The molecular mechanisms, diagnosis and management of congenital hyperinsulinism

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ABSTRACT

Congenital hyperinsulinism (CHI) is the result of unregulated insulin secretion from the pancreatic β-cells leading to severe hypoglycaemia. In these patients it is important to make an accurate diagnosis and initiate the appropriate management so as to avoid hypoglycemic episodes and prevent the potentially associated complications like epilepsy, neurological impairment and cerebral palsy. At a genetic level abnormalities in eight different genes (*ABCC8, KCNJ11, GLUD1, GCK, HADH, SLC16A1, HNF4A* and *UCP2*) have been reported with CHI. Loss of function mutations in *ABCC8/KCNJ11* lead to the most severe forms of CHI which are usually medically unresponsive. At a histological level there are two major subgroups, diffuse and focal, each with a different genetic etiology. The focal form is sporadic in inheritance and is localized to a small region of the pancreas whereas the diffuse form is inherited in an autosomal recessive (or dominant) manner. Imaging using a specialized positron emission tomography scan with the isotope fluroine-18 L-3, 4-dihydroxyphenyalanine (18F-DOPA-PET-CT) is used to accurately locate the focal lesion pre-operatively and if removed can cure the patient from hypoglycemia. Understanding the molecular mechanisms, the histological basis, improvements in imaging modalities and surgical techniques have all improved the management of patients with CHI.

Key words: Diazoxide, hypoglycemia, hyperinsulinism

INTRODUCTION

Congenital hyperinsulinism (CHI) is an important cause of hyperinsulinaemic hypoglycaemia (HH) in the neonatal, infancy and childhood periods. It occurs due to the unregulated secretion of insulin form pancreatic β -cells leading to severe and persistent hypoglycemia. The early recognition, diagnosis and immediate management are important as delay in the diagnosis and inappropriate management can lead to hypoglycemic brain injury.^[1] CHI is associated with an increased risk of brain injury secondary to the metabolic actions of insulin. Unregulated insulin secretion leads to suppression of glucose production by

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inhibition of glycogenolysis and gluconeogenesis.^[2] In the absence of glucose as energy source, brain tends to utilise the ketone bodies as an alternative substrate. However, in CHI, brain is deprived of both glucose and ketones as insulin inhibits lipolysis and ketogenesis.

The other forms of hyperinsulinaemic hypoglycaemia (HH) include transient hyperinsulinism due to risk factors such as birth asphyxia, intra-uterine growth retardation (IUGR) and maternal diabetes mellitus (gestational or insulin dependent), several developmental syndromes (such as Beckwith-Wiedemann) and rare metabolic conditions such as CDG syndromes.^[3]

CHI is extremely heterogeneous in terms of clinical presentation, histological subgroups and underlying molecular biology. The incidence of CHI can vary from 1 in 35,000-40,000 in the general population^[4] to 1 in 2500 in some communities with high rates of consanguinity.^[5] The molecular basis of CHI involves genetic defects in eight different genes (*ABCC8, KCNJ11, GLUD1, GCK*,

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HADH, SLC16A1, HNF4A and UCP2) that are involved in regulating insulin secretion from β -cells.^[6]

At a histological level two major subtypes of CHI (diffuse and focal) have been described.^[7] The diffuse form is inherited in an autosomal recessive (or dominant) manner whereas the focal form is sporadic in inheritance. The differentiation of these two histological subtypes is important from the management point of view. The diffuse from may require a near total pancreatectomy (with the risk of diabetes mellitus and pancreatic exocrine insufficiency) whereas the focal form will only require a limited focal lesionectomy. Recently a specialized positron emission tomography scan using the isotope Fluorine 18 L-3, 4-dihydroxyphenyalanine (¹⁸F-DOPA-PET) has become the gold standard for the localisation of focal lesions within the pancreas.^[8]

The aim of this review is to provide an overview of pancreatic β -cell physiology and insulin secretion, to review the different molecular mechanisms of CHI, to outline the clinical presentation and describe the recent advances in imaging and management.

