

Protein processing in endoplasmic reticulum

Antigen processing and presentation

Tuberculosis

Salmonella infection

Th17 cell differentiation

HTLV-I infection

pos
(15)

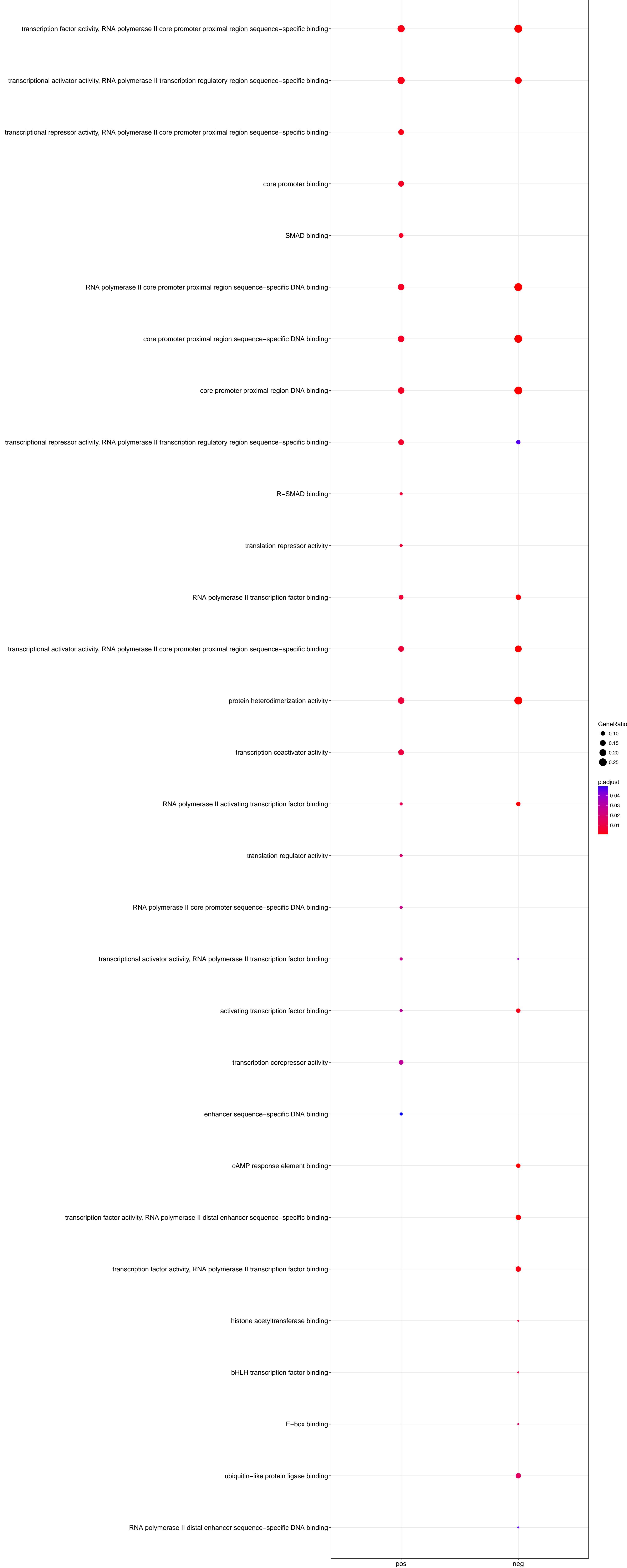
neg
(20)

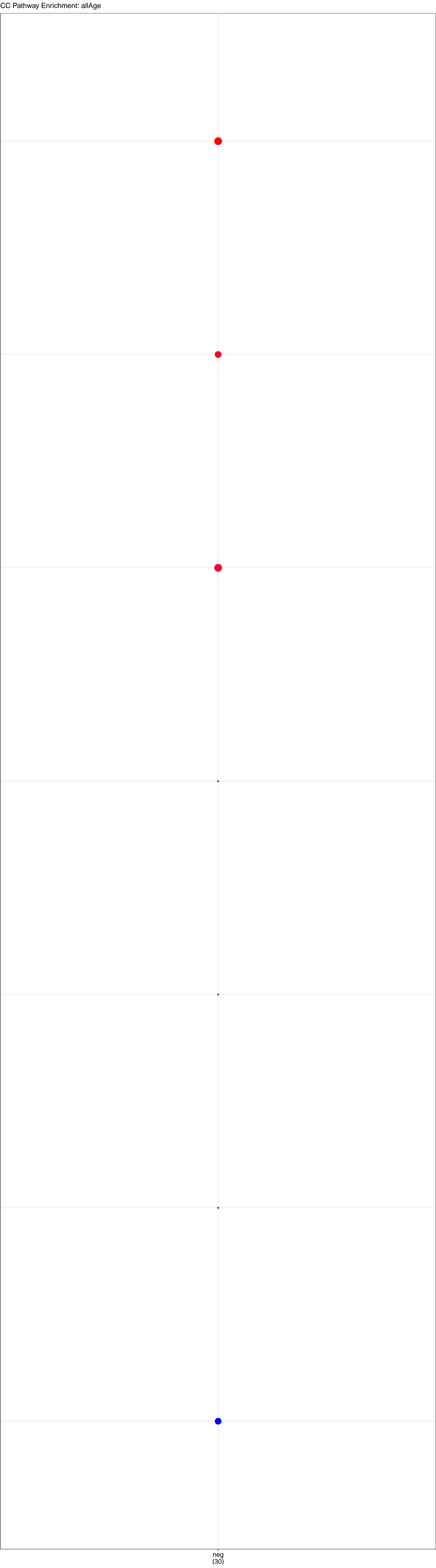
GeneRatio

- 0.15
- 0.18
- 0.21
- 0.24

p.adjust

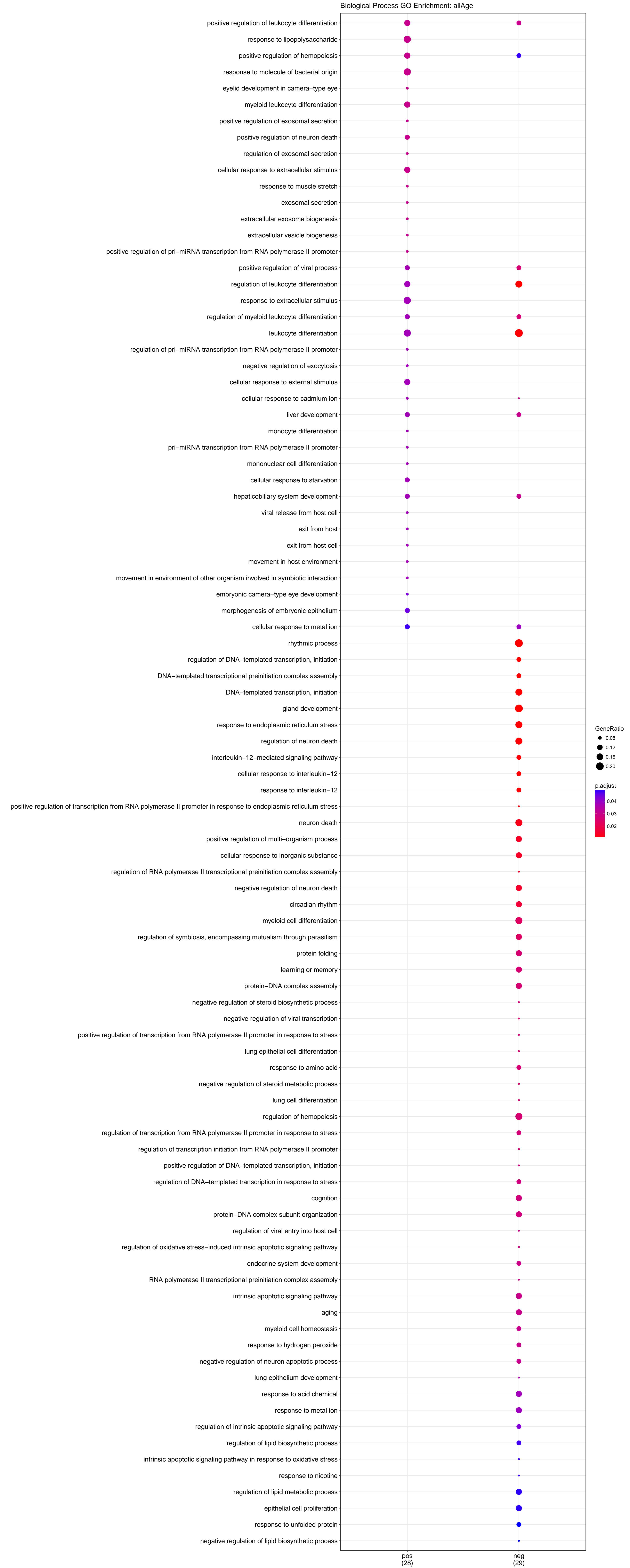
- 0.044
- 0.042
- 0.040





neg
(30)

