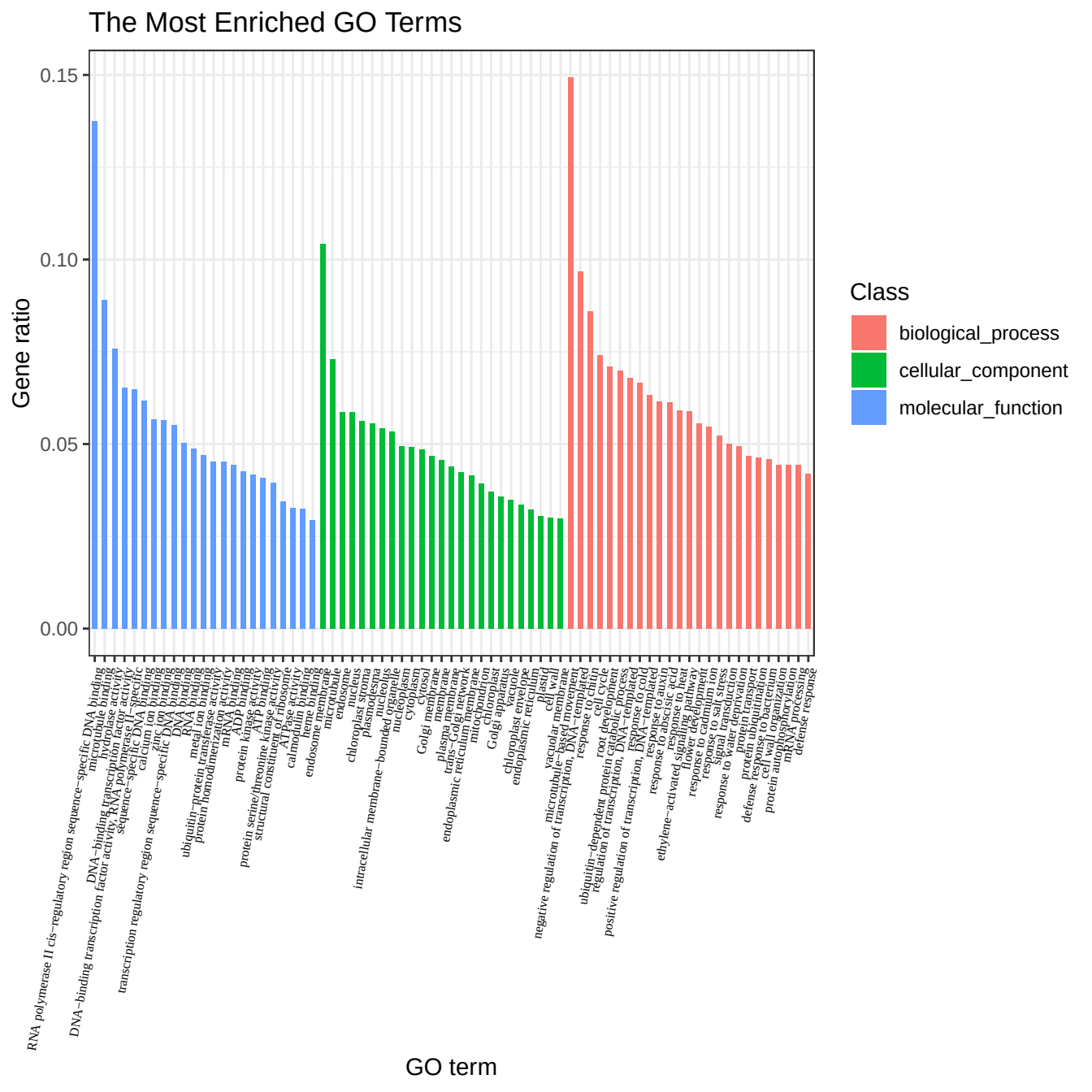
The Most Enriched GO Terms
Gene ratio
0.15
0.10
0.05
0.00
GO term
Class
biological_process
cellular_component
molecular_function
RNA polymerase II cis−regulatory region sequence−specific DNA binding
DNA−binding transcription factor activity, RNA polymerase II−specific
transcription regulatory region sequence−specific DNA binding
ubiquitin−protein transferase activity
protein serine/threonine kinase activity
structural constituent of ribosome
endosome membrane
intracellular membrane−bounded organelle
negative regulation of transcription, DNA−templated
ubiquitin−dependent protein catabolic process
positive regulation of transcription, DNA−templated
ethylene−activated signaling pathway
defense response
protein autophosphorylation
mRNA processing