Pancreatic β -Cell Physiology, Glucose Metabolism and Insulin Secretion

Pancreatic β -cells play a key role in regulating insulin production and secretion. Insulin secretion is tightly linked to glucose metabolism by pancreatic β -cell ATP-sensitive K^+ channels. These ATP-sensitive K^+ channel (K_{ATP} channel) plays a fundamental role in glucose homeostasis by linking glucose metabolism to electrical excitability of the β -cell and insulin secretion.^[9] The β -cell K_{ATD} channel is a hetero-octameric complex (arranged in a 4:4 stoichiometry) composed of two types of subunits: Four inward-rectifying potassium channel pore-forming (Kir6.2) subunits and four high-affinity sulfonylurea receptor 1 (SUR 1) subunits.^[10] The Kir6.2 forms the core of the channel whilst the SUR1 (an ATP binding cassette transporter) acts as a regulatory subunit. KATP channels can only function if they are assembled and correctly transported to the cell membrane surface (trafficking). The assembly and trafficking of K_{ATP} channels are intricately linked processes. Only octameric KATP channel complexes are capable of expressing on the cell membrane surface. For example both Kir6.2 and SUR1 possess an endoplasmic reticulum (ER) retention signal (RKR) that prevents the trafficking of each subunit to the plasma membrane in the absence of the other subunit.^[11] Co expression of two subunits masks these retention signals, allowing them to traffic to the plasma membrane. The retention signal is present in the C terminal region of Kir6.2 and in an intracellular loop between TM11 and NBF-1 in SUR1. Truncation of the C-terminus of Kir6.2 deletes its retention signal, allowing functional expression of Kir6.2 in the absence of SUR1 subunit.^[12]

 K_{ATP} channels are regulated by adenine nucleotides to convert changes in cellular metabolic levels into membrane excitability.^[13] The Kir6.2 subunit determines the biophysical properties of the channel complex including K⁺ selectivity, rectification, and inhibition by ATP and activation by acyl-CoAs.^[12] The sulfonylurea receptors endow K_{ATP} channels with sensitivity to the stimulatory actions of Mg-nucleotides and K_{ATP} channel openers like diazoxide and the inhibitory effects of sulfonylureas.^[14]

Glucose metabolism in the β -cells regulates insulin secretion. Metabolism raises the intracytosolic ATP/ADP ratio which inhibits the plasma membrane sulfonylurea receptor 1 (SUR 1). This results in the closure of the K_{ATP} channel which in turn leads to cell membrane depolarisation and Ca²⁺ influx into the cell via voltage gated calcium channels. The increase in the intracellular concentration of calcium triggers the release of insulin. When glucose levels are low, K_{ATP} channels are open and potassium diffusing via these channels maintains the resting membrane potential at a hyperpolarized level.

Glucose stimulates insulin secretion by triggering (KATP channel dependent) and amplifying (augmentation pathway) signals in beta-cells. The triggering pathway is well characterized involving the pancreatic beta-cells. The augmentation pathway is less well defined but involves an increase in the efficacy of Ca²⁺ on the exocytosis of insulin granules. The amplification pathway serves to optimize the secretory response not only to glucose but also to non-glucose stimuli.

Aetiology of Congenital Hyperinsulinism [Figure 1]

So far at a genetic level, abnormalities in eight different genes have been described which lead to unregulated insulin secretion. The most common forms are due to defects in the genes which encode the components of the pancreatic K_{ATP} channel. Below is a brief description of the different forms of CHI.

Pancreatic β-cell K_{ATP} channel defects

The Kir6.2 and SUR1 subunits are encoded by the genes KCNJ11 and ABCC8 (localized to chromosome 11p15.1), respectively. Recessive mutations in ABCC8 and KCNJ11 are the most common causes of CHI.^[15,16] The inactivating mutations in ABCC8/KCNJ11 reduce or completely abolish the activity of the K_{ATP} channel, leading to unregulated insulin release despite severe hypoglycemia.^[15] Germline