Biological Process GO Enrichment: allAge



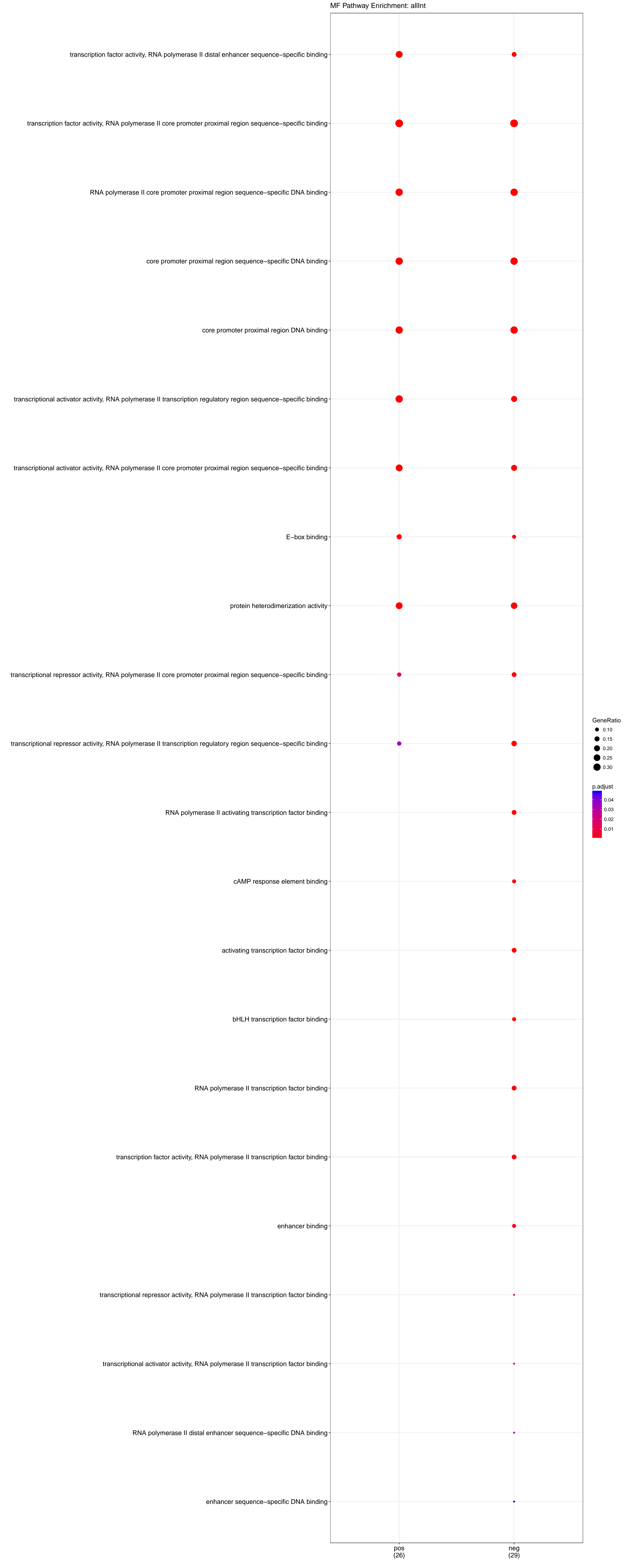


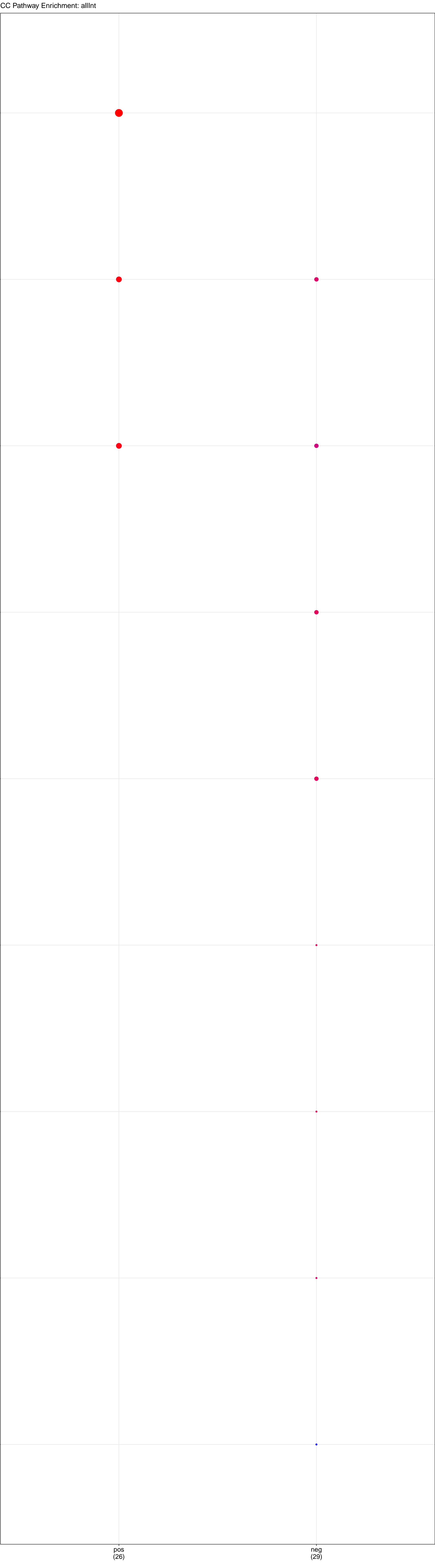
GeneRatio

- 0.14
- 0.16
- 0.18
- 0.20
- 0.22

p.adjust

- 0.03
- 0.02
- 0.01





transcription factor complex

RNA polymerase II transcription factor complex

nuclear transcription factor complex

melanosome

pigment granule

dendrite cytoplasm

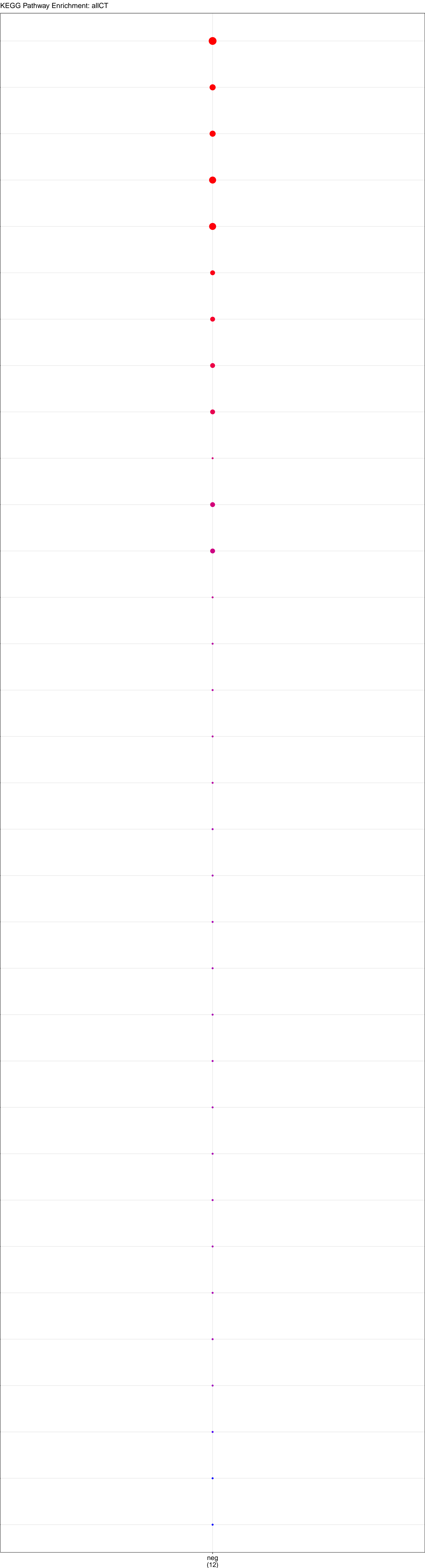
nuclear euchromatin

euchromatin

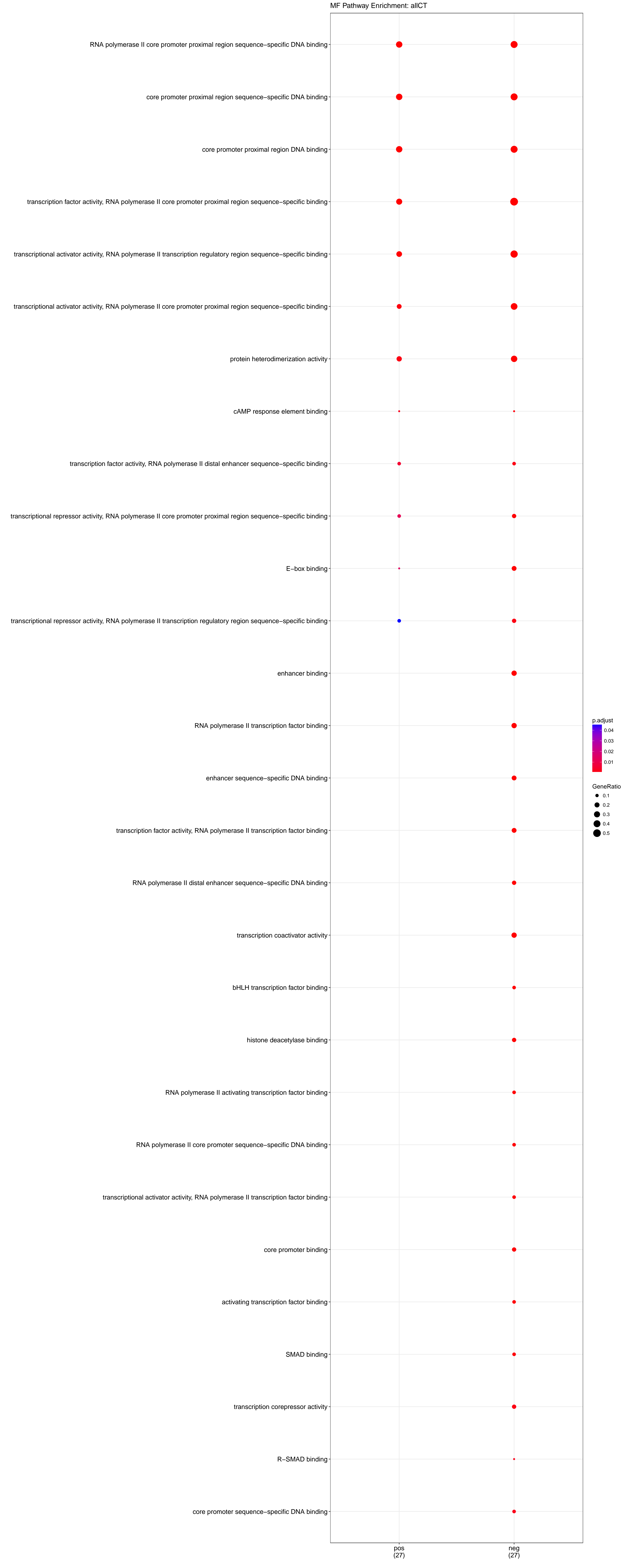
cell projection cytoplasm

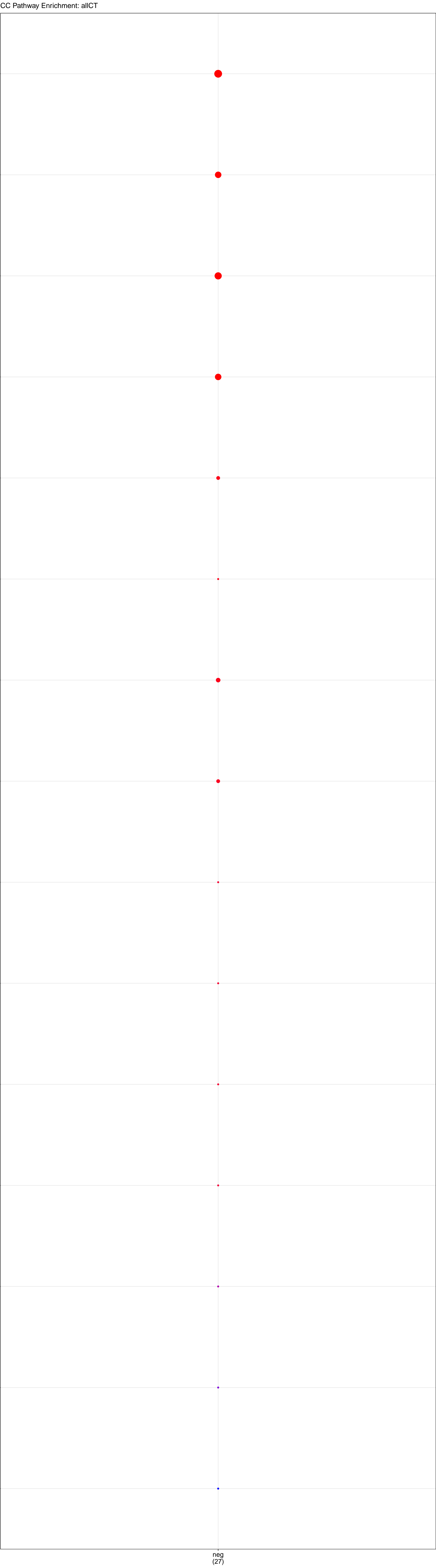
Biological Process GO Enrichment: allInt





neg
(12)





