

Inborn Errors of Metabolism

GLYCOGEN STORAGE DISEASES

LIVER GLYCOGENOSIS

- O Glycogen synthase 2.4.1.11
- I Glucose-6-phosphatase 3.1.3.9 (von Gierke)
- III Amylo-1,6-glucosidase (debrancher) 3.2.1.33 (Cori)
- IV 1,4-Glucan branching 2.4.1.18 (Andersen)
- VII Liver phosphorylase b kinase 2.7.1.38 (Hug)
- VI Liver phosphorylase 2.4.1.11 (Hers)

MUSCLE GLYCOGENOSIS

- V Muscle phosphorylase 2.4.1.11 (McArdle)
- II α-1,4-Glucosidase 3.2.1.20 (Pompe)
- VII Muscle phosphorylase kinase 2.7.1.11

MONO- AND DI-SACCHARIDE METABOLISM

- #### GALACTOSE
- β-Galactosidase 3.2.1.23
 - Galactokinase 2.7.1.6
 - Galactose-1-phosphate uridylyltransferase 2.7.1.12
 - UDP-Galactose 4-epimerase 5.1.3.2
 - Sorbitol dehydrogenase deficiency 1.1.1.14
 - Primary sucrose-isomaltase deficiency 3.2.1.48
 - Sucrose 3.2.1.48
 - Pentosa 1.1.1.10
 - L-Xylose reductase 1.1.1.10
- #### FRUCTOSE
- Essential fructosuria 3.2.1.13
 - Ketohexokinase 2.7.1.3
 - Fructokinase deficiency 2.7.1.4
 - Fructose-1,6-bis-phosphatase deficiency 3.1.3.11
 - Fructose intolerance 3.1.3.11
 - Fructose-1,6-bis-phosphatase aldolase (isozyme B) 4.1.2.13

PHOTOSYNTHESIS

- #### PENTOSE PHOSPHATE PATHWAY
- Phosphogluconate dehydrogenase (decarboxylating) 1.1.1.44
 - Glucose-6-phosphate dehydrogenase 1.1.1.49
- #### GLUCONEOGENESIS
- Glucose-6-phosphatase 3.1.3.9
 - Fructose-1,6-bis-phosphatase 3.1.3.11
 - Phosphoenolpyruvate carboxykinase (GTP) 4.1.1.32
 - Pyruvate carboxylase 6.4.1.1

NON-SPHEROCYTIC HAEMOLYTIC ANEMIAS DUE TO ERYTHROCYTE ENZYMOPATHIES

- Hexokinase 2.7.1.1
- Glucose-6-phosphate isomerase 5.3.1.9
- 6-Phosphofruktokinase 2.7.1.11
- Fructose-1,6-bis-phosphatase aldolase 4.1.2.13
- Tissue phosphate isomerase 5.3.1.1
- Glyceraldehyde-3-phosphate dehydrogenase 1.2.1.12
- Adenylate kinase 2.7.4.3
- Phosphoglycerate kinase 2.7.2.3
- Phosphoglycerate mutase 5.4.2.1
- Pyruvate kinase 2.7.1.40

LIPID METABOLISM

- Type II Glycolic aciduria 2.6.1.44
- Acylcarnitine hydrolase 3.1.1.28
- Acid lipase 3.1.1.3
- Triacylglycerol lipase 3.1.1.3
- Pancreatic lipase 3.1.1.3
- Long-chain acyl-CoA dehydrogenase 1.3.99.3
- Medium-chain acyl-CoA dehydrogenase 1.3.99.2
- Short-chain acyl-CoA dehydrogenase 1.3.99.2
- 3-Hydroxyacyl-CoA dehydrogenase 1.1.1.35
- Acetyl-CoA C-acetyltransferase 2.3.1.16
- Hydroxymethylglutaryl-CoA synthase 4.1.3.5
- Hydroxymethylglutaryl-CoA lyase 4.1.3.4

