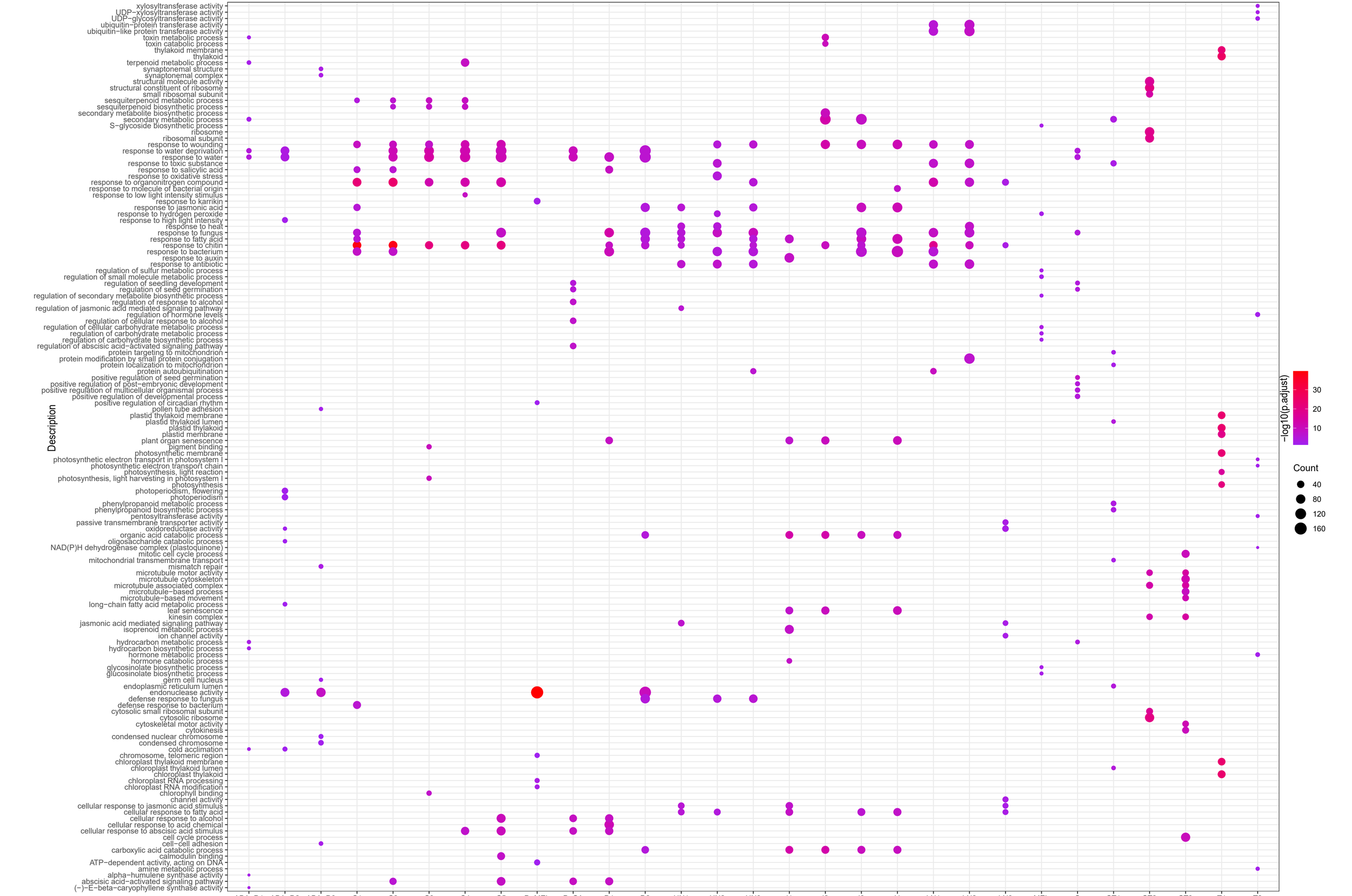
xylosyltransferase activity
UDP−xylosyltransferase activity
UDP−glycosyltransferase activity
ubiquitin−protein transferase activity
ubiquitin−like protein transferase activity
toxin metabolic process
toxin catabolic process
thylakoid membrane
thylakoid
terpenoid metabolic process
synaptonemal structure
synaptonemal complex
structural molecule activity
structural constituent of ribosome
small ribosomal subunit
sesquiterpenoid metabolic process
sesquiterpenoid biosynthetic process
secondary metabolite biosynthetic process
secondary metabolic process
S−glycoside biosynthetic process
ribosome
ribosomal subunit
response to wounding
response to water deprivation
response to water
response to toxic substance
response to salicylic acid
response to oxidative stress
response to organonitrogen compound
response to molecule of bacterial origin
response to low light intensity stimulus
response to karrikin
response to jasmonic acid
response to hydrogen peroxide
response to high light intensity
response to heat
response to fungus
response to fatty acid
response to chitin
response to bacterium
response to auxin
response to antibiotic
regulation of sulfur metabolic process
regulation of small molecule metabolic process
regulation of seedling development
regulation of seed germination
regulation of secondary metabolite biosynthetic process
regulation of response to alcohol
regulation of jasmonic acid mediated signaling pathway
regulation of hormone levels
regulation of cellular response to alcohol
regulation of cellular carbohydrate metabolic process
regulation of carbohydrate metabolic process
regulation of carbohydrate biosynthetic process
regulation of abscisic acid−activated signaling pathway
protein targeting to mitochondrion
protein modification by small protein conjugation
protein localization to mitochondrion
protein autoubiquitination
positive regulation of seed germination
positive regulation of post−embryonic development
positive regulation of multicellular organismal process
positive regulation of developmental process
positive regulation of circadian rhythm
pollen tube adhesion
plastid thylakoid membrane
plastid thylakoid lumen
plastid thylakoid
plastid membrane