transcription factor complex

RNA polymerase II transcription factor complex

nuclear chromatin

nuclear transcription factor complex

histone methyltransferase complex

Set1C/COMPASS complex

protein-DNA complex

methyltransferase complex

nuclear euchromatin

MLL1/2 complex

MLL1 complex

euchromatin

SWI/SNF superfamily-type complex

transcriptional repressor complex

ATPase complex

GeneRatio

0.1

0.2

0.3

0.4

p.adjust

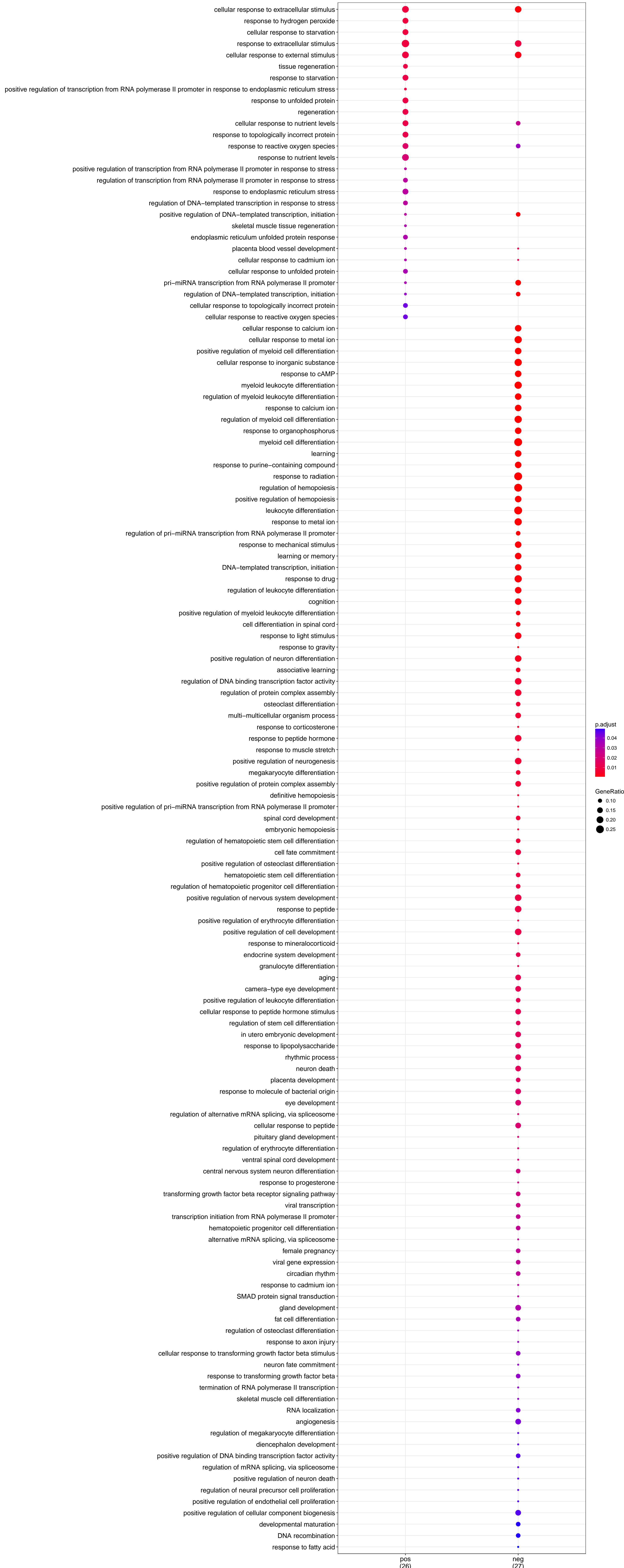
0.03

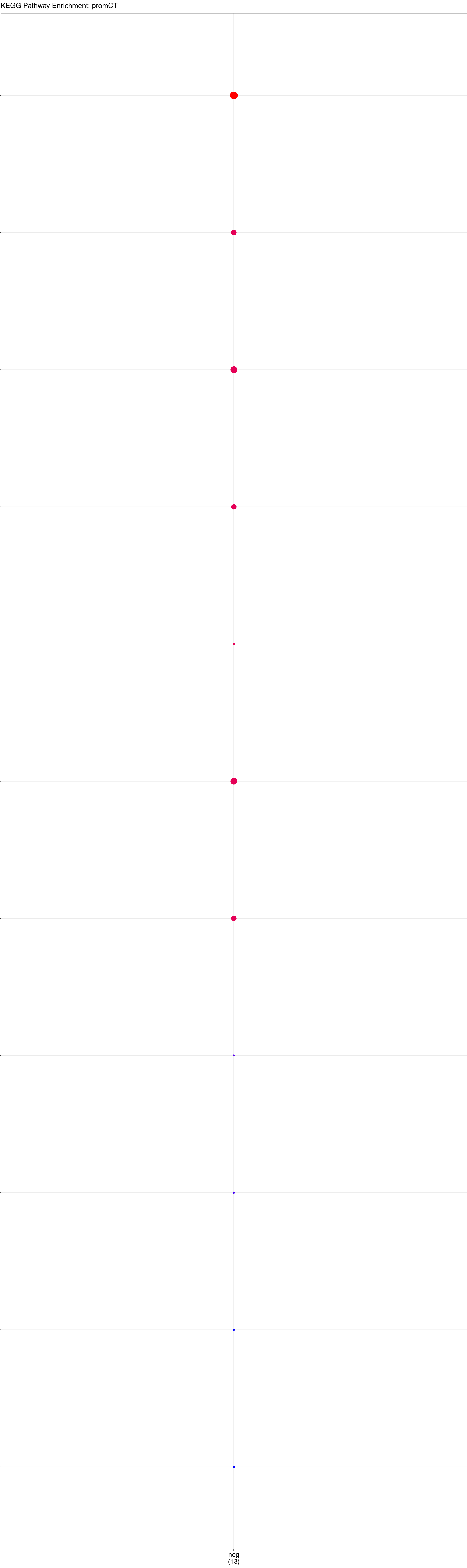
0.02

0.01

neg
(27)

Biological Process GO Enrichment: allCT

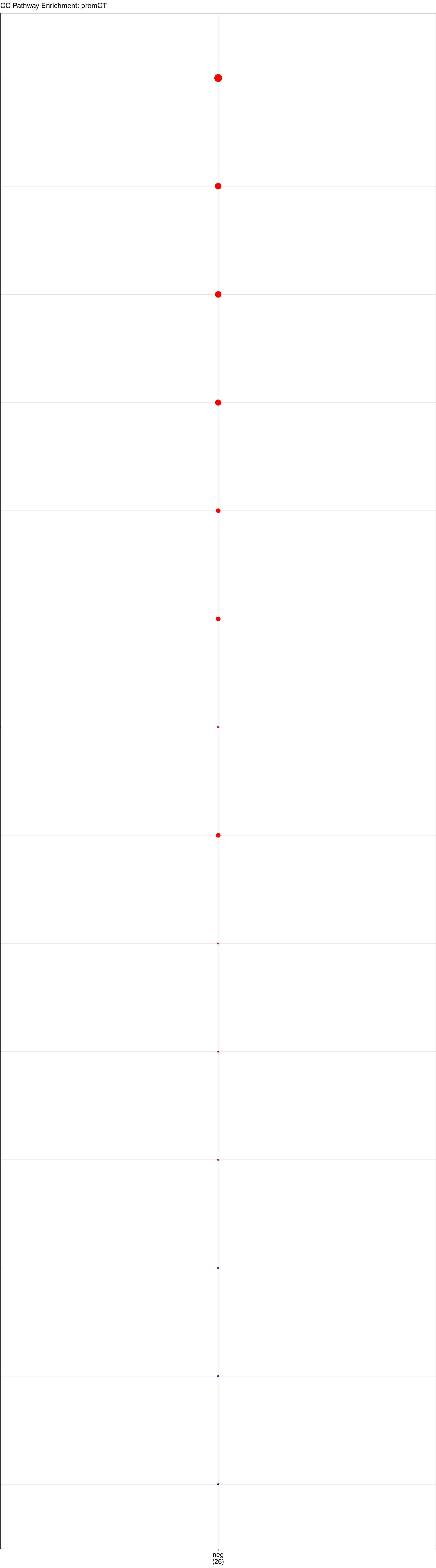




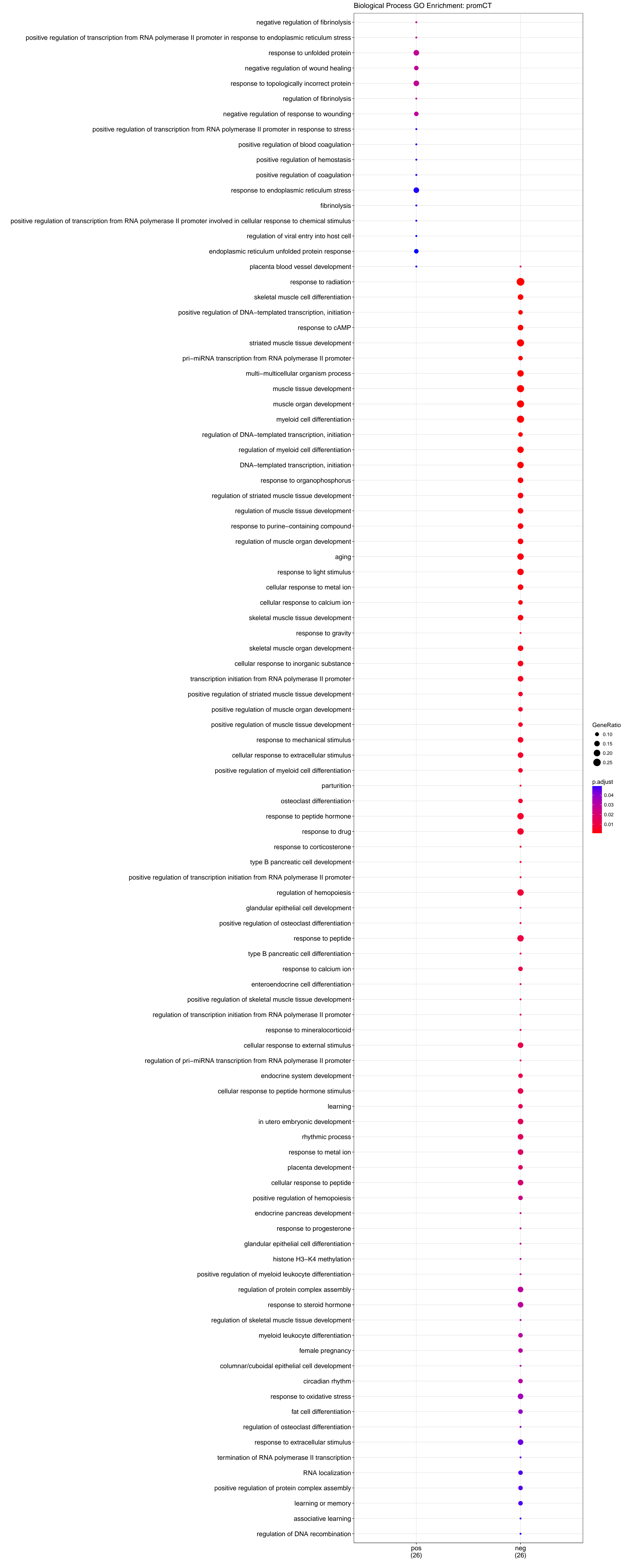
neg
(13)

