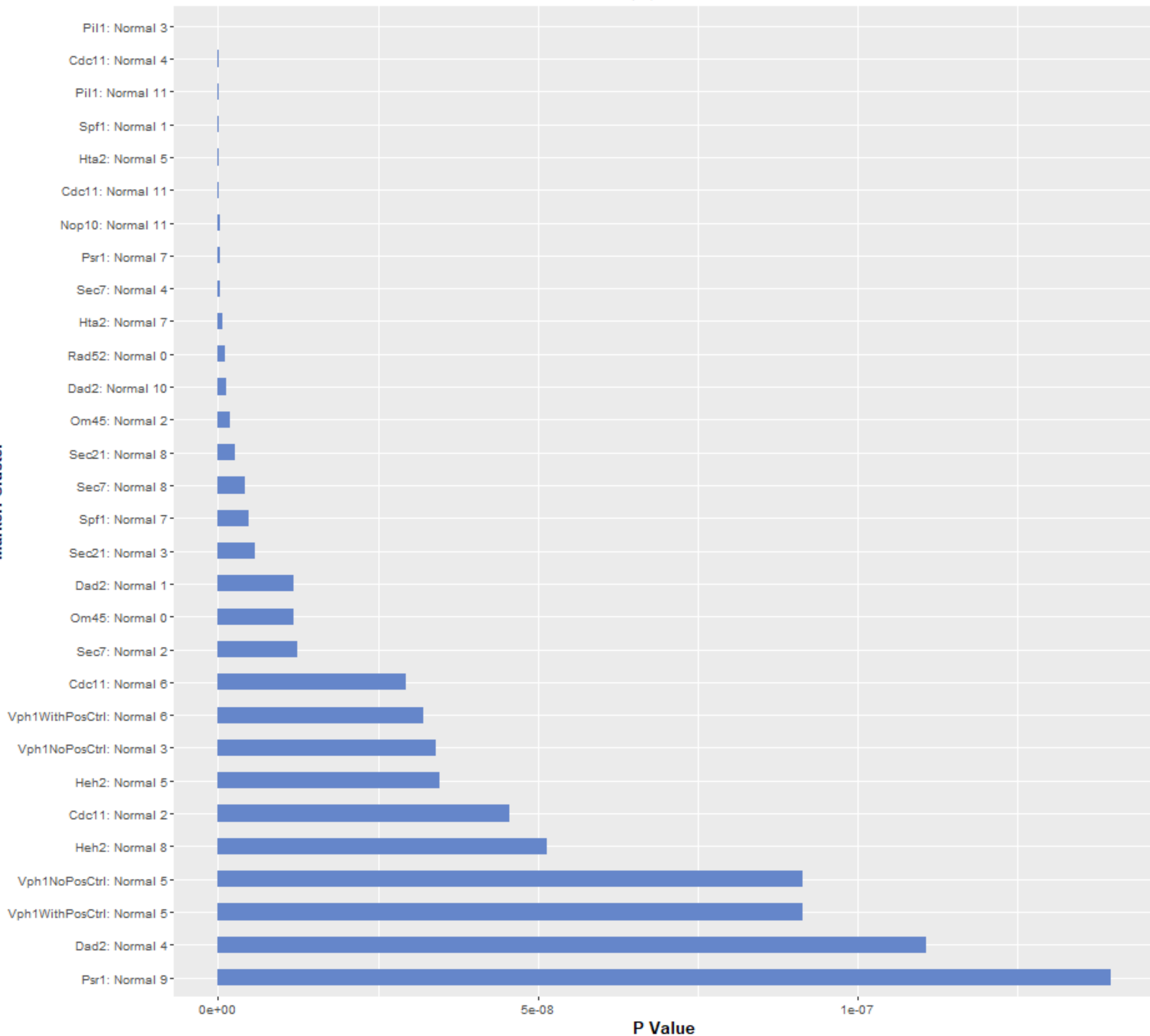
Marker: Cluster



exocytosis

With AreaShape | Unordered Gene Set

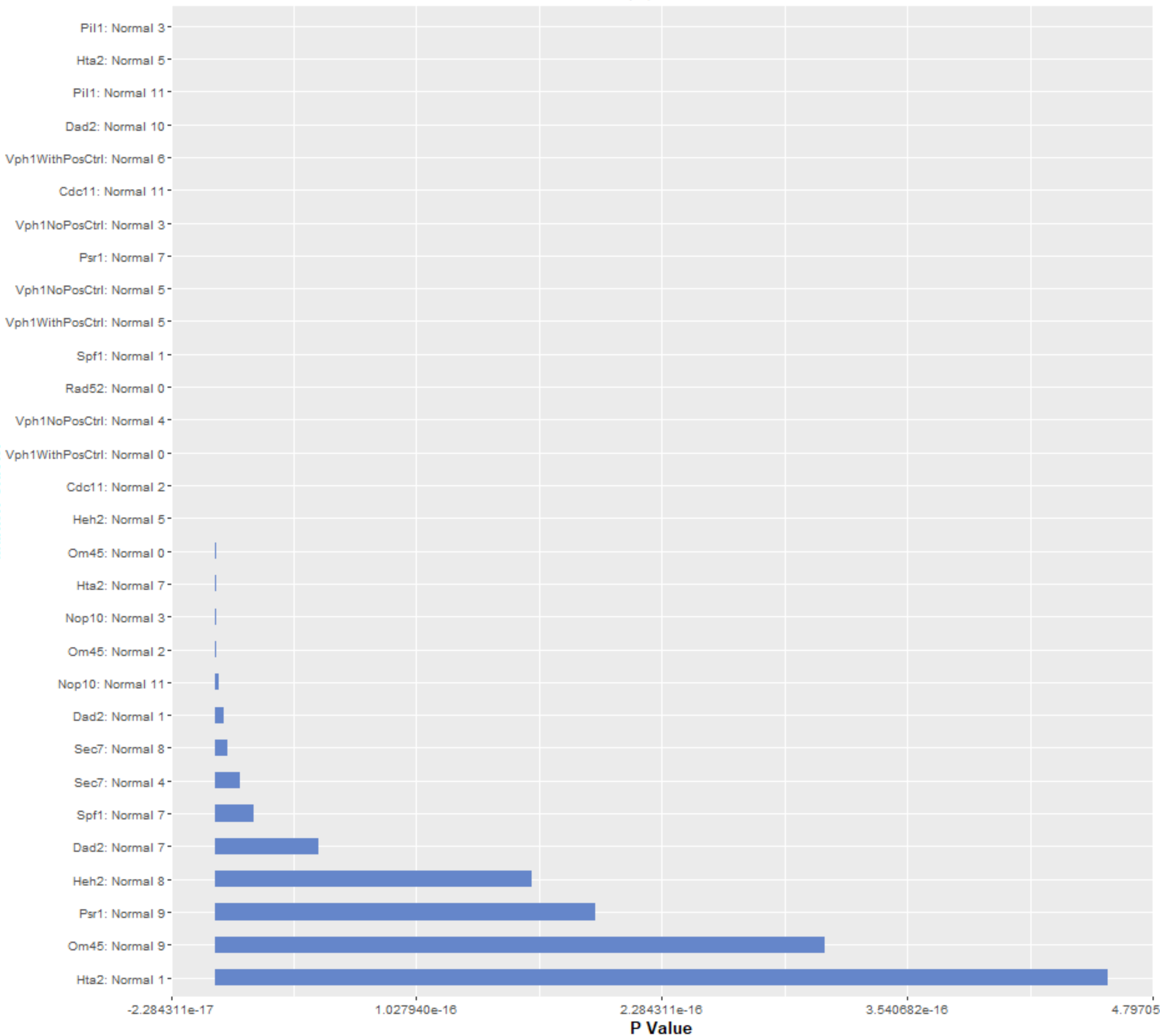
Marker: Cluster



Golgi vesicle transport

With AreaShape | Unordered Gene Set

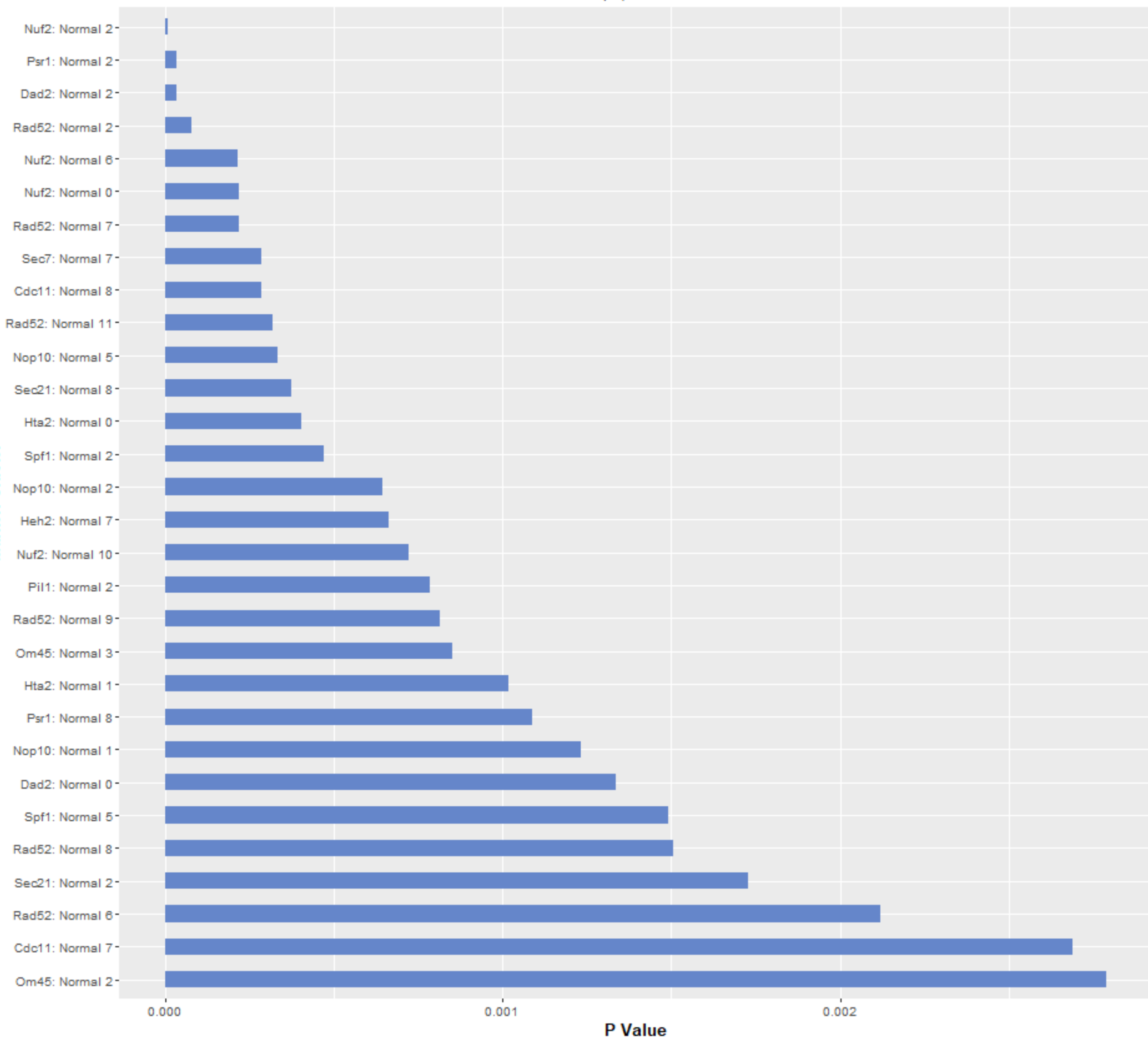
Marker: Cluster



histone modification

With AreaShape | Unordered Gene Set

Marker: Cluster



invasive growth in response to glucose limitation

With AreaShape | Unordered Gene Set

Marker: Cluster

Psr1: Normal 0

Rad52: Normal 1

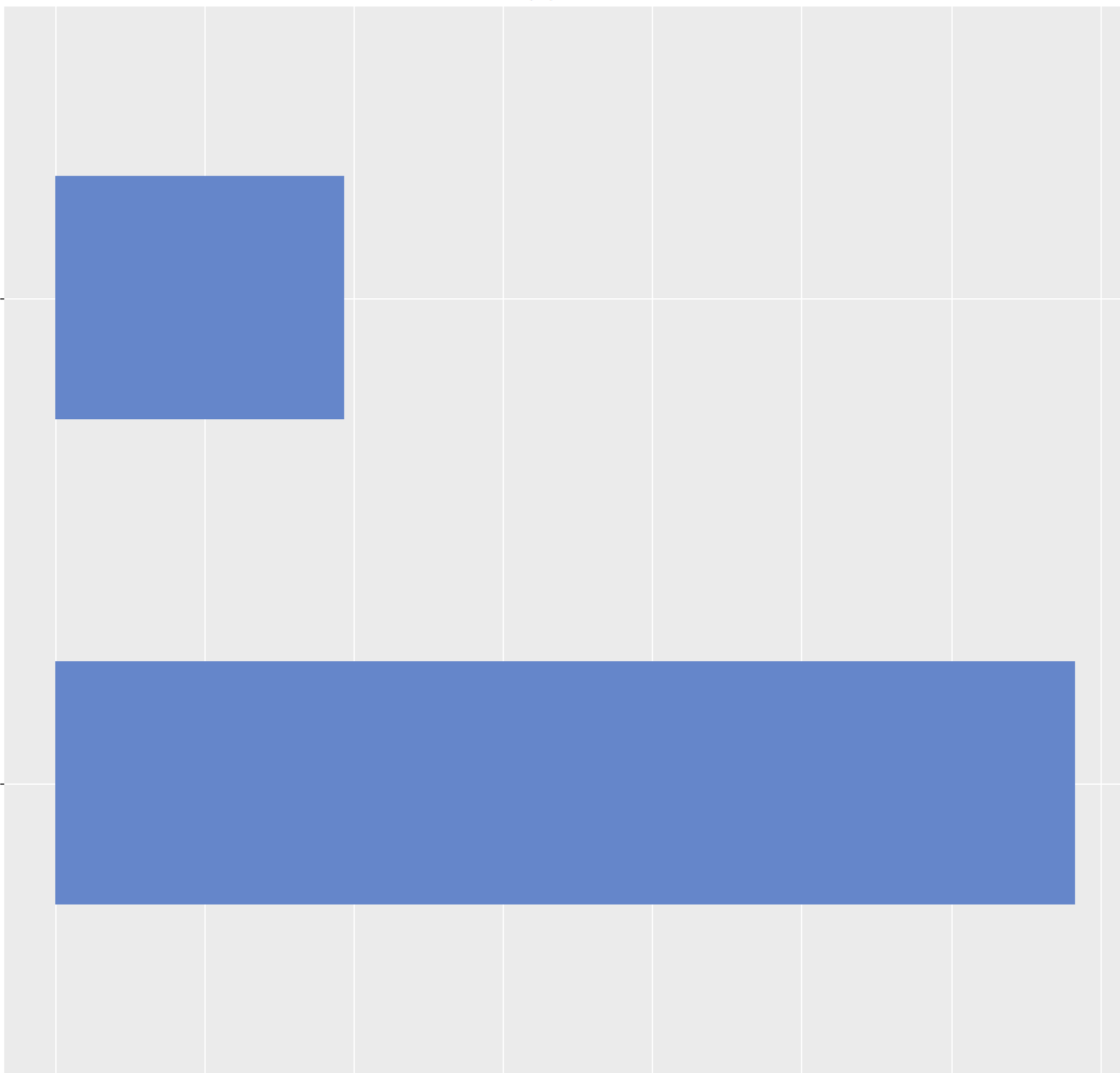
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0.0025

