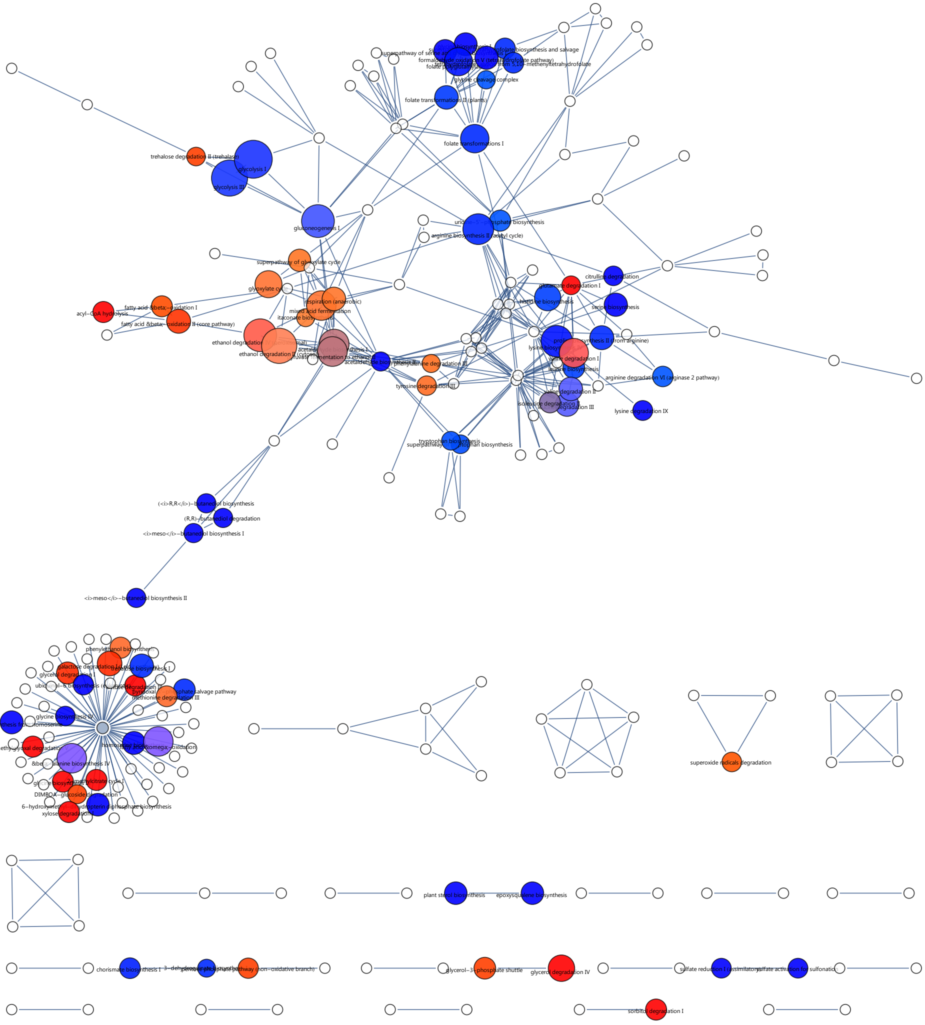
trehalose degradation II (trehalase)
glycolysis I
glycolysis III
gluconeogenesis I
superpathway of glyoxylate cycle
glyoxylate cycle
fatty acid β-oxidation I
acyl-CoA hydrolysis
fatty acid β-oxidation II (core pathway)
respiration (anaerobic)
mixed acid fermentation
ethanol degradation II (cytosol)
phenylalanine degradation III
tyrosine degradation III
folate transformations II (plants)
folate transformations I
glycine cleavage complex
formaldehyde oxidation V (tetrahydrofolate pathway)
arginine biosynthesis II (acetyl cycle)
citrulline degradation
glutamate degradation I
histidine biosynthesis
serine biosynthesis
arginine degradation VI (arginase 2 pathway)
lysine degradation IX
tryptophan biosynthesis
(<i>R,R</i>)-butanediol biosynthesis
(R,R)-butanediol degradation
<i>meso</i>-butanediol biosynthesis I
<i>meso</i>-butanediol biosynthesis II
glycine biosynthesis IV
methionine degradation III
β-alanine biosynthesis IV
xylose degradation
superoxide radicals degradation
plant sterol biosynthesis
epoxysqualene biosynthesis
chorismate biosynthesis I
glycerol-3-phosphate shuttle
glycerol degradation IV
sorbitol degradation I