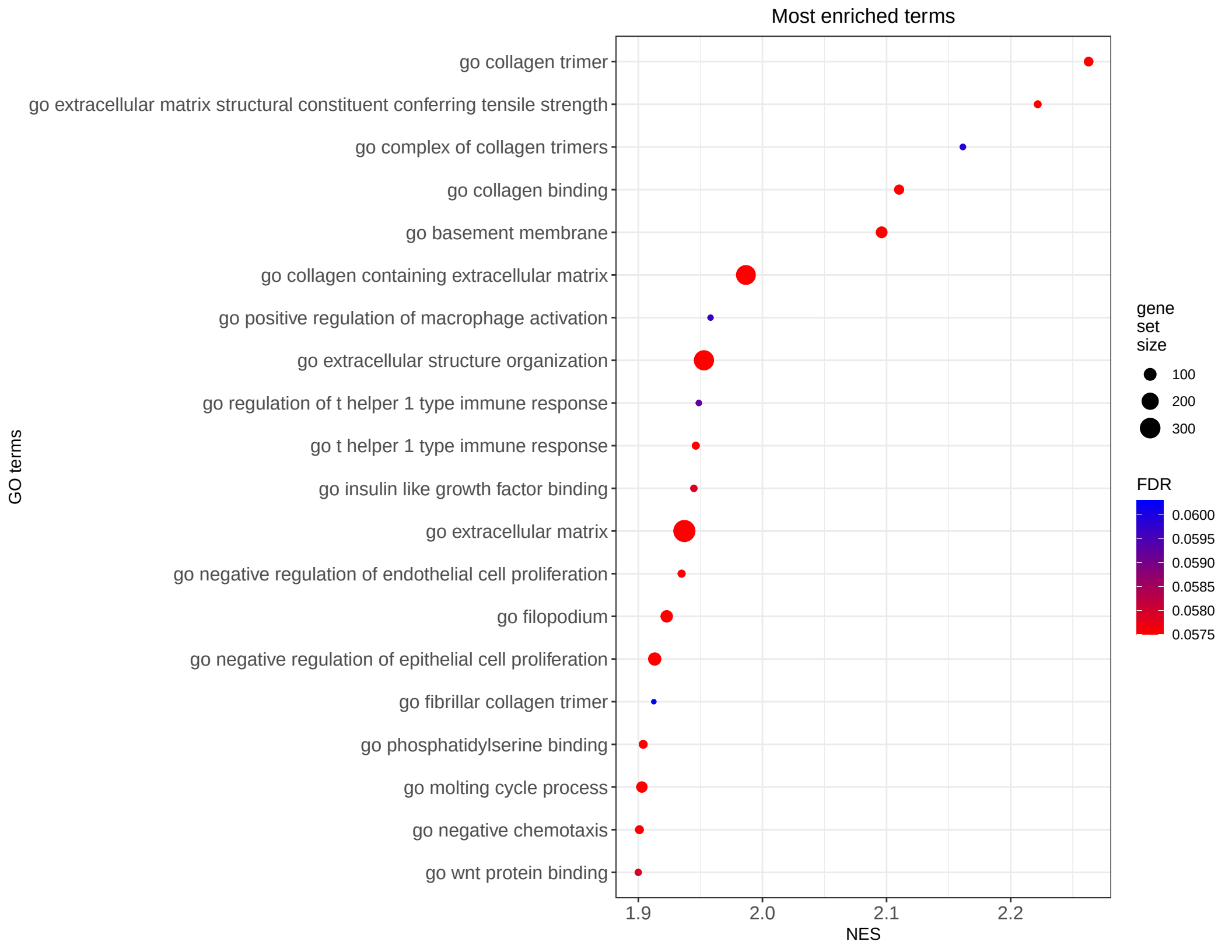
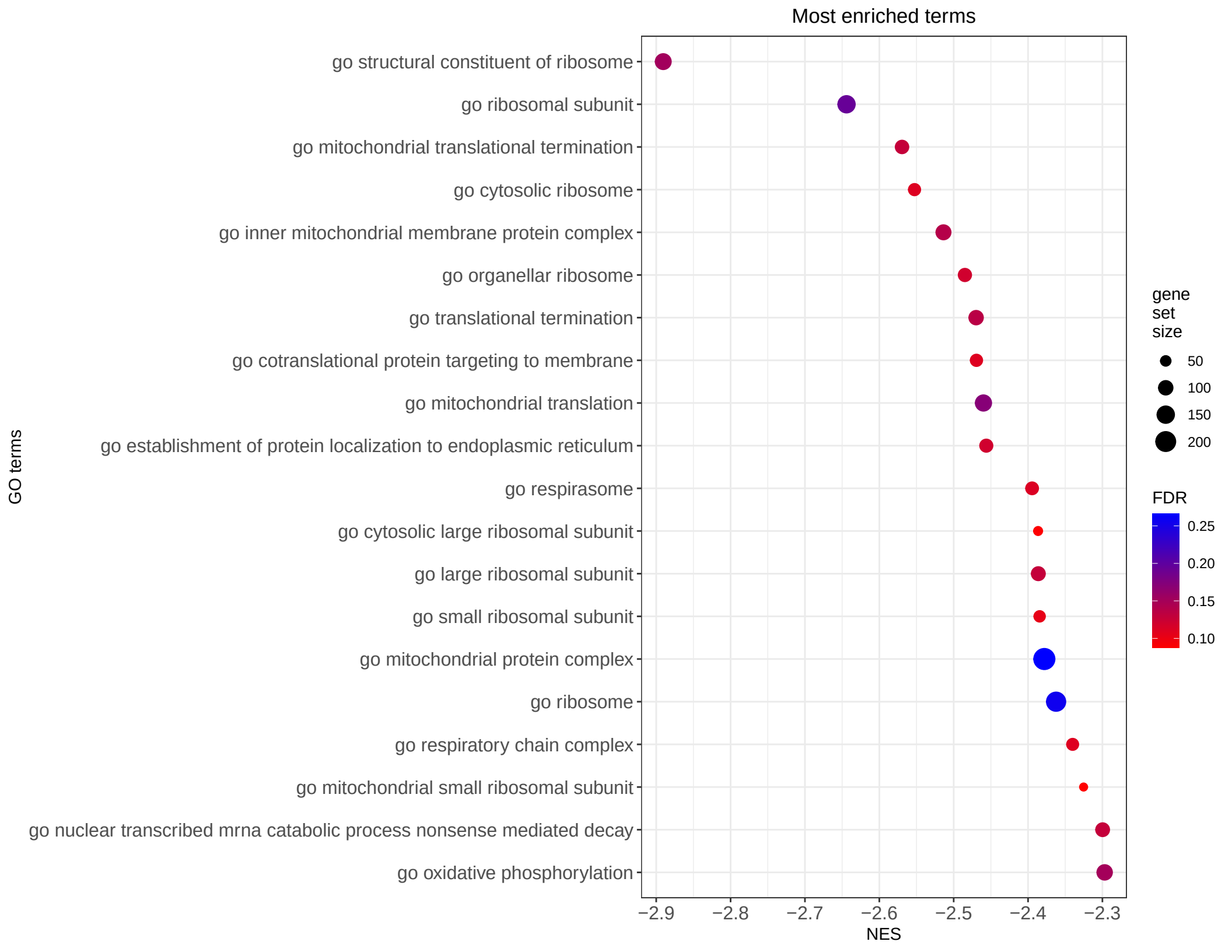
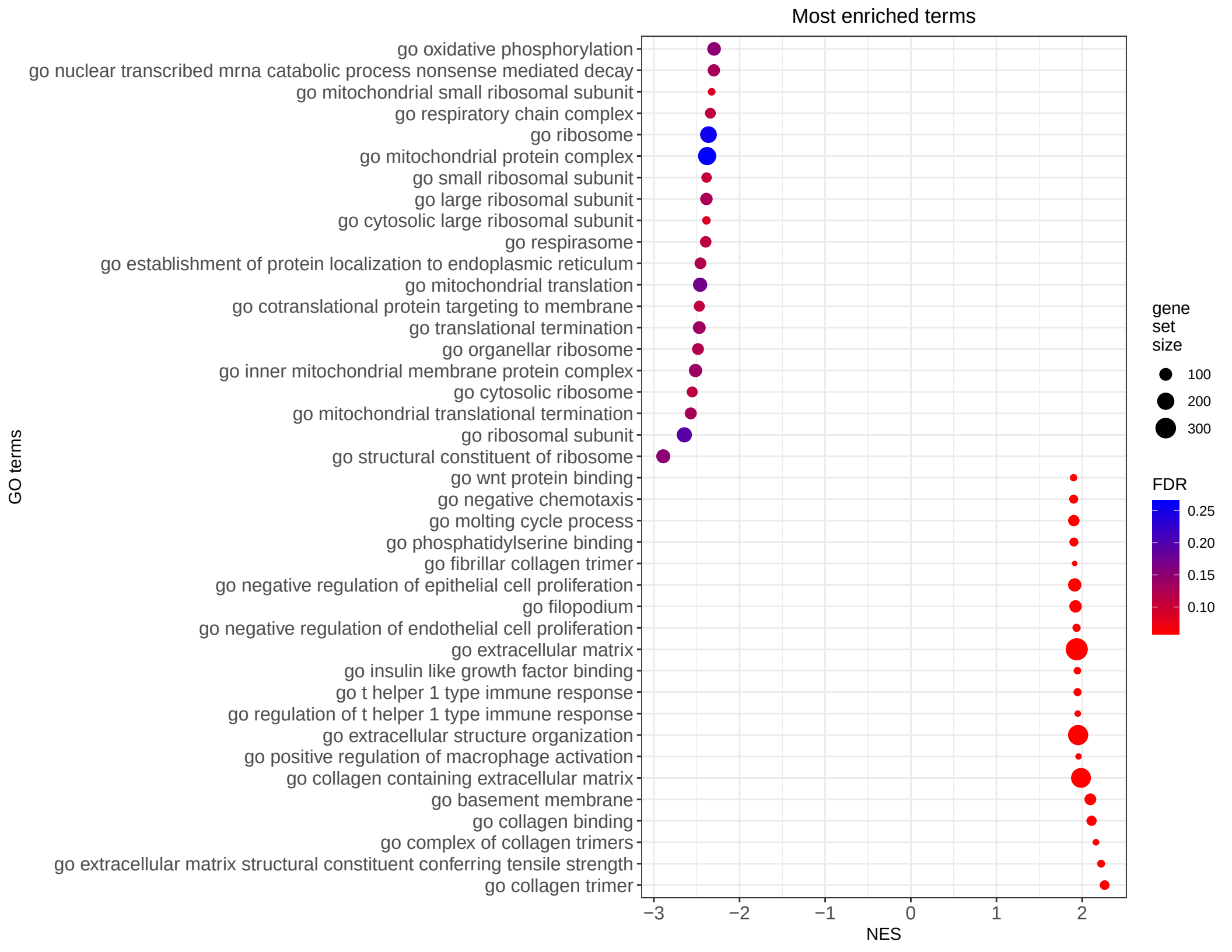

Most enriched terms
go collagen trimer
go extracellular matrix structural constituent conferring tensile strength
go complex of collagen trimers
go collagen binding
go basement membrane
go collagen containing extracellular matrix
go positive regulation of macrophage activation
go extracellular structure organization
go regulation of t helper 1 type immune response
go t helper 1 type immune response
go insulin like growth factor binding
go extracellular matrix
go negative regulation of endothelial cell proliferation
go filopodium
go negative regulation of epithelial cell proliferation
go fibrillar collagen trimer
go phosphatidylserine binding
go molting cycle process
go negative chemotaxis
go wnt protein binding
GO terms
1.9
2.0
2.1
2.2
NES
gene set size
100
200
300
FDR
0.0600
0.0595
0.0590
0.0585
0.0580
0.0575


Most enriched terms
go structural constituent of ribosome