0.0050

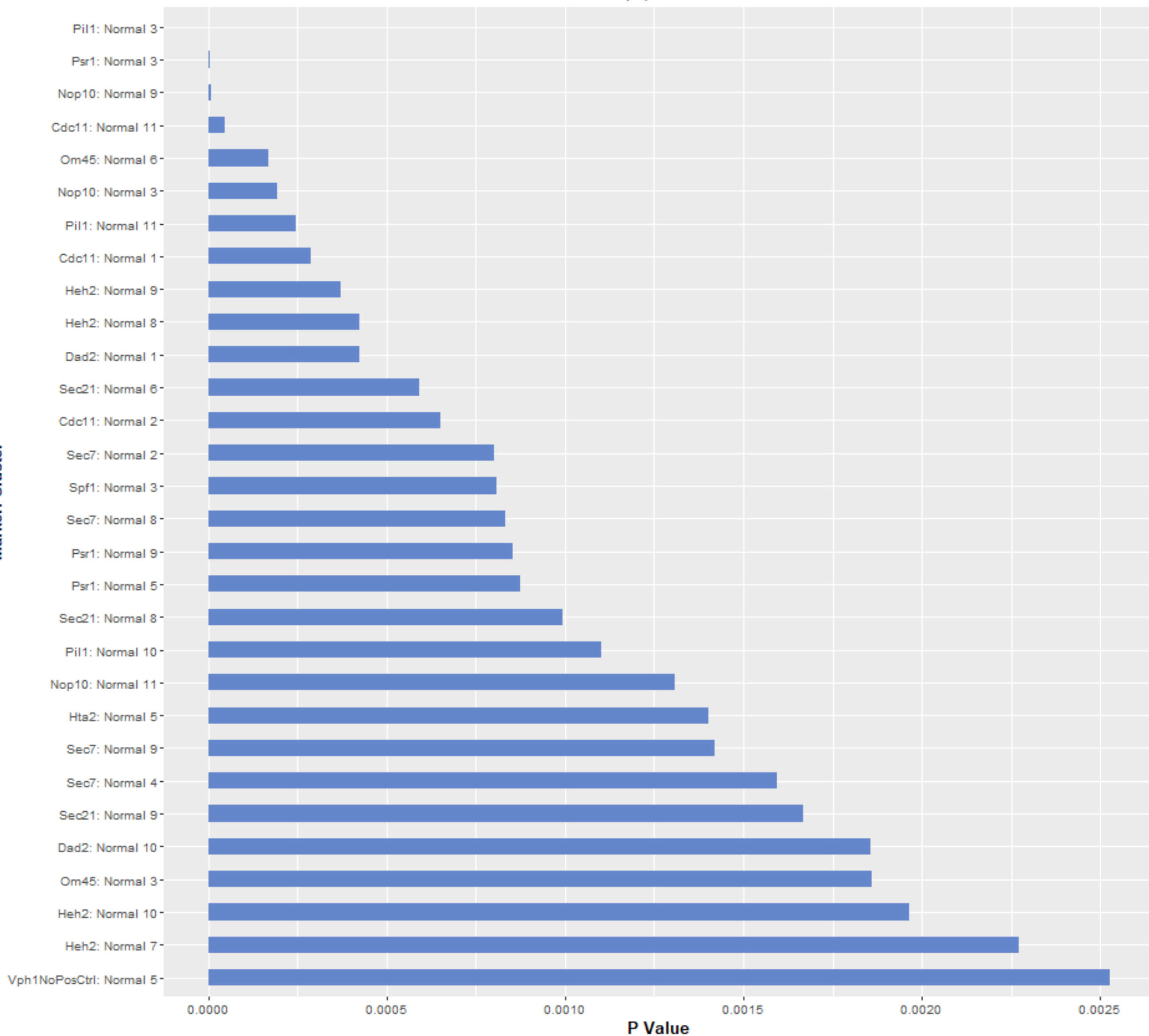
0.0075

P Value

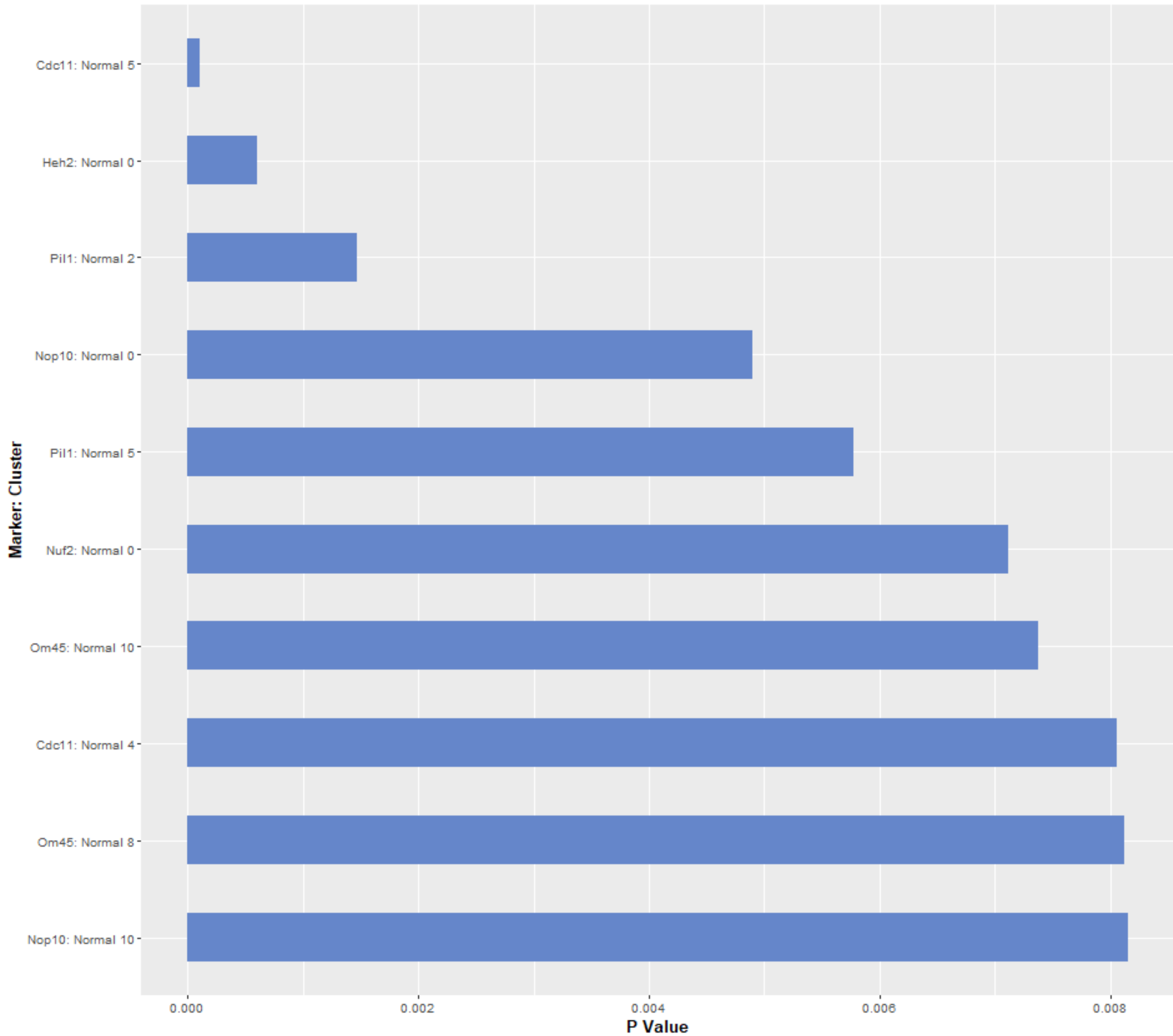


lipid metabolic process
With AreaShape | Unordered Gene Set

Marker: Cluster



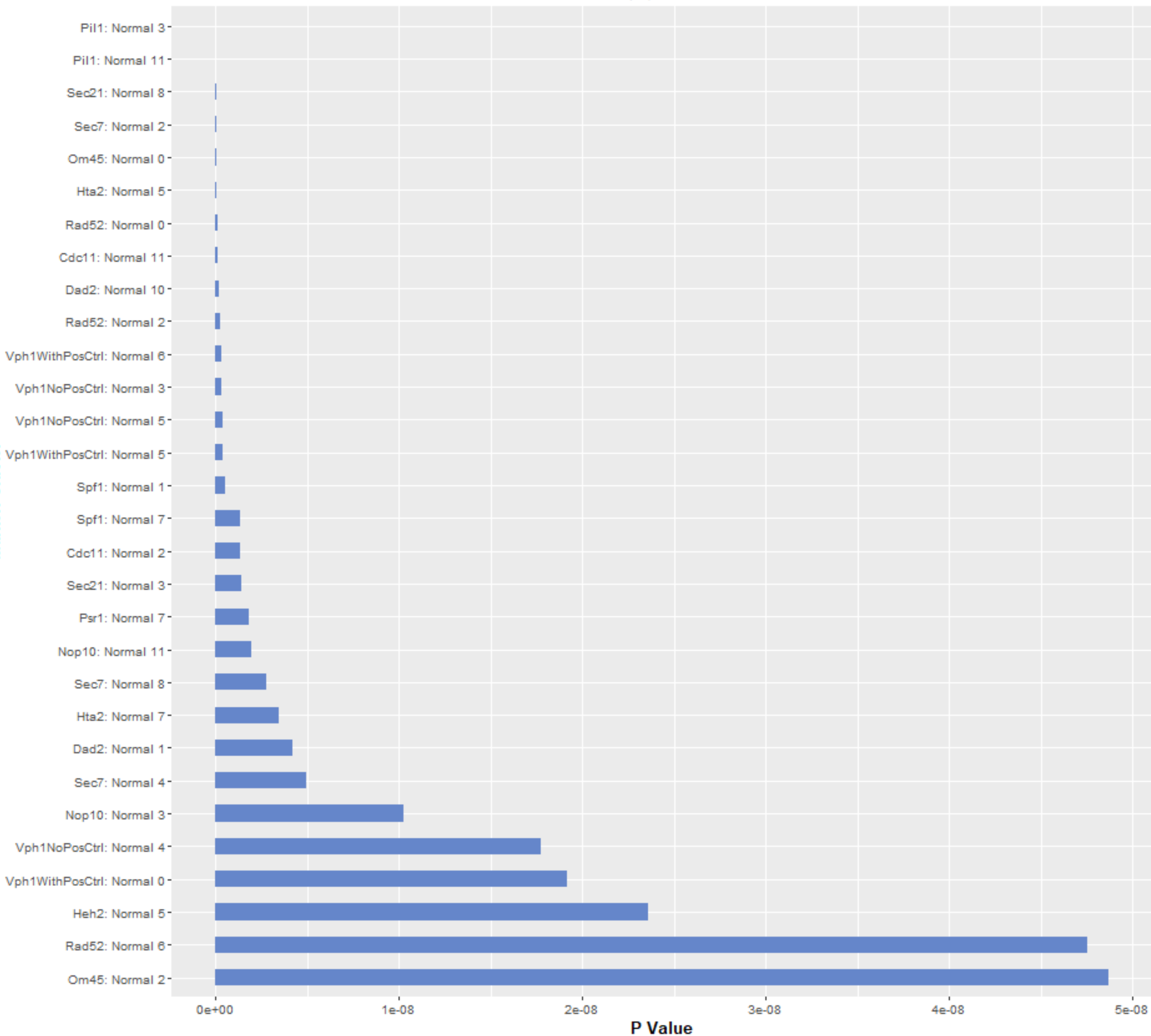
meiotic cell cycle
With AreaShape | Unordered Gene Set



membrane fusion

With AreaShape | Unordered Gene Set

Marker: Cluster



mitochondrion organization

With AreaShape | Unordered Gene Set

Marker: Cluster

Om45: Normal 3

0e+00

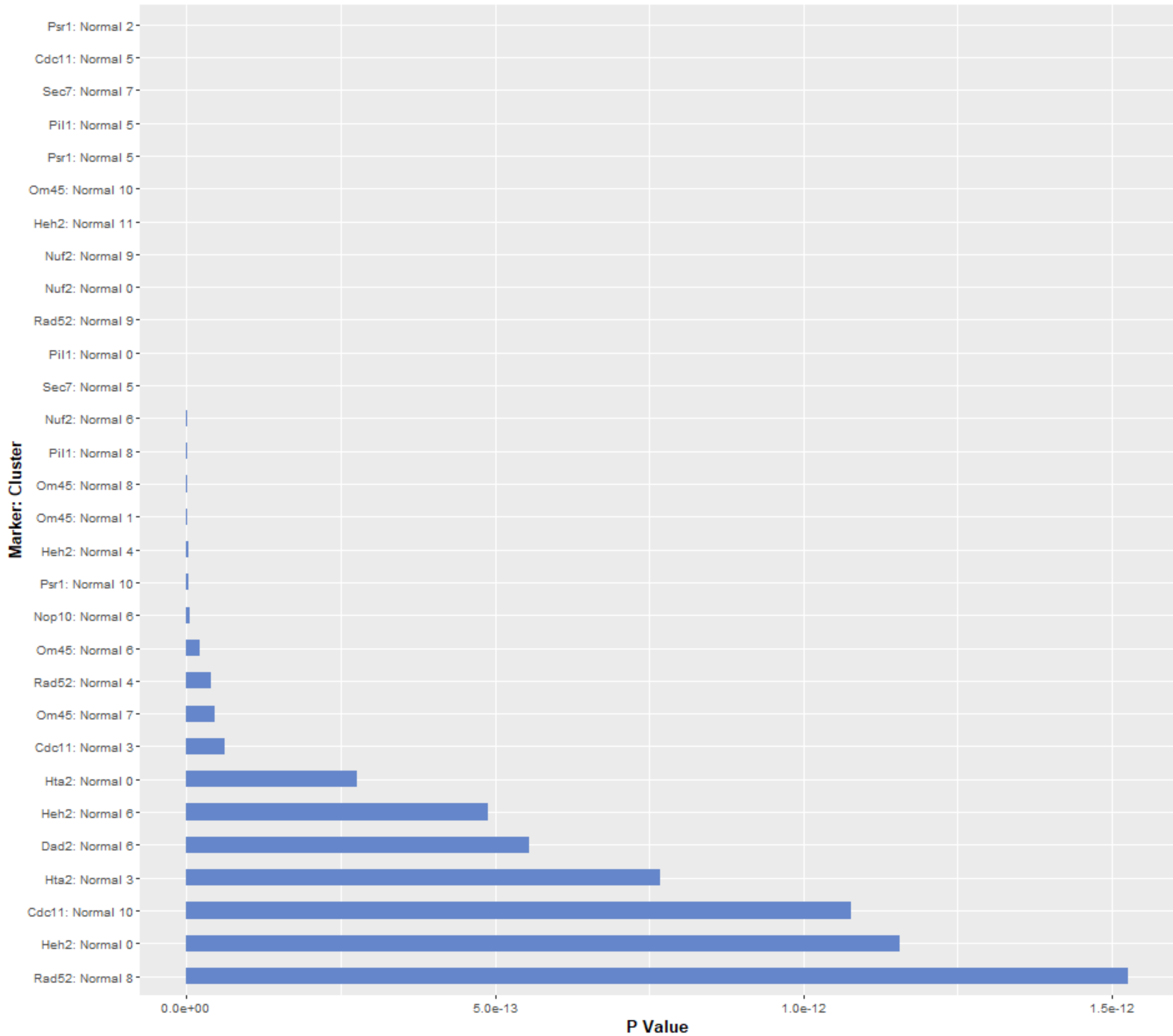
5e-04

1e-03

P Value

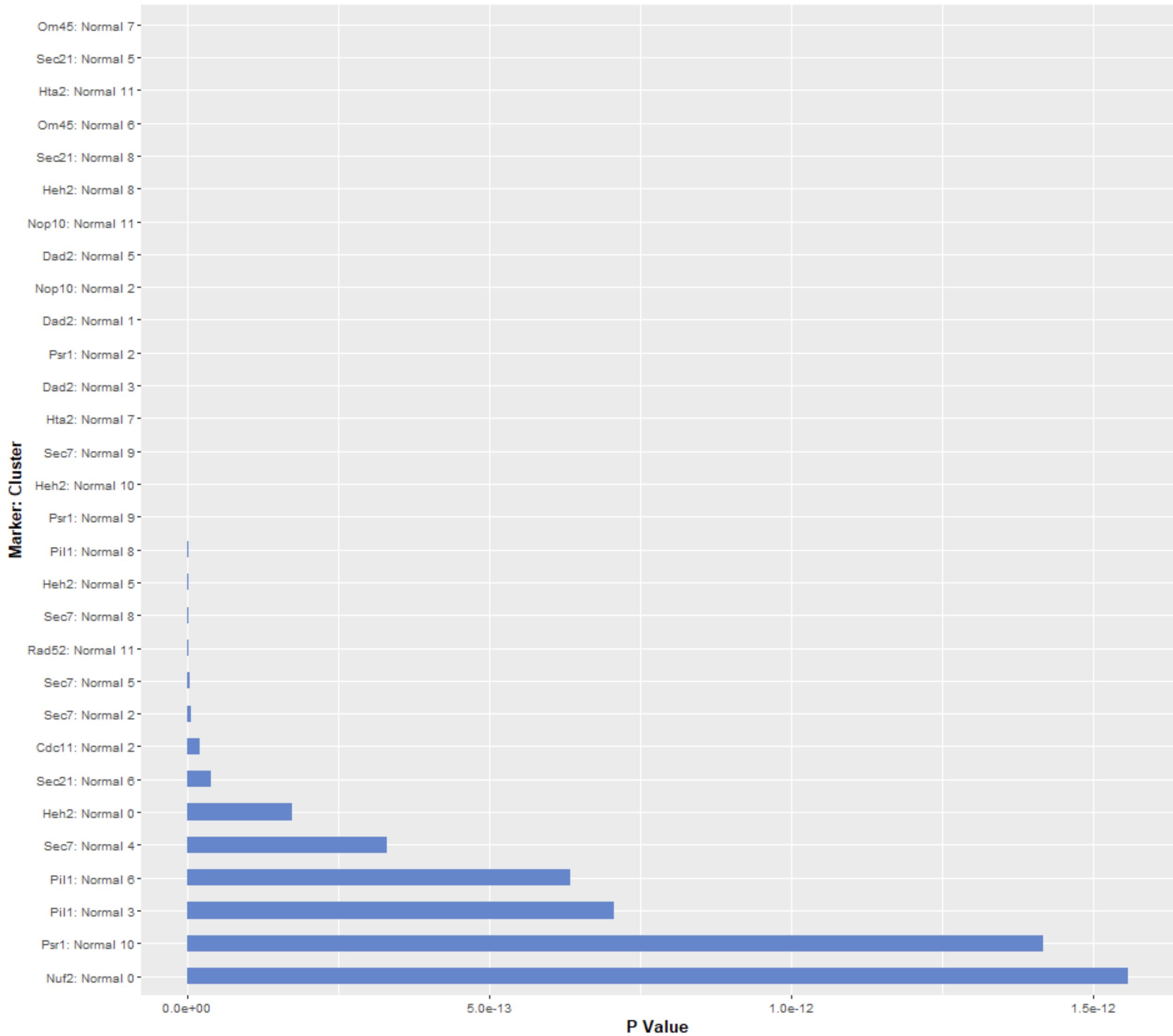


mitotic cell cycle
With AreaShape | Unordered Gene Set



mRNA processing

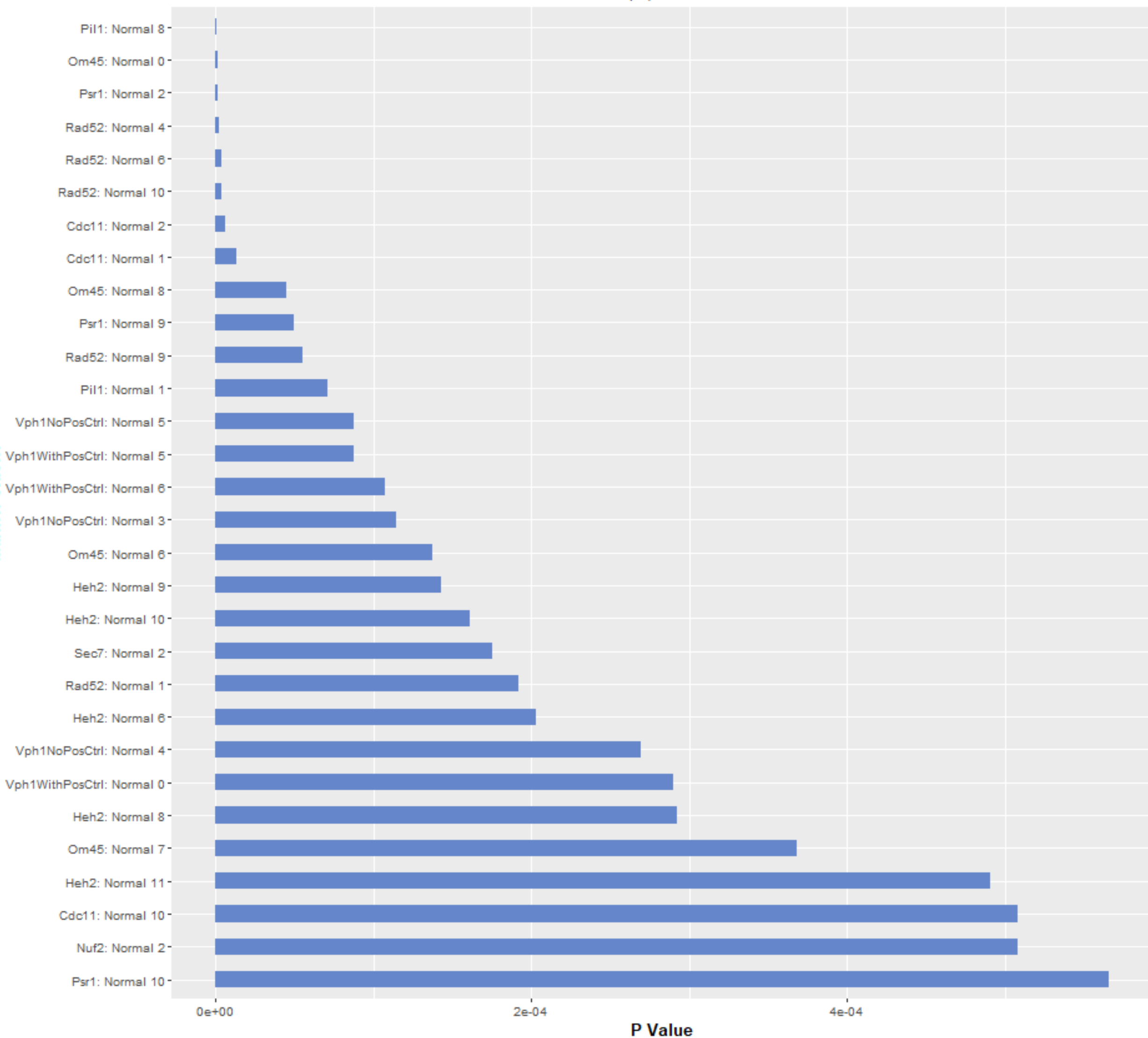
With AreaShape | Unordered Gene Set



nuclear transport

With AreaShape | Unordered Gene Set

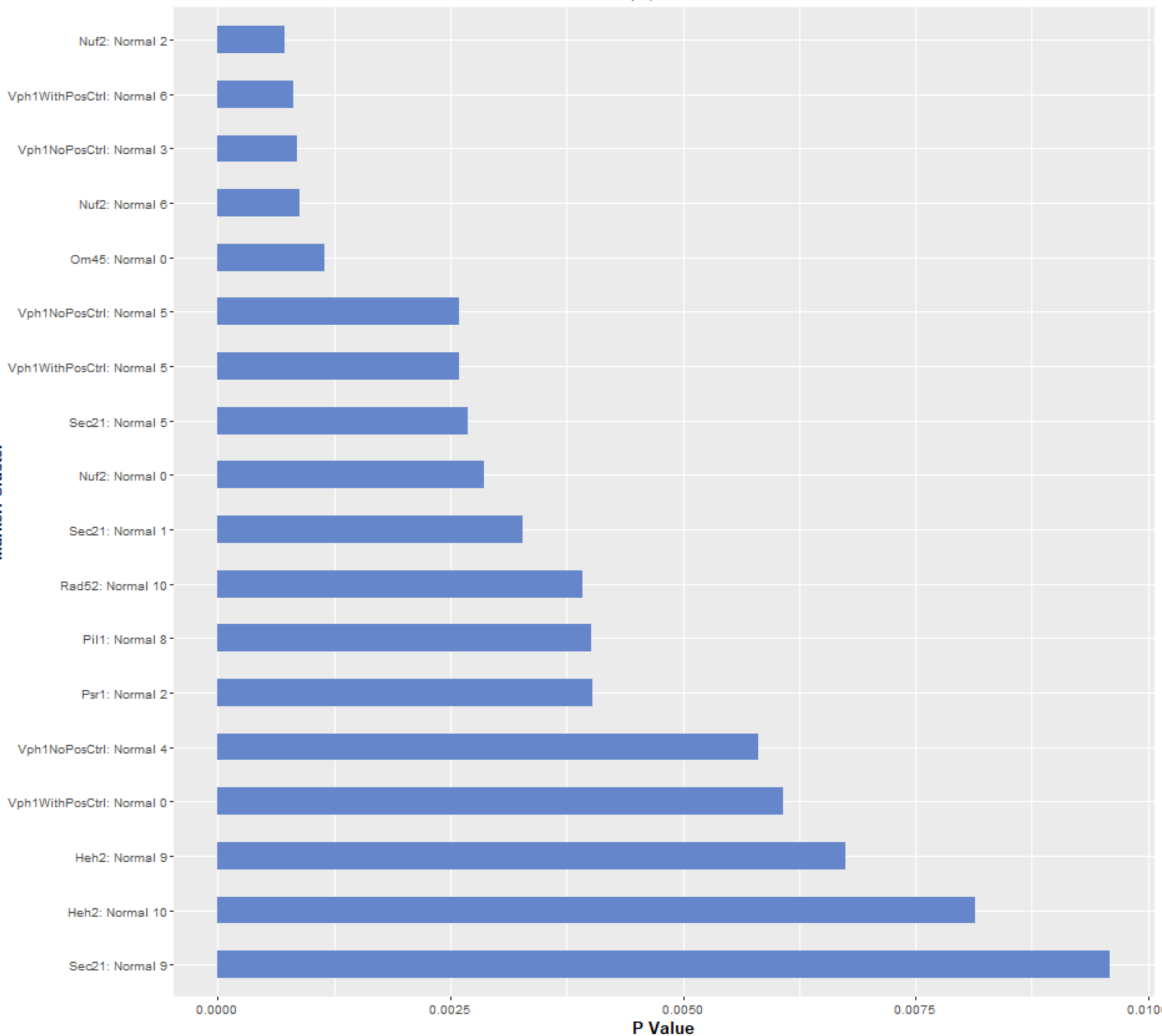
Marker: Cluster



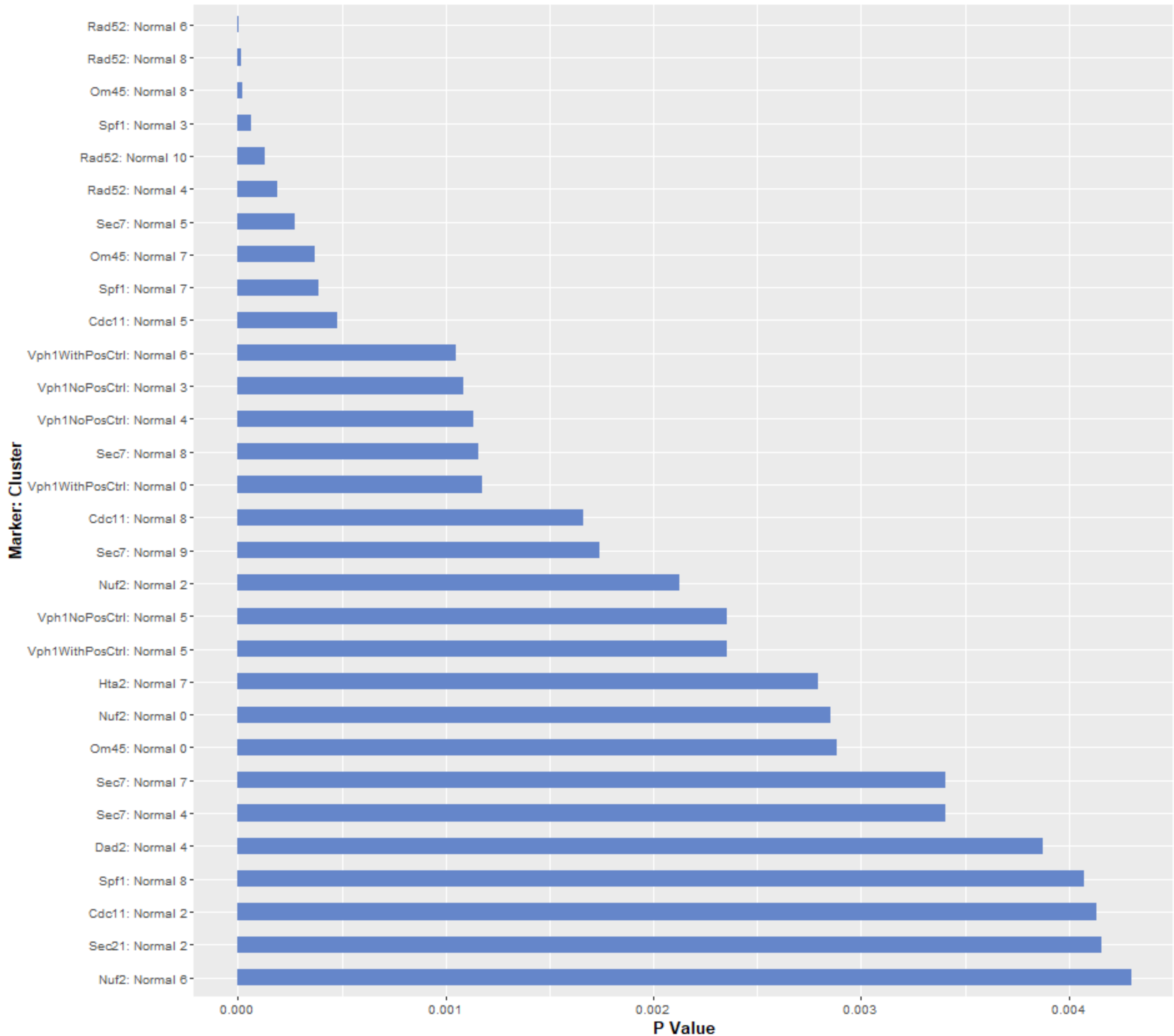
nucleobase-containing compound transport