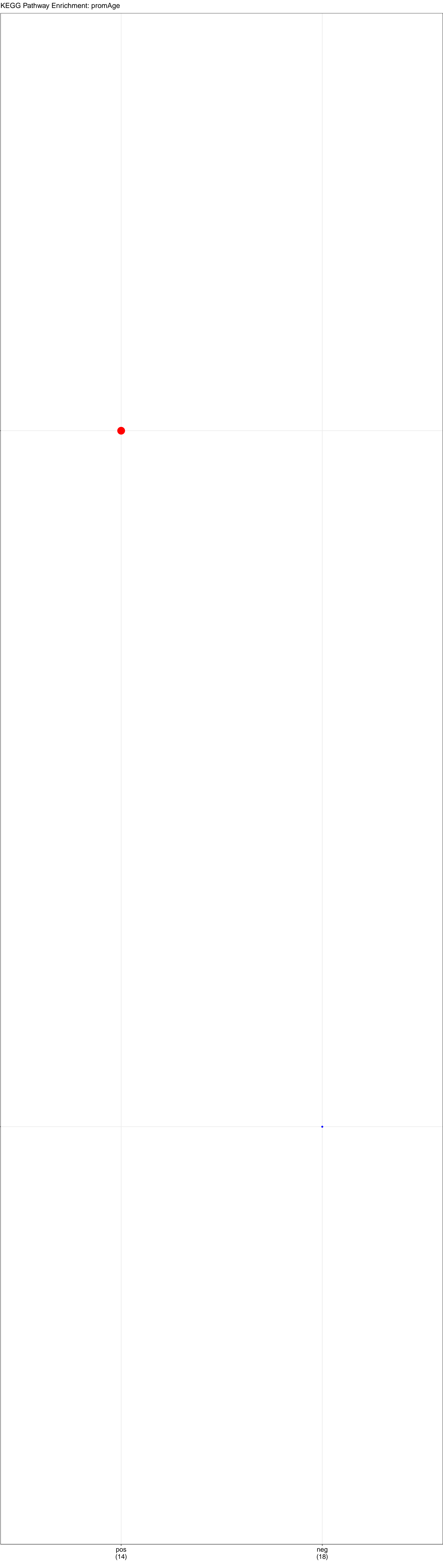
MF Pathway Enrichment: promCT





neg
(26)





Non-alcoholic fatty liver disease (NAFLD)

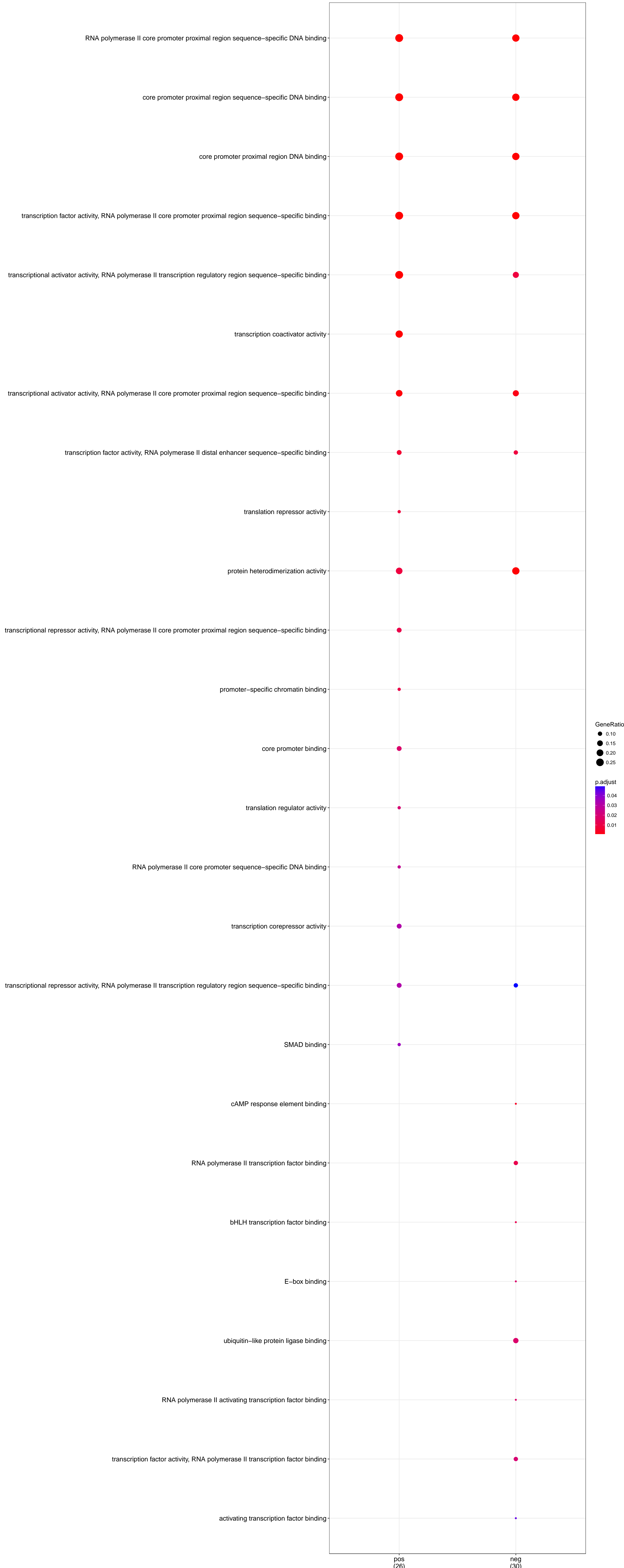
Protein processing in endoplasmic reticulum

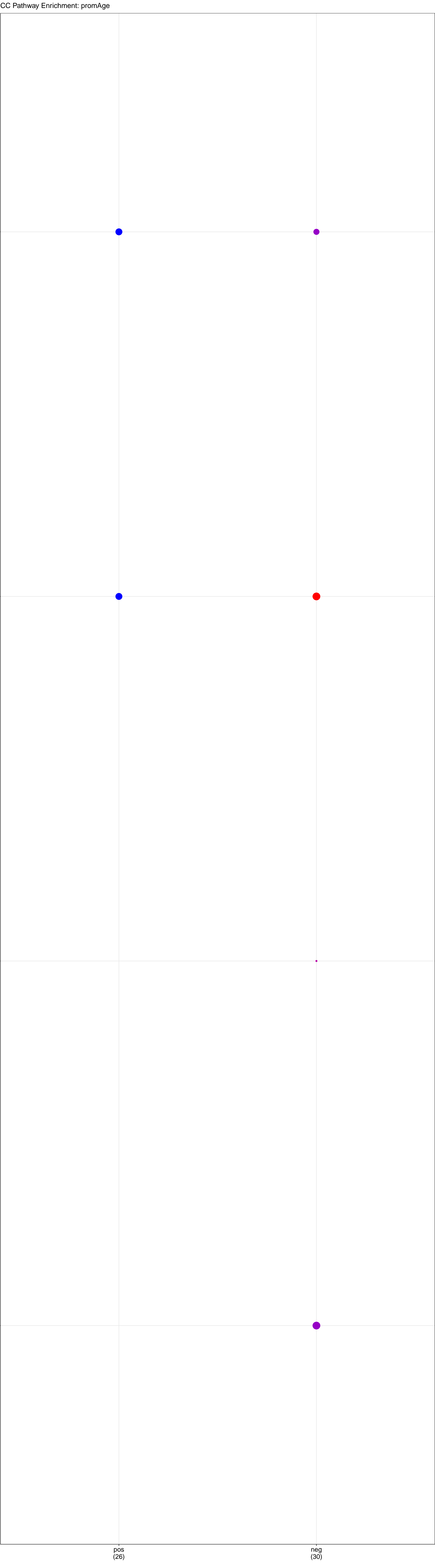
pos

(14)

neg

(18)





RNA polymerase II transcription factor complex

nuclear transcription factor complex

dendrite cytoplasm

transcription factor complex

pos

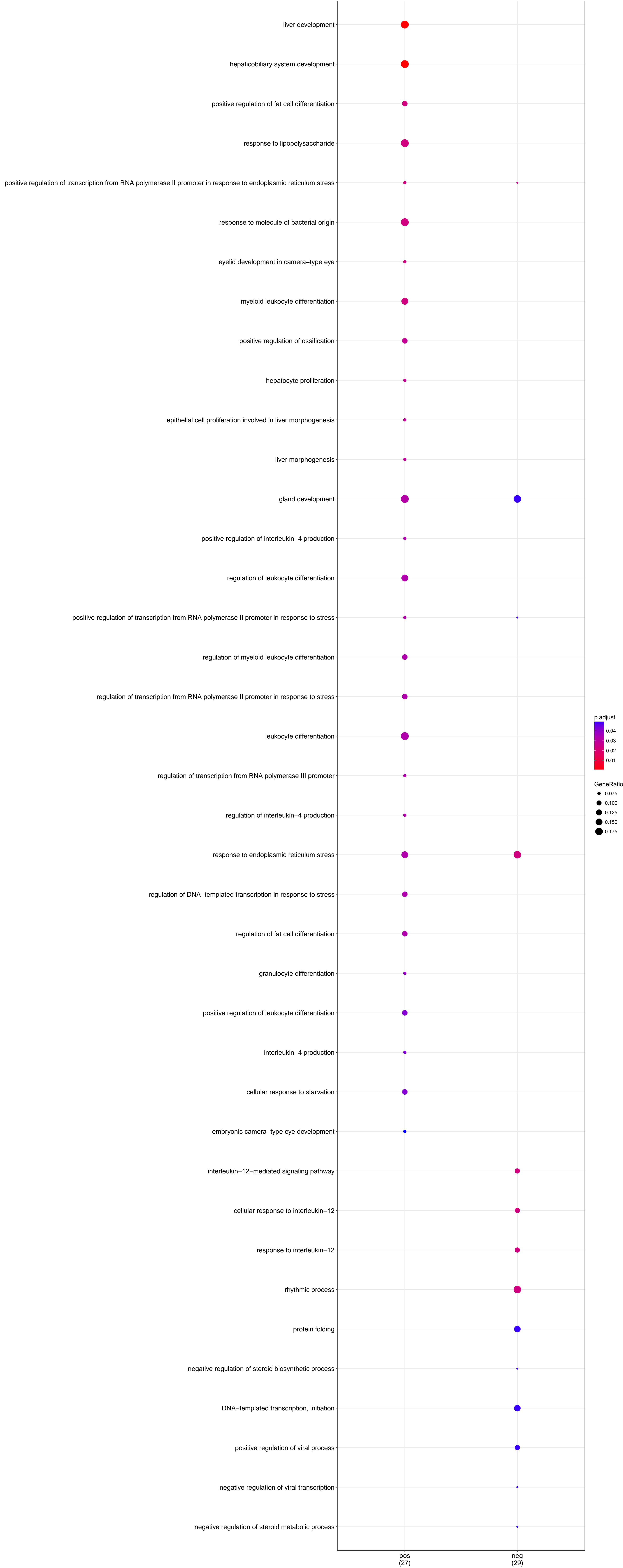
(26)

