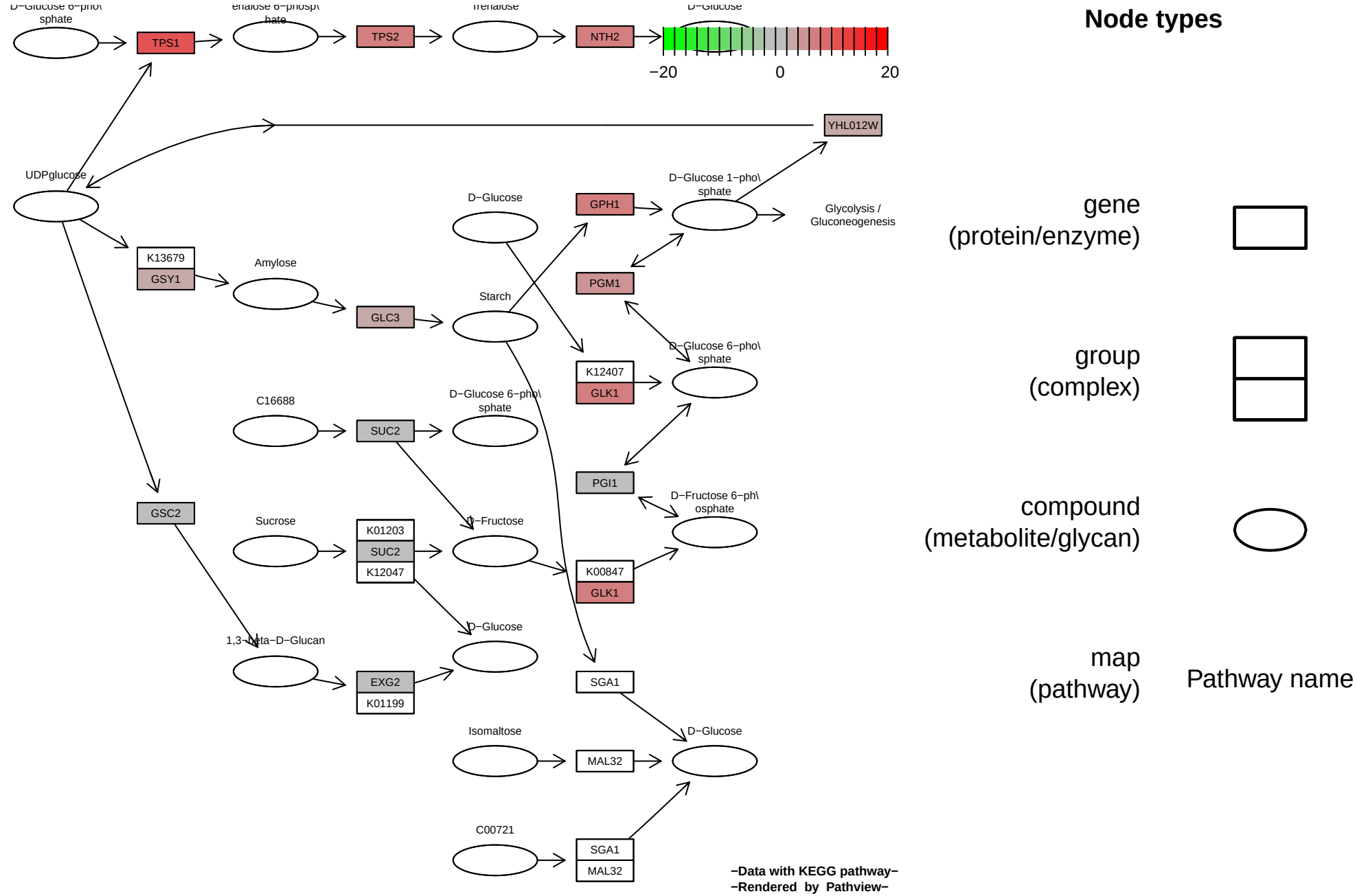
D−Glucose 6−pho\
sphate
TPS1
erulose 6−phosp\
hate
TPS2
Trehalose
NTH2
D−Glucose
−20
0
20
YHL012W
UDPglucose
D−Glucose 1−pho\
sphate
Glycolysis /
Gluconeogenesis
D−Glucose
GPH1
K13679
GSY1
Amylose
PGM1
GLC3
Starch
D−Glucose 6−pho\
sphate
K12407
GLK1
C16688
D−Glucose 6−pho\
sphate
SUC2
PGI1
D−Fructose 6−ph\
osphate
GSC2
Sucrose
K01203
SUC2
K12047
D−Fructose
K00847
GLK1
D−Glucose
1,3−beta−D−Glucan
EXG2
K01199
SGA1
Isomaltose
D−Glucose
MAL32
C00721
SGA1
MAL32
−Data with KEGG pathway−
−Rendered by Pathview−
Node types
gene
(protein/enzyme)
group
(complex)
compound
(metabolite/glycan)
map
(pathway)
Pathway name