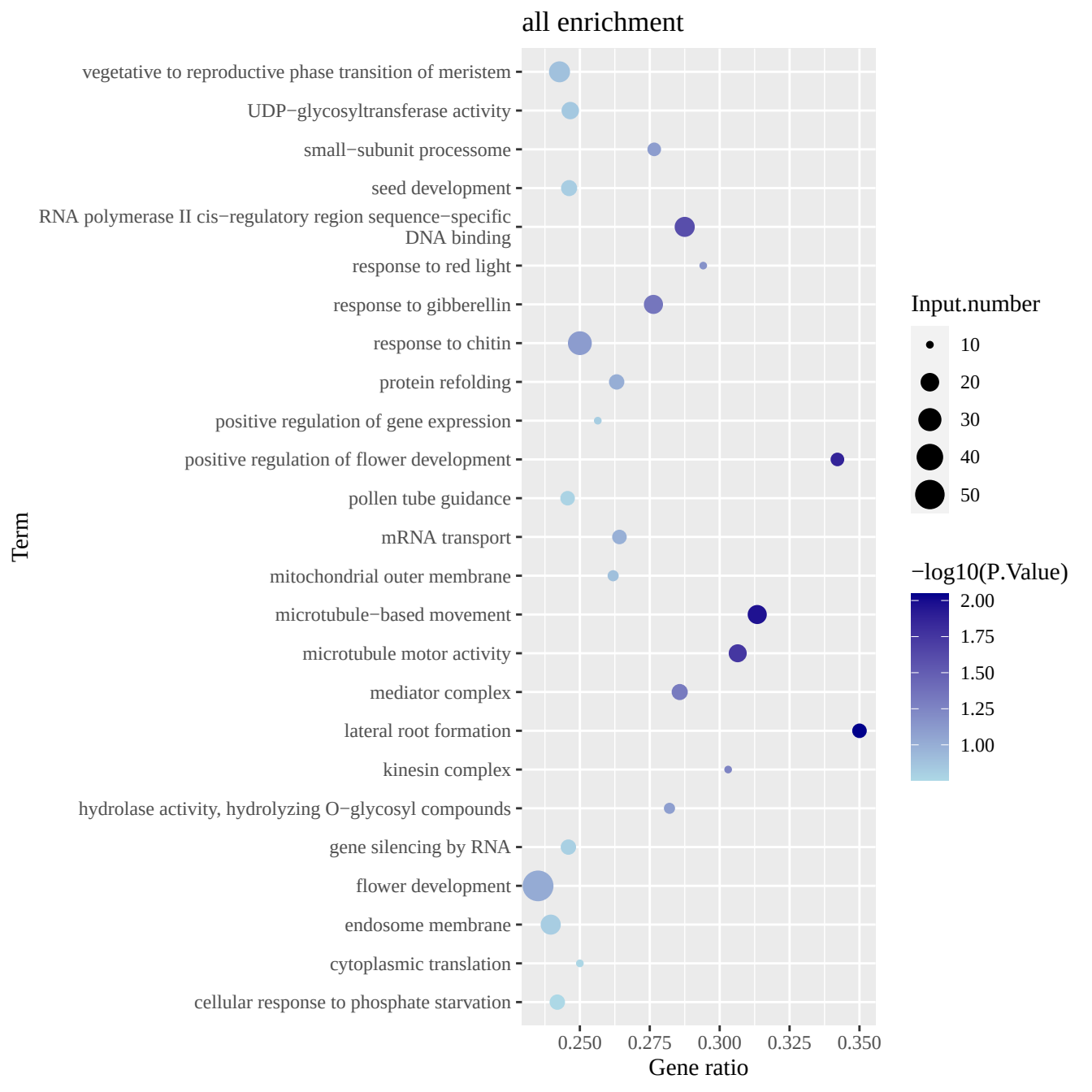
all enrichment
vegetative to reproductive phase transition of meristem
UDP−glycosyltransferase activity
small−subunit processome
seed development
RNA polymerase II cis−regulatory region sequence−specific DNA binding
response to red light
response to gibberellin
response to chitin
protein refolding
positive regulation of gene expression
positive regulation of flower development
pollen tube guidance
mRNA transport
mitochondrial outer membrane
microtubule−based movement
microtubule motor activity
mediator complex
lateral root formation
kinesin complex
hydrolase activity, hydrolyzing O−glycosyl compounds
gene silencing by RNA
flower development
endosome membrane
cytoplasmic translation
cellular response to phosphate starvation
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