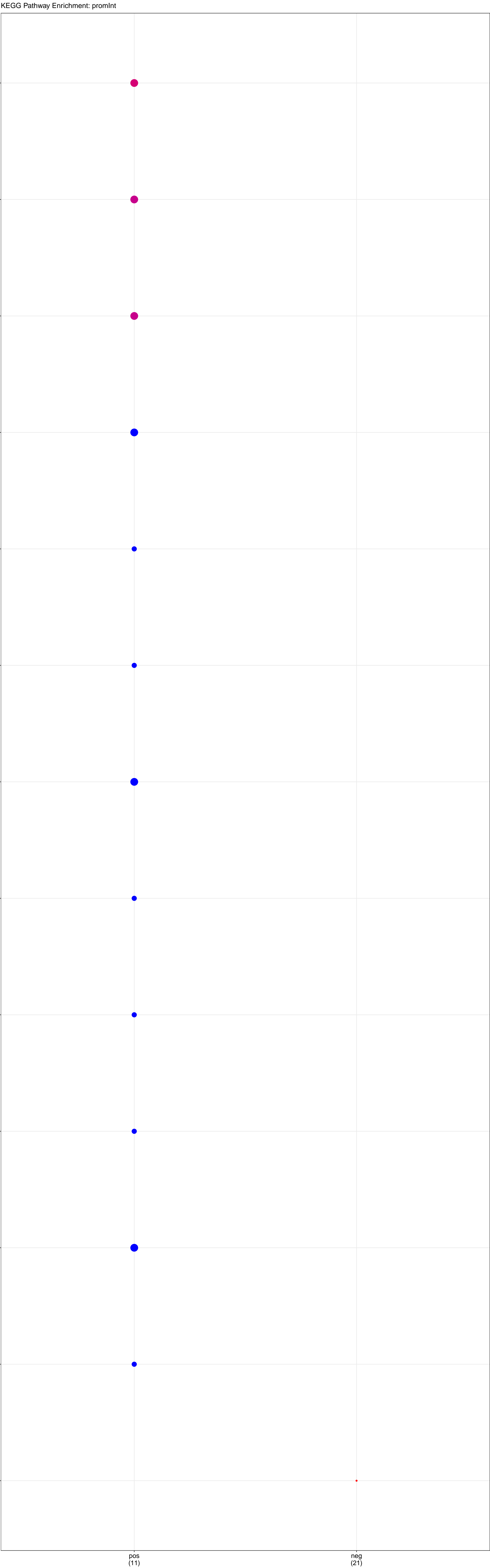
neg

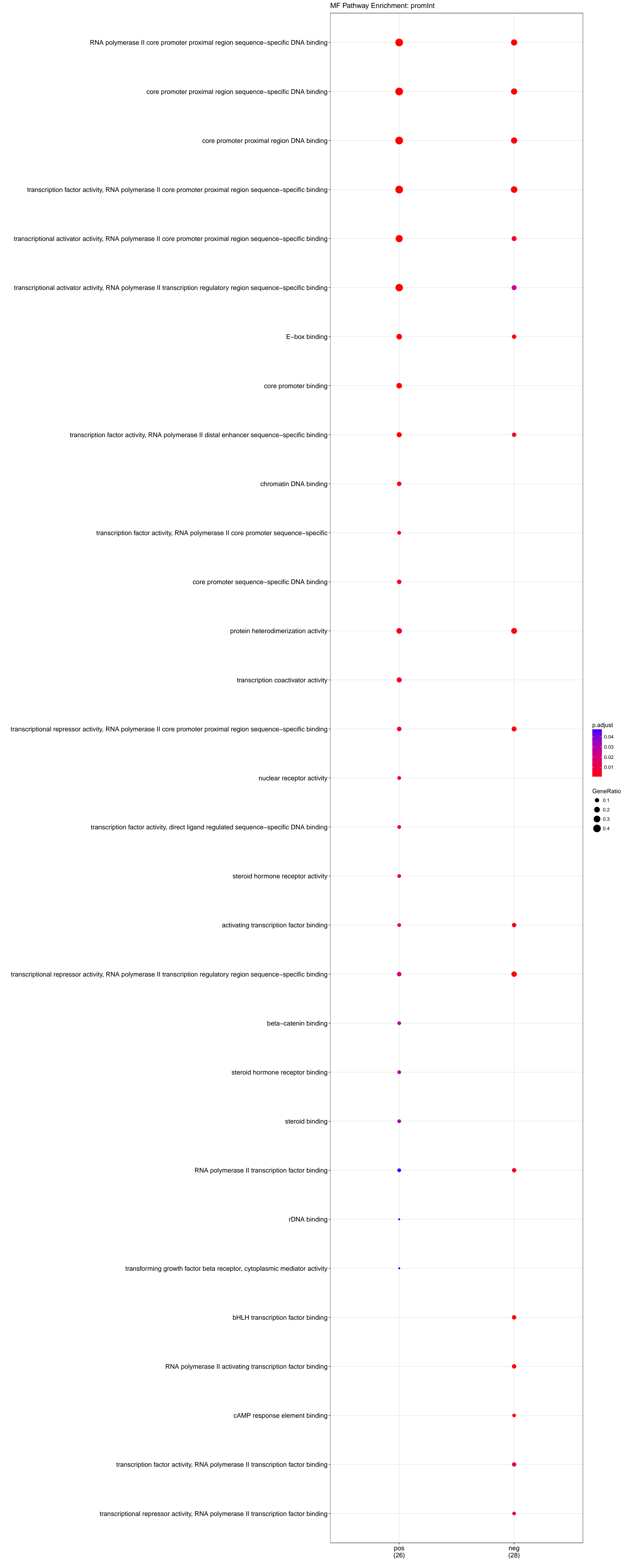
(30)

