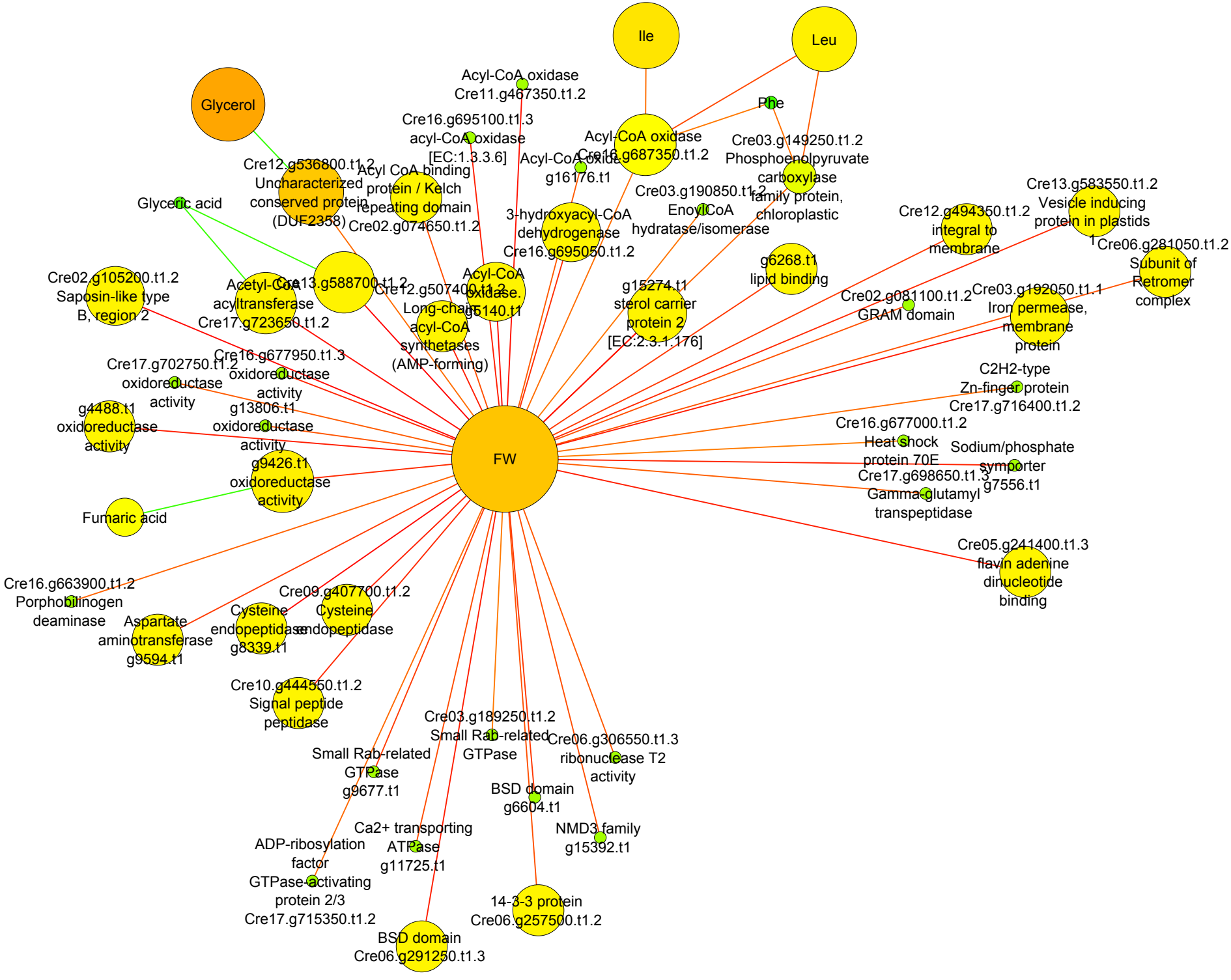
Ile
Leu
Glycerol
Acyl-CoA oxidase
Cre11.g467350.t1.2
Phe
Cre16.g695100.t1.3
acyl-CoA oxidase
[EC:1.3.3.6]
Acyl-CoA oxidase
Cre16.g687350.t1.2
Cre03.g149250.t1.2
Phosphoenolpyruvate
carboxylase
family protein,
chloroplastic
Acyl-CoA oxidase
g16176.t1
Cre12.g536800.t1.2
Uncharacterized
conserved protein
(DUF2358)
Acyl CoA binding
protein / Kelch
repeating domain
Cre02.g074650.t1.2
Cre03.g190850.t1.2
Enoyl-CoA
hydratase/isomerase
Glyceric acid
3-hydroxyacyl-CoA
dehydrogenase
Cre16.g695050.t1.2
Cre13.g583550.t1.2
Vesicle inducing
protein in plastids
1
Cre12.g494350.t1.2
integral to
membrane
Cre06.g281050.t1.2
Subunit of
Retromer
complex
g6268.t1
lipid binding
Cre02.g105200.t1.2
Saposin-like type
B, region 2
Acetyl-CoA
acyltransferase
Cre17.g723650.t1.2
Cre13.g588700.t1.2
Cre12.g507400.t1.2
Long-chain
acyl-CoA
synthetases
(AMP-forming)
Acyl-CoA
oxidase.
g5140.t1
g15274.t1
sterol carrier
protein 2
[EC:2.3.1.176]
Cre02.g081100.t1.2
GRAM domain
Cre03.g192050.t1.1
Iron permease,
membrane
protein
Cre17.g702750.t1.2
oxidoreductase
activity
Cre16.g677950.t1.3
oxidoreductase
activity
C2H2-type
Zn-finger protein
Cre17.g716400.t1.2
g4488.t1
oxidoreductase
activity
g13806.t1
oxidoreductase
activity
Cre16.g677000.t1.2
Heat shock
protein 70E
Sodium/phosphate
symporter
g7556.t1
FW
g9426.t1
oxidoreductase
activity
Cre17.g698650.t1.3
Gamma-glutamyl
transpeptidase
Fumaric acid
Cre05.g241400.t1.3
flavin adenine
dinucleotide
binding
Cre16.g663900.t1.2
Porphobilinogen
deaminase
Aspartate
aminotransferase
g9594.t1
Cysteine
endopeptidase
g8339.t1
Cre09.g407700.t1.2
Cysteine
endopeptidase
Cre10.g444550.t1.2
Signal peptide
peptidase
Cre03.g189250.t1.2
Small Rab-related
GTPase
Cre06.g306550.t1.3
ribonuclease T2
activity
Small Rab-related
GTPase
g9677.t1
BSD domain
g6604.t1
Ca2+ transporting
ATPase
g11725.t1
NMD3 family
g15392.t1
ADP-ribosylation
factor
GTPase-activating
protein 2/3
Cre17.g715350.t1.2
14-3-3 protein
Cre06.g257500.t1.2
BSD domain
Cre06.g291250.t1.3