With AreaShape | Unordered Gene Set

Marker: Cluster

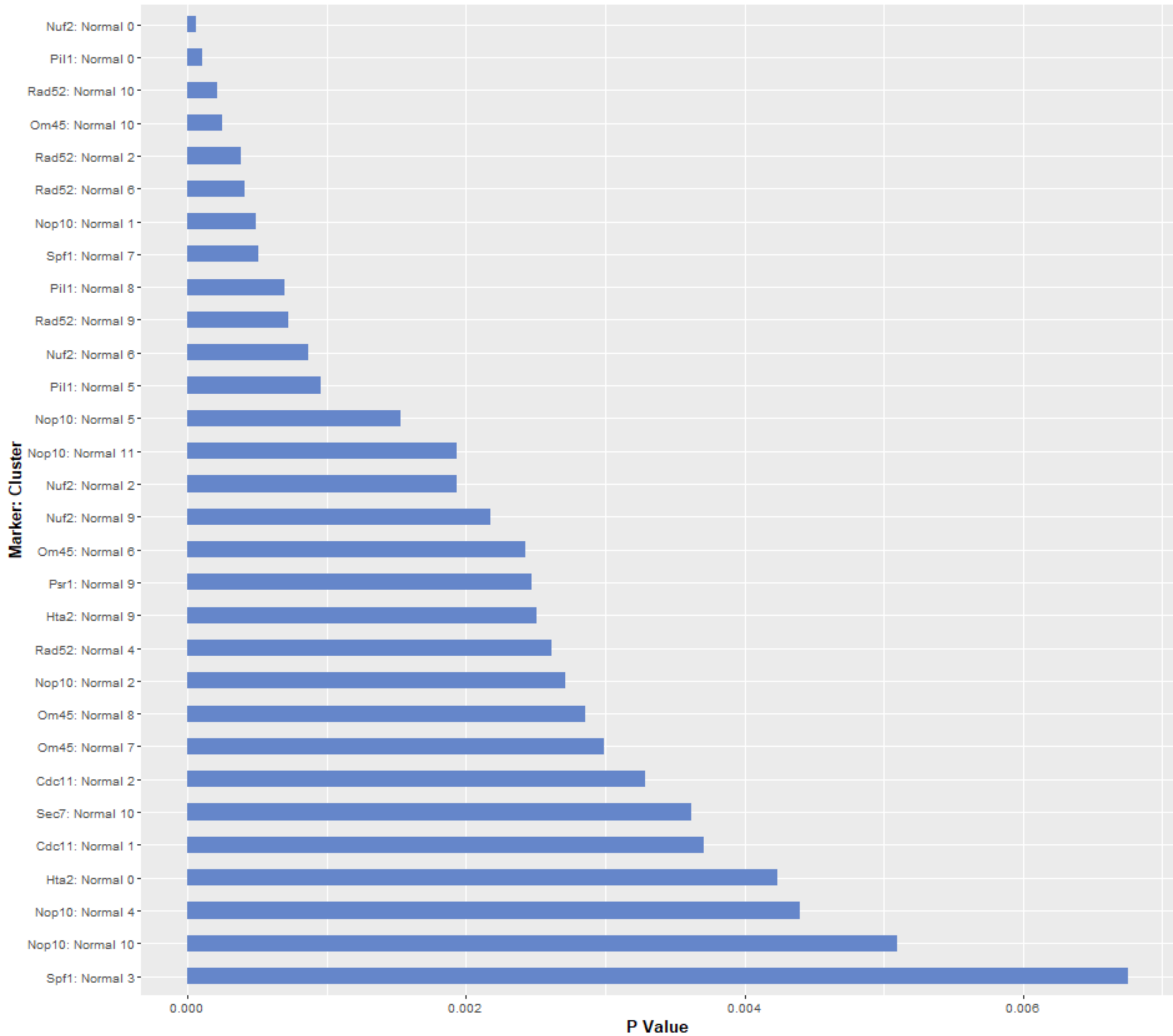


nucleus organization
With AreaShape | Unordered Gene Set

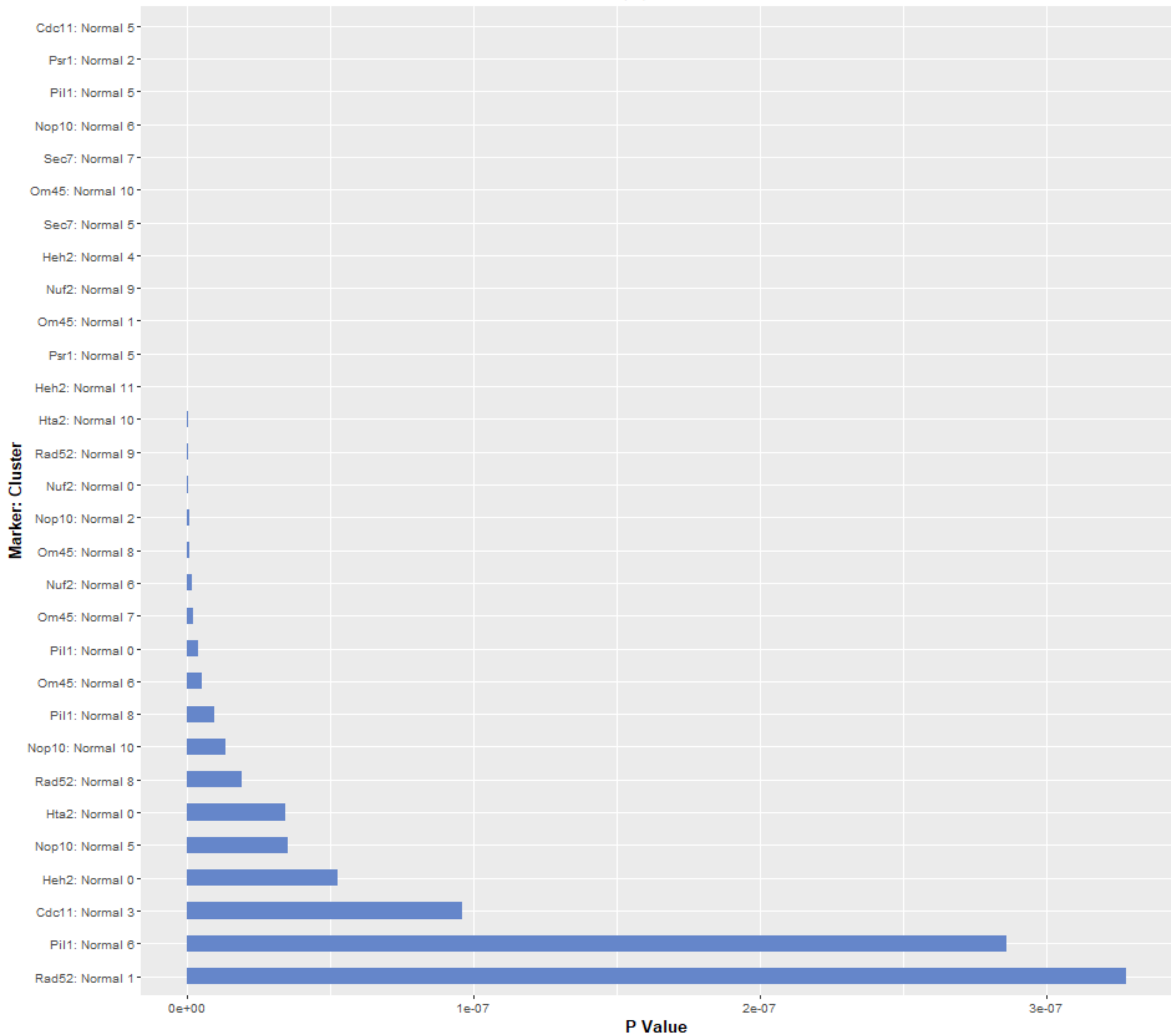


organelle assembly

With AreaShape | Unordered Gene Set



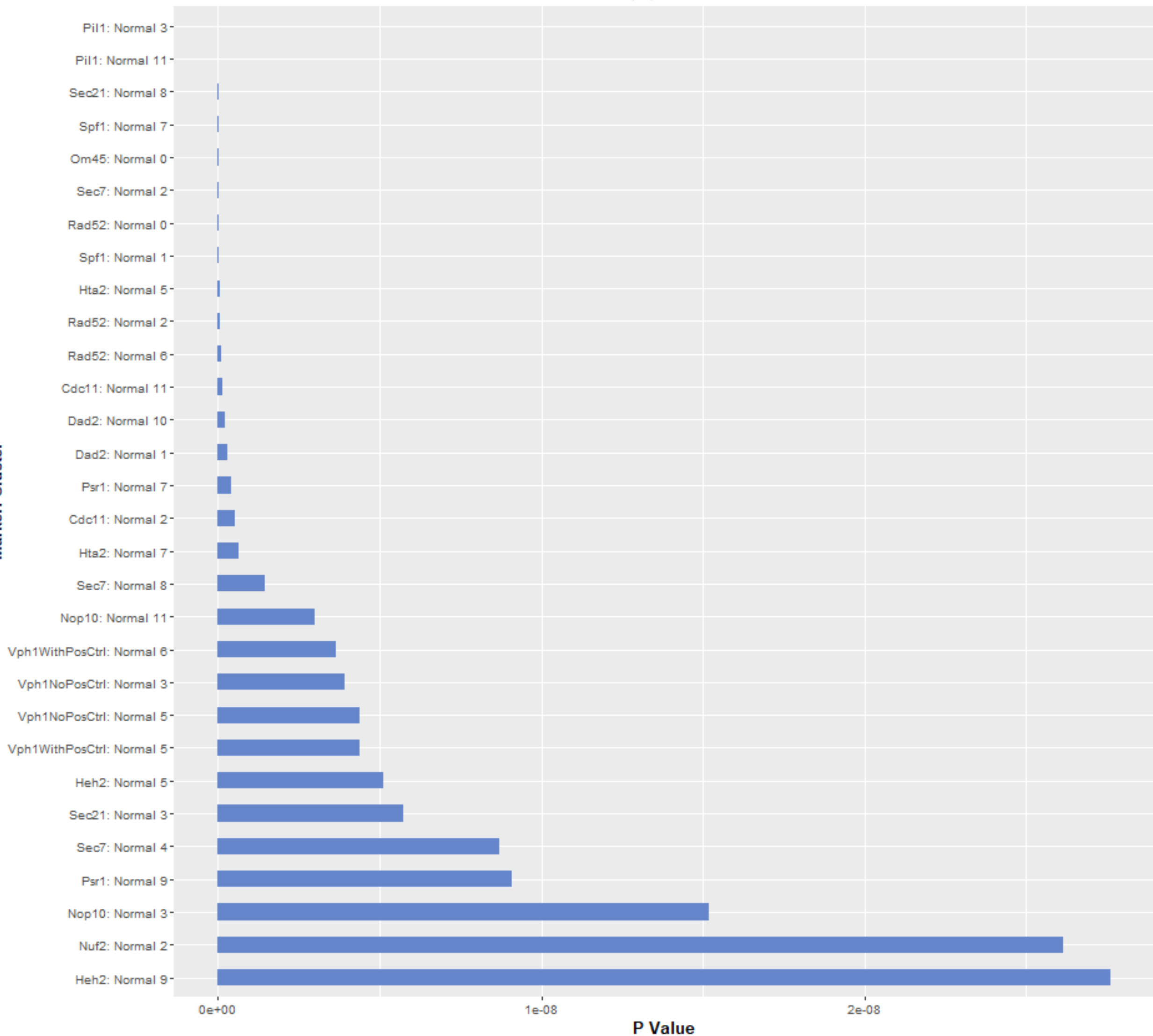
organelle fission
With AreaShape | Unordered Gene Set



organelle fusion

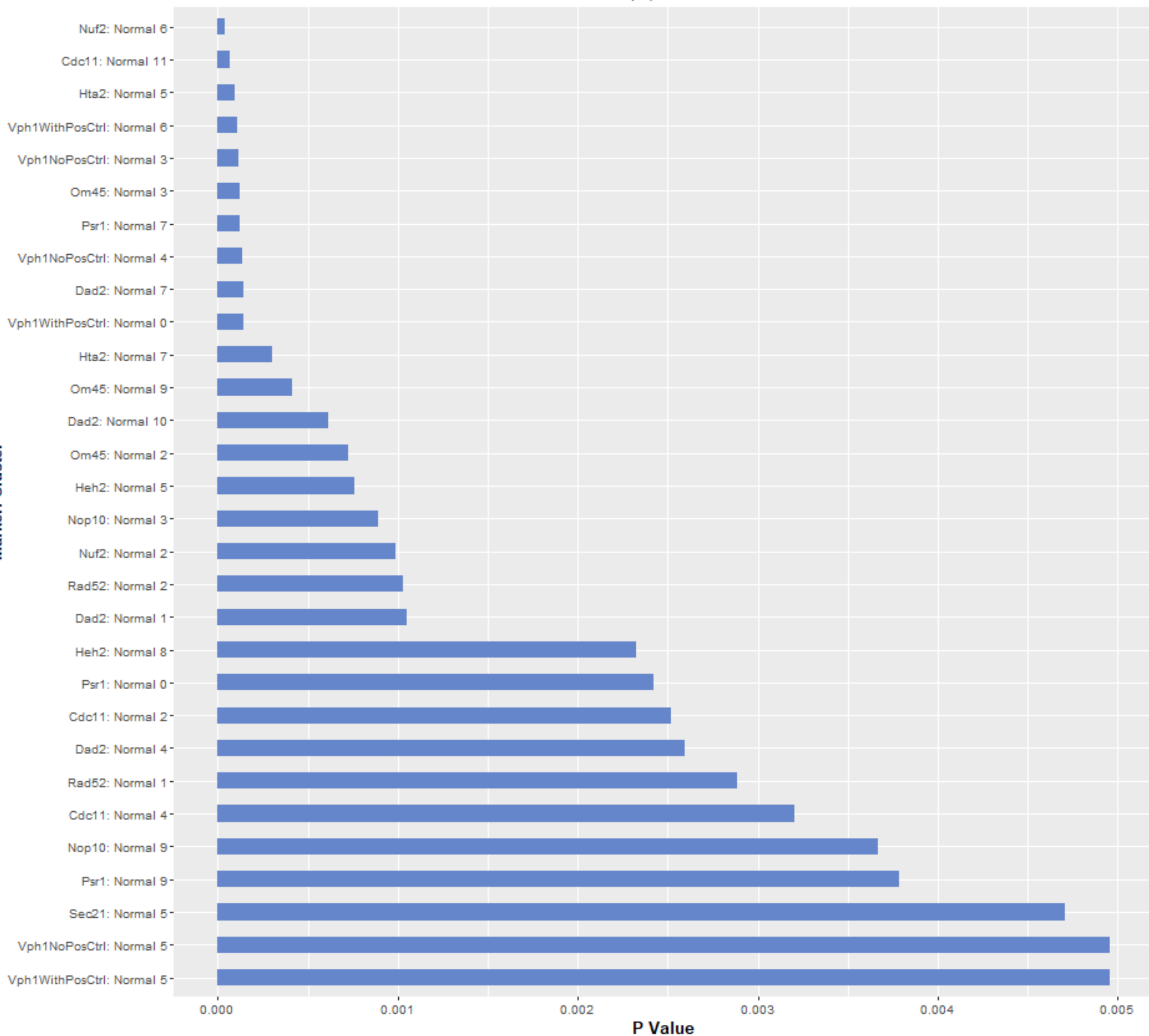
With AreaShape | Unordered Gene Set

Marker: Cluster



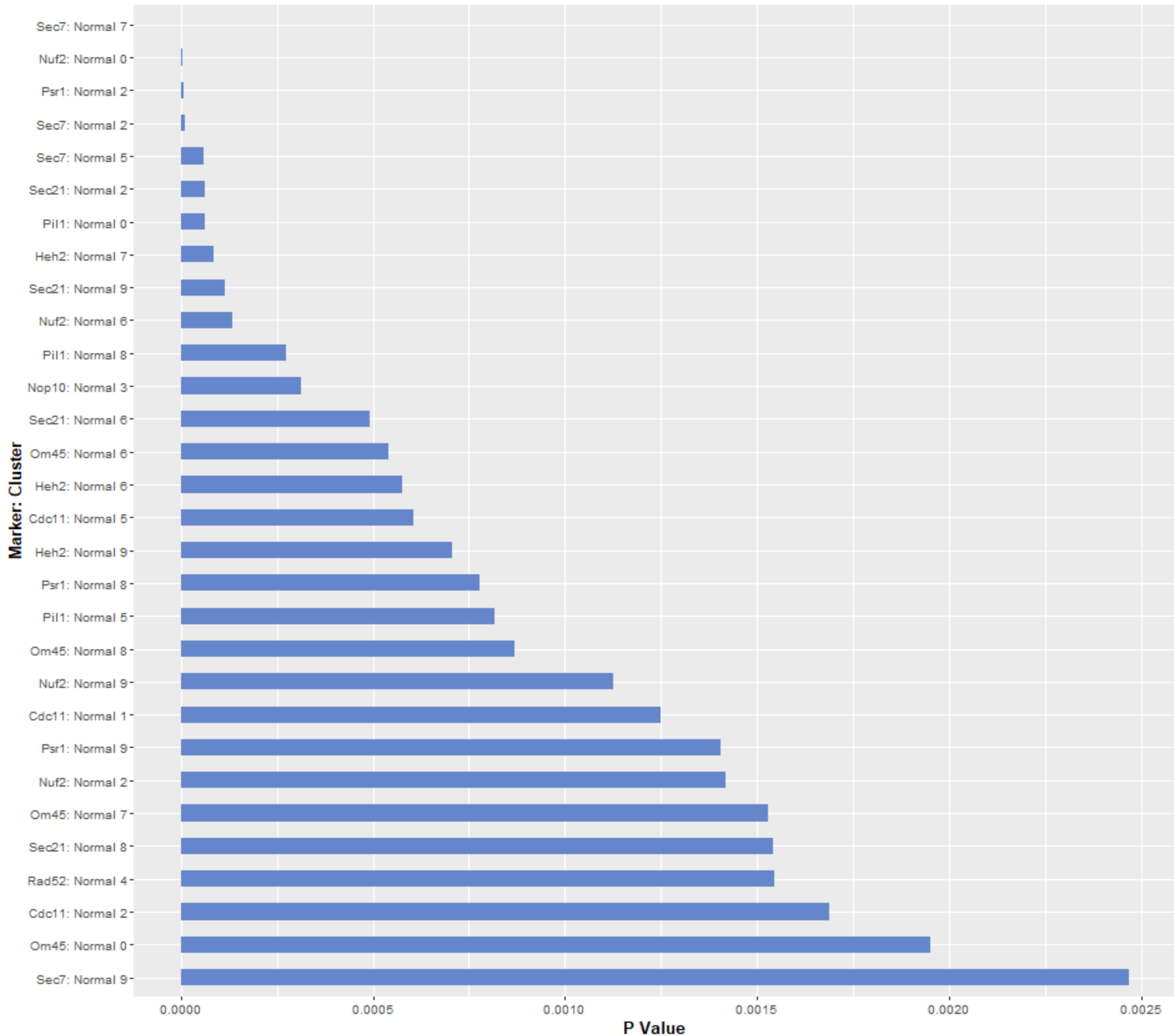
organelle inheritance
With AreaShape | Unordered Gene Set

