

Two patients with *HADH* (SCHAD) Hyperinsulinism in part without detectable 3-Hydroxybutyrylcarnitine/ 3-Hydroxyglutarate

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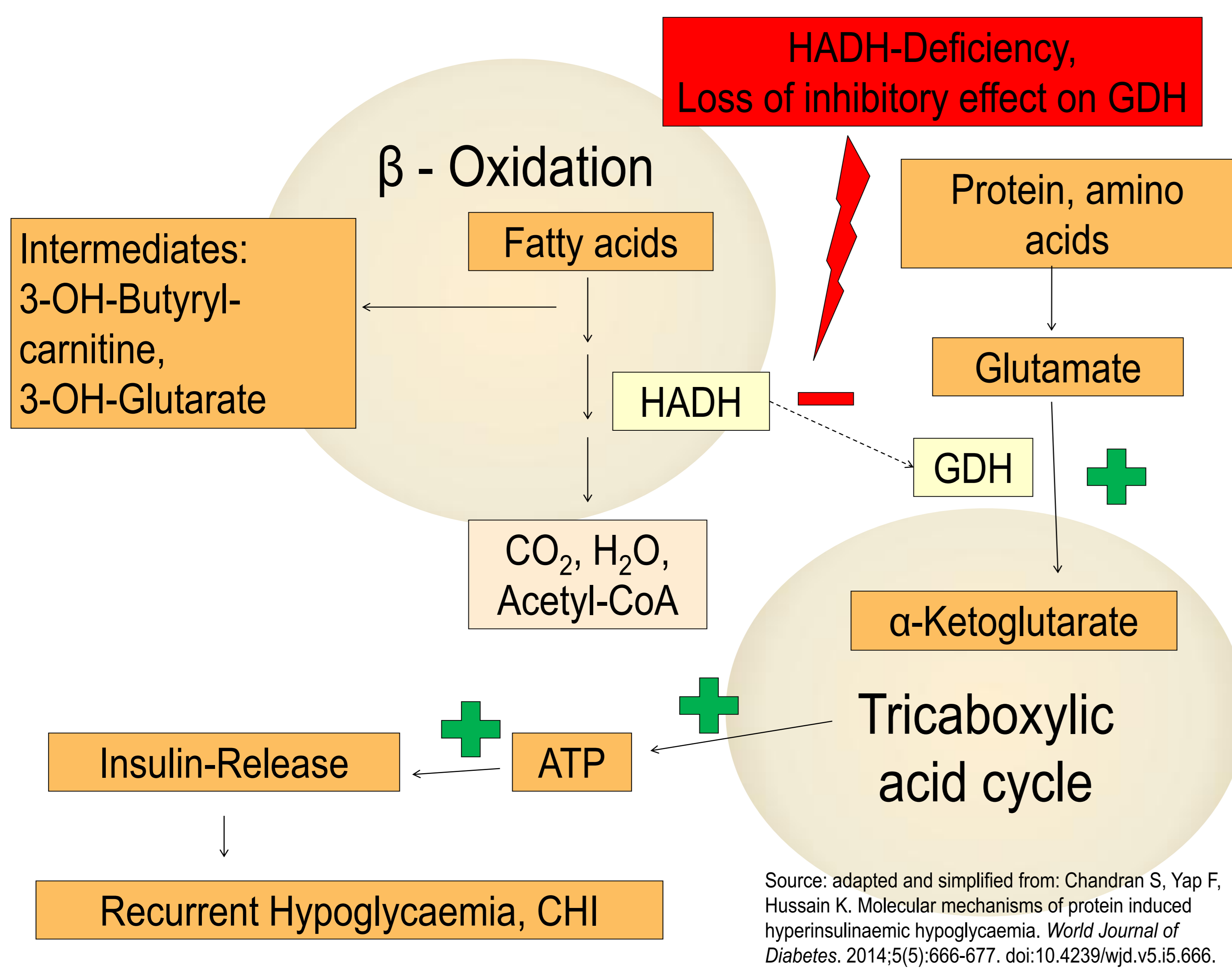
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Introduction

Congenital hyperinsulinism of infancy (CHI) is the most common cause for persisting hypoglycaemia in infancy. The most common genetic causes are mutations in *ABCC8* or *KCNJ11* (coding for K^+ _{ATP}-channel subunits), less frequently mutations in *GCK* or *GLUD1*. Further genetic analysis is often performed only if phenotypic aspects point to other specific genes, such as the very rare short chain 3-Hydroxyacyl-CoA dehydrogenase (*HADH*/*SCHAD*) deficiency. This disorder is usually characterized by an accumulation of 3-hydroxybutyrylcarnitine in plasma and 3-hydroxyglutarate in the urine.

Pathophysiology in beta cells of patients with *HADH* CHI



Patients: clinical presentation

Two patients of Turkish origin, consanguineous parents.

Patient 1: male, 17 years, postnatal macrosomy, recurrent hyperinsulinaemic hypoglycaemia since birth, **diazoxide responsive** (still ongoing, 4,5 mg/kg/d, side effect: hypertrichosis), mild cognitive impairment, obesity, older sister and grandmother also suffering from CHI.

Patient 2: female, 9 years, recurrent hyperinsulinaemic hypoglycaemia since birth, **diazoxide responsive** (still ongoing, 5mg/kg/d), PET-CT: diffuse form of CHI, normal cognitive function, grandfather and cousin also suffering from CHI.

Methods

Patient 1: No mutations in *ABCC8*, *KCNJ11*. Therefore extensive next-generation sequencing (NGS) was performed.

Patient 2: Sanger-sequencing of *ABCC8*/*KCNJ11*, *GLUD1*, *GCK* did not lead to a conclusive genetic diagnosis, and was followed by further Sanger sequencing of *HADH* and *HNF4A*.

Biochemical analysis: Acylcarnitine-profile (blood) and organic acids (urine) via tandem mass spectrometry in patient 1.

Results

Age	Sex	Gene	Mutation	Biochemical Analysis
17 J	m	HADH	HOM c.428T>A (p.Ile143Asn), new mutation	no detection of significant concentrations of 3-OH-Butyrylcarnitine or 3-OH-Glutarate.
9 J	w	HADH	HOM c.706C>T (p.Arg236*) mutation published before ¹	no analysis.

Conclusions

HADH deficiency should be considered in patients with CHI who are negative for *ABCC8* and *KCNJ11*, and might be more frequent than known so far. Its specific biochemical markers are not necessarily present in individual patients or situations, and should not be regarded as a prerequisite for sequencing of *HADH* gene.

These data underline the broad clinical and genetic heterogeneity of CHI, and the value of extensive sequencing, e.g. using NGS, to detect the molecular cause of the disease.

Acknowledgements and literature

Written and informed consent was obtained from parents before inclusion in the study.

Literature:

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