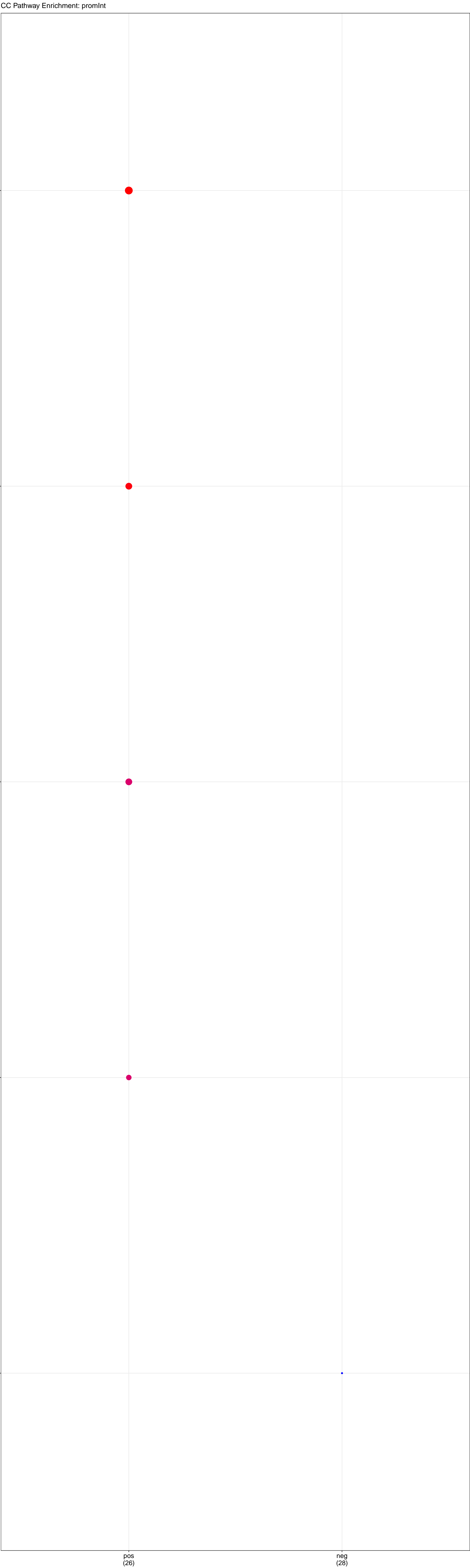


Biological Process GO Enrichment: promAge









p.adjust

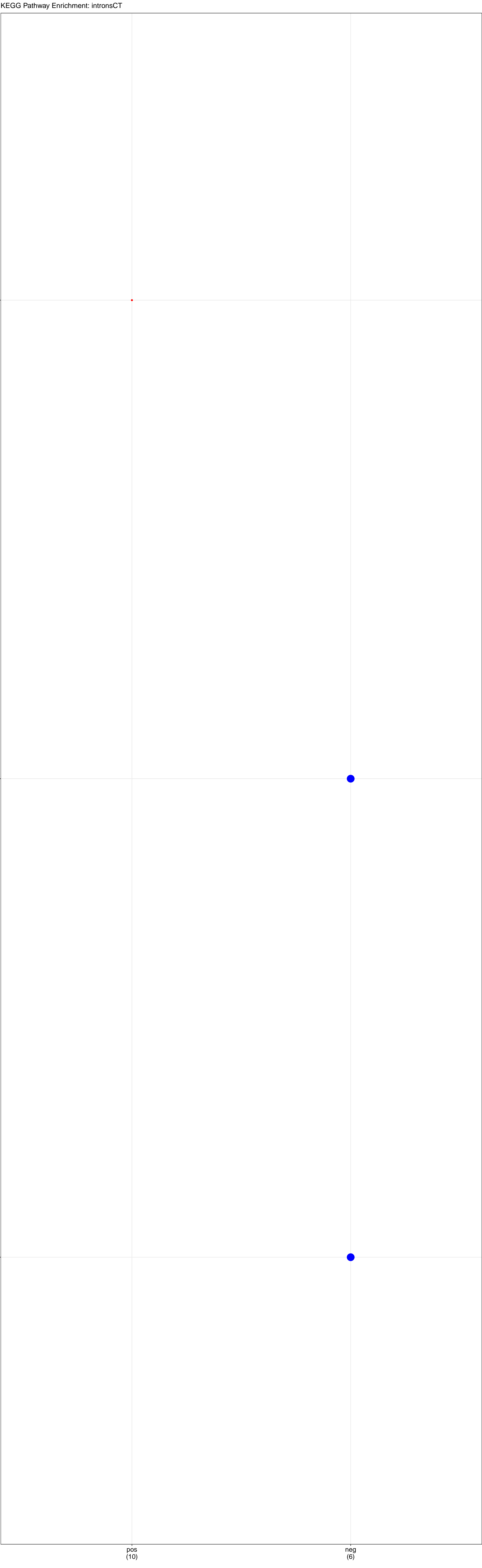
0.04
0.03
0.02
0.01

GeneRatio

0.075
0.100
0.125
0.150
0.175

Biological Process GO Enrichment: promInt





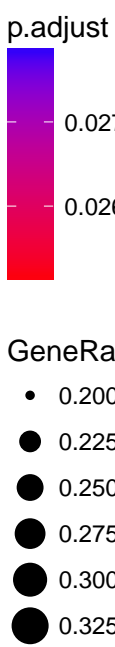
Base excision repair

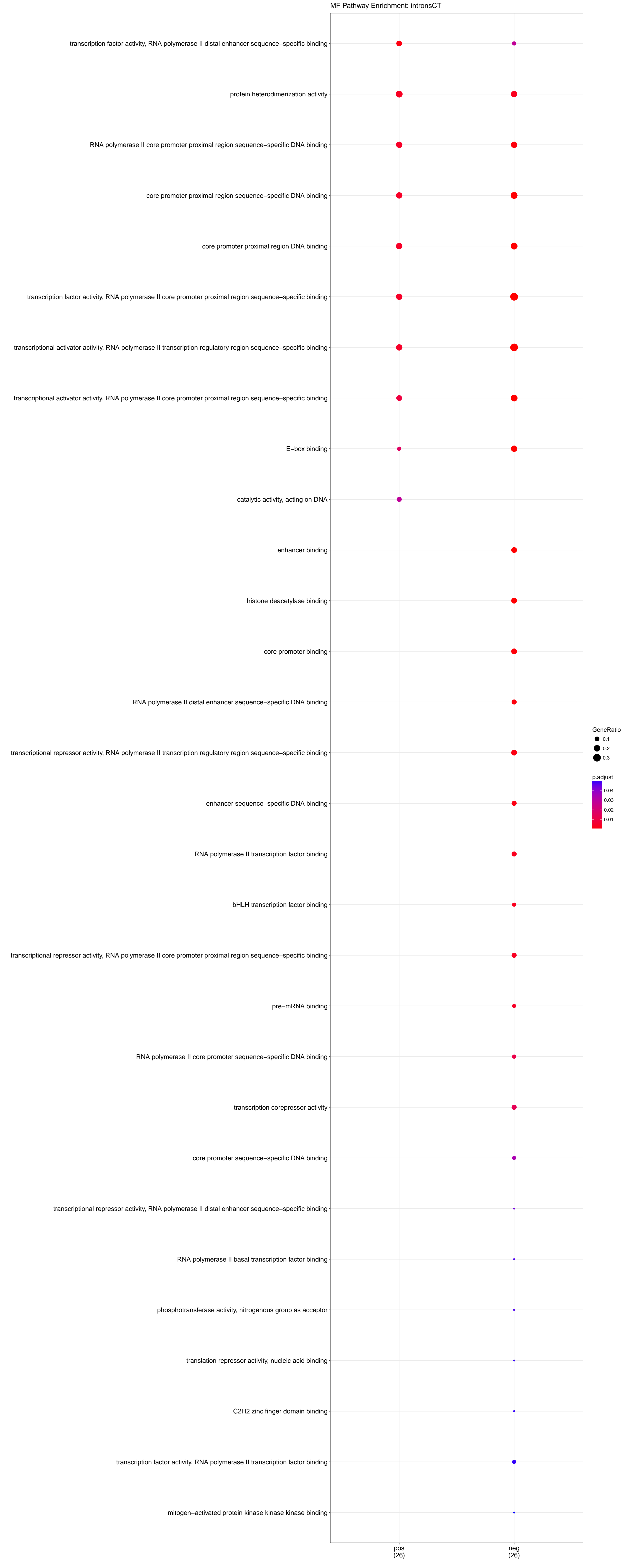
Apelin signaling pathway

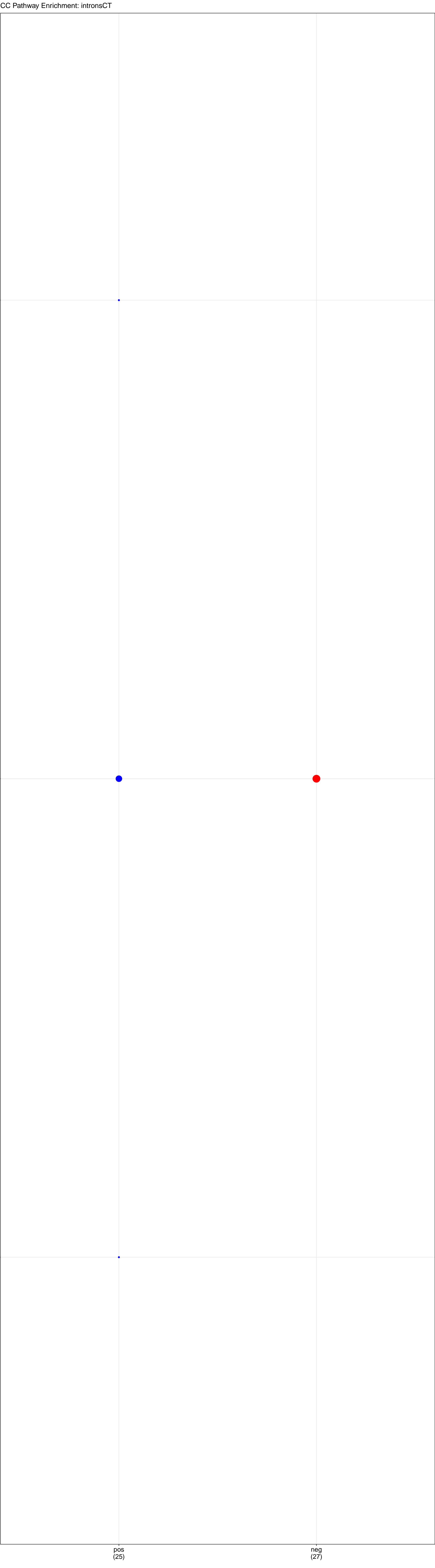
cGMP-PKG signaling pathway

pos
(10)

neg
(6)







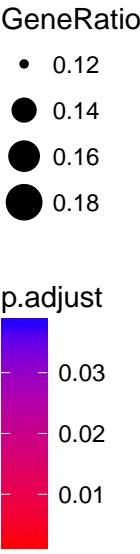
RNA polymerase II transcription factor complex

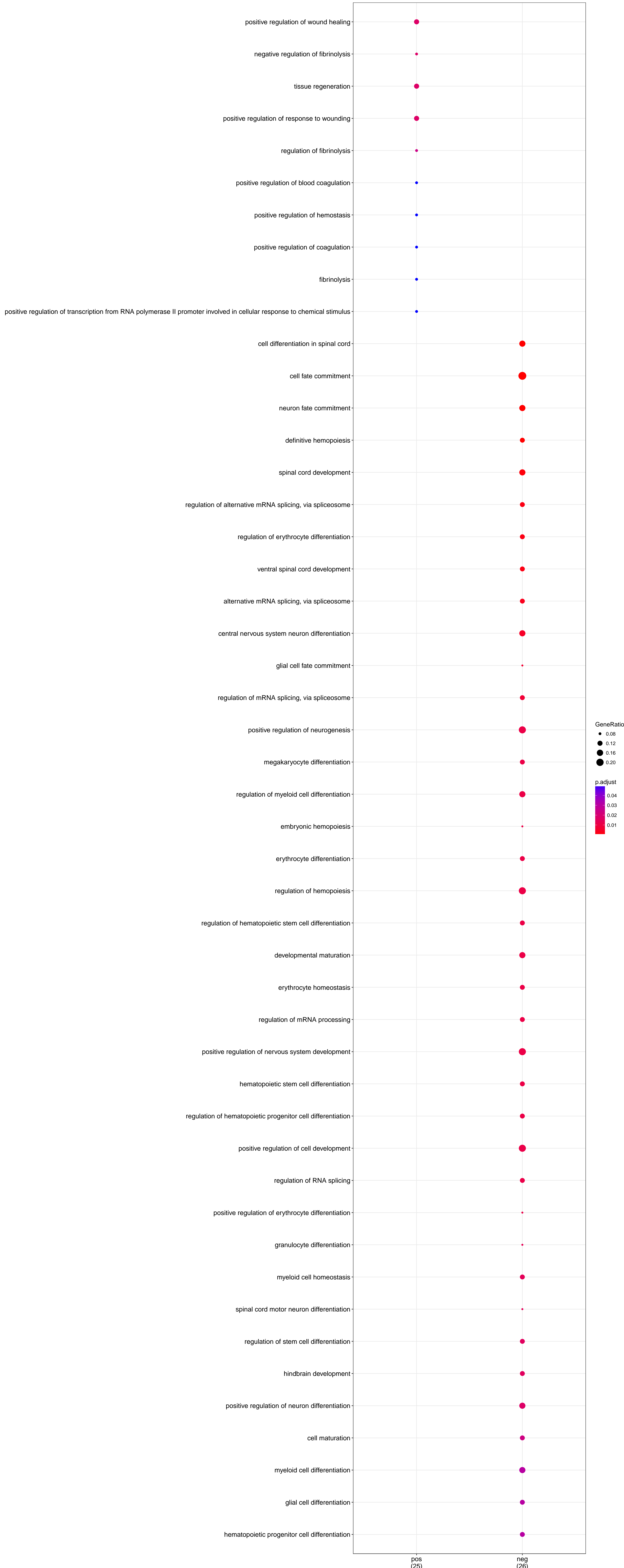
transcription factor complex

nuclear transcription factor complex

pos
(25)

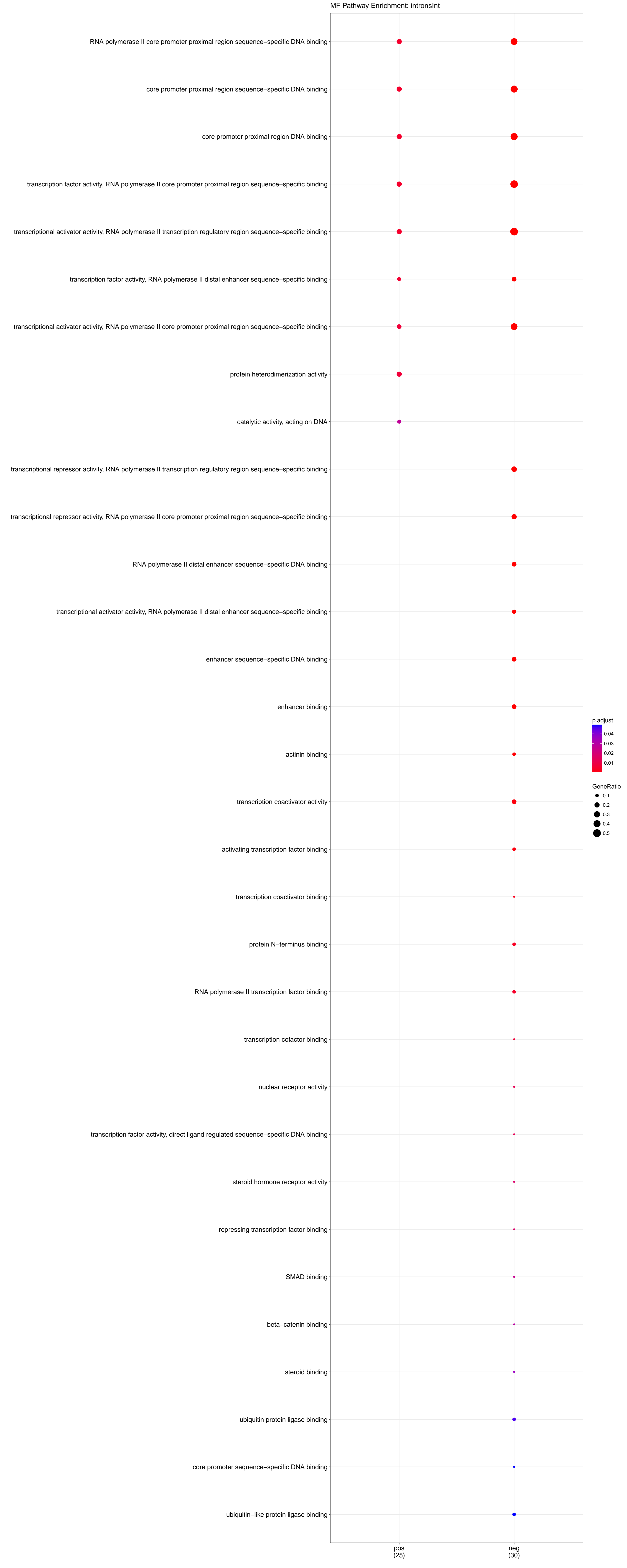
neg
(27)