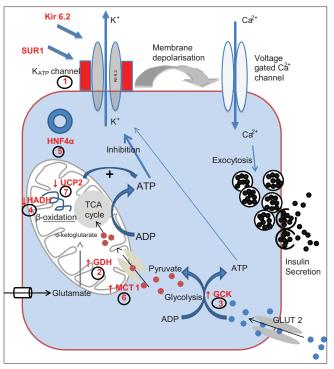


Figure 1: Common mutations associated with CHI. (1) ATP gated K⁺ channel (KATP) encoded by ABCC8 and KCNJ11; (2) Glutamate Dehydrogenase (GDH) encoded by GLUD1; (3) Glucokinase (GCK) encoded by GCK gene; (4) L-3-hyroxyacyl-coenzyme A dehydrogenase (HADH) encoded by HADH; (5) Hepatocyte Nuclear Factor 4α (HNF 4α) encoded by HNF4A gene; (6) The moncarboxylate transporter (MCT1) encoded by SLC16A1; (7) Uncoupling Protein 2 (UCP2)

mutations in *ABCC8/KCNJ11* are found in approximately 50% of CHI patients.^[16] To date about 150 homozygous, compound heterozygous and heterozygous inactivating mutations in *ABCC8* and around 24 *KCNJ11* mutations have been reported.^[16]

The recessive inactivating mutations in *ABCC8* and *KCNJ11* usually cause severe CHI which is unresponsive to medical treatment with diazoxide. The molecular basis of recessive inactivating *ABCC8* and *KCNJ11* mutations involves defects in K_{ATP} channel biogenesis and turnover,^[17] channel trafficking from the ER and Golgi apparatus^[18,19] to the plasma membrane and alterations of channels in response to nucleotide regulation and open state frequency.^[20] Dominant inactivating mutations in *ABCC8* and *KCNJ11* usually cause CHI with a milder phenotype^[21,22] although medically unresponsive forms have recently been reported.^[23]

Hyperinsulinism-hyperammonaemia syndrome

Hyperinsulinism-hyperammonemia syndrome (HI/HA), the second most common form of (CHI) is associated with activating missense mutations in the *GLUD1* gene, which encodes the mitochondrial matrix enzyme, glutamate dehydrogenase (GDH). Patients present with recurrent symptomatic postprandial hypoglycemia following protein-rich meals (leucine-sensitive hypoglycemia) as well as fasting hypoglycemia accompanied by asymptomatic elevations of plasma ammonia. GDH is expressed in liver, kidney, brain, and pancreatic β -cells. Mutations in *GLUD1* lead to a gain of enzyme function by reducing its sensitivity to allosteric inhibition by the high-energy phosphates such as GTP and ATP and allowing activation by the amino acid leucine.^[24]

GDH catalyses the reversible oxidative deamination of glutamate to alpha-ketoglutarate and ammonia using NAD or NADP as co-factors. The increased GDH activity leads to inappropriate insulin secretion in pancreatic β -cells, as well as to excessive ammonia production and reduced urea synthesis in the liver. Recent animal studies have suggested the role of renal ammoniagenesis due to activation of GDH as a source of hyperammonemia in these patients.^[25] The phenotype is characterized by recurrent postprandial hypoglycemia following protein-rich meals as well as fasting hypoglycemia accompanied by asymptomatic hyperammonemia.^[26]

Though hyperammonemia has remained the most consistent feature of HI/HA, there is a rare group of patients with *GLUD1* mutations who demonstrate leucine hypersensitivity but have normal serum ammonia level.^[27] These patients may be mosaic for GDH enzyme activity (with normal GDH activity in the liver but elevated activity in the pancreas), however, this remains to be proven.

The phenotype of HI/HA is reported to be milder in contrast to other forms of CHI, thus escaping recognition for the first few months of life.^[27] Patients with HI/HA syndrome have more neurological complications such as epilepsy and learning disabilities.^[28] However, in contrast to hyperanmonemia patients due to urea cycle disorders, patients with HI/HA syndrome do not experience lethargy, headaches, or manifest CNS symptoms that might be expected for their degree of hyperanmonemia, and are resistant to ammonia scavenging agents or protein restriction. Routine measurement of plasma ammonia concentrations in all patients with hypoglycemia is an essential screening test for the disorder.

