BP Ontology Enrichment

Gene number
- 100
- 200
- 300

−log10(weight)
- 30
- 20
- 10

Y-axis categories (top to bottom):
- pollen tube reception
- abscisic acid−activated signaling pathwa...
- regulation of defense response by callos...
- transport of virus in host, tissue to ti...
- lipopolysaccharide−mediated signaling pa...
- response to salicylic acid
- negative regulation of defense response
- detection of lipopolysaccharide
- cellular response to heat
- endoplasmic reticulum unfolded protein r...
- immune system process
- regulation of double fertilization formi...
- response to lipid
- cell wall macromolecule catabolic proces...
- detoxification
- jasmonic acid metabolic process
- salicylic acid metabolic process
- response to auxin
- negative regulation of catalytic activit...
- response to bacterium
- defense response to fungus
- cytidine deamination
- lipid localization
- cell surface receptor signaling pathway
- xyloglucan metabolic process
- detection of calcium ion
- recognition of pollen
- response to organic cyclic compound
- toxin metabolic process
- response to toxic substance
- defense response
- response to biotic stimulus
- regulation of pH
- glutathione metabolic process
- sequestering of actin monomers
- response to amino acid
- negative regulation of molecular functio...
- protein deubiquitination
- sulfation
- N−terminal protein myristoylation
- NADH oxidation
- auxin metabolic process
- cell wall modification
- intracellular signal transduction
- calcium−mediated signaling
- dephosphorylation
- pectin catabolic process
- protein phosphorylation

X-axis categories: size_2, size_3_5, size_6_10, size_gt_10