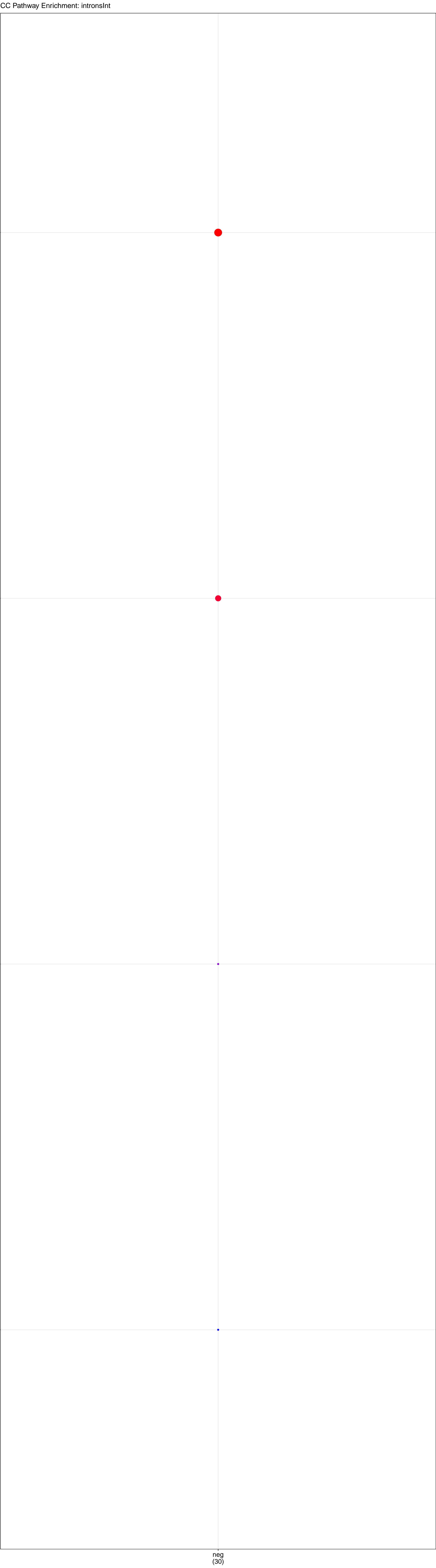




KEGG Pathway Enrichment: intronsInt







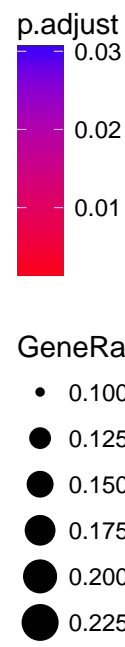
nuclear chromatin

transcription factor complex

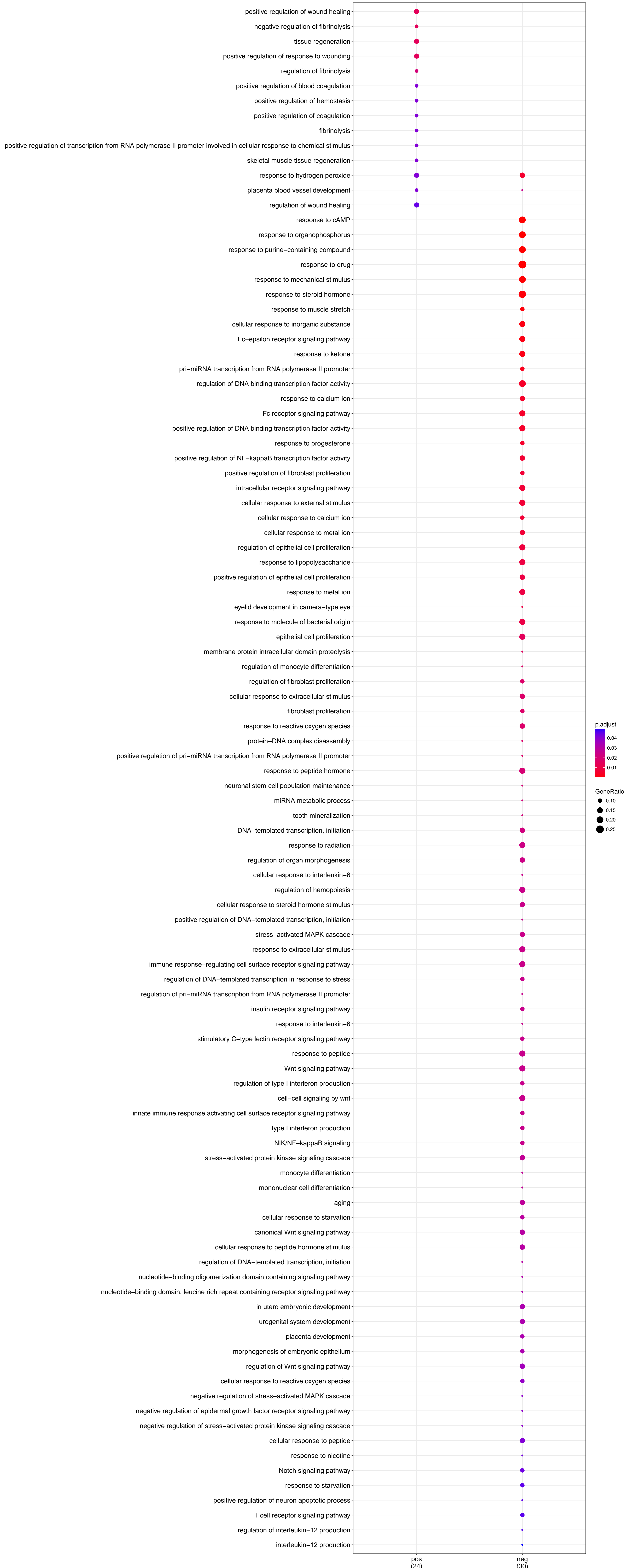
RNA polymerase II transcription factor complex

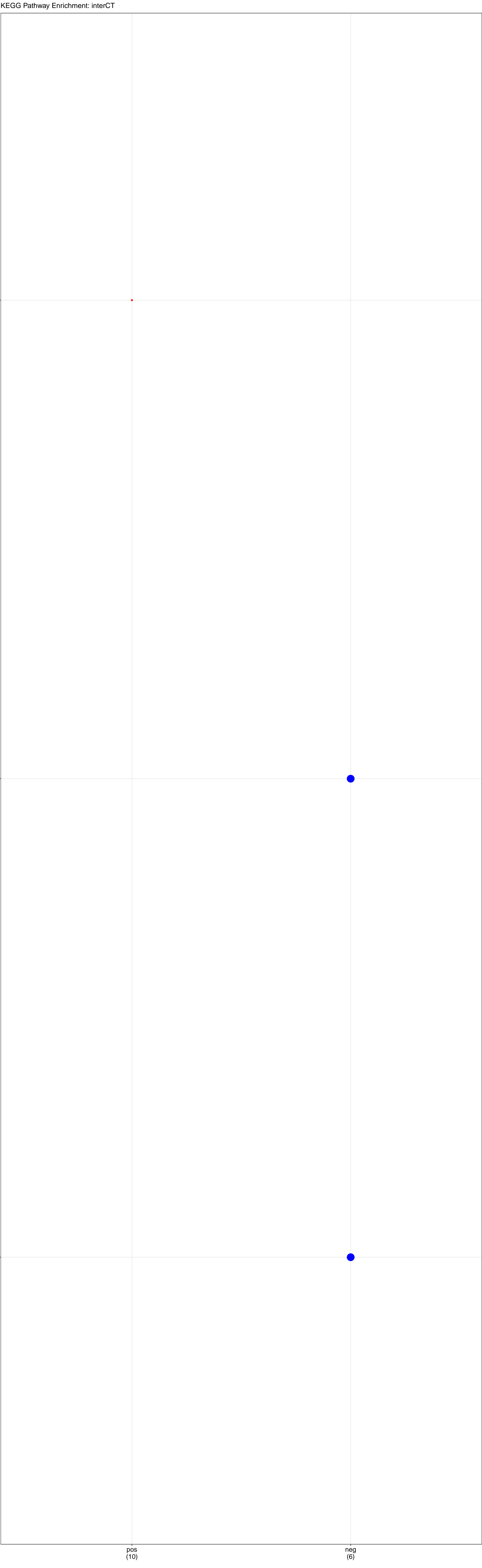
nuclear transcription factor complex

neg
(30)