HADH and hyperinsulinism

HADH encodes for the enzyme 3-hydroxyacyl-coenzyme A dehydrogenase (HADH) and catalyses the penultimate reaction in the beta-oxidation of fatty acids the NAD⁺-dependent conversion of L-3-hydroxyacyl-CoA to 3-ketoacyl-CoA. Mutations in the mitochondrial HADH gene (encoding the enzyme L-3-hydroxyacyl-Coenzyme A dehydrogenase, HADH), are a rare cause of CHI.^[29] This

enzyme catalyses the conversion of L3-hydroxyacyl CoAs of variable chain length to their corresponding 3-ketoacyl CoAs and exerts highest activity to 3-hydroxybutyryl-CoA. HADH gene has been mapped to chromosome 4q22- $26^{[30]}$ and is expressed in most tissues, although the enzyme activity is high in the pancreas, particularly in the islets of Langerhans.^[31] HADH expression is regulated by transcription factors such as Foxa2, which are essential for β -cell differentiation and mice which have Foxa2 knocked out, showed a 3-fold down regulation of HADH mRNA and severe HH.^[32]

HADH gene mutations can lead either to severe neonatal HH or to mild late onset HH.^[32,33] All patients reported so far have responded to diazoxide and some had abnormal acylcarnitine metabolites (raised plasma hydroxybutyrylcarnitine and urinary 3-hydroxyglutarate levels). Protein sensitivity has been demonstrated in patients with HADH mutations^[34] and this has been confirmed in the HADH knockout mouse.^[35] However, the precise mechanism of dysregulated insulin secretion in patients with a HADH deficiency is not understood but might involve an interaction between GDH and HADH.^[35] Genetic analysis for *HADH* gene is recommended in patients with diazoxide responsive HH from consanguineous families, who are negative for mutations in the K_{ATP} channels.^[36]

HNF4A and hyperinsulinism

HNF4A gene encodes for the transcription factor hepatocyte nuclear factor 4α (HNF-4α), which belongs to the nuclear hormone receptor superfamily and has been shown to control the expression of genes involved in glucose stimulated insulin secretion.^[37] Heterozygous mutations in the *HNF4A* gene causes maturity-onset diabetes of the young type 1 (MODY1), which is characterized by progressive β-cell dysfunction and failure of glucose induced insulin secretion.^[38]

Recently heterozygous mutations in the *HNF4A* gene were also reported to result in transient^[39] or persistent HH.^[40] The phenotype of these patients is characterized by macrosomia and neonatal HH.^[41,42] The severity of HH in these patients varies from diet-controlled neonatal hypoglycemia to persistent HH requiring diazoxide treatment. In a recent series of 11 patients with HNF4A mutations, the HH was noted to range from three months to eight years with on-going need for diazoxide therapy.^[43] Interestingly, only few of the parents of children in this series had diabetes, suggesting that the absence of a history of diabetes in the parents should not preclude sequencing of the *HNF4A* gene. HNF4A mutation has been noted to have variable penetrance with only a minority of HNF4A mutation carriers developing HH. The precise mechanism by which HNF4A mutations cause HH is not clear but might involve a reduction in expression of the potassium channel subunit Kir6.2^[37] or reduction in the levels of PPAR α .^[44] PPAR α is a transcription factor that is known to control the expression of genes encoding enzymes of the beta oxidation pathway of fatty acids. Low levels of PPAR α are reported in HNF-4 α deficient β -cells.^[37] It can be postulated that HNF-4 α deficiency causes lower levels of PPAR α and a decrease in beta-oxidation of fatty acids resulting in the accumulation of lipids (such as malonyl-CoA) in the cytoplasm. Increased malonyl-CoA is thought to inhibit the enzyme carnitine-palmitoyltransferase1 thereby increasing cytosolic long-chain acyl-CoA levels, which signals insulin release.^[45] In support of this hypothesis PPARa null mice develop fasting HH suggesting that PPAR α is important for regulated insulin secretion during fasting.^[44]

Exercise-induced hyperinsulinism (SLC16A1)