Marker: Cluster

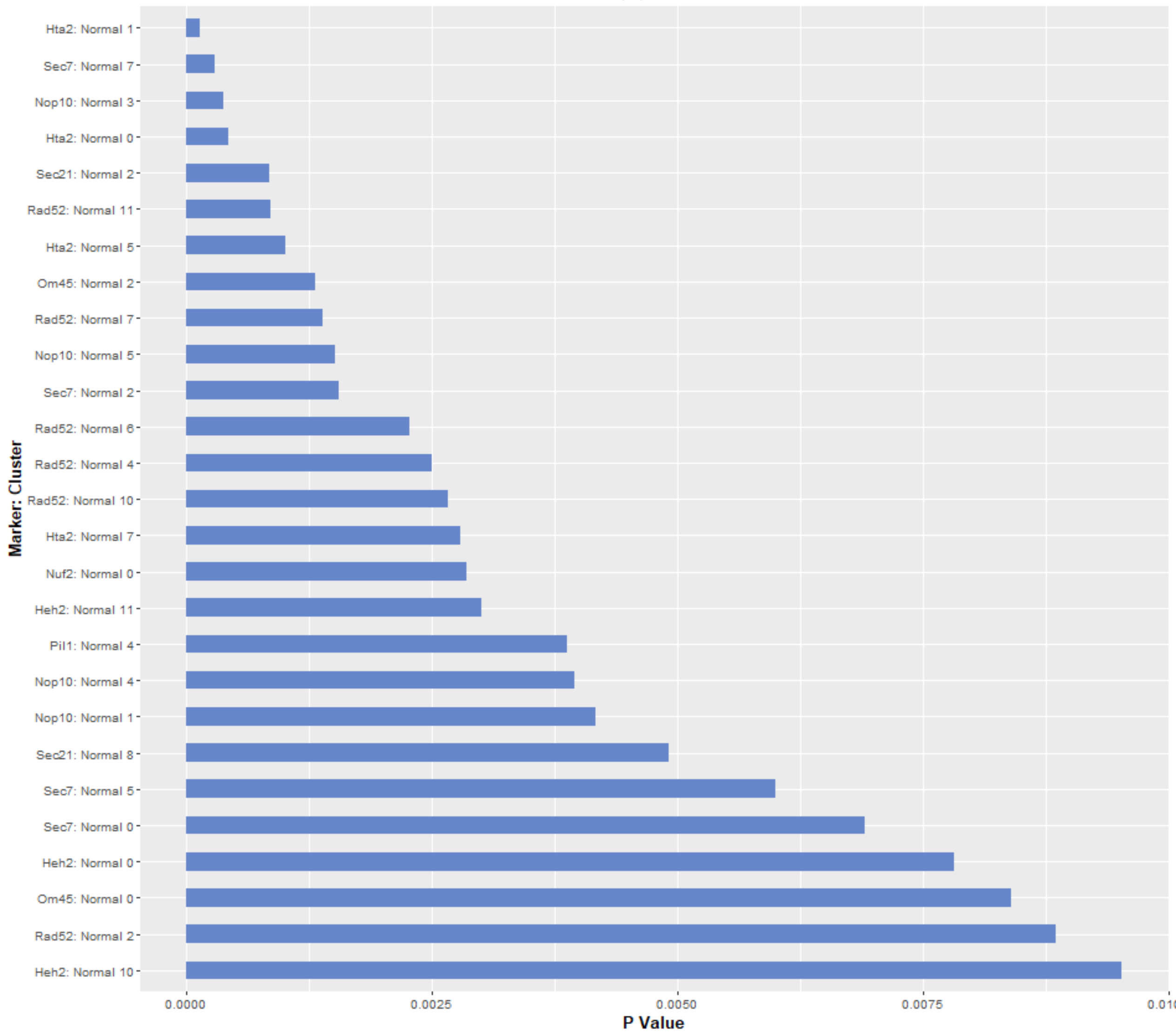


peptidyl-amino acid modification

With AreaShape | Unordered Gene Set



protein acylation
With AreaShape | Unordered Gene Set



protein alkylation

With AreaShape | Unordered Gene Set

Marker: Cluster

Dad2: Normal 2

Sec21: Normal 1

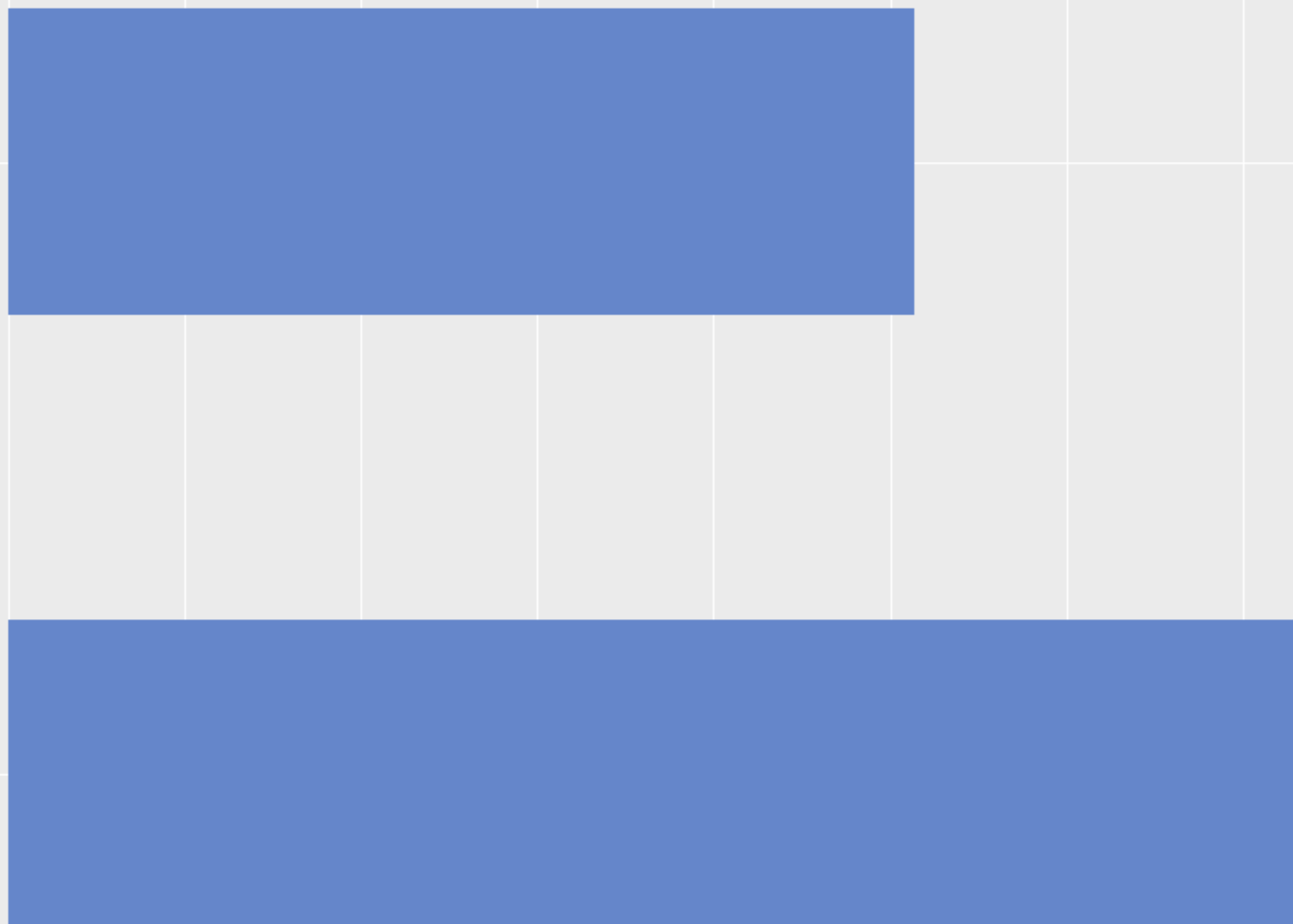
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0.002

0.004

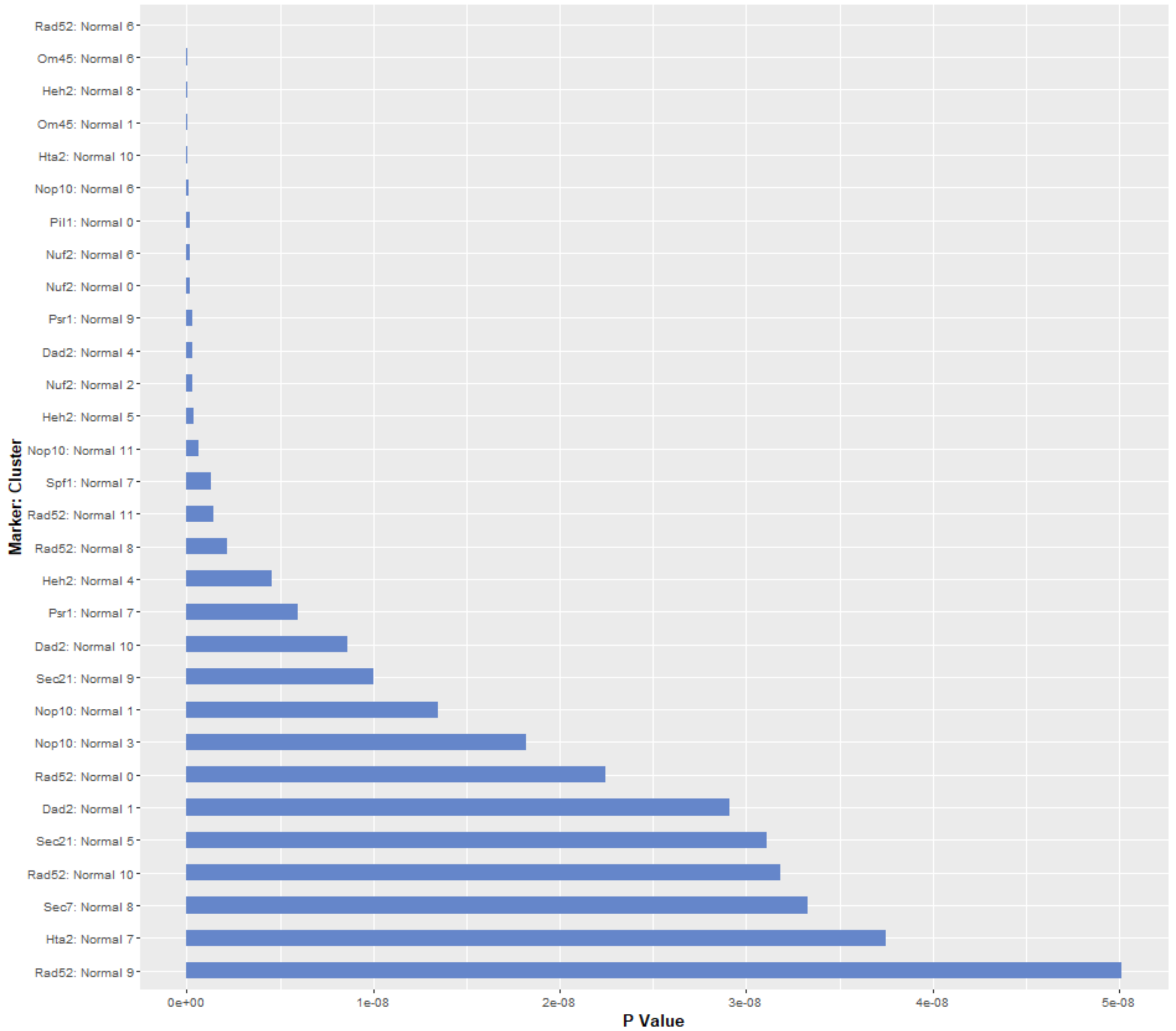
0.006

P Value

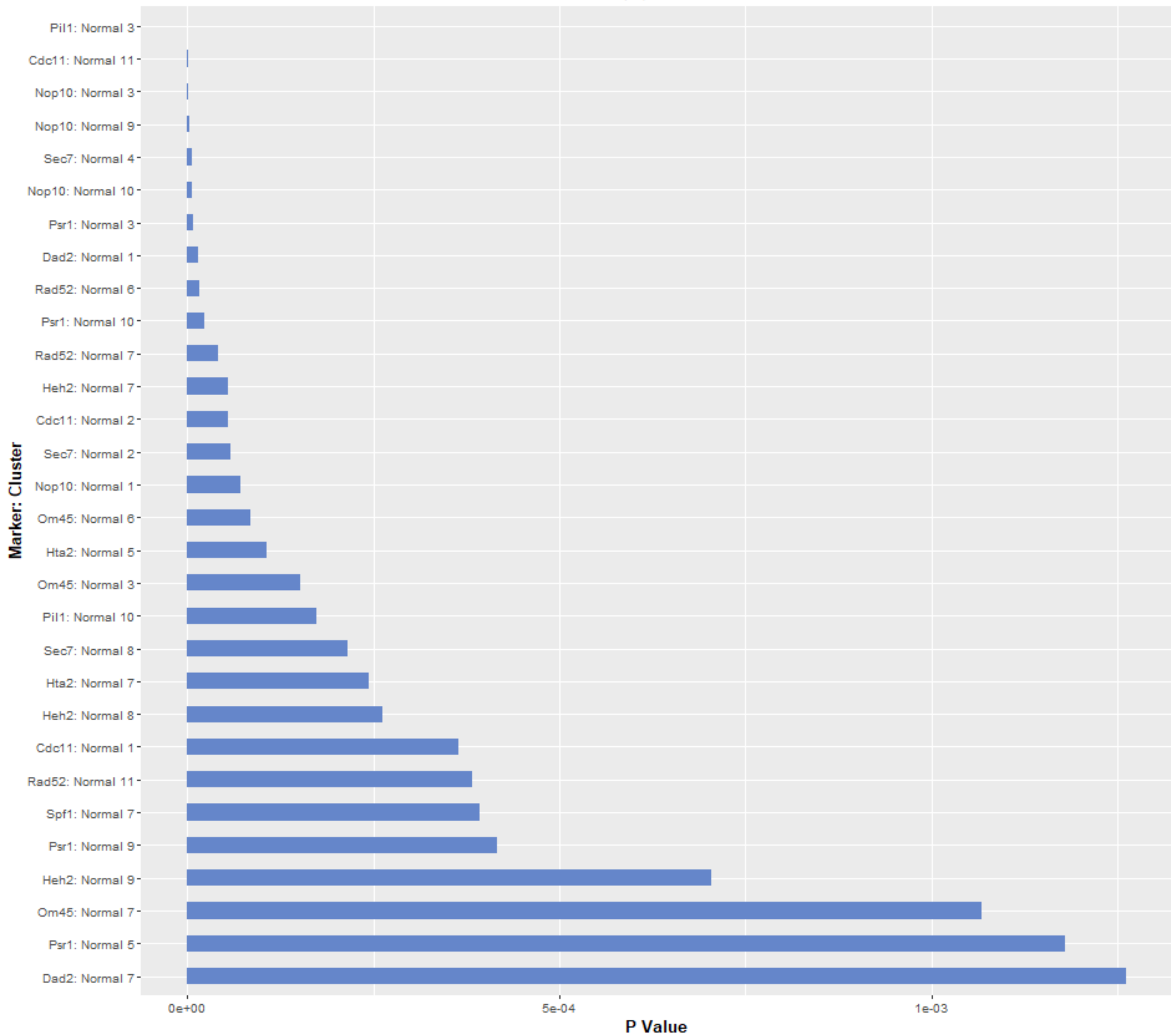


protein complex biogenesis

With AreaShape | Unordered Gene Set



protein lipidation
With AreaShape | Unordered Gene Set



protein maturation

With AreaShape | Unordered Gene Set

Marker: Cluster

Psr1: Normal 0

0.000

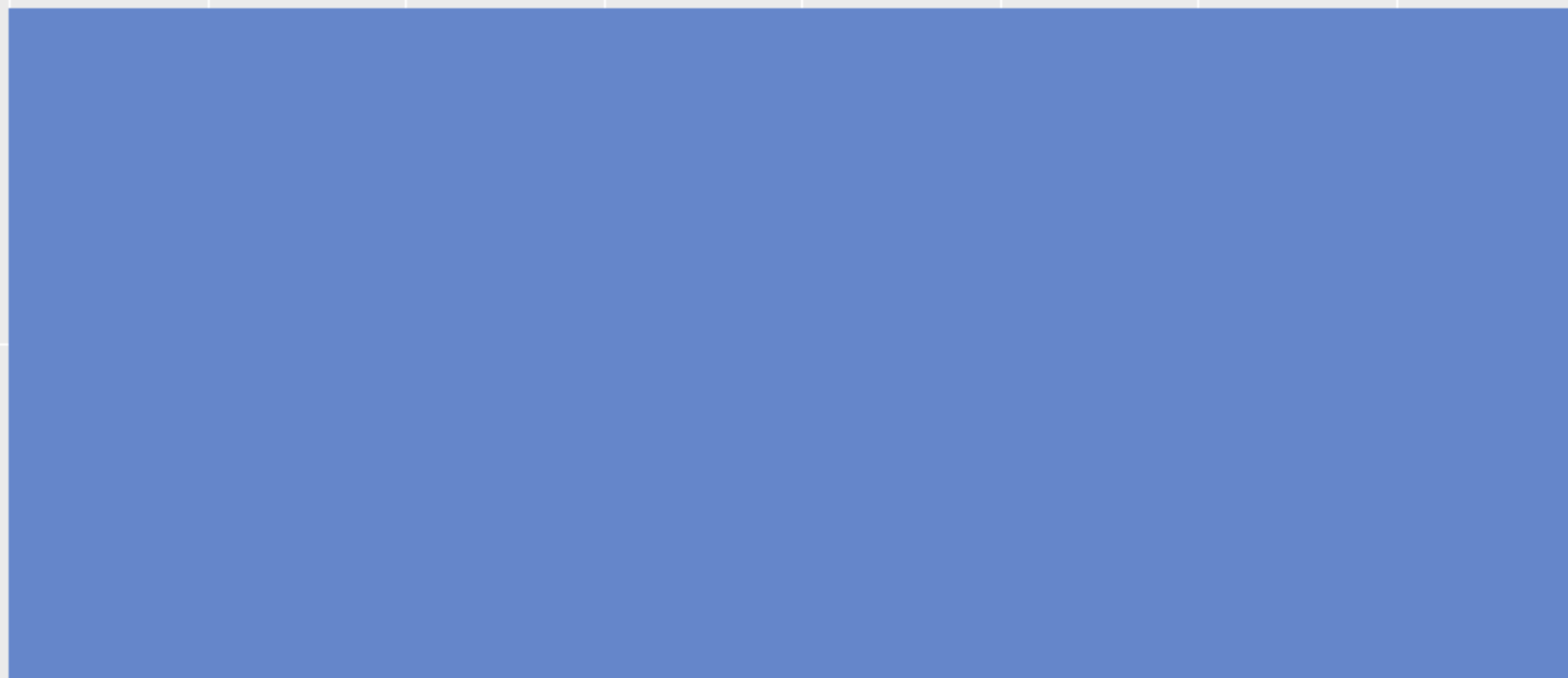
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0.004

0.006

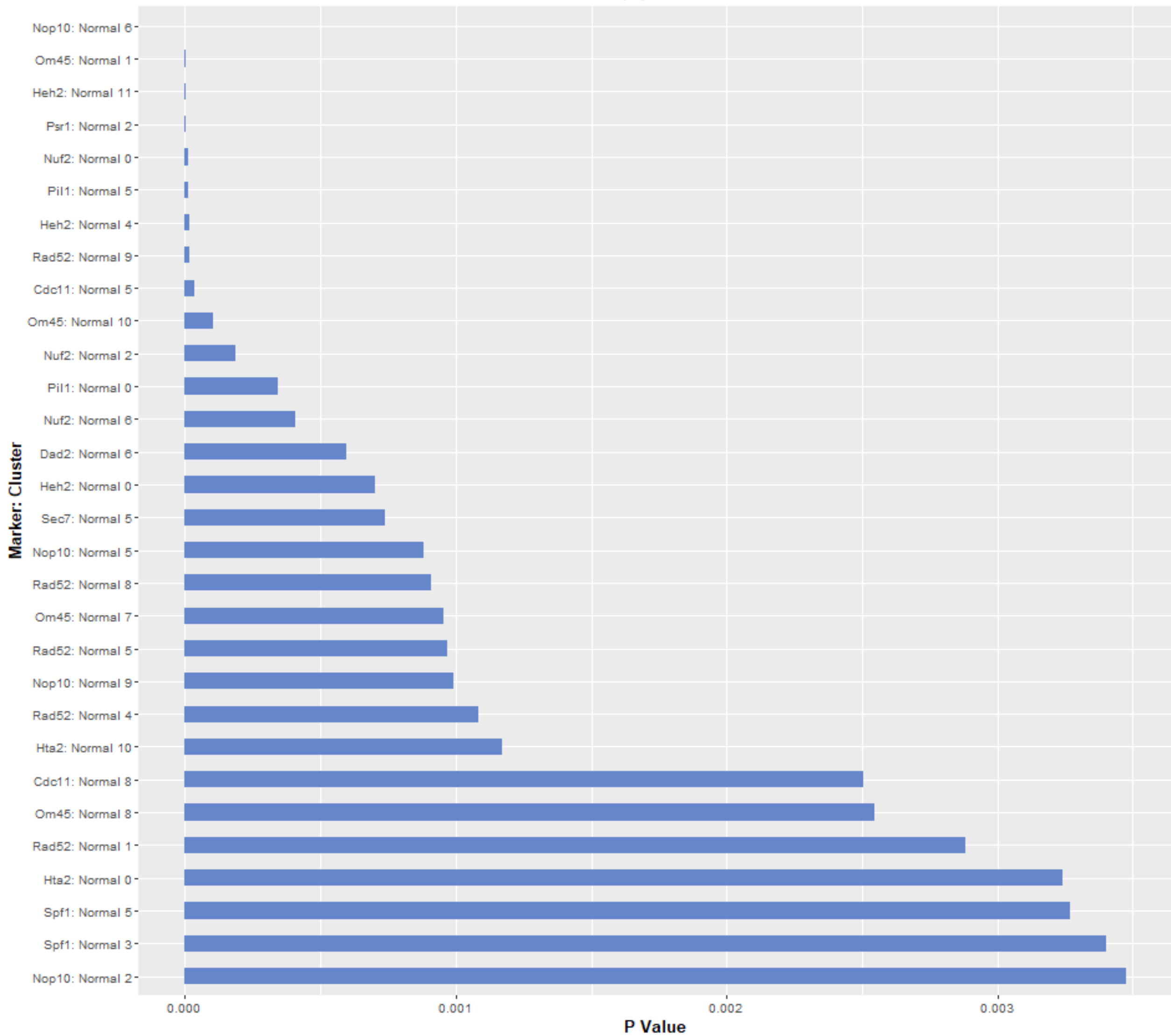
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P Value