plant organ senescence
pigment binding
photosynthetic membrane
photosynthetic electron transport in photosystem I
photosynthetic electron transport chain
photosynthesis, light reaction
photosynthesis, light harvesting in photosystem I
photosynthesis
photoperiodism, flowering
photoperiodism
phenylpropanoid metabolic process
phenylpropanoid biosynthetic process
pentosyltransferase activity
passive transmembrane transporter activity
oxidoreductase activity
organic acid catabolic process
oligosaccharide catabolic process
NAD(P)H dehydrogenase complex (plastoquinone)
mitotic cell cycle process
mitochondrial transmembrane transport
mismatch repair
microtubule motor activity
microtubule cytoskeleton
microtubule associated complex
microtubule−based process
microtubule−based movement
long−chain fatty acid metabolic process
leaf senescence
kinesin complex
jasmonic acid mediated signaling pathway
isoprenoid metabolic process
ion channel activity
hydrocarbon metabolic process
hydrocarbon biosynthetic process
hormone metabolic process
hormone catabolic process
glycosinolate biosynthetic process
glucosinolate biosynthetic process
germ cell nucleus
endoplasmic reticulum lumen
endonuclease activity
defense response to fungus
defense response to bacterium
cytosolic small ribosomal subunit
cytosolic ribosome
cytoskeletal motor activity
cytokinesis
condensed nuclear chromosome
condensed chromosome
cold acclimation
chromosome, telomeric region
chloroplast thylakoid membrane
chloroplast thylakoid lumen
chloroplast thylakoid
chloroplast RNA processing
chloroplast RNA modification
chlorophyll binding
channel activity
cellular response to jasmonic acid stimulus
cellular response to fatty acid
cellular response to alcohol
cellular response to acid chemical
cellular response to abscisic acid stimulus
cell cycle process
cell−cell adhesion
carboxylic acid catabolic process
calmodulin binding
ATP−dependent activity, acting on DNA
amine metabolic process
alpha−humulene synthase activity
abscisic acid−activated signaling pathway
(−)−E−beta−caryophyllene synthase activity
Description
ABA_D1
ABA_D2
ABA_D3
C1
C2
C3
C4
C5
D_MEL
D_SA
D1
D2
HN1
HN2
HN3
I1
I2
I3
I4
LN1
LN2
LN3
MEL
S
ST1
ST2
ST3
T1
T2
group
−log10(p.adjust)
30
20
10
Count
40
80
120
160