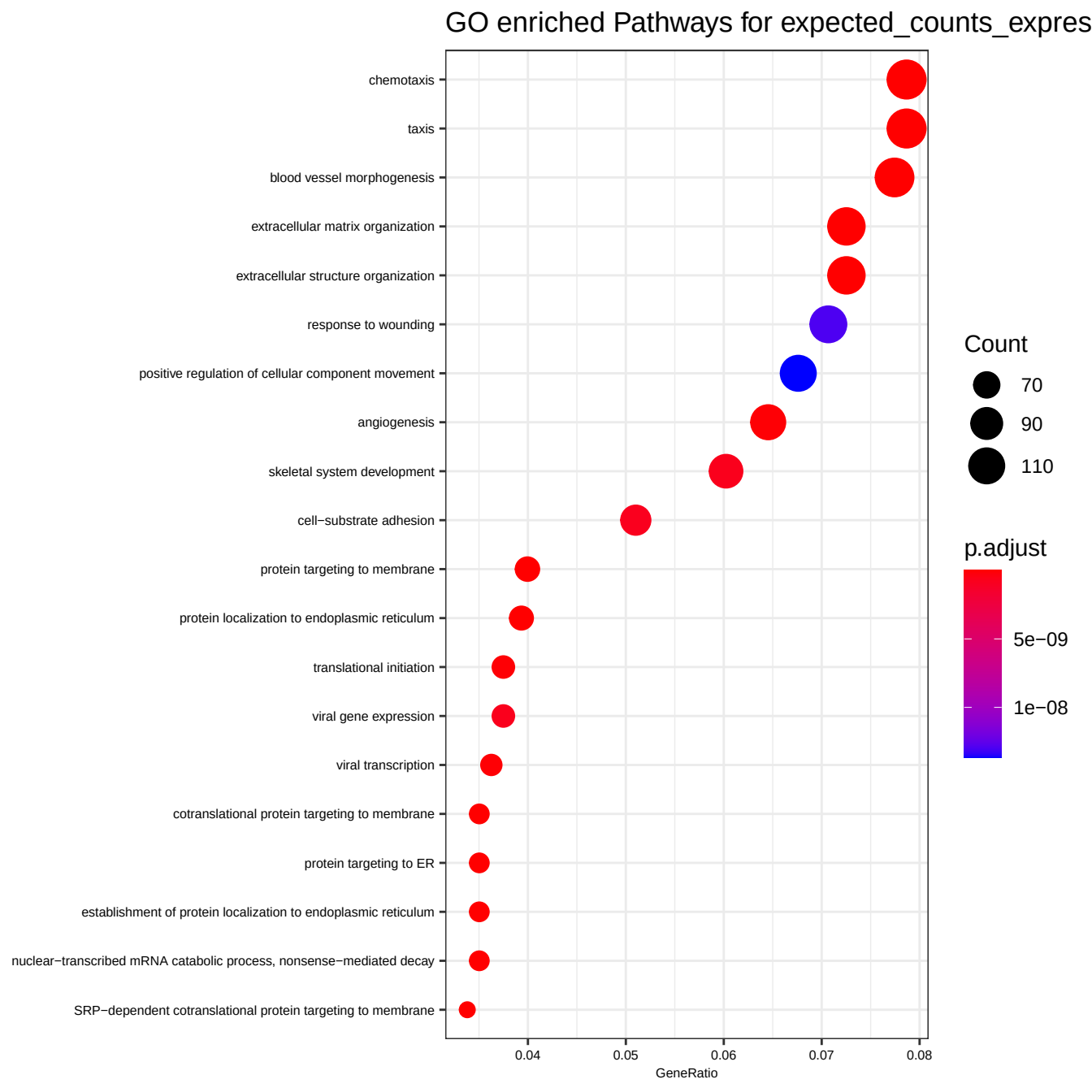
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chemotaxis
taxis
blood vessel morphogenesis
extracellular matrix organization
extracellular structure organization
response to wounding
positive regulation of cellular component movement
angiogenesis
skeletal system development
cell−substrate adhesion
protein targeting to membrane
protein localization to endoplasmic reticulum
translational initiation
viral gene expression
viral transcription
cotranslational protein targeting to membrane
protein targeting to ER
establishment of protein localization to endoplasmic reticulum
nuclear−transcribed mRNA catabolic process, nonsense−mediated decay
SRP−dependent cotranslational protein targeting to membrane