Exercise-induced hyperinsulinism (EIHI) is a dominantly inherited and characterized by the inappropriate insulin secretion during anaerobic exercise or on pyruvate load.^[46-48] To date, 13 patients have been reported, 12 from two Finnish pedigrees and one unrelated patient.^[49] Affected patients become hypoglycemic typically 30-45 minutes after a period of intensive anaerobic exercise.^[47]

The transport of lactate and pyruvate into the β -cells are mediated by monocarboxylate transporter 1 (MCT1) which is encoded by the *SLC16A1* (solute carrier family 16, member 1) gene. Under normal physiological conditions lactate and pyruvate concentrations are low in β -cells and do not stimulate insulin secretion.^[50] However, promoter-activating mutations in *SLC16A1* induce the expression of MCT1 in β -cells (where this gene is not usually transcribed) permitting pyruvate uptake and pyruvate-stimulated insulin release despite ensuing hypoglycemia.^[49] This novel disease mechanism results from the failure of cell-specific transcriptional silencing of a gene (*SLC16A1*) that is highly expressed in other tissues. During strenuous anaerobic exercise there is accumulation of lactate and pyruvate which then act as insulin secretagogues.

Glucokinase induced hyperinsulinism

Glucokinase is a key glycolytic enzyme that plays a pivotal role as a glucose sensor in the pancreatic β -cell and appears to have a similar role in entero-endocrine cells, hepatocytes, and hypothalamic neurons.^[51] In β -cells, glucokinase a rate-limiting enzyme for glucose metabolism governs glucose-stimulated insulin secretion. Heterozygous activating mutations of glucokinase lead to CHI.^[52] These glucokinase mutations result in increased affinity of the enzyme for glucose, resulting in an increase in the ATP: ADP ratio in the pancreatic β -cell, closure of K_{ATP} channel, and inappropriate insulin secretion. The activating glucokinase mutations are inherited in an autosomal dominant manner with the severity of symptoms varying markedly within and between families.^[53] The age of presentation of glucokinase induced hyperinsulinism can range widely from infancy to adulthood.^[54] Affected children have fasting hypoglycemia and may manifest variable responsiveness to medical treatment. Some appear to respond well to pharmacologic intervention with diazoxide, whilst others require more intensive medical management including octreotide and even surgery.^[55]

Mutations in the UCP2 gene and HH

Uncoupling protein 2 (UCP2) is expressed in several tissues, and acts in the protection against oxidative stress, in fatty acid metabolism and in the negative regulation of insulin secretion by beta cells.^[56] UCP2 knockout mice exhibit HH, suggesting an involvement of UCP2 in insulin secretion. A few patients have been described with mild HH due to loss of function mutations in the UCP2 gene.^[57] Parental-inherited heterozygous UCP2 variants encoding amino-acid changes were found in two unrelated children with CHI.^[57] Functional assays in yeast and in insulin-secreting cells revealed an impaired activity of UCP2 mutants.^[57]

HISTOLOGICAL SUBTYPES OF CONGENITAL HYPERINSULINISM

There are two major histological subtypes of CHI; diffuse and focal.^[58] The diffuse form consists of hyper-functioning pancreatic β -cells and affects the whole of pancreas. The most common causes of diffuse CHI are the recessive and dominant mutations in *ABCC8* and *KCNJ11*. Patients with diffuse disease due to recessive mutations in *ABCC8* and *KCNJ11* do not usually respond to diazoxide.

The second histological subtype of CHI is the focal disease, which involves a small localized region of pancreas (2-10 mm in diameter). It is characterized by nodular hyperplasia of islet-like cell clusters, including ductuloinsular complexes and giant β -cell nuclei surrounded by a histologically and functionally normal pancreatic tissue.^[59] The focal lesions can sometimes be deeply embedded within the pancreatic tissue.

The focal form has a distinctive genetic etiology from that of the diffuse disease and involves two independent events, the first of which is the inheritance of a paternal mutation in *ABCC8* or *KCNJ11*.^[60] The second event is the somatic loss of the maternal 11p allele (11p15.1 to 11p15.5) involving the *ABCC8* and *KCNJ11* region within the focal lesion.^[61] This paternal uniparental disomy unmasks the paternally inherited K_{ATP} channel mutation, which leads to altered expression of a number of imprinted genes, including the maternally expressed tumour suppressor genes *H19* and *CDKN1C*, and the paternally expressed growth factor *IGF2*.^[61] These events eventually give rise to the increase in proliferation of β -cells evolving into a focal adenomatous hyperplasia. The focal disease is always sporadic in origin.

