

Ref: 262904

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|---|---|
| - Metabolic pathways (map01100) | - Biosynthesis of secondary metabolites (map01110) |
| - ABC transporters (map02010) | - Biosynthesis of amino acids (map01230) |
| - Biosynthesis of cofactors (map01240) | - Aminoacyl-tRNA biosynthesis (map00970) |
| - 2-Oxocarboxylic acid metabolism (map01210) | - Galactose metabolism (map00052) |
| - Biosynthesis of various plant secondary metabolites (map00999) | - Tryptophan metabolism (map00380) |
| - Phenylalanine metabolism (map00360) | - D-Amino acid metabolism (map00470) |
| - Arginine and proline metabolism (map00330) | - Glucosinolate biosynthesis (map00966) |
| - Tyrosine metabolism (map00350) | - Glyoxylate and dicarboxylate metabolism (map00630) |
| - Carbon metabolism (map01200) | - Diterpenoid biosynthesis (map00904) |
| - Phenylalanine, tyrosine and tryptophan biosynthesis (map00400) | - Arachidonic acid metabolism (map00590) |
| - Alanine, aspartate and glutamate metabolism (map00250) | - Arginine biosynthesis (map00220) |
| - Lysine degradation (map00310) | - Glycine, serine and threonine metabolism (map00260) |
| - Flavonoid biosynthesis (map00941) | - Indole alkaloid biosynthesis (map00901) |
| - Citrate cycle (TCA cycle) (map00020) | - Purine metabolism (map00230) |
| - Cyanoamino acid metabolism (map00460) | - Pyrimidine metabolism (map00240) |
| - Nucleotide metabolism (map01232) | - alpha-Linolenic acid metabolism (map00592) |
| - Biosynthesis of various alkaloids (map00996) | |
| - Tropane, piperidine and pyridine alkaloid biosynthesis (map00960) | |
| - Glutathione metabolism (map00480) | |
| - Fructose and mannose metabolism (map00051) | |
| - Linoleic acid metabolism (map00591) | |

subject to the condition that the original source is acknowledged by citing at least one KEGG paper.

Permission granted:

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