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0.08
GeneRatio
Count
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90
110
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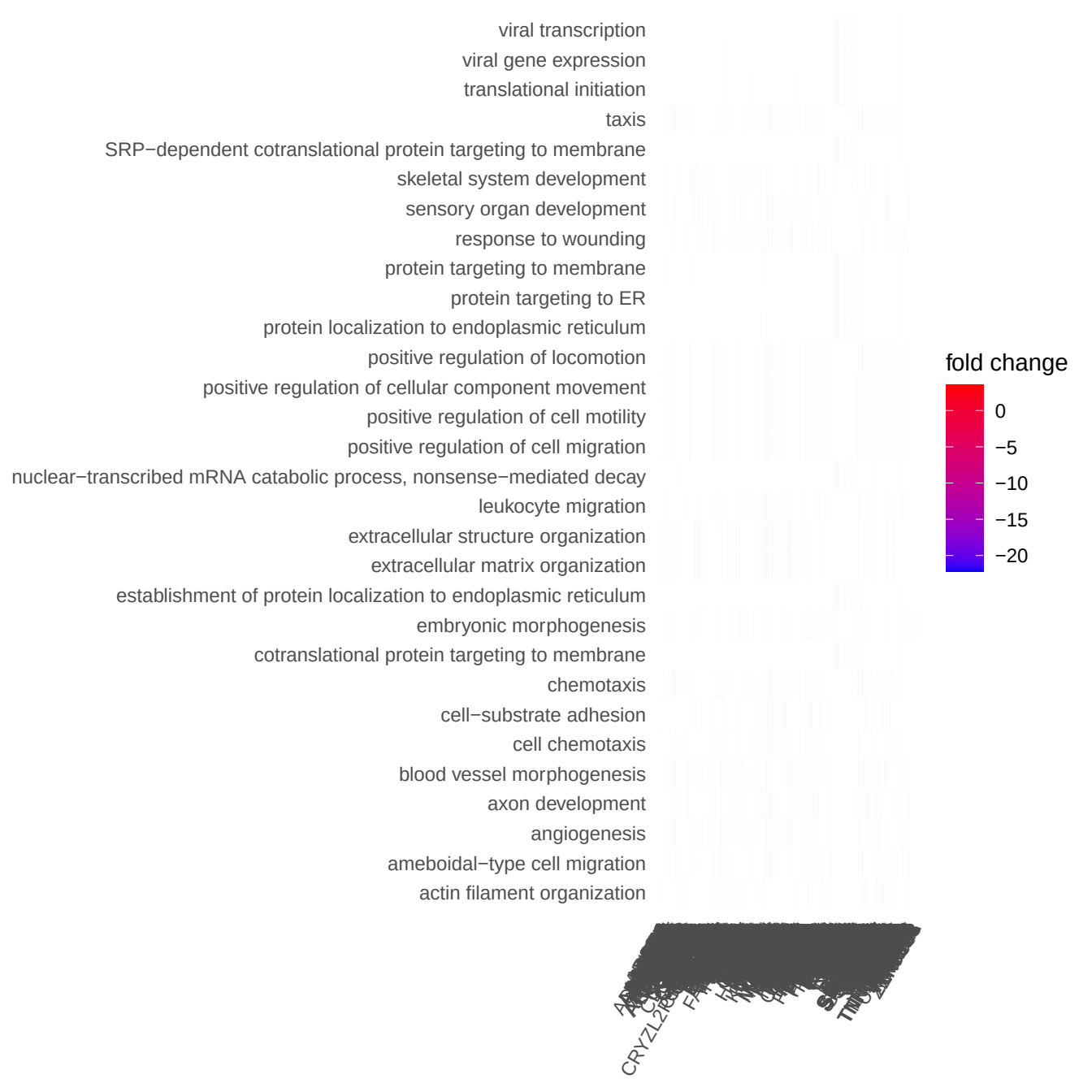
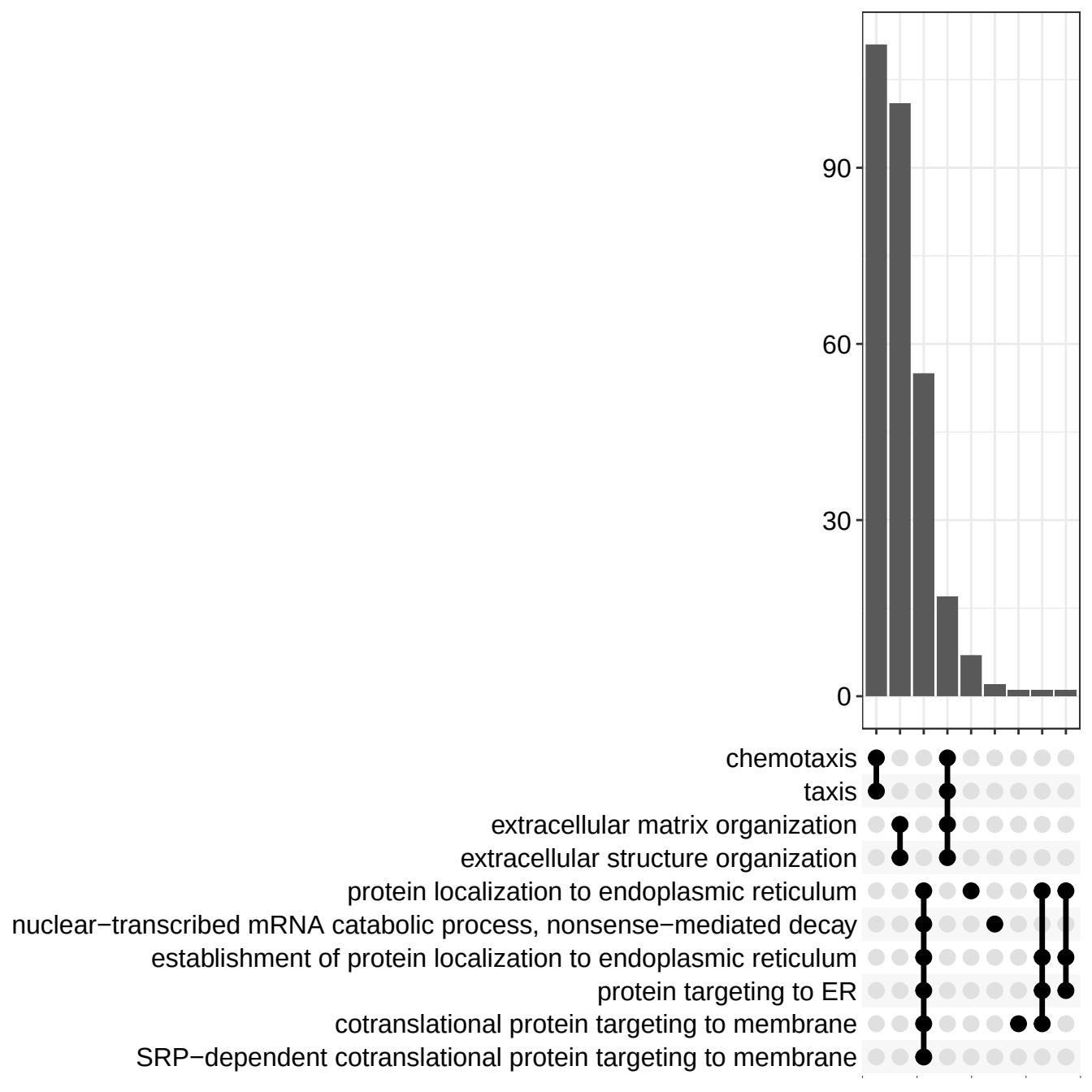

viral transcription
viral gene expression
translational initiation
taxis
SRP−dependent cotranslational protein targeting to membrane
skeletal system development
sensory organ development
response to wounding
protein targeting to membrane
protein targeting to ER
protein localization to endoplasmic reticulum
positive regulation of locomotion
positive regulation of cellular component movement
positive regulation of cell motility
positive regulation of cell migration
