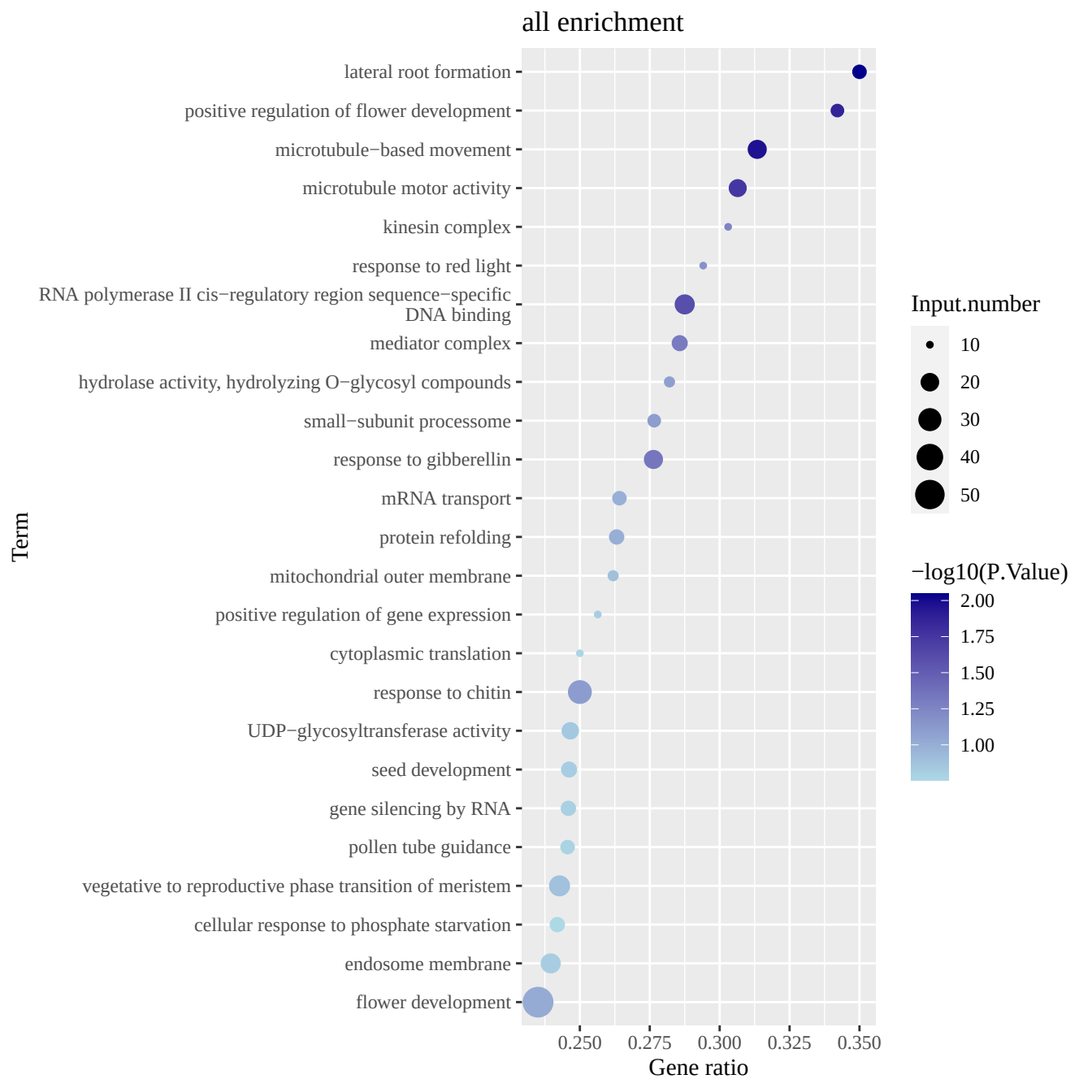
all enrichment
lateral root formation
positive regulation of flower development
microtubule−based movement
microtubule motor activity
kinesin complex
response to red light
RNA polymerase II cis−regulatory region sequence−specific DNA binding
mediator complex
hydrolase activity, hydrolyzing O−glycosyl compounds
small−subunit processome
response to gibberellin
mRNA transport
protein refolding
mitochondrial outer membrane
positive regulation of gene expression
cytoplasmic translation
response to chitin
UDP−glycosyltransferase activity
seed development
gene silencing by RNA
pollen tube guidance
vegetative to reproductive phase transition of meristem
cellular response to phosphate starvation
endosome membrane
flower development
Term
0.250
0.275
0.300
0.325
0.350
Gene ratio
Input.number
10
20
30
40
50
−log10(P.Value)
2.00
1.75
1.50
1.25
1.00