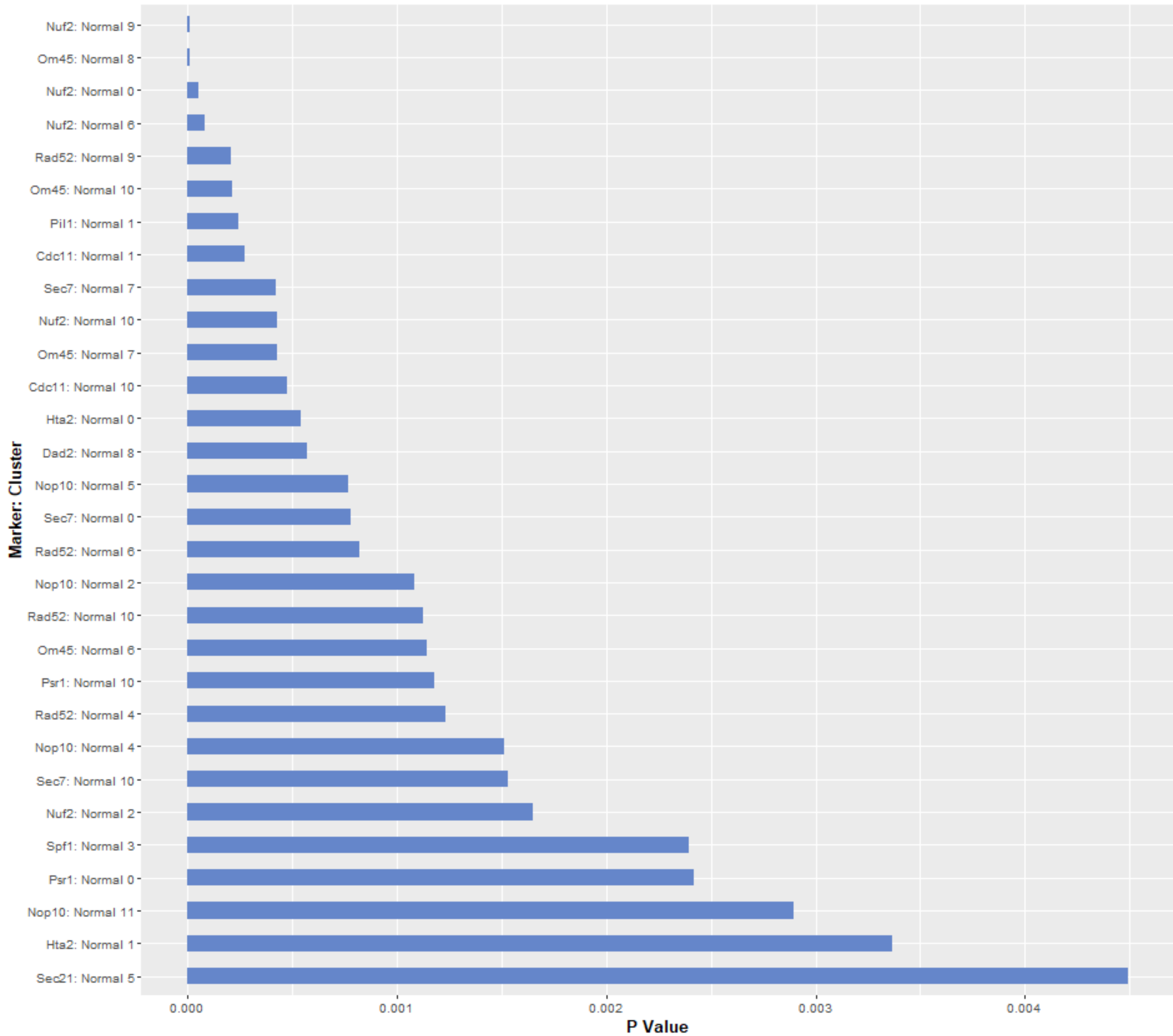


protein modification by small protein conjugation or removal

With AreaShape | Unordered Gene Set



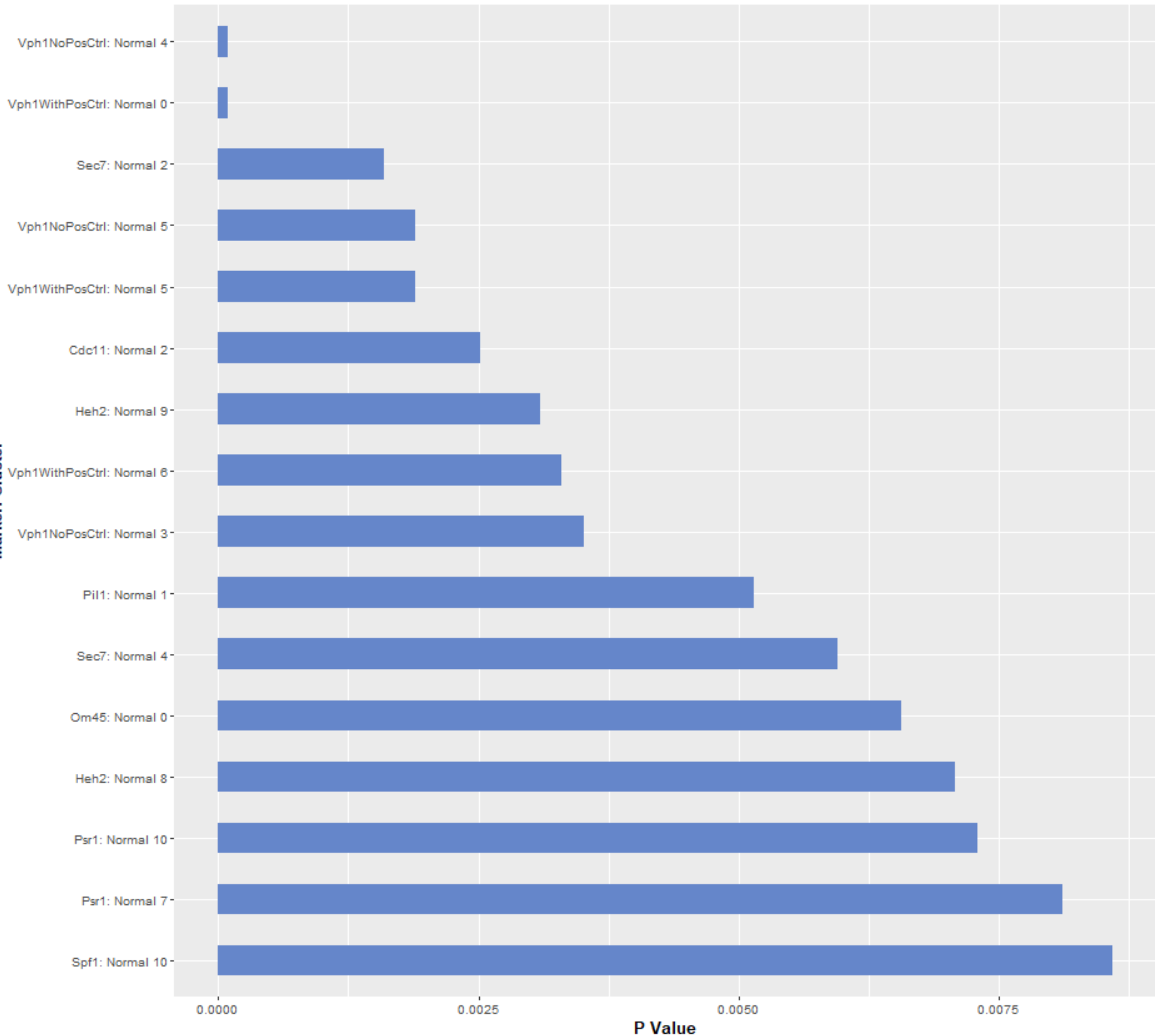
protein phosphorylation
With AreaShape | Unordered Gene Set



protein targeting

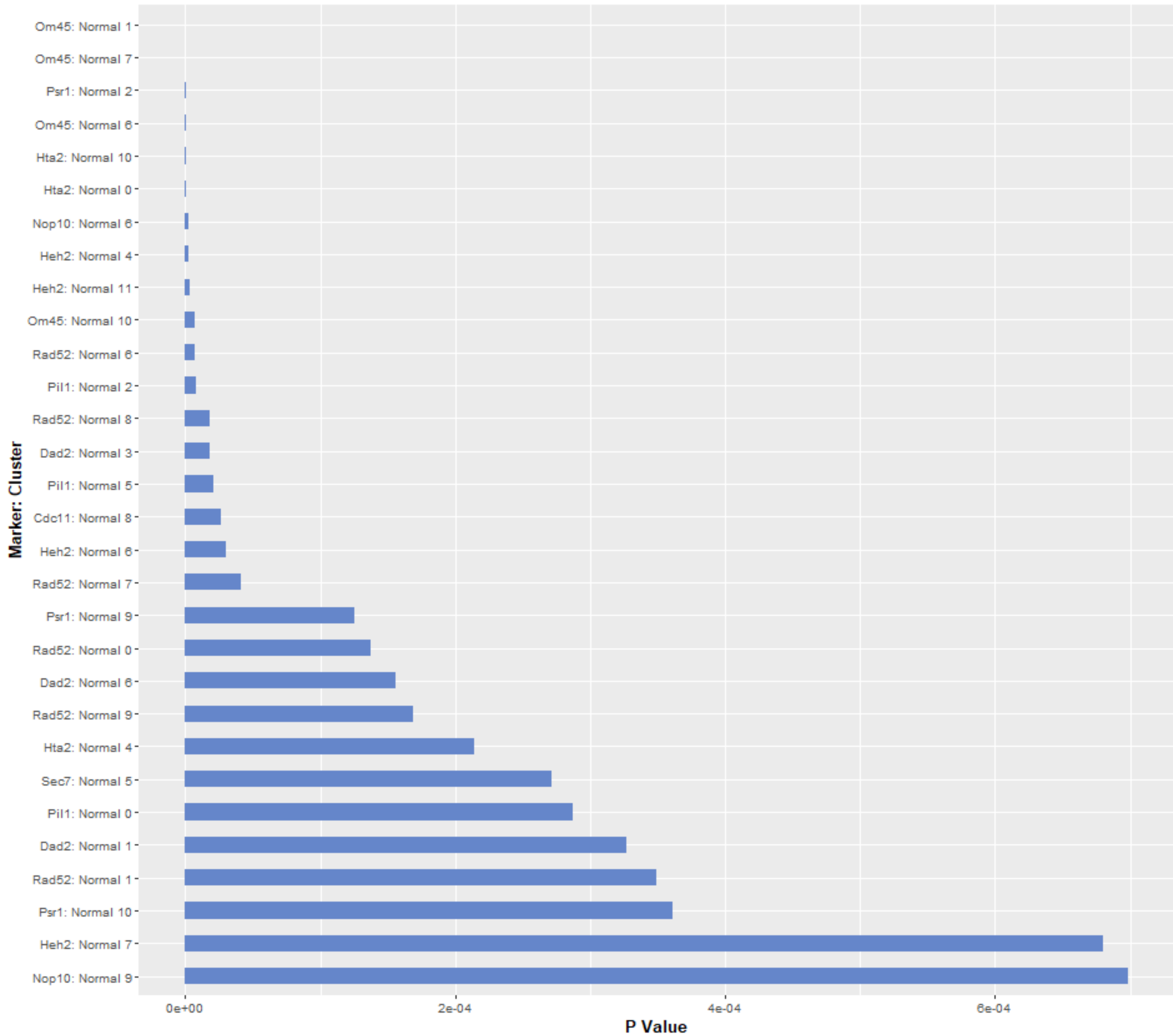
With AreaShape | Unordered Gene Set

Marker: Cluster



proteolysis involved in cellular protein catabolic process

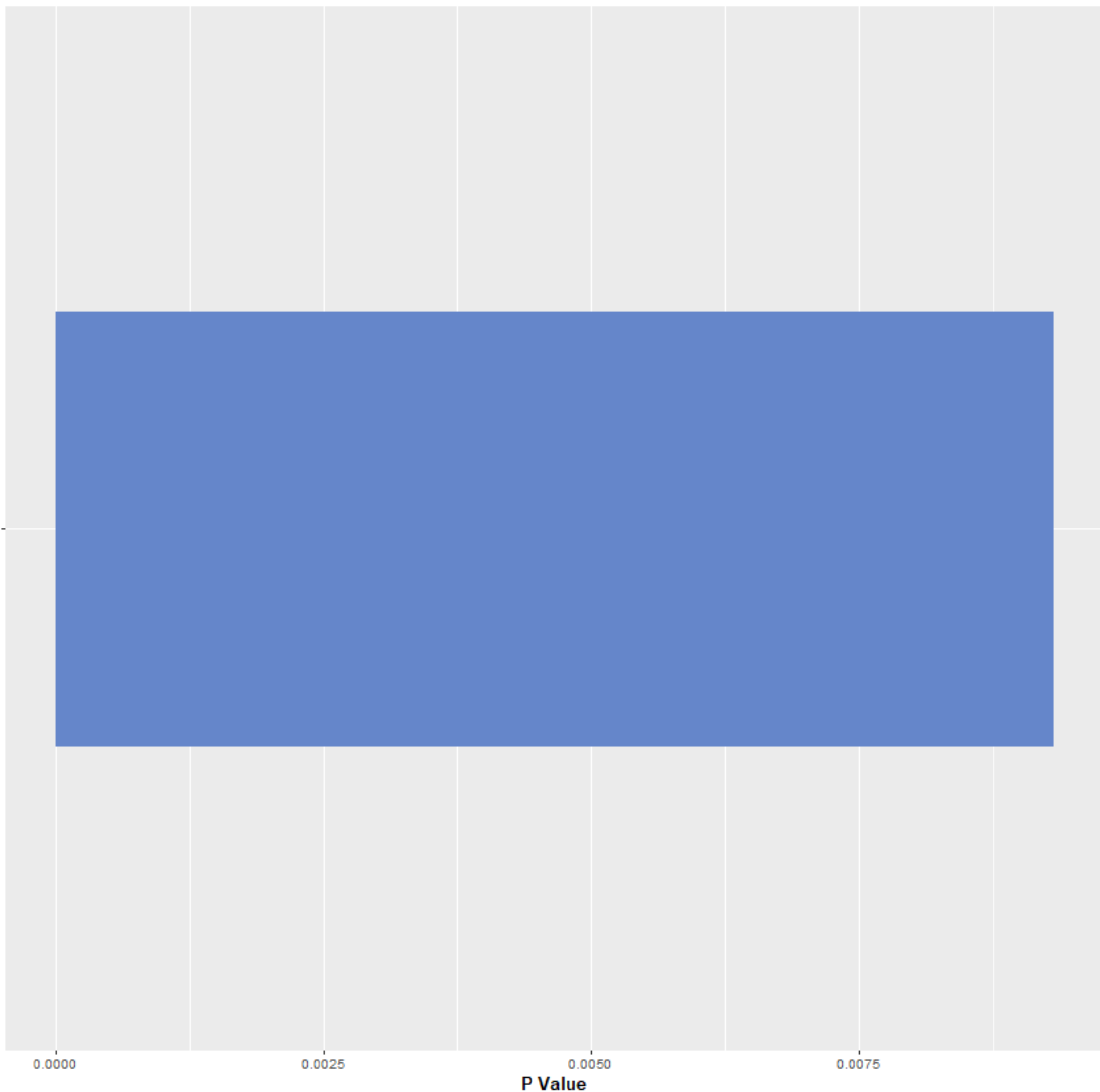
With AreaShape | Unordered Gene Set



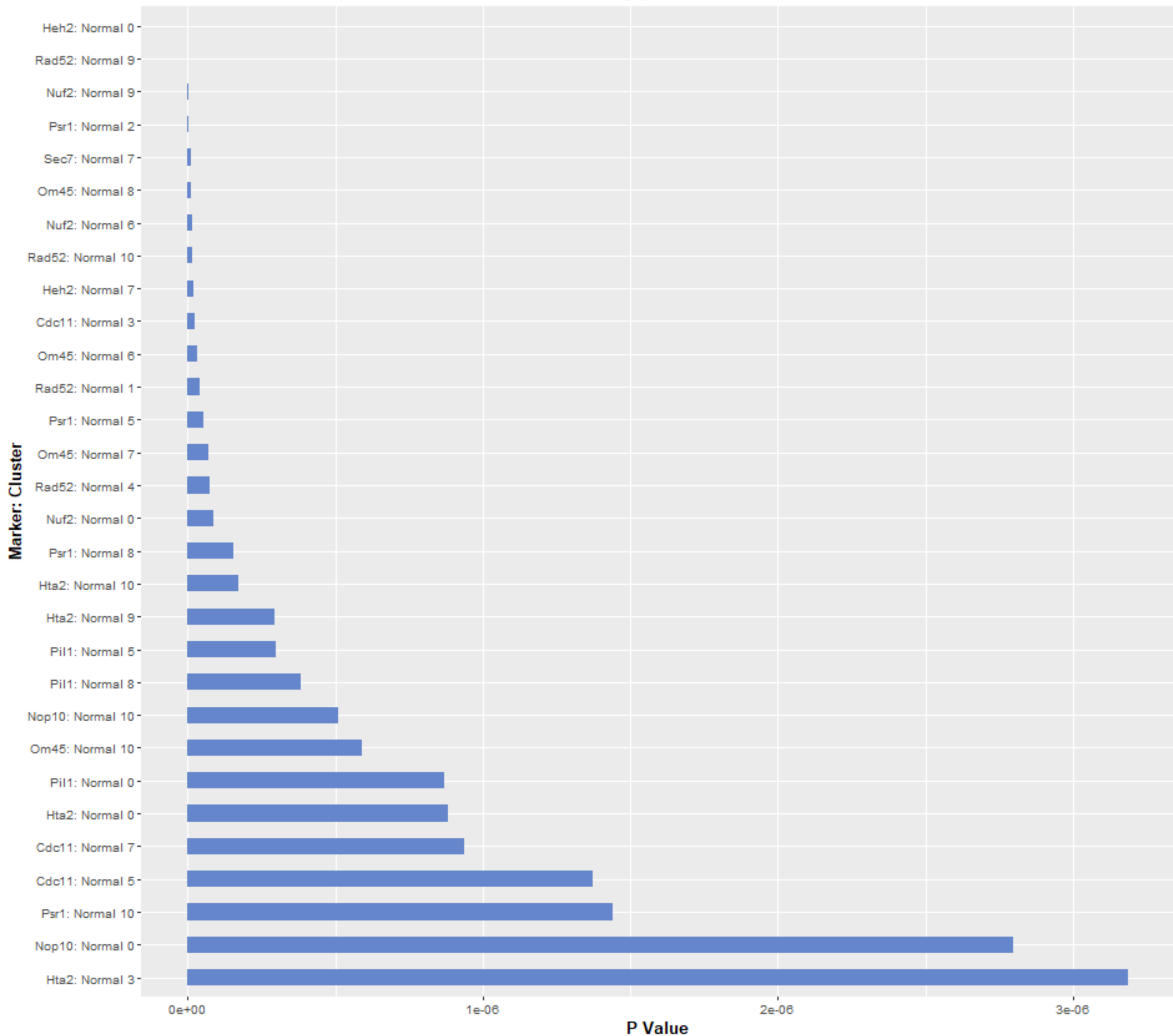
pseudohyphal growth
With AreaShape | Unordered Gene Set

Marker: Cluster

Rad52: Normal 9



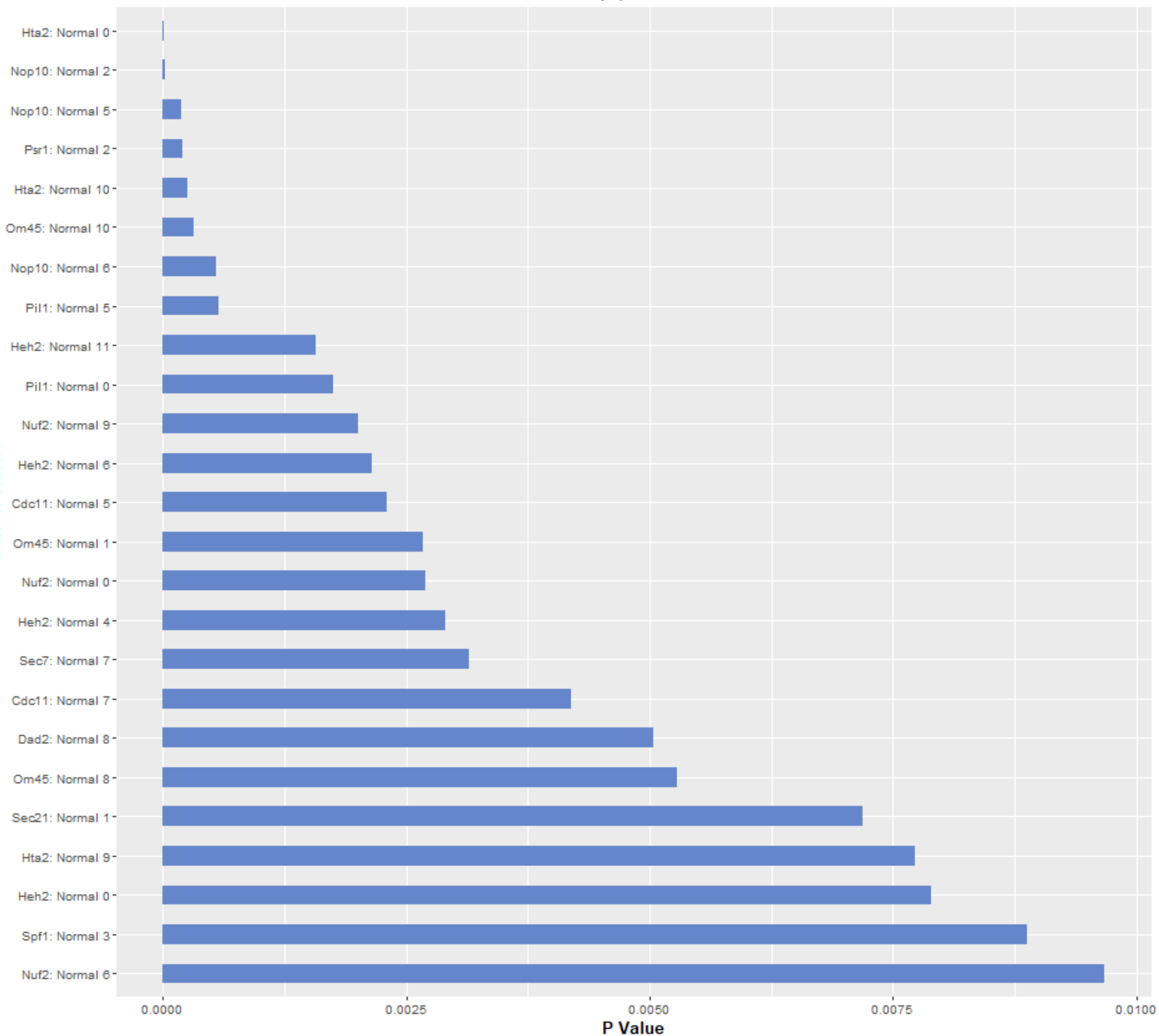
regulation of cell cycle
With AreaShape | Unordered Gene Set