DIFFERENTIAL DIAGNOSIS

Transient HH

Transient HH generally refers to the group of patients in whom HH resolves spontaneously within few days after birth. Transient HH is associated with certain risk factors such as maternal diabetes mellitus (gestational or insulin dependent), intra-uterine growth retardation, perinatal asphyxia, erythroblastosis fetalis, maternal use of drugs like sulphonylureas, and intravenous maternal glucose infusions during labor. Occasionally, some patients with intra-uterine growth retardation and perinatal asphyxia have a protracted form of HH which resolves over several months and may require treatment with diazoxide.^[62] The mechanism of transient HH is not clear.

Metabolic conditions associated with HH Congenital disorders of glycosylation

Congenital disorders of glycosylation (CDG) are a rapidly evolving family of inherited multisystem disorders resulting from defects in the synthesis of the glycan moiety of glycoconjugates (mainly glycoproteins or glycolipids) or in the attachment of glycans to macromolecules.^[63] CDG type Ia is the most common and is caused by mutations in the phosphomannomutase 2 gene (PMM2) gene. The clinical spectrum of CDG type Ia has expanded since its initial description to now include rare features such as HH congenital nephrotic syndrome and obstructive cardiomyopathy.^[64] A milder disease with single organ involvement, presenting as isolated HH has been reported in a female patient.^[65]

HH as a leading symptom has been described predominantly in CDG type Ib (phosphomannose-isomerase deficiency).^[64] The phenotype is characterized by protein-losing enteropathy, congenital hepatic fibrosis, and coagulopathy without overt neurologic manifestations that are commonly seen in other CDGs. Early diagnosis is essential, because patients can be successfully treated with oral mannose.^[66] HH has also been described in patients with CDG Id.^[67]

The precise mechanism for insulin dysregulation is unknown, but the rapid resolution of HH in CDG type Ib patients with oral mannose supplementation suggests a role of glycosylation in maintenance of normoglycemia, perhaps at the level of the sulfonylurea receptor. CDGs should be considered in patients with HH of undiagnosed etiology.

Postprandial forms of HH

Postprandial hyperinsulinemic hypoglycemia (PPHH) refers to the development of hypoglycemia within a few hours of meal ingestion. It is associated with inappropriate insulin secretion in response to the meal. The most common cause is due to the "dumping" syndrome in infants who have undergone gastro-oesophageal surgery.^[68] It has also been observed in adult patients who have undergone gastric bypass surgery for morbid obesity.^[69] It has been noted that children with PPHH after Nissen fundoplication have abnormally exaggerated secretion of glucagon like peptide-1 (GLP-1) which may contribute to the exaggerated insulin surge and resultant hypoglycemia.^[70] PPHH is also observed in the insulin autoimmune syndrome which is characterized by the presence of insulin-binding autoantibodies in subjects who have not been previously exposed to exogenous insulin.^[71]

A syndrome of autosomal dominant PPHH with onset in adolescence to adulthood and linked to a mutation (Arg1174Gln) in the insulin receptor kinase gene has been reported.^[72] Impaired insulin clearance and insulin resistance due to mutations in insulin receptor have been hypothesized to affect insulin action differently in various tissues leading to hypoglycemia.

In adults, a syndrome of "non-insulinoma pancreatogenous" PPHH has been recognized.^[73] These patients demonstrate neuroglycopenic episodes from hypoglycemia within 4hrs of meal ingestion and have negative 72 hrs controlled fasts. The exact mechanism of hypoglycemia is not clear.

Syndromes associated with HH

A large number of developmental syndromes may present in the newborn period with HH [Table 1]. The most common syndrome associated with HH is Beckwith-Wiedemann syndrome (BWS).^[74] This syndrome is characterized by prenatal and/or postnatal overgrowth, macroglossia, anterior abdominal wall defects, organomegaly, hemihypertrophy, ear lobe creases, helical pits, and renal tract abnormalities. HH is observed in about 50% of patients with BWS and in the vast majority of patients with BWS, HH is usually transient and resolves spontaneously in a few days.^[74] However, a small number of patients (5% of cases), have persistent HH requiring medical therapy or even sub-total pancreatectomy.