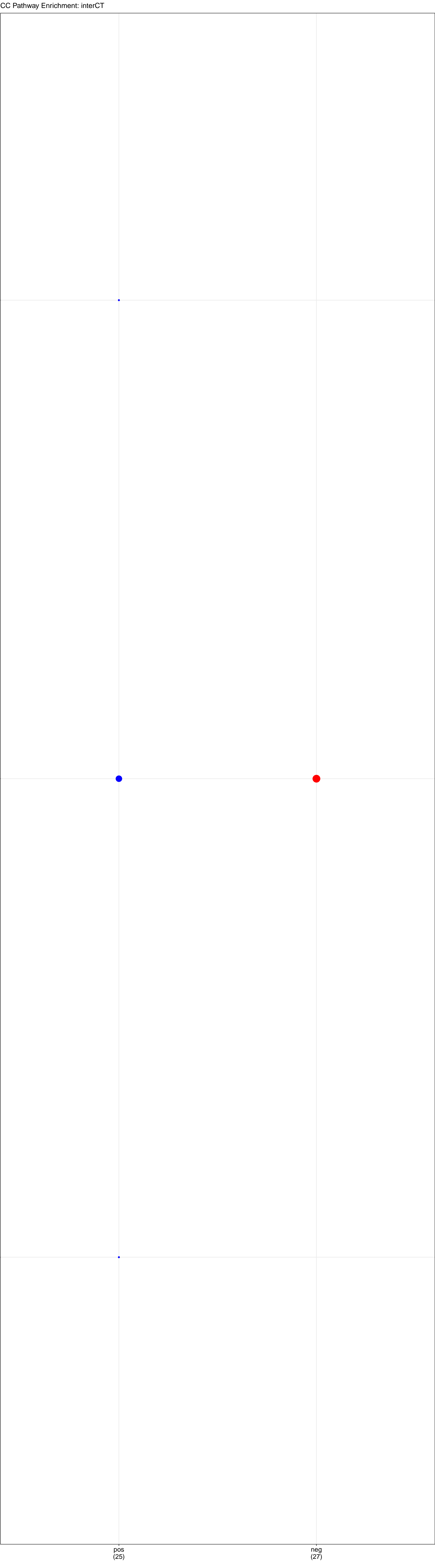


Biological Process GO Enrichment: intronsInt









RNA polymerase II transcription factor complex

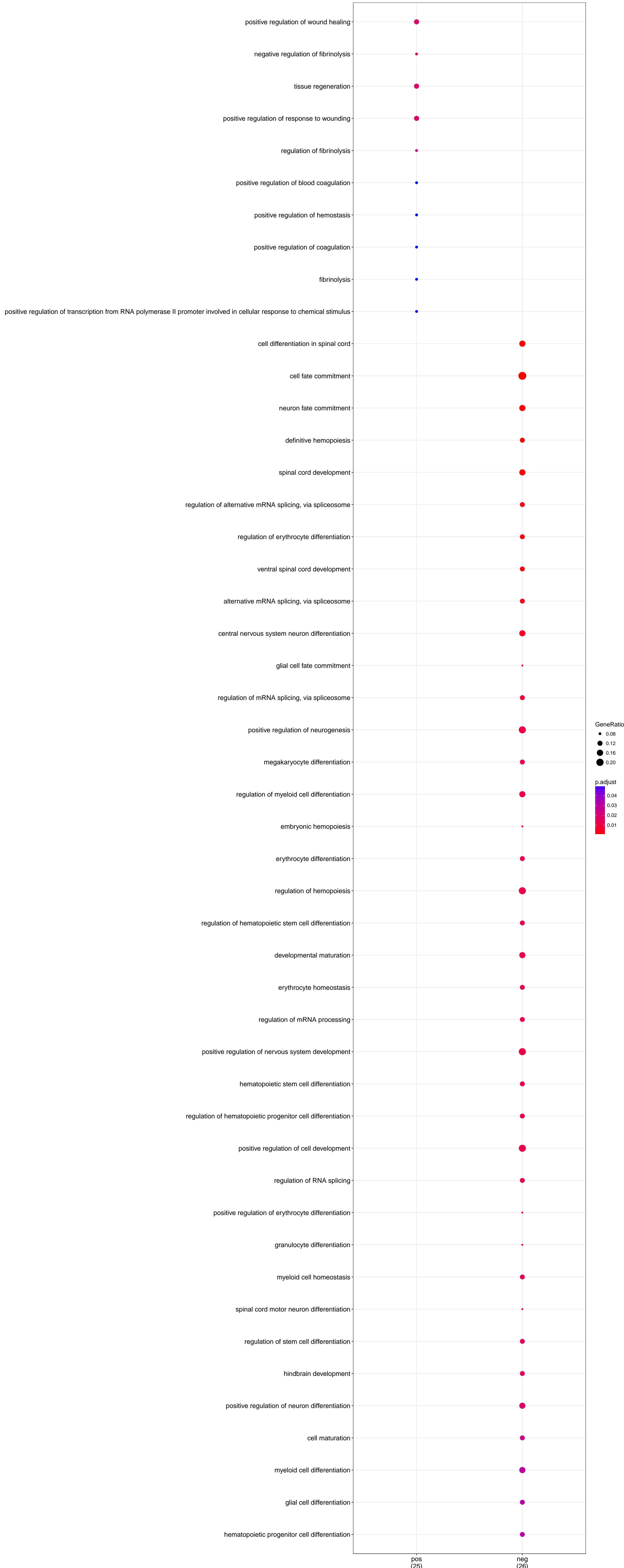
transcription factor complex

nuclear transcription factor complex

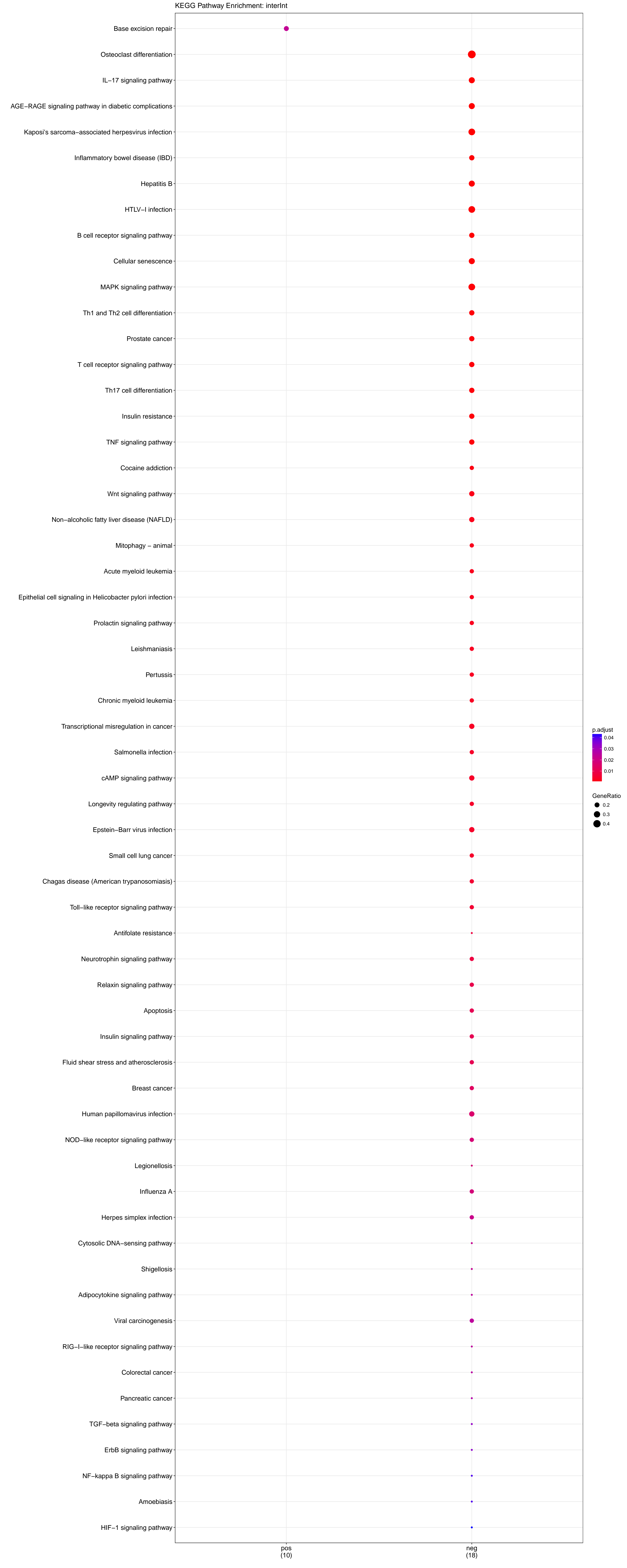
pos
(25)

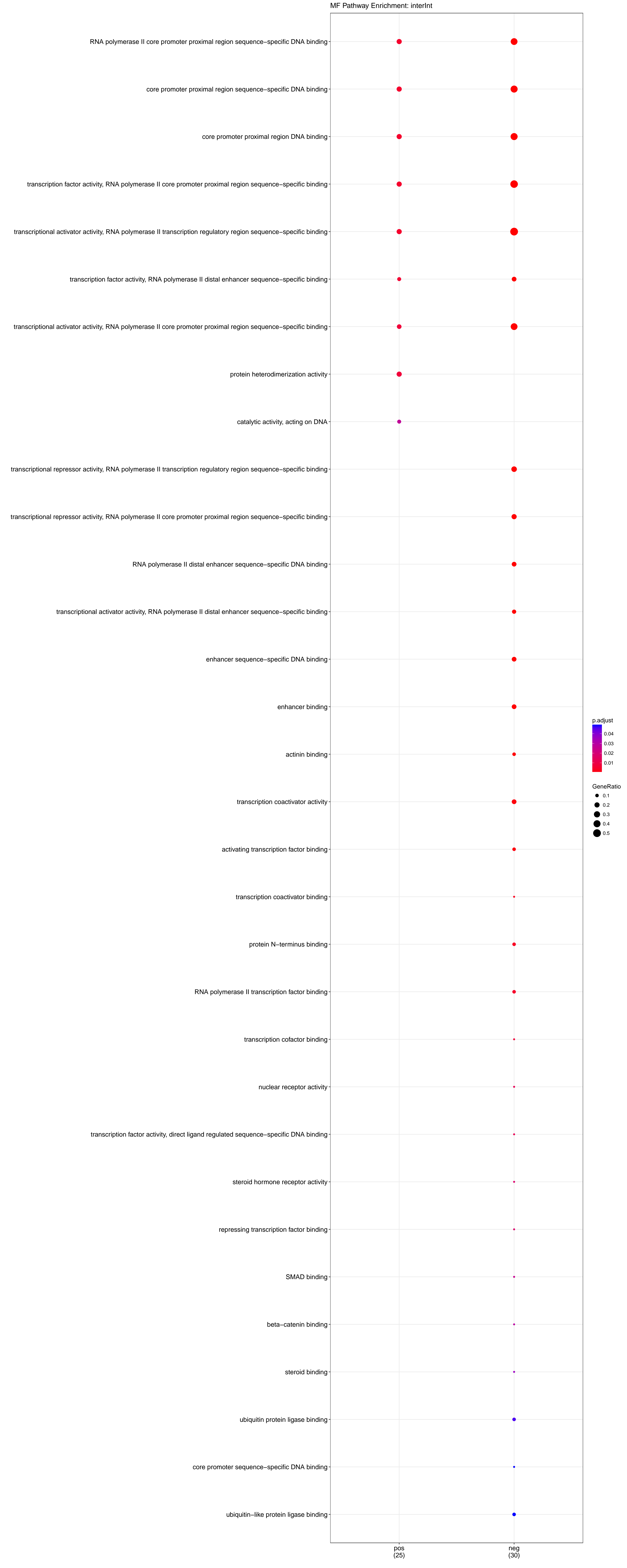
neg
(27)

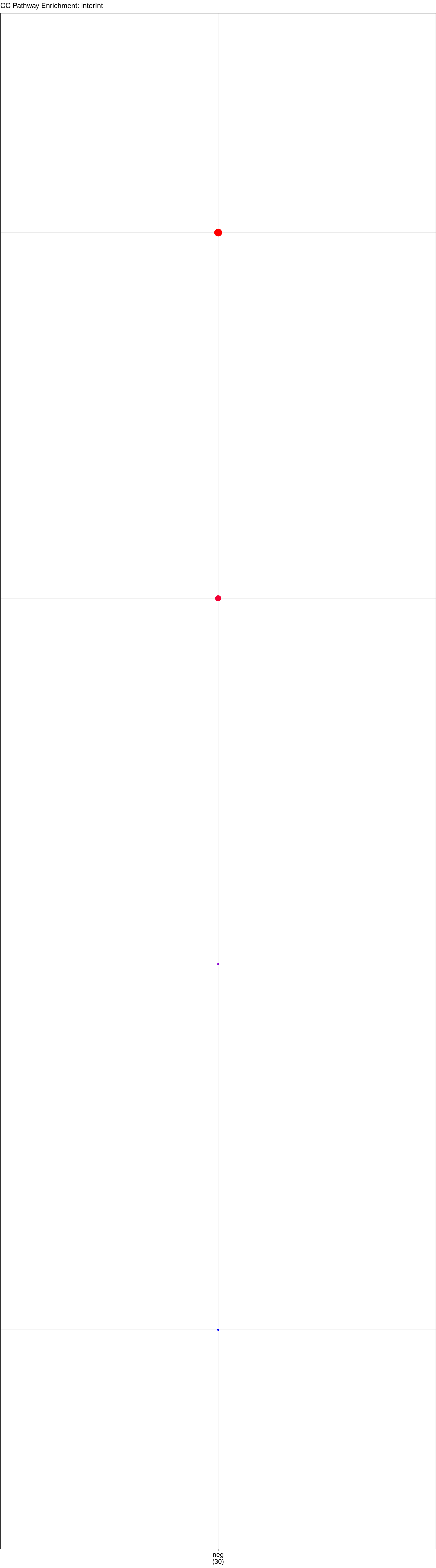




KEGG Pathway Enrichment: interInt







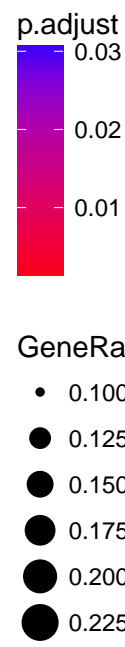
nuclear chromatin

transcription factor complex

RNA polymerase II transcription factor complex

nuclear transcription factor complex

neg
(30)



Biological Process GO Enrichment: interlnt

