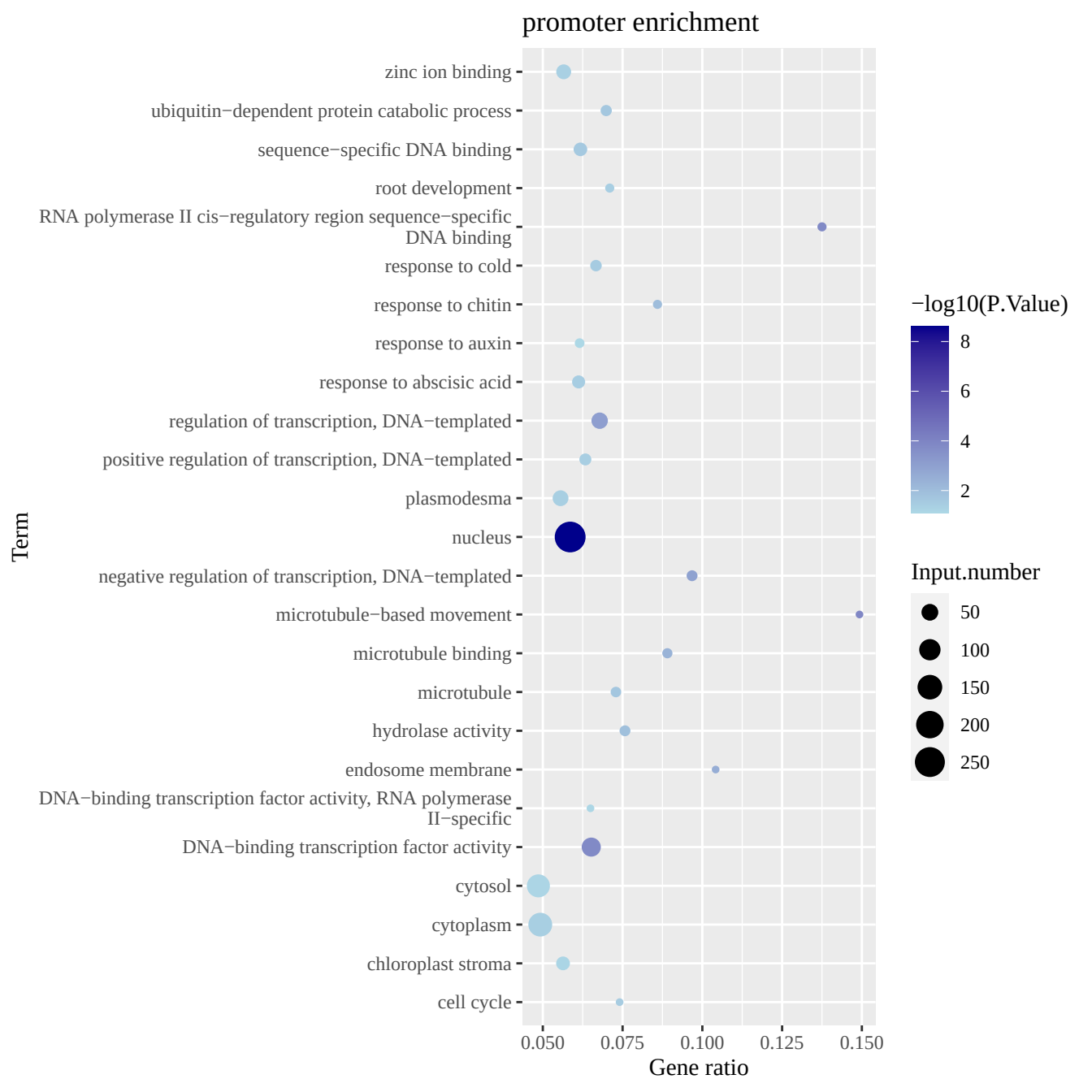
promoter enrichment
zinc ion binding
ubiquitin−dependent protein catabolic process
sequence−specific DNA binding
root development
RNA polymerase II cis−regulatory region sequence−specific DNA binding
response to cold
response to chitin
response to auxin
response to abscisic acid
regulation of transcription, DNA−templated
positive regulation of transcription, DNA−templated
plasmodesma
nucleus
negative regulation of transcription, DNA−templated
microtubule−based movement
microtubule binding
microtubule
hydrolase activity
endosome membrane
DNA−binding transcription factor activity, RNA polymerase II−specific
DNA−binding transcription factor activity
cytosol
cytoplasm
chloroplast stroma
cell cycle
Term
0.050
0.075
0.100
0.125
0.150
Gene ratio
−log10(P.Value)
8
6
4
2
Input.number
50
100
150
200
250