nuclear−transcribed mRNA catabolic process, nonsense−mediated decay
leukocyte migration
extracellular structure organization
extracellular matrix organization
establishment of protein localization to endoplasmic reticulum
embryonic morphogenesis
cotranslational protein targeting to membrane
chemotaxis
cell−substrate adhesion
cell chemotaxis
blood vessel morphogenesis
axon development
angiogenesis
ameboidal−type cell migration
actin filament organization
fold change
0
−5
−10
−15
−20


90
60
30
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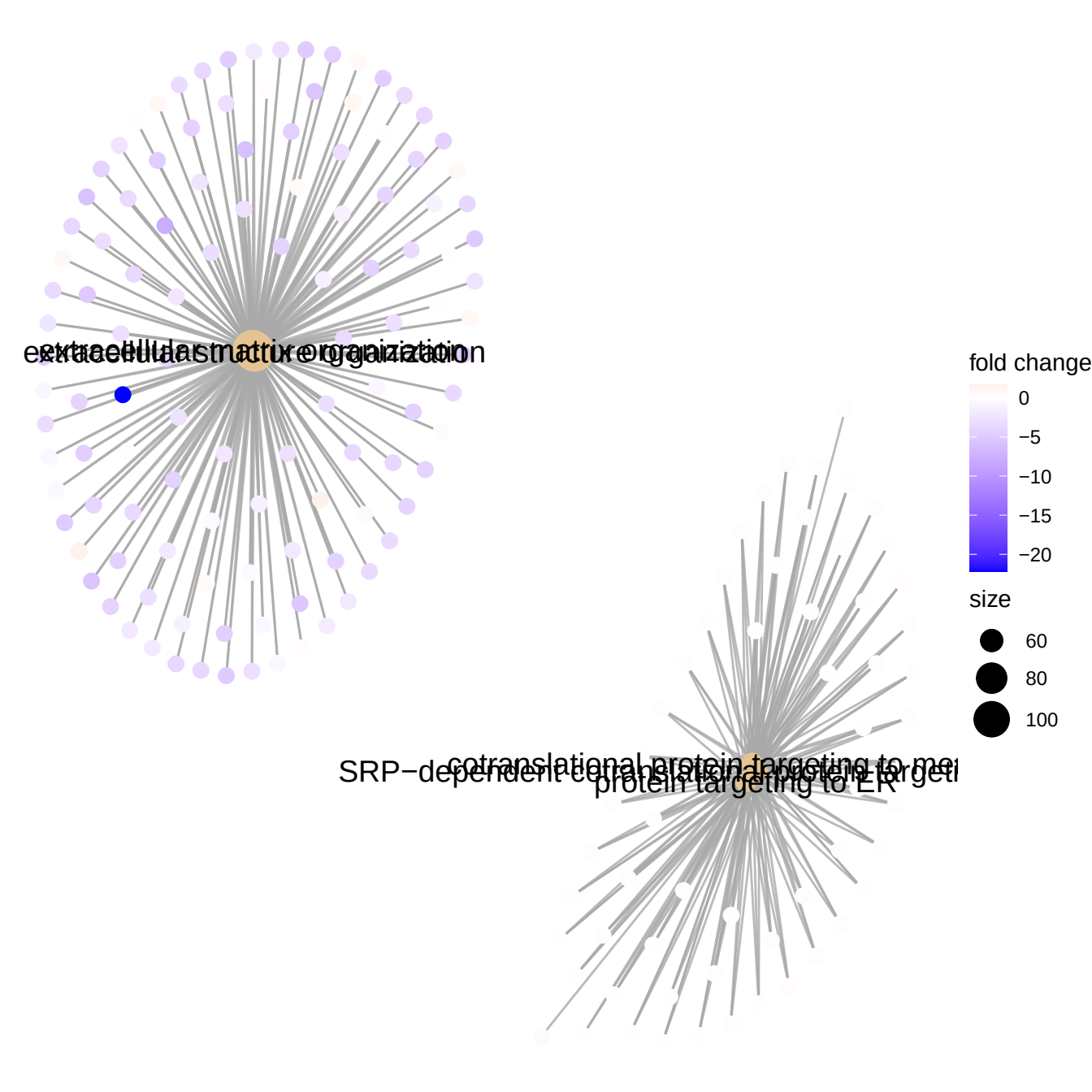
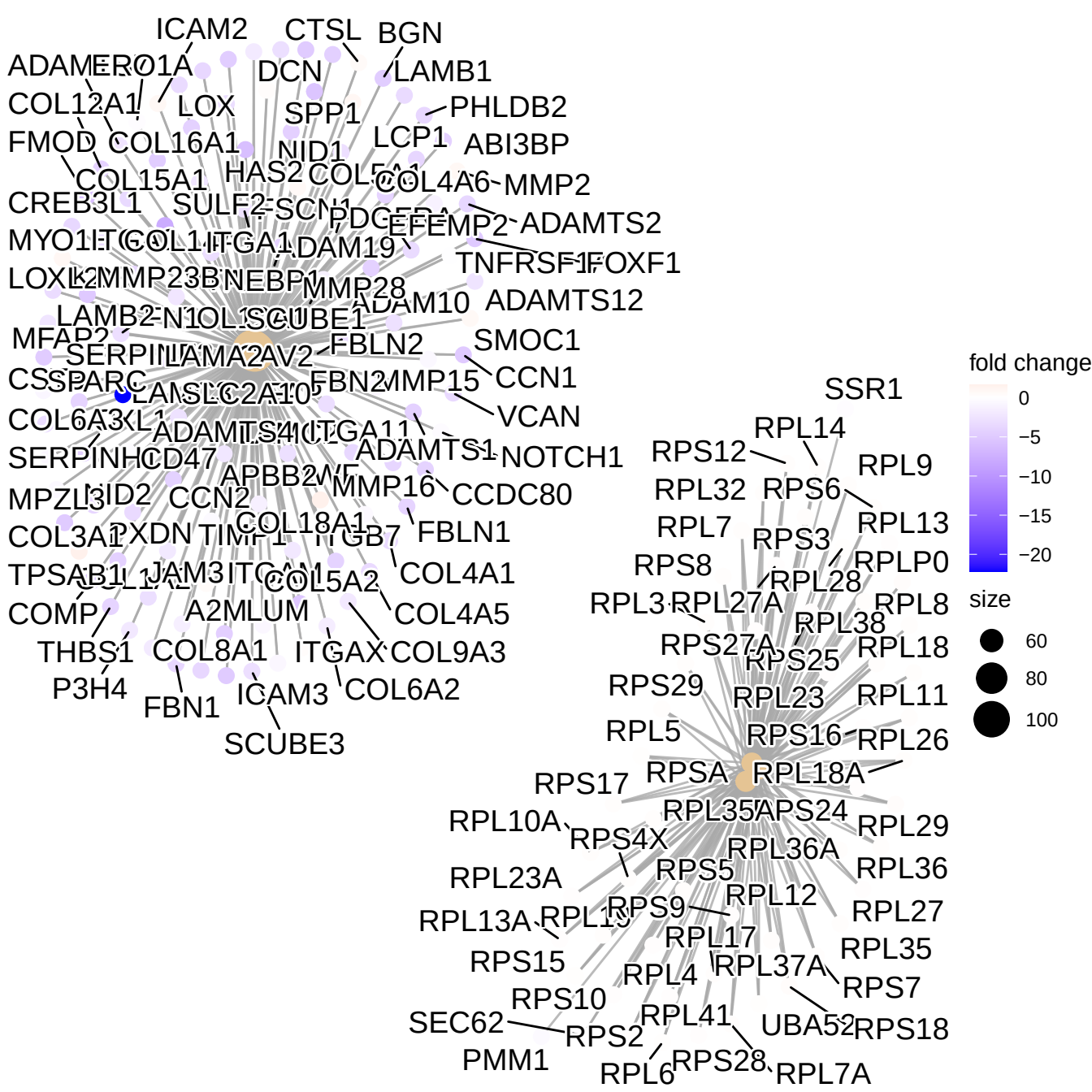
chemotaxis
taxis
extracellular matrix organization
extracellular structure organization
protein localization to endoplasmic reticulum