regulation of DNA metabolic process

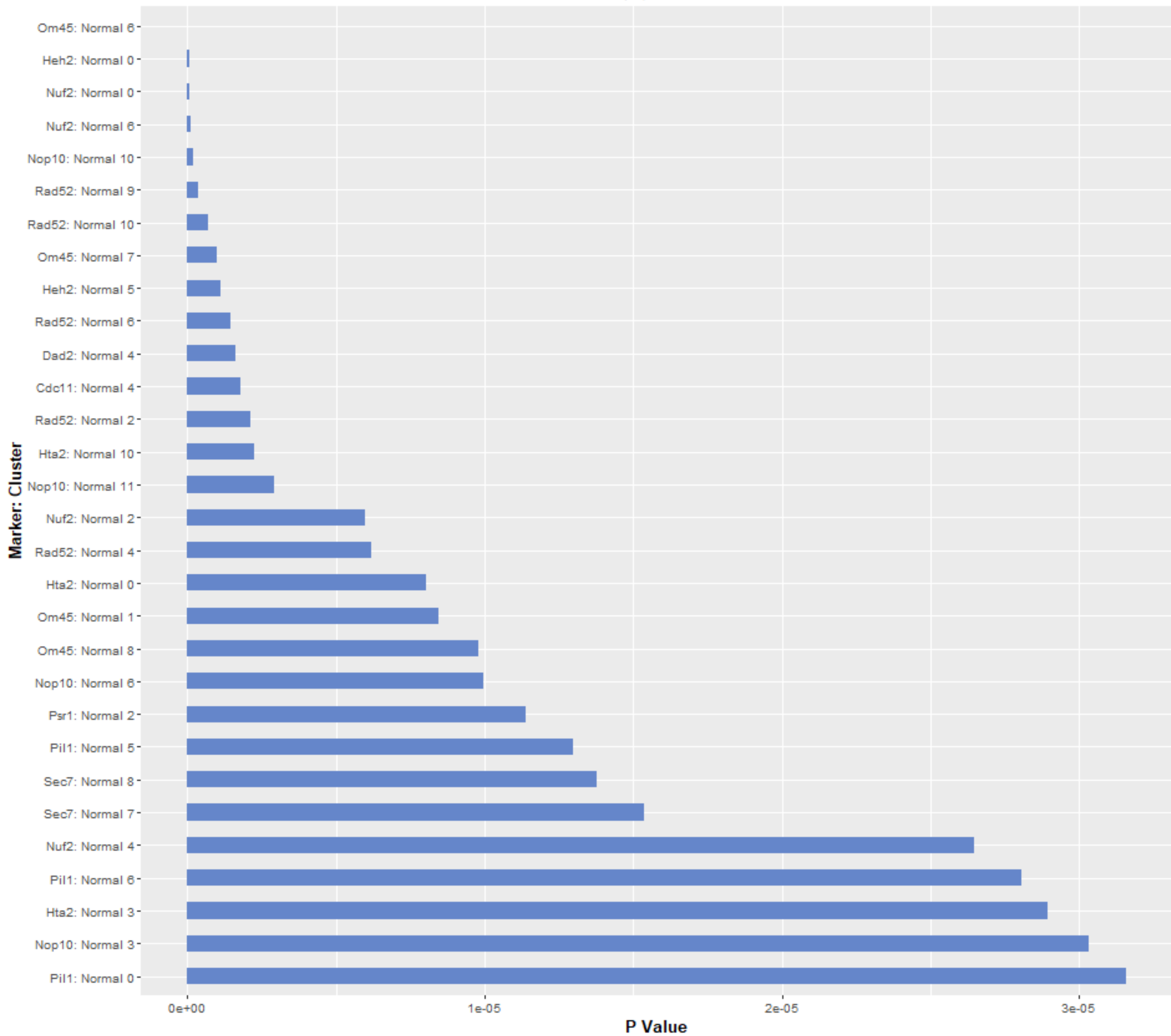
With AreaShape | Unordered Gene Set

Marker: Cluster



regulation of organelle organization

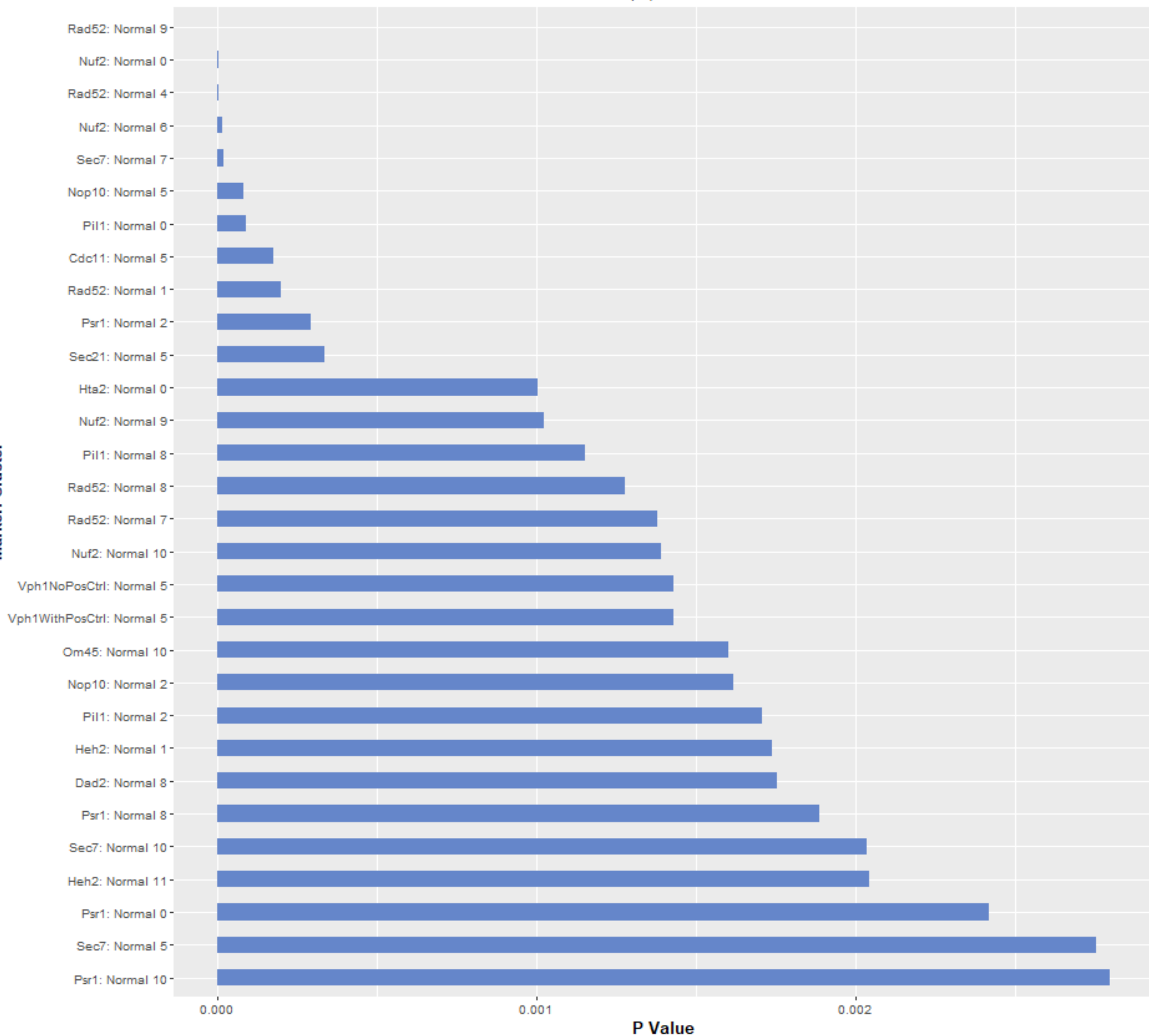
With AreaShape | Unordered Gene Set



regulation of protein modification process

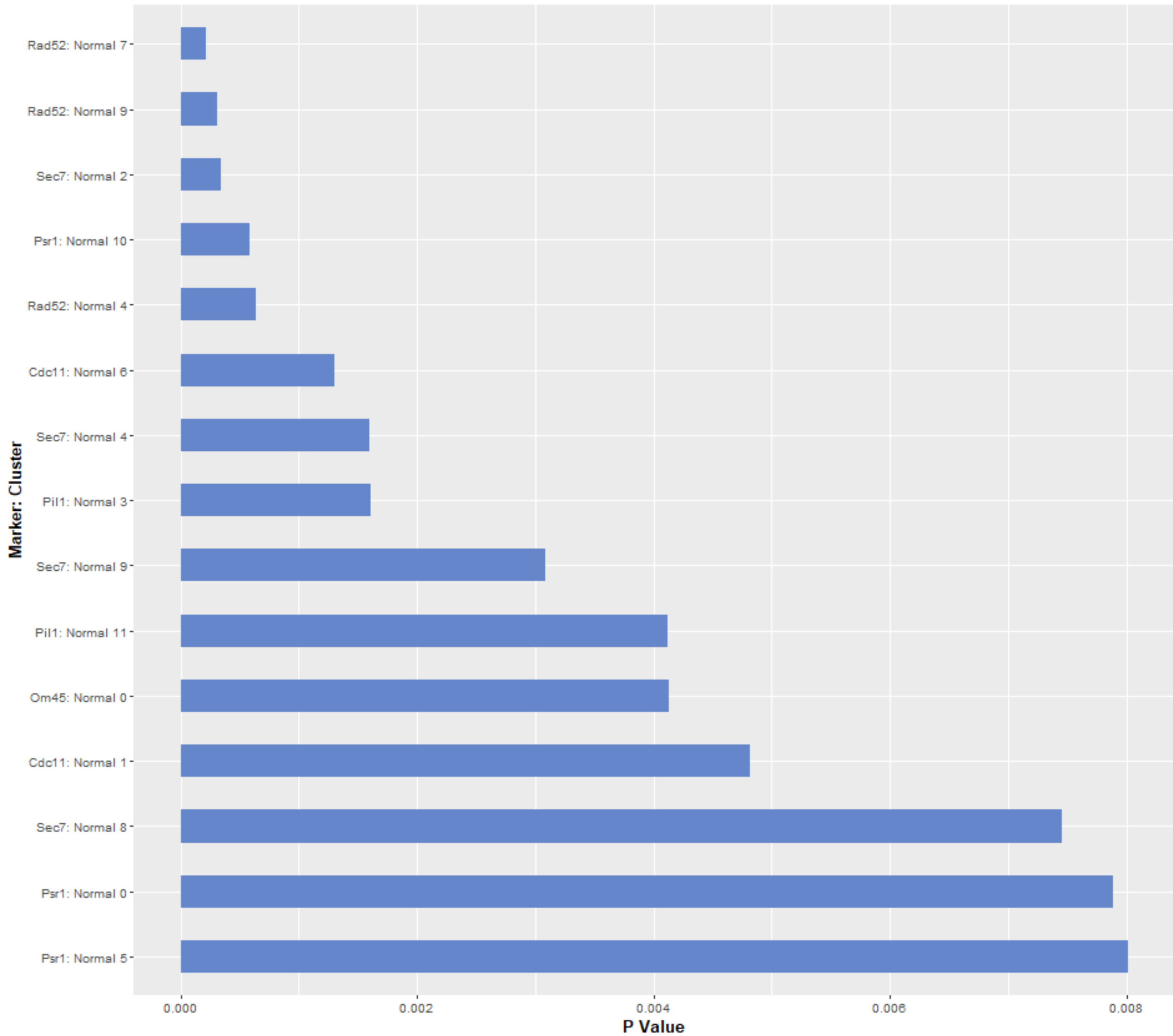
With AreaShape | Unordered Gene Set

Marker: Cluster



regulation of transport

With AreaShape | Unordered Gene Set



response to heat

With AreaShape | Unordered Gene Set

Marker: Cluster

Psr1: Normal 0

0.000

0.002

0.004

0.006

0.008

P Value



ribosomal large subunit biogenesis

With AreaShape | Unordered Gene Set

Marker: Cluster

Pil1: Normal 10

Psr1: Normal 9

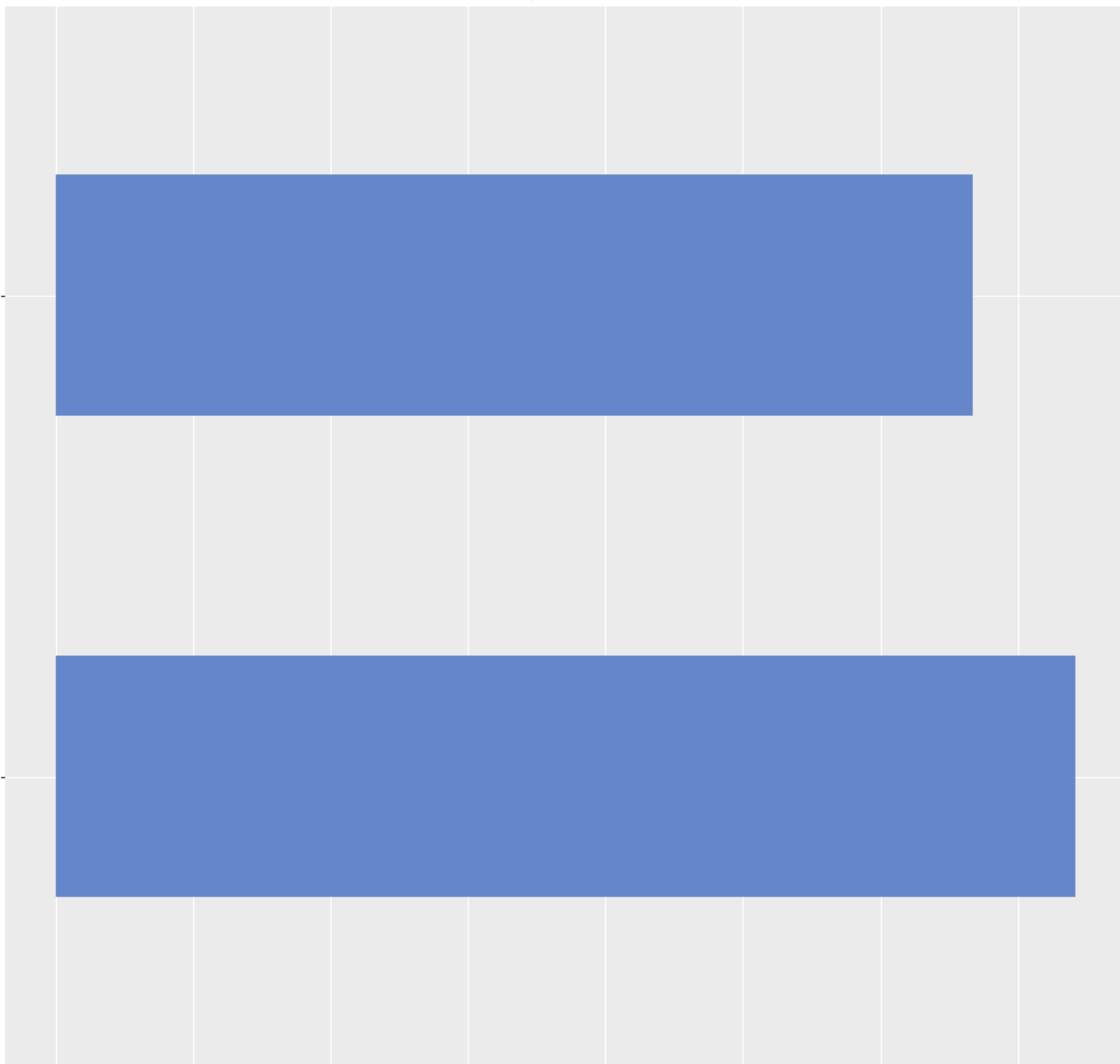
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0.0050

