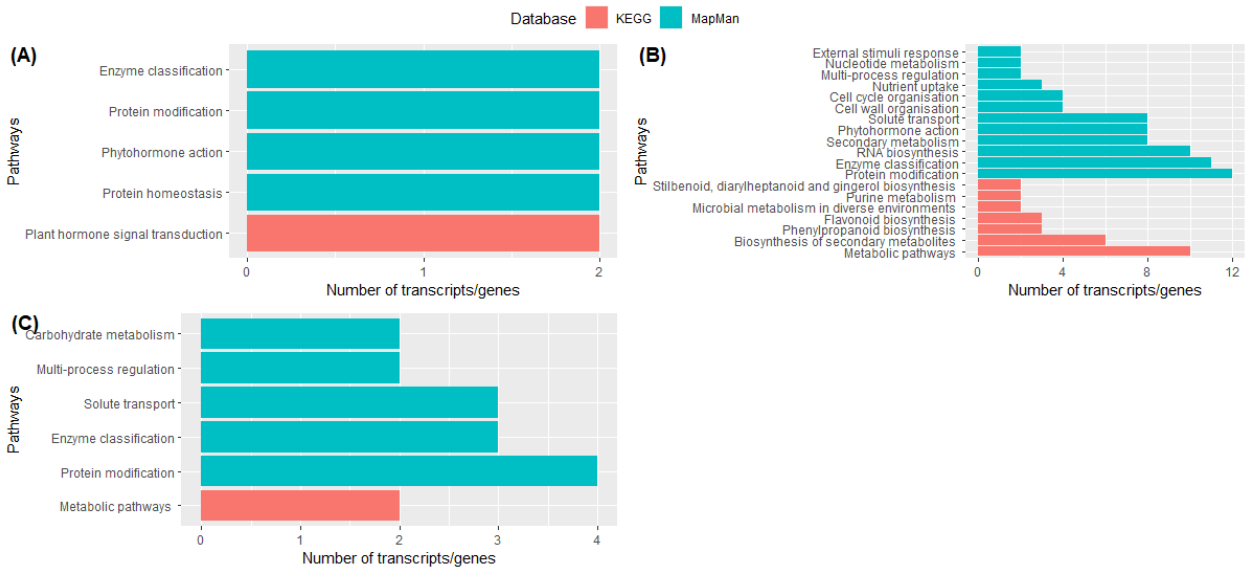
Database
KEGG
MapMan
(A)
Enzyme classification
Protein modification
Phytohormone action
Protein homeostasis
Plant hormone signal transduction
Pathways
0
1
2
Number of transcripts/genes
(B)
External stimuli response
Nucleotide metabolism
Multi-process regulation
Nutrient uptake
Cell cycle organisation
Cell wall organisation
Solute transport
Phytohormone action
Secondary metabolism
RNA biosynthesis
Enzyme classification
Protein modification
Stilbenoid, diarylheptanoid and gingerol biosynthesis
Purine metabolism
Microbial metabolism in diverse environments
Flavonoid biosynthesis
Phenylpropanoid biosynthesis
Biosynthesis of secondary metabolites
Metabolic pathways
Pathways
0
4
8
12
Number of transcripts/genes
(C)
Carbohydrate metabolism
Multi-process regulation
Solute transport
Enzyme classification
Protein modification
Metabolic pathways
Pathways
0
1
2
3
4
Number of transcripts/genes