nuclear−transcribed mRNA catabolic process, nonsense−mediated decay
establishment of protein localization to endoplasmic reticulum
protein targeting to ER
cotranslational protein targeting to membrane
SRP−dependent cotranslational protein targeting to membrane


extracellular matrix organization
SRP−dependent cotranslational protein targeting to membrane
protein targeting to ER
fold change
0
−5
−10
−15
−20
size
60
80
100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LCP1
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HAS2
COL5A1
COL4A6
MMP2
CREB3L1
SULF2
SCN1
PDGFRA
EFEMP2
ADAMTS2
MYO1B
ITGA1
ADAM19
TNFRSF11B
FOXF1
LOXL2
MMP23B
NEBP1
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ADAM10
ADAMTS12
LAMB2
N1OL1
SCUBE1
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FBLN2
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SERPINH1
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FBN2
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TIMP1
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FBLN1
TPSAB1
JAM3
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A2MLUM
COL4A5
THBS1
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COL6A2
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RPL5
RPS16
RPL26
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RPL19
RPS9
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RPS15
RPL35
RPL4
RPL37A
RPS7
RPS10
RPL41
UBA52
RPS18
SEC62
RPS2
PMM1
RPL6
RPS28
RPL7A
fold change
0
−5
−10
−15
−20
size
60
80
100