SPHINGOLIPIDOSES

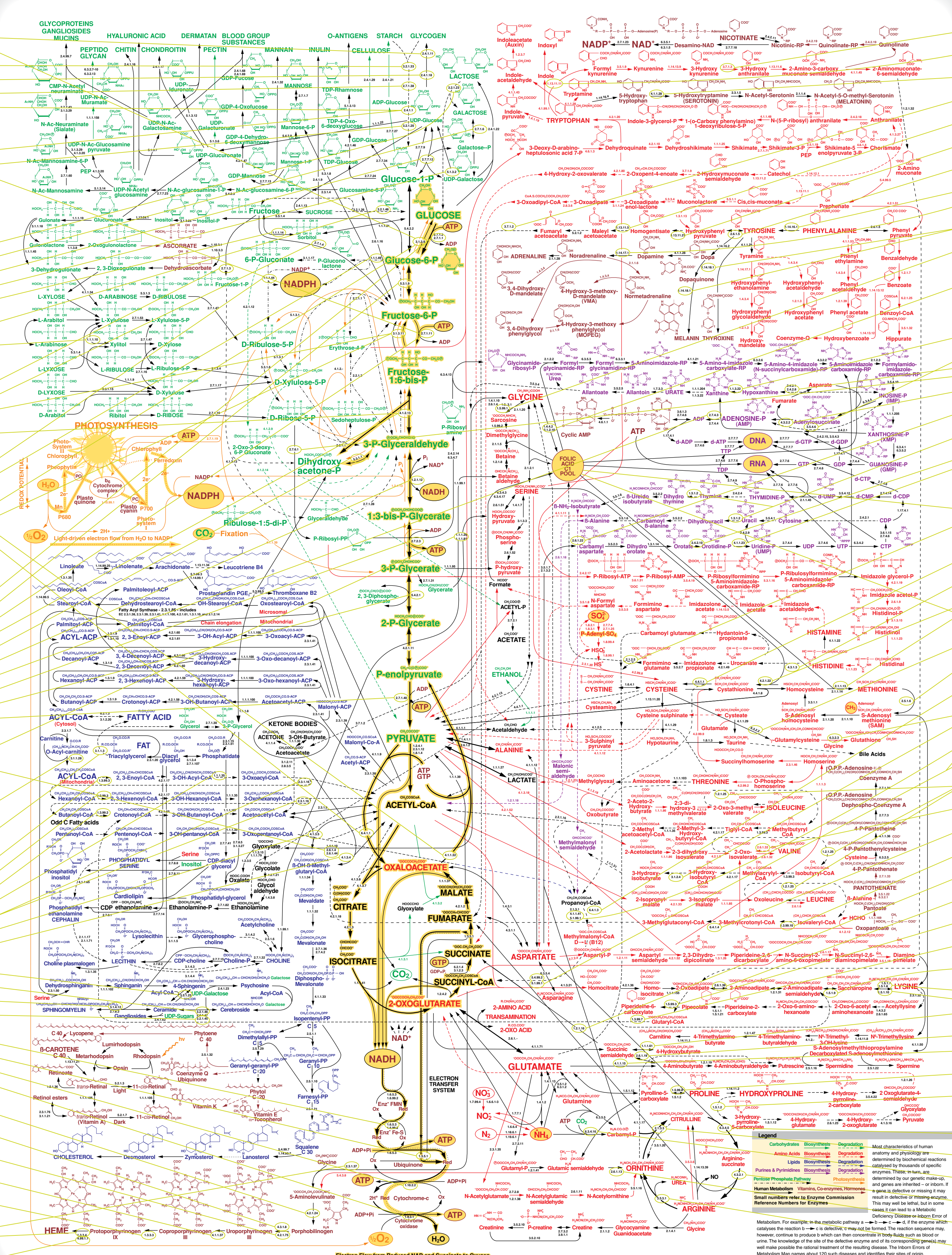
- Acylsphingosine deacylase (Ceramide) 3.5.1.23 (Farber)
- Galactosyl ceramidase 3.2.1.46 (Krabbe)
- Sphingomyelin phosphodiesterase 3.1.4.12 (Niemann-Pick)
- Glucosyl ceramidase 3.2.1.45 (Gaulcher)

RESPIRATORY CHAIN

- NADH dehydrogenase (Ubiquinone) 1.6.5.3
- Succinic dehydrogenase (Ubiquinone) 1.3.5.1
- Ubiquinol-cytochrome c reductase 1.10.2.2
- Cytochrome c oxidase 1.9.3.1

PORPHYRIAS

- Porphobilinogen synthase deficiency 4.2.1.24
- Acute intermittent porphyria (AIP) 4.3.1.8
- Porphyria cutanea tarda (PCT) 4.3.1.8
- Congenital erythropoietic porphyria 4.1.1.37
- Uroporphyrinogen III synthase 4.2.1.25
- Porphyria cutanea tarda (PCT) 4.1.1.37
- Uroporphyrinogen decarboxylase 4.1.1.37
- Hereditary coproporphyrin (HCP) 4.1.1.37
- Coproporphyrin oxidase 1.3.3.3
- Variegated Porphyria (VP) 4.1.1.37
- 5-Aminolevulinic acid synthase 2.3.1.37
- Protoporphyrin oxidase 1.3.3.4
- Erythropoietic Protoporphyrin 4.9.1.14



AMINO ACID METABOLISM

- Xanthurenic aciduria 3.7.1.3
- Tryptophanuria 1.13.11.11
- Hereditary Tyrosinemia (Type I) 4.1.1.28
- Fumarylacetoacetate 3.7.1.2
- Alcaptonuria 1.13.11.5
- Hawkinsinuria 1.13.11.7
- Hereditary Tyrosinemia (Type II) 2.6.1.5
- Tyrosine aminotransferase 2.6.1.5
- Phenylketonuria 1.14.16.1
- Dopamine hydroxylase 1.14.17.1
- Tyrosine-negative Albinism 1.14.18.1
- Hypersarcosinemia 1.5.99.1
- Non-ketotic hyperglycinemia 1.4.4.2
- Glycine decarboxylase (decarboxylating) 4.1.1.4
- Aminomethyltransferase 2.1.1.10

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