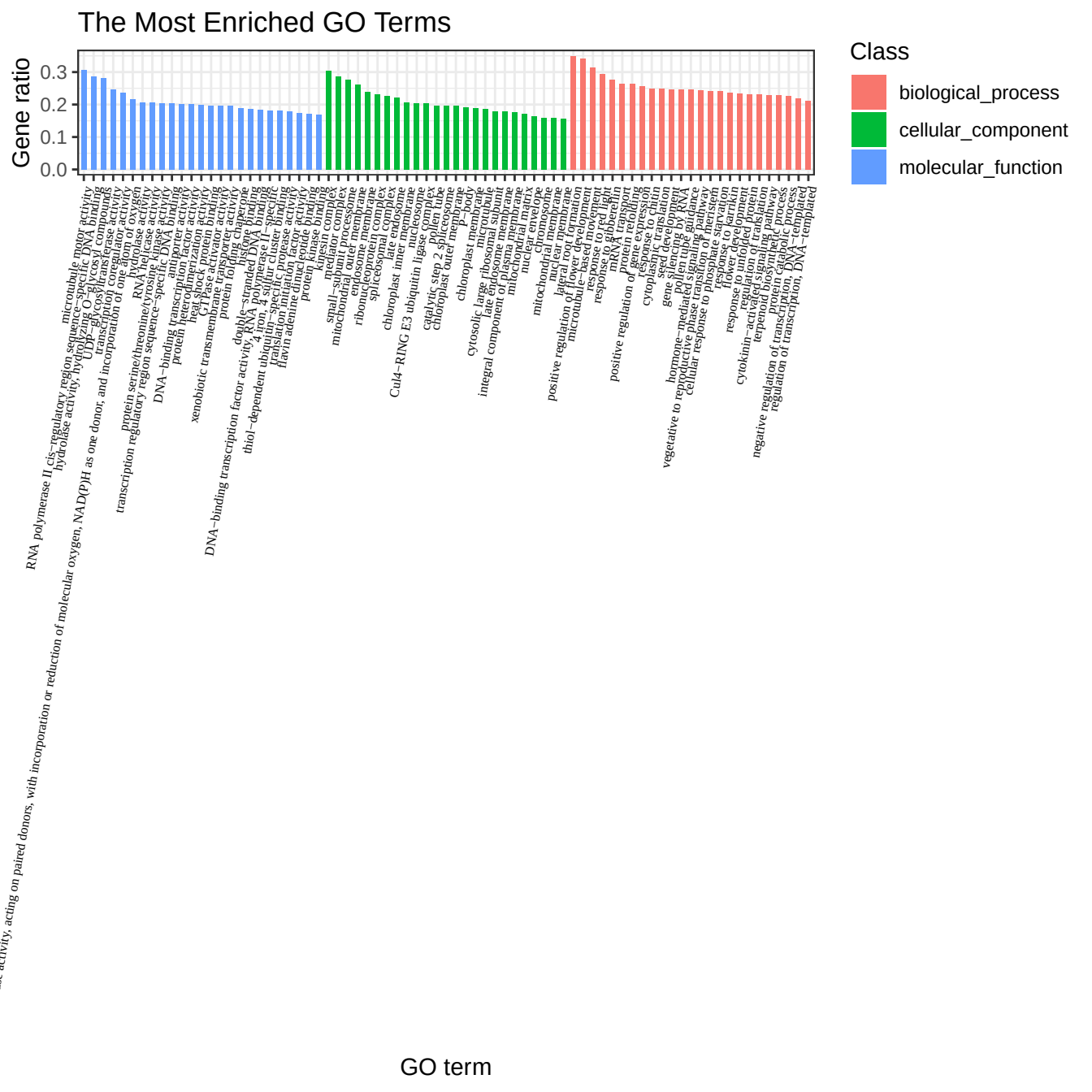
The Most Enriched GO Terms
Gene ratio
0.0
0.1
0.2
0.3
GO term
Class
biological_process
cellular_component
molecular_function
RNA polymerase II cis−regulatory region sequence−specific DNA binding
hydrolase activity, hydrolyzing O−glycosyl compounds
UDP−glycosyltransferase activity
glycosyltransferase activity
transcription coregulator activity
oxidoreductase activity, acting on paired donors, with incorporation or reduction of molecular oxygen, NAD(P)H as one donor, and incorporation of one atom of oxygen
hydrolase activity
RNA helicase activity
protein serine/threonine/tyrosine kinase activity
transcription regulatory region sequence−specific DNA binding
DNA−binding transcription activator activity
protein heterodimerization activity
antiporter activity
heterodimerization activity
protein transporter activity
heat shock protein binding
xenobiotic transmembrane transporter activity
protein folding chaperone
histone binding
DNA−binding transcription factor activity
double−stranded DNA binding
thiol−dependent ubiquitin−specific protease activity
4 iron, 4 sulfur cluster binding
translation factor activity, RNA binding
flavin adenine dinucleotide binding
protein kinase binding
kinesin binding
small−subunit processome
mediator complex
mitochondrial outer membrane translocase complex
endosome membrane
ribonucleoprotein complex
spliceosomal complex
late endosome
chloroplast inner membrane
nucleosome
Cul4−RING E3 ubiquitin ligase complex
catalytic step 2 spliceosome
pollen tube
chloroplast outer membrane
plastid
chloroplast membrane
microtubule
cytosolic large ribosomal subunit
late endosomal membrane
integral component of plasma membrane
mitochondrial matrix
nuclear envelope
chromosome
mitochondrial membrane
nuclear membrane
positive regulation of flower development
integral component of membrane
microtubule−based movement
response to pollen
response to gibberellin
mRNA transport
positive regulation of gene expression
protein refolding
response to chitin
cytoplasmic translation
seed development
gene silencing by RNA
hormone−mediated signaling pathway
reproductive shoot system development
pollen tube guidance
vegetative to reproductive phase transition of meristem
cellular response to phosphate starvation
response to karrikin
flower development
response to unfolded protein
cytokinin−activated signaling pathway
regulation of translation
terpenoid biosynthetic process
negative regulation of transcription, DNA−templated
protein catabolic process
regulation of transcription, DNA−templated