go ribosomal subunit
go mitochondrial translational termination
go cytosolic ribosome
go inner mitochondrial membrane protein complex
go organellar ribosome
go translational termination
go cotranslational protein targeting to membrane
go mitochondrial translation
go establishment of protein localization to endoplasmic reticulum
go respirasome
go cytosolic large ribosomal subunit
go large ribosomal subunit
go small ribosomal subunit
go mitochondrial protein complex
go ribosome
go respiratory chain complex
go mitochondrial small ribosomal subunit
go nuclear transcribed mrna catabolic process nonsense mediated decay
go oxidative phosphorylation
GO terms
−2.9
−2.8
−2.7
−2.6
−2.5
−2.4
−2.3
NES
gene set size
50
100
150
200
FDR
0.25
0.20
0.15
0.10


Most enriched terms
go oxidative phosphorylation
go nuclear transcribed mrna catabolic process nonsense mediated decay
go mitochondrial small ribosomal subunit
go respiratory chain complex
go ribosome
go mitochondrial protein complex
go small ribosomal subunit
go large ribosomal subunit
go cytosolic large ribosomal subunit
go respirasome
go establishment of protein localization to endoplasmic reticulum
go mitochondrial translation
go cotranslational protein targeting to membrane
go translational termination
go organellar ribosome
go inner mitochondrial membrane protein complex
go cytosolic ribosome
go mitochondrial translational termination
go ribosomal subunit
go structural constituent of ribosome
go wnt protein binding
go negative chemotaxis
go molting cycle process
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go regulation of t helper 1 type immune response
go extracellular structure organization
go positive regulation of macrophage activation
go collagen containing extracellular matrix
go basement membrane
go collagen binding
go complex of collagen trimers
go extracellular matrix structural constituent conferring tensile strength
go collagen trimer
GO terms
−3
−2
−1
0
1
2
NES
gene set size
100
200
300
FDR
0.25
0.20
0.15
0.10