0.0075

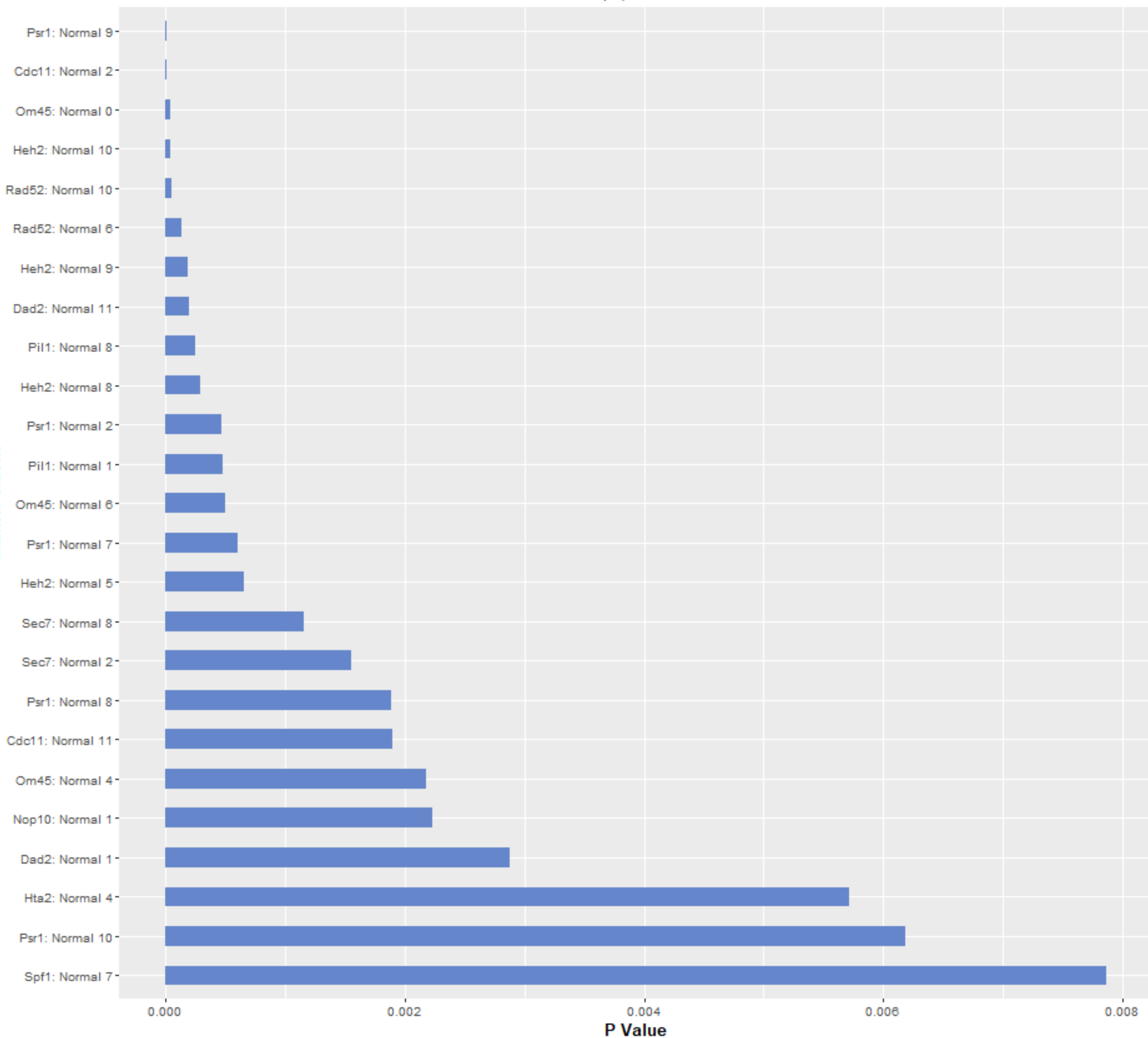
P Value



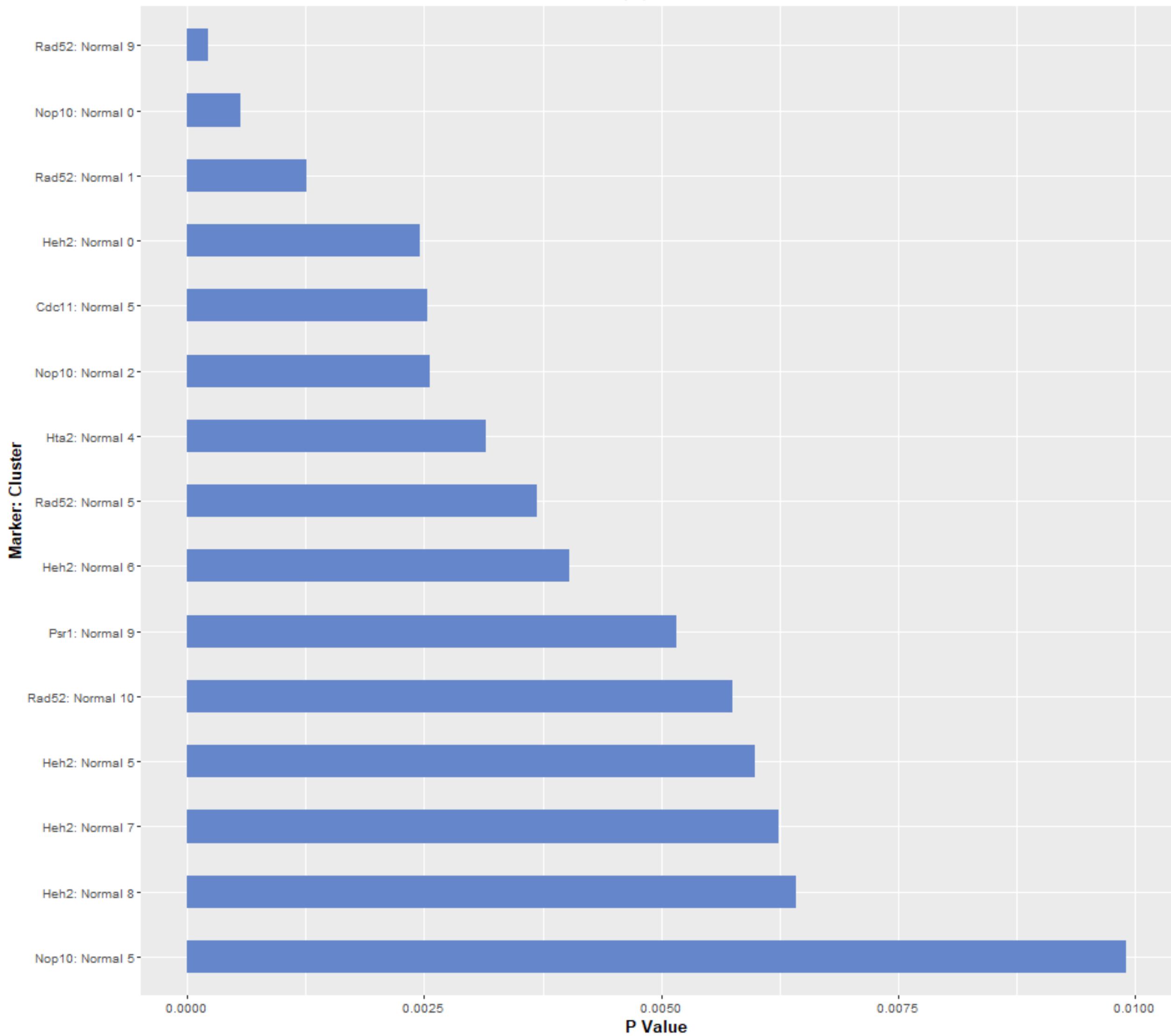
ribosomal subunit export from nucleus

With AreaShape | Unordered Gene Set

Marker: Cluster

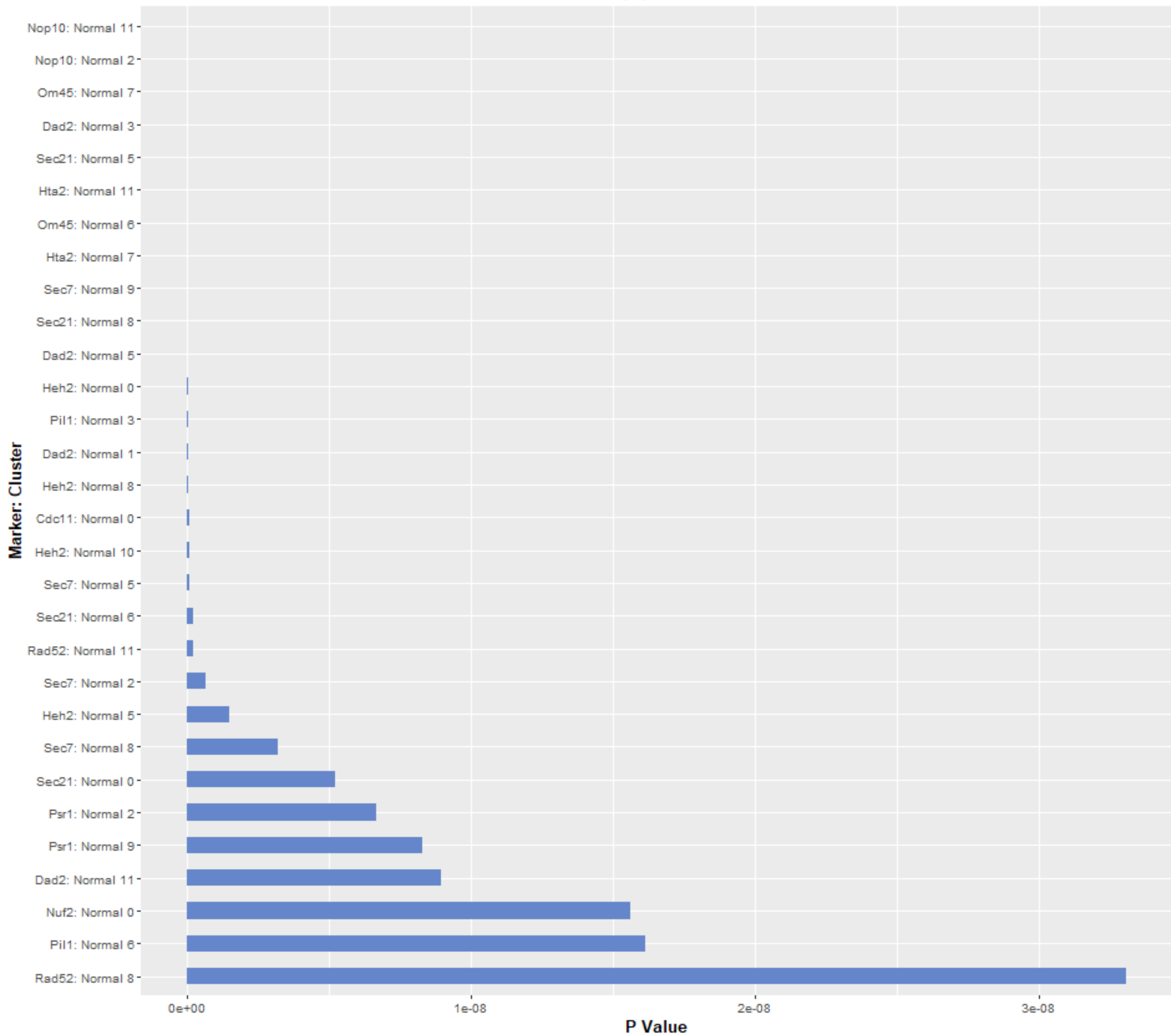


RNA catabolic process
With AreaShape | Unordered Gene Set

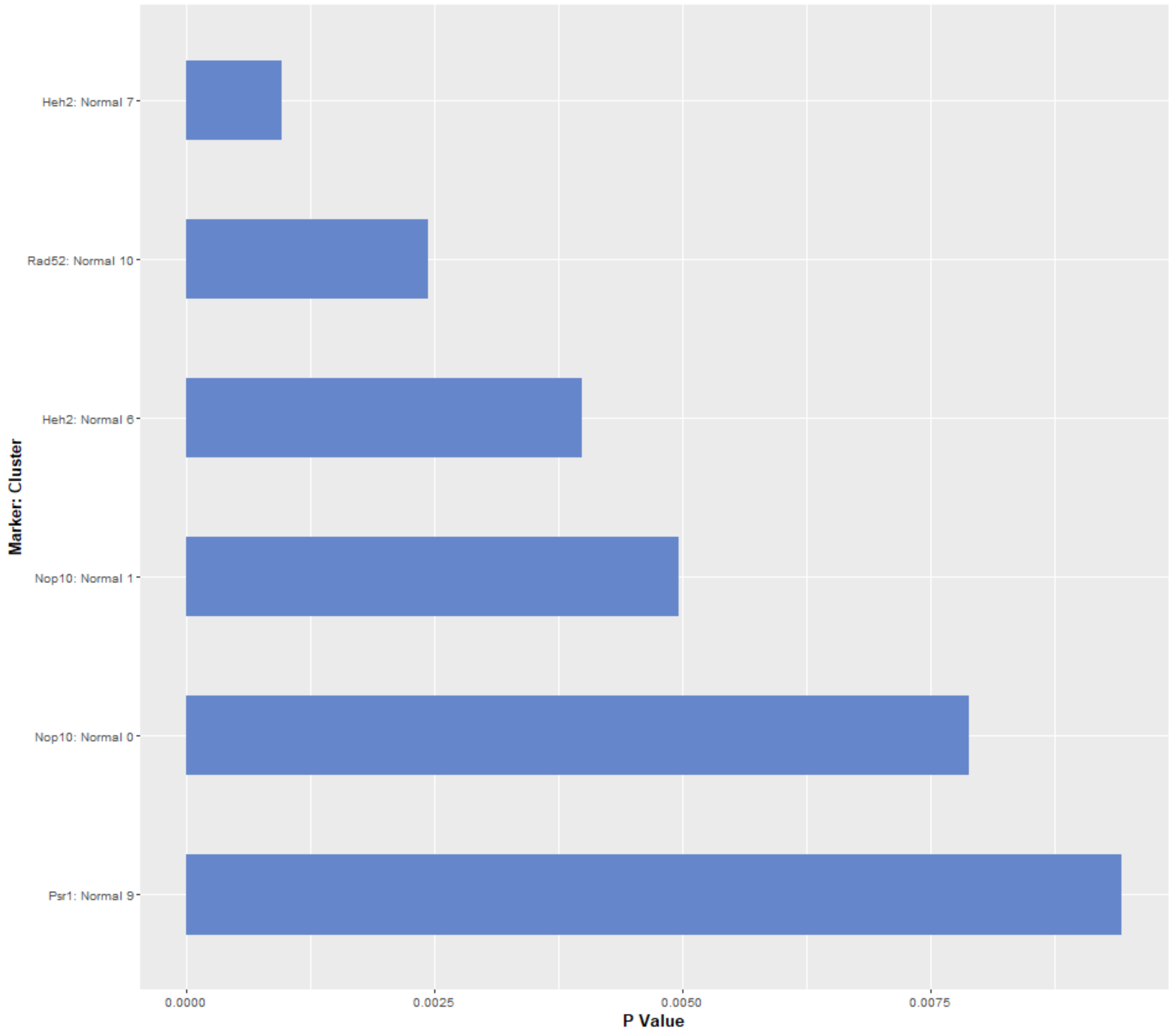


RNA splicing

With AreaShape | Unordered Gene Set



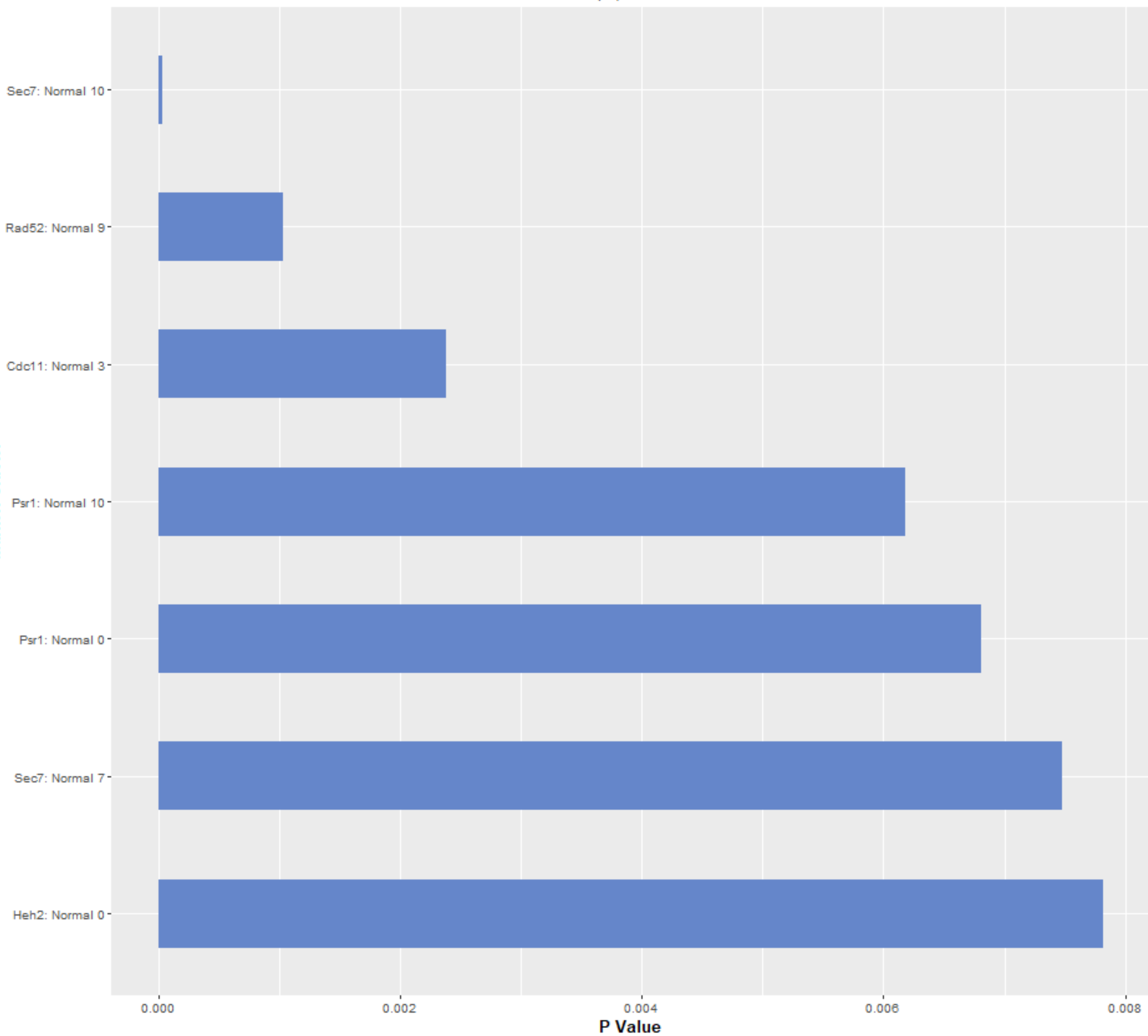
rRNA processing
With AreaShape | Unordered Gene Set



signaling

With AreaShape | Unordered Gene Set

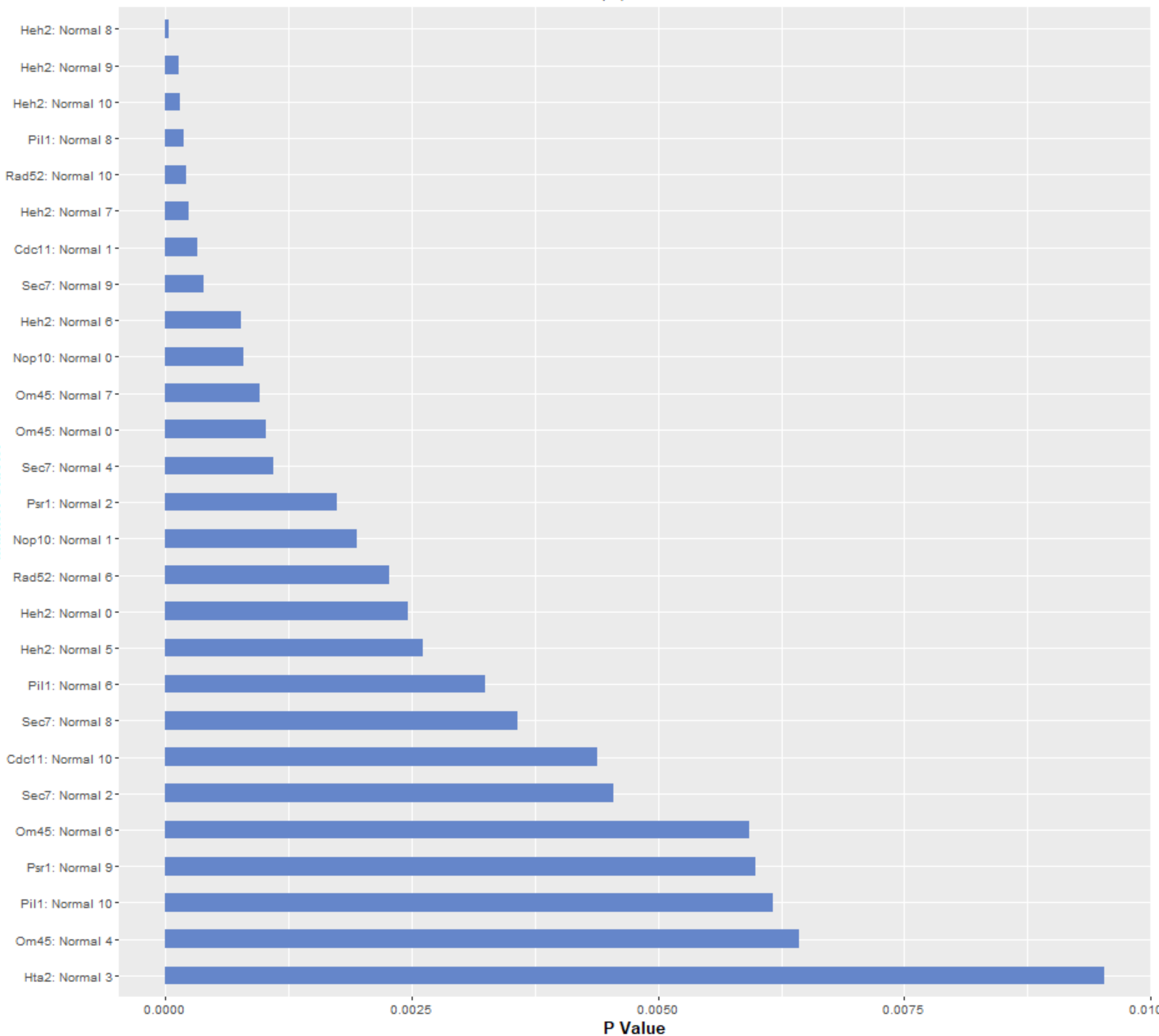
Marker: Cluster



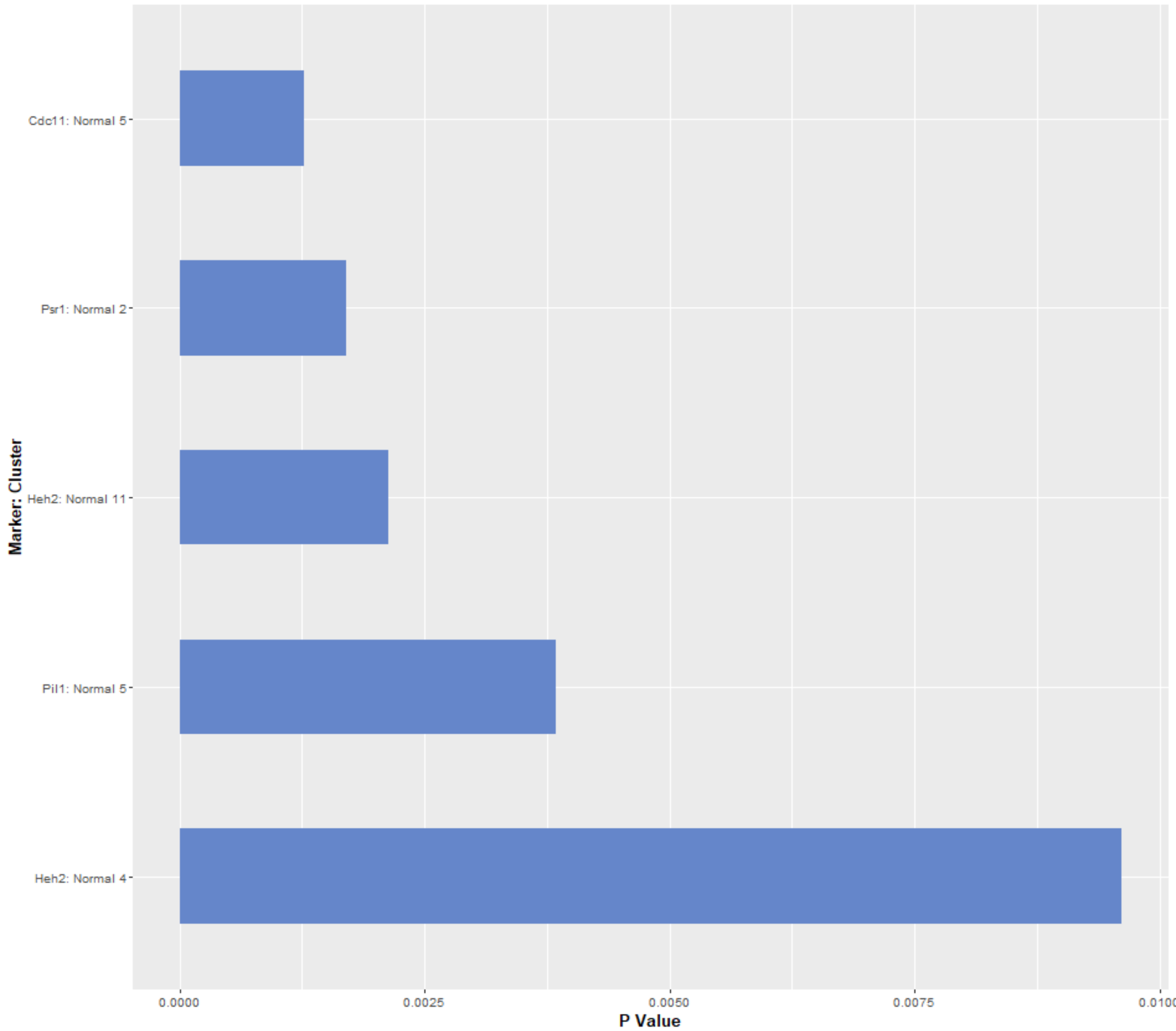
snoRNA processing

With AreaShape | Unordered Gene Set

Marker: Cluster



telomere organization
With AreaShape | Unordered Gene Set



transcription from RNA polymerase I promoter

With AreaShape | Unordered Gene Set

Marker: Cluster

Pil1: Normal 1

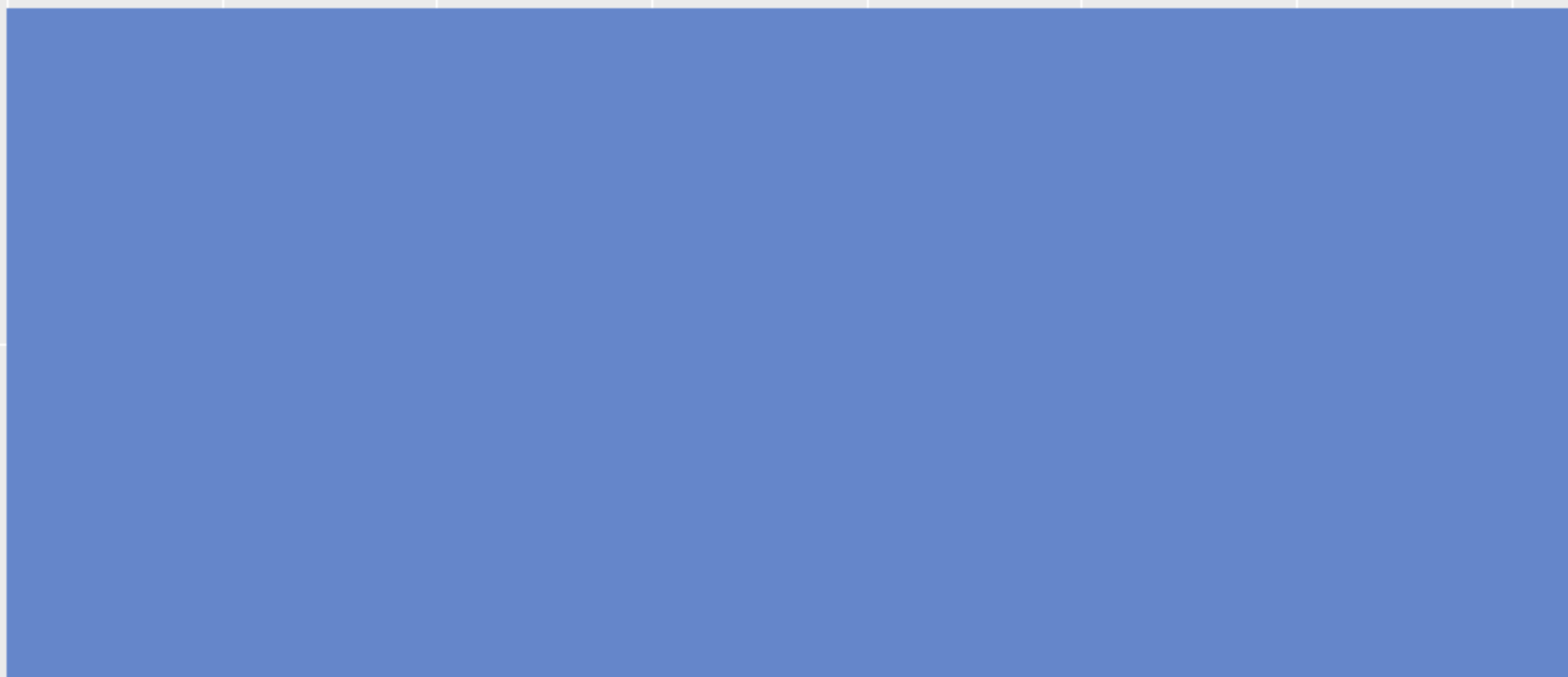
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0.002

0.004

0.006

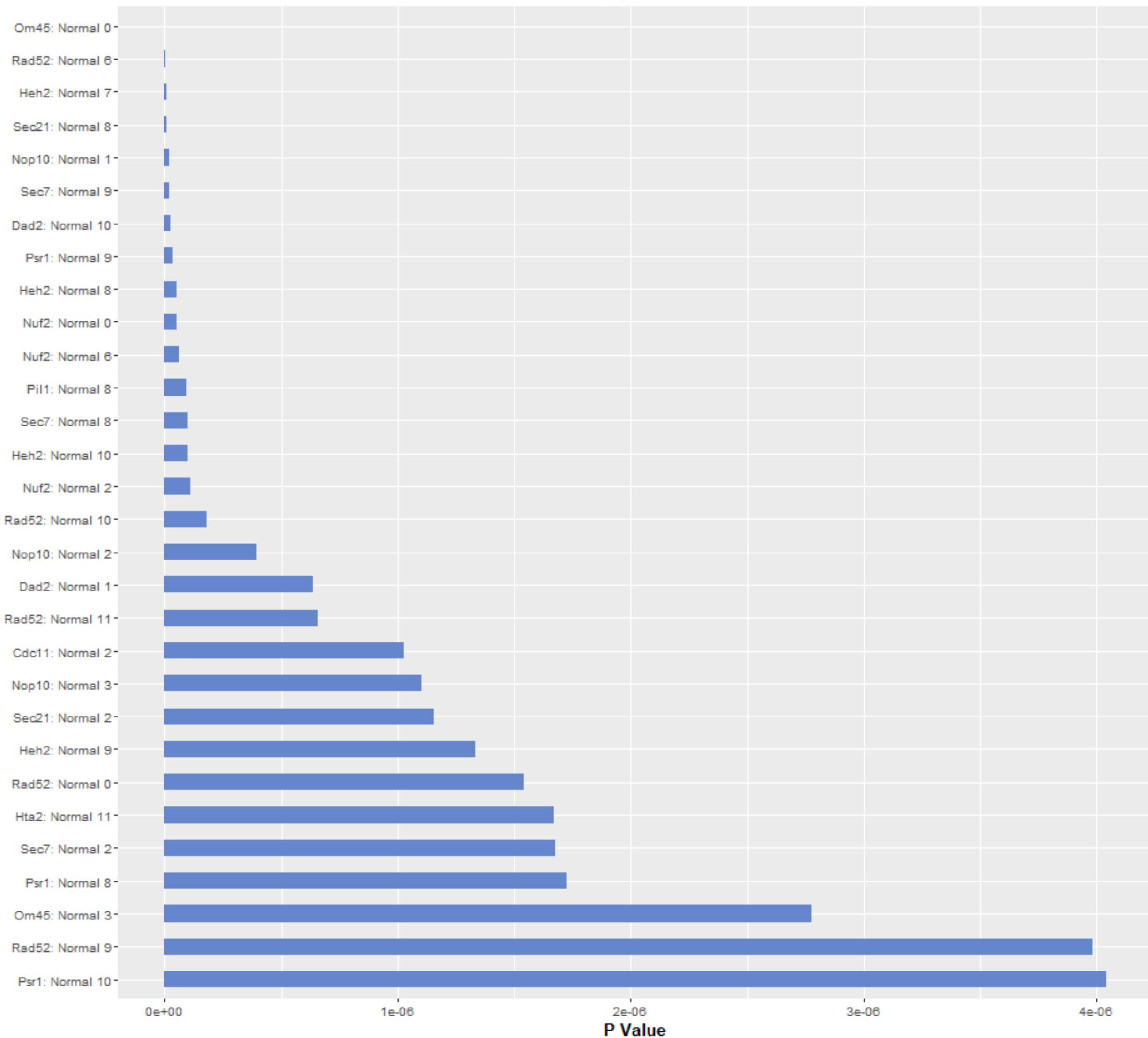
P Value



transcription from RNA polymerase II promoter

With AreaShape | Unordered Gene Set

Marker: Cluster



transcription from RNA polymerase III promoter

With AreaShape | Unordered Gene Set

Marker: Cluster

Hta2: Normal 9

Heh2: Normal 7

Hta2: Normal 4

0.000

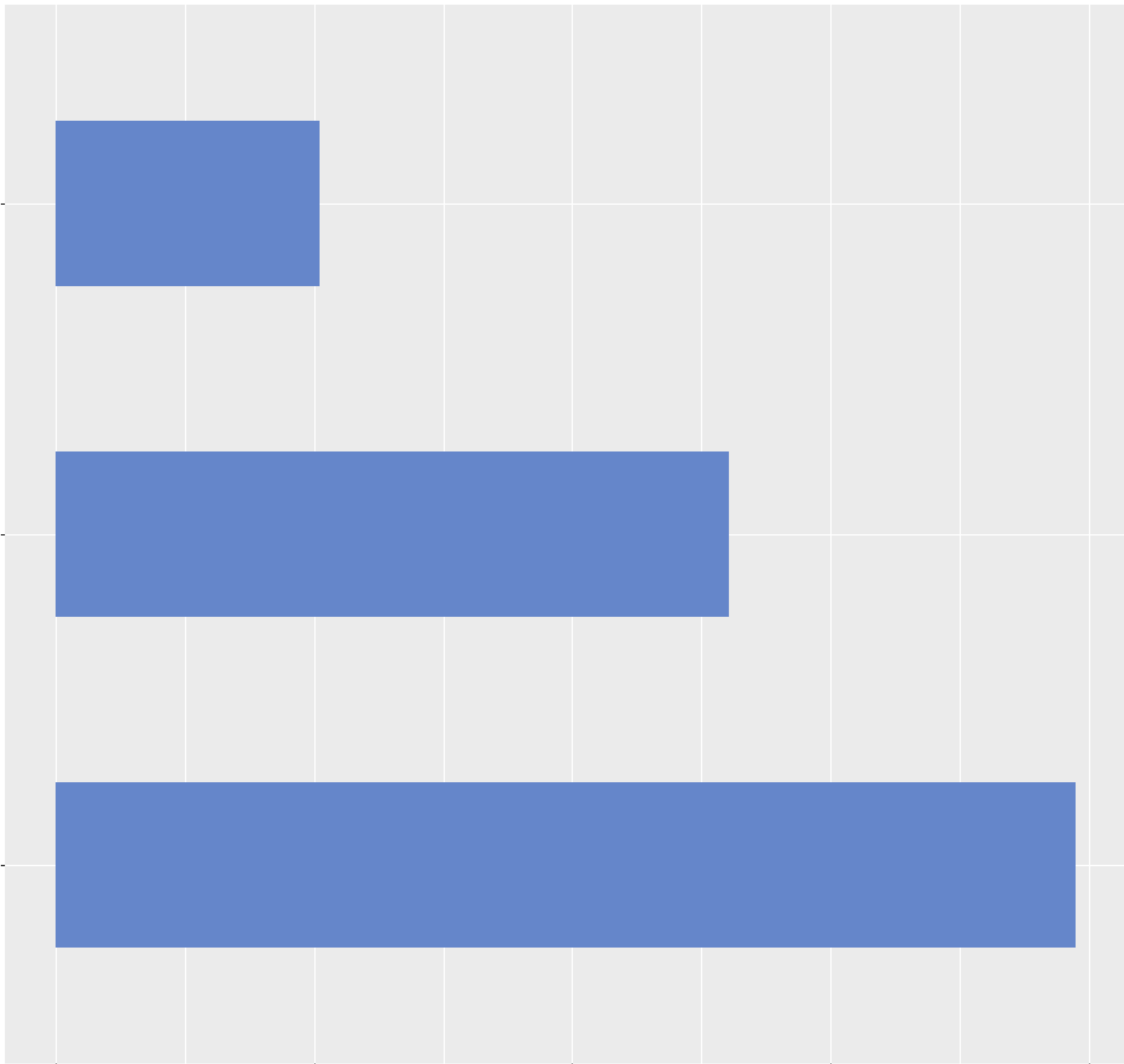
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0.002

0.003

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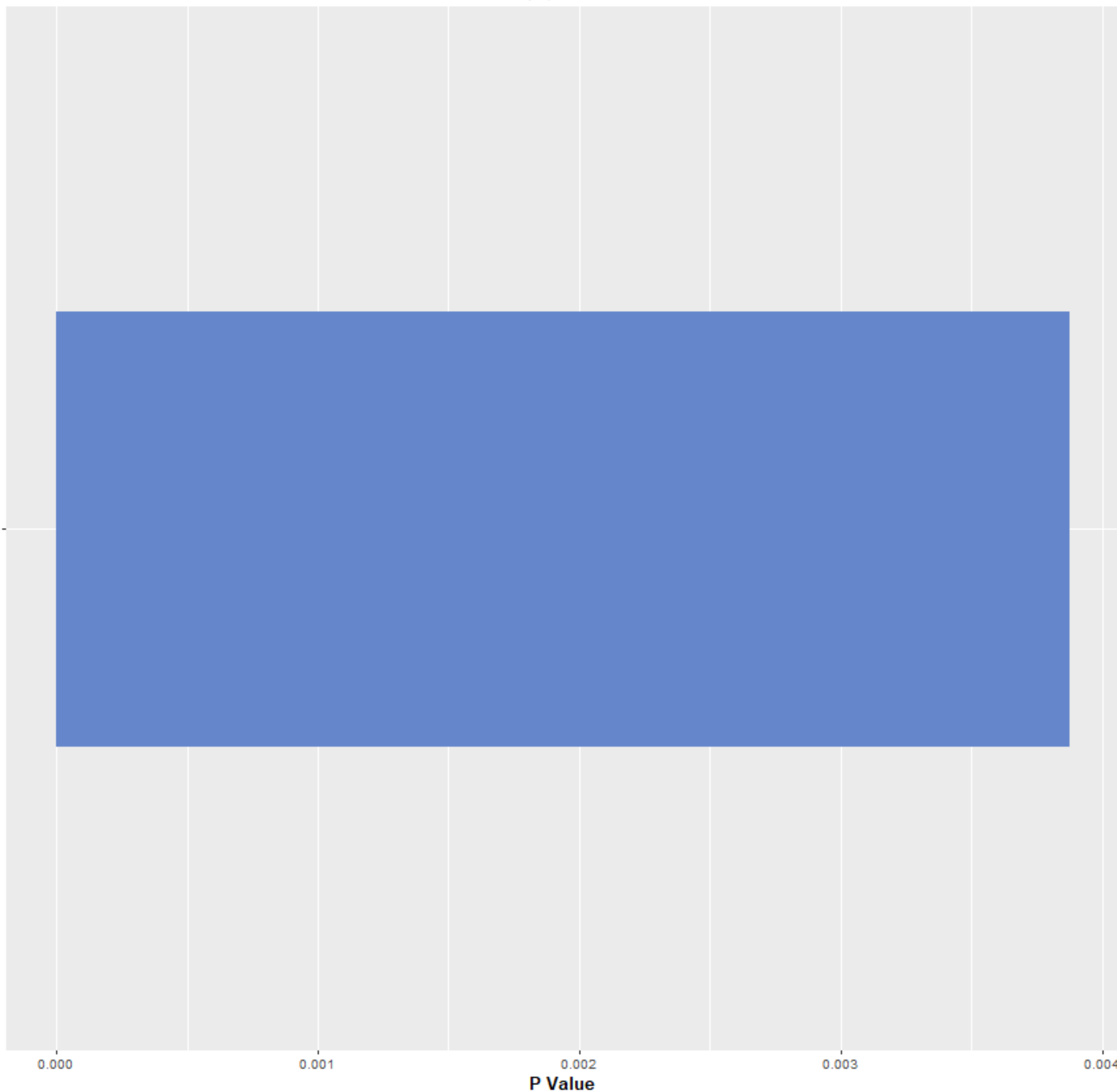
P Value



translational elongation
With AreaShape | Unordered Gene Set

Marker: Cluster

Pil1: Normal 4



translational initiation

With AreaShape | Unordered Gene Set

Marker: Cluster

Pil1: Normal 1

Hta2: Normal 9

0.000

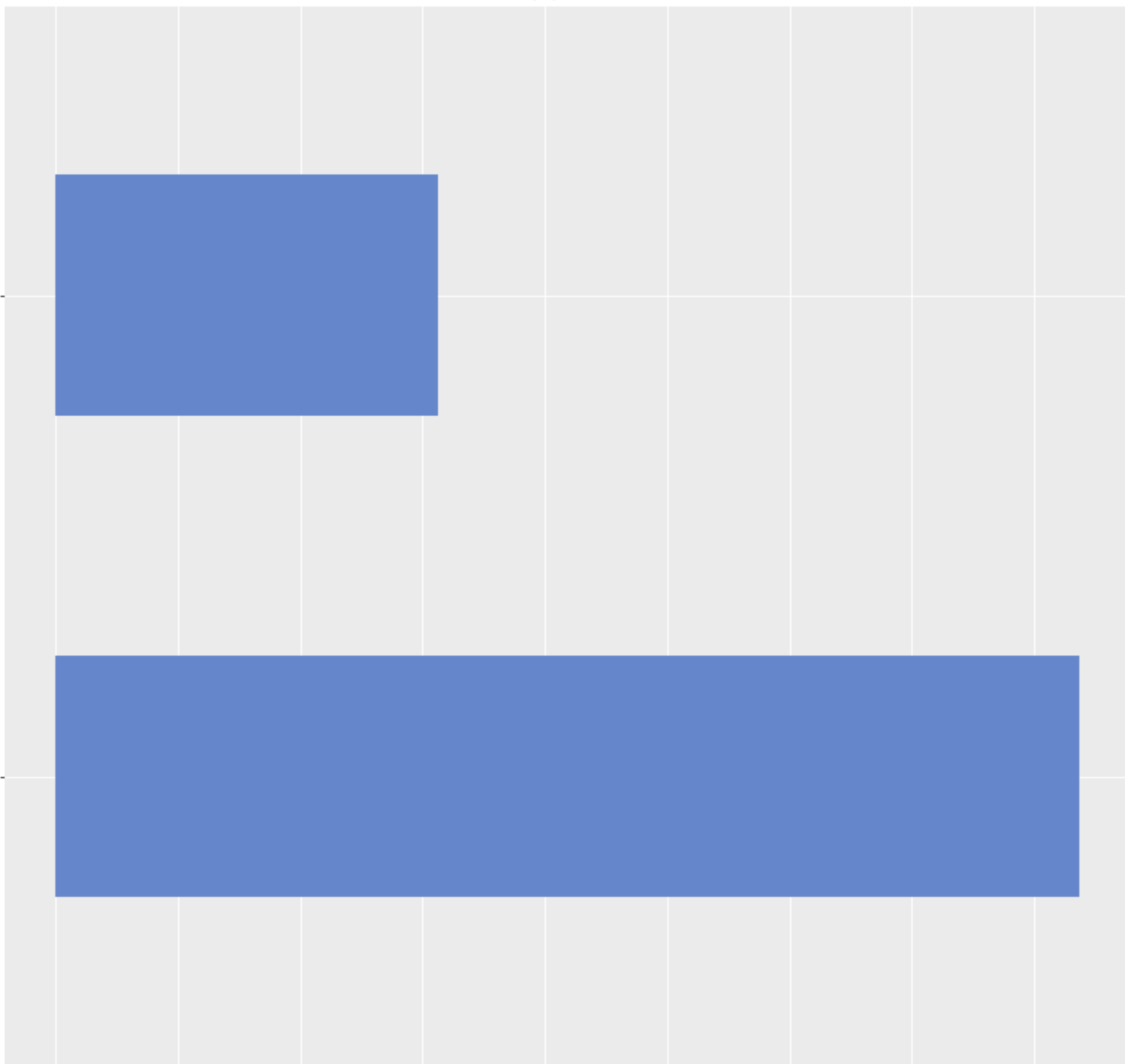
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0.002

0.003

0.004

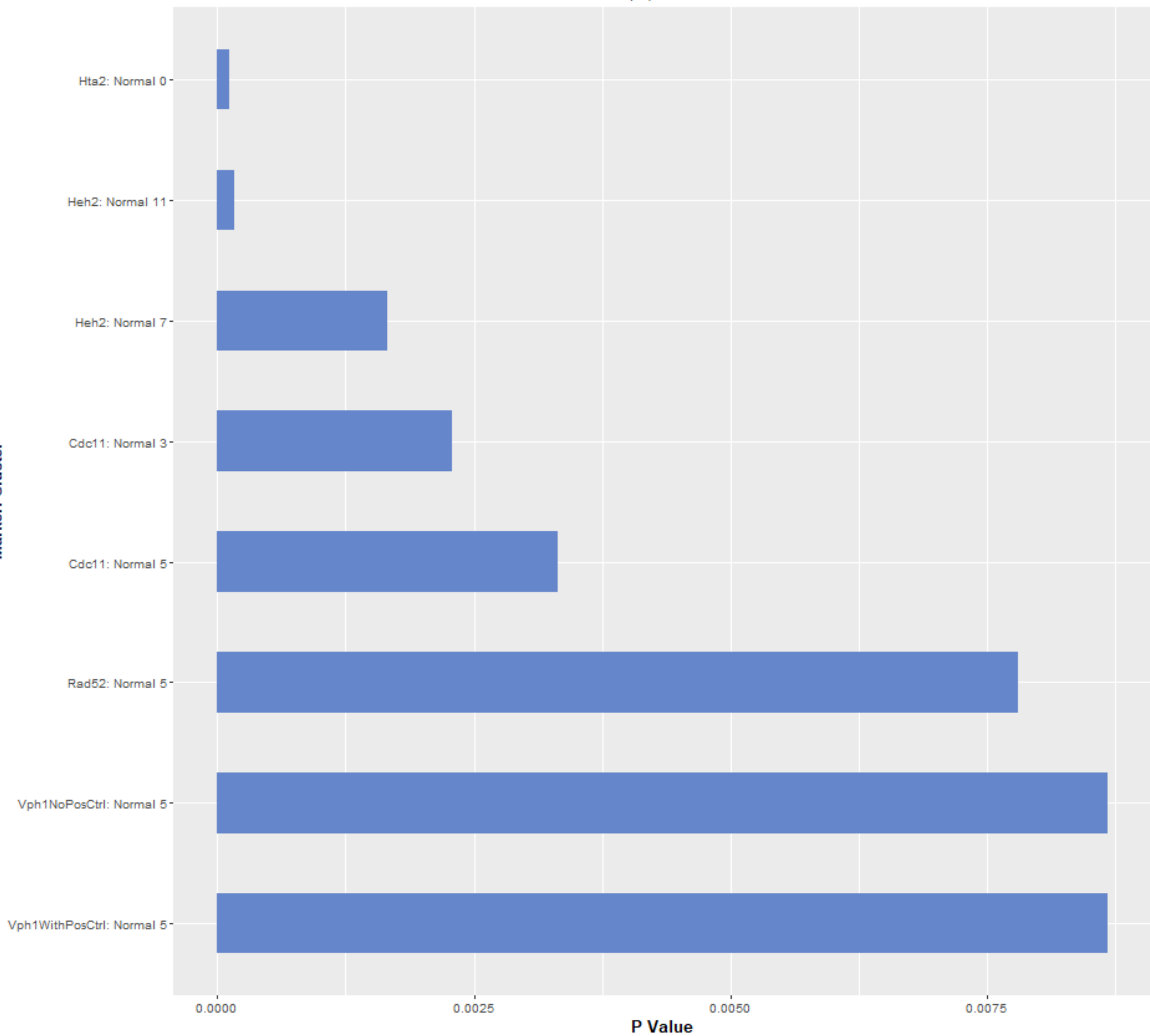
P Value



transposition

With AreaShape | Unordered Gene Set

Marker: Cluster



tRNA aminoacylation for protein translation

With AreaShape | Unordered Gene Set

Marker: Cluster

Om45: Normal 8

Heh2: Normal 9

Heh2: Normal 8

Pil1: Normal 8

Om45: Normal 9

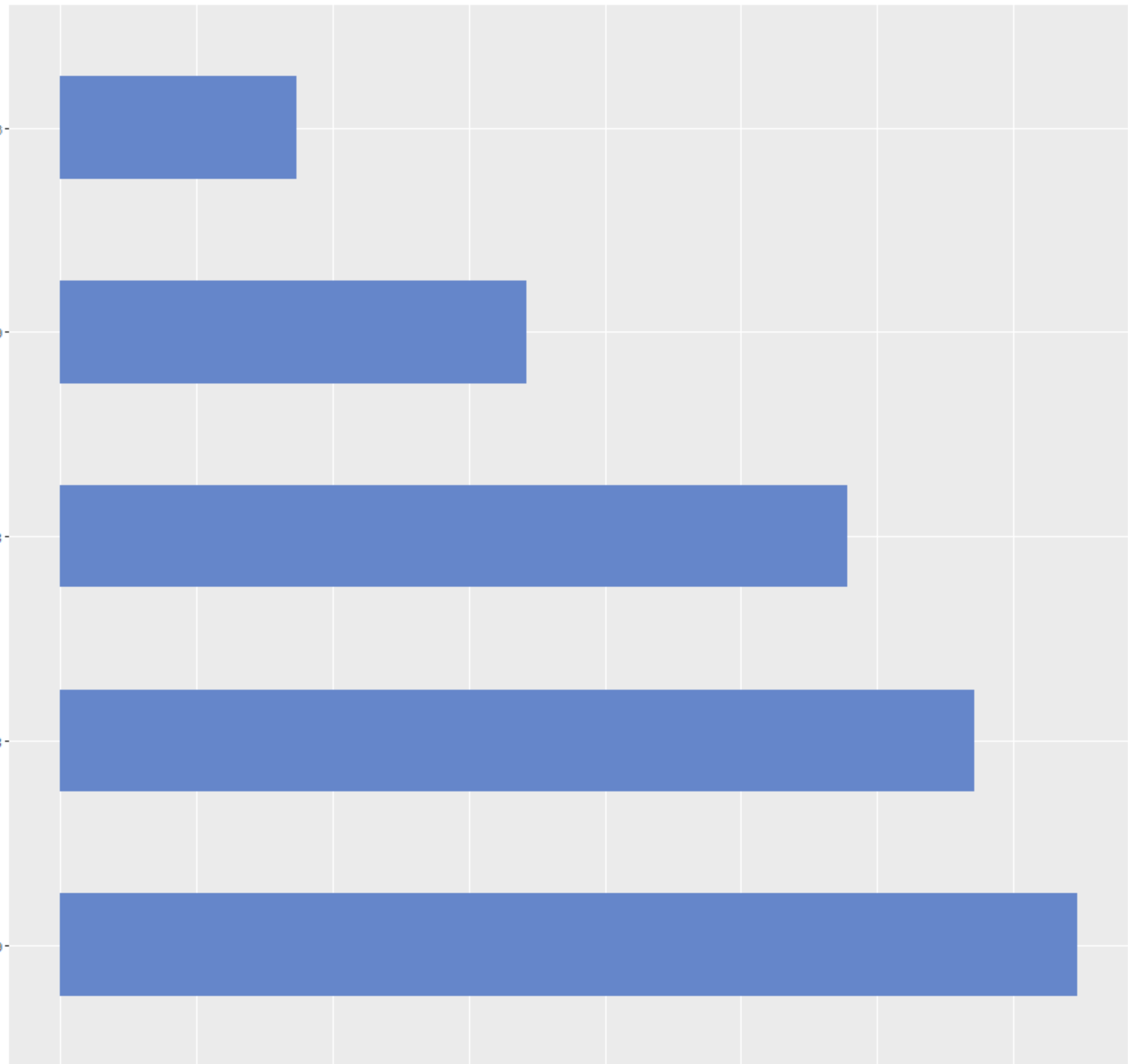
0.000

0.002

0.004

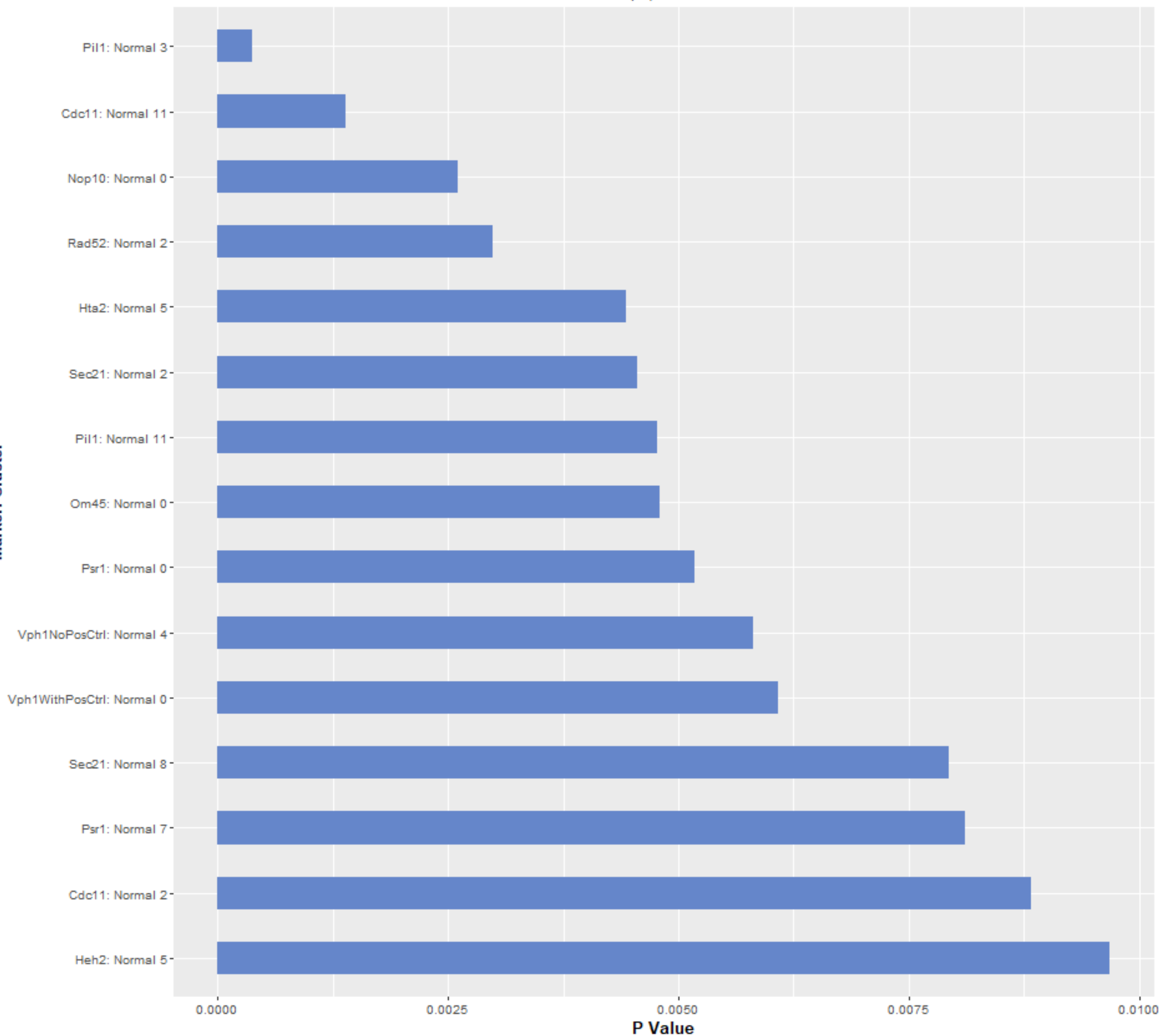
0.006

P Value



vacuole organization
With AreaShape | Unordered Gene Set

Marker: Cluster



vesicle organization
With AreaShape | Unordered Gene Set

Marker: Cluster