Other causes of HH

An insulinoma must be considered in older children or adolescents with HH.^[75] Insulinoma may be a part of multiple endocrine neoplasia syndrome type 1 (MEN1)

Table 1: Summary of the syndromes associated with hyperinsulinaemic hypoglycaemia

Developmental syndromes

Pre and post-natal overgrowth syndromes Beckwith-Wiedemann syndrome
Sotos syndrome
Simpson Golabi Behmel syndrome
Chromosomal abnormality syndromes
Trisomy 13 (Patau syndrome)
Mosaic Turner syndrome
Postnatal growth failure syndromes
Kabuki syndrome
Costello syndrome
Contiguous gene deletion affecting the ABCC8 gene
Usher syndrome
Syndromes with abnormal calcium homeostasis
Timothy syndrome
Congenital disorders of Glycosylation syndromes (CDG)
Congenital disorder of glycosylation 1a
Congenital disorder of glycosylation 1b
Congenital disorder of glycosylation 1d
Others
Congenital central hypoventilation syndrome

and hence a family history may provide a diagnostic clue in the familial cases. Munchausen by proxy can present as factitious HH due to administration of insulin or anti diabetic drugs such as sulphonylureas. In some cases, this has led to misdiagnosis and consequent pancreatectomy.^[76]

CLINICAL PRESENTATION

CHI most commonly presents in the newborn but it can also present during infancy childhood and rarely in adults. The clinical presentation of CHI is most severe in the newborn and may be quite subtle in the infancy and childhood periods. The CHI due to recessive mutations in *ABCC8/KCNJ11* genes is usually refractory to oral feeds and requires high concentrations of intravenous glucose to maintain normoglycemia.^[1] However, the milder forms may be able to maintain normoglycemia on oral feeds. Hypoglycemic symptoms may vary from being non-specific (such as poor feeding, lethargy and irritability) to severe (such as apnea, seizures or coma).

As a result of the foetal hyperinsulinemia, newborns with CHI may be macrosomic, however; the absence of macrosomia does not exclude CHI. Hypertrophic cardiomyopathy and hepatomegaly is observed in some patients with CHI. The mechanism of cardiomyopathy and hepatomegaly in these patients is unclear but might be related to the effect of fetal hyperinsulinemia.^[1] In some patients, hypoglycemia and associated symptoms manifest only following protein ingestion or exercise. In dumping syndrome, hypoglycemia occurs typically post feed. Hence it is important to ask these specific questions in the history to identify the aetiology.

DIAGNOSIS OF CONGENITAL Hyperinsulinism

It is essential to make an early diagnosis of CHI so as to avoid hypoglycemic brain injury hence clinicians should have a low threshold for recognizing these patients. Any patient with recurrent or persistent hypoglycemia can potentially have CHI. CHI is the only condition, wherein hypoglycemia can persist despite continuous intravenous administration of glucose. An intravenous glucose infusion rate of >8 mg/kg/min (normal 4-6 mg/kg/min) is virtually diagnostic of CHI.^[1] Factors like duration of fasting, protein sensitivity (hypoglycemia precipitated by meals) and hypoglycemia induced by exercise aid the diagnosis of milder forms of CHI.

The key feature of CHI is an inappropriate concentration of serum insulin (and/or c-peptide) for the level of blood glucose (spontaneous or provoked). Any detectable insulin during an episode of hypoglycemia is abnormal. However, some assays (Delphia) will detect insulin even when the blood glucose is low making a cut off more difficult. The other biochemical features are inappropriately low levels of serum ketone bodies and fatty acids during the hypoglycemic episode, which are secondary to the metabolic effects of in appropriate insulin levels. There is no correlation between the serum insulin concentration and the severity of the hypoglycemia.^[77] The level of hypoglycaemia is not only dependent on augmented insulin secretion, but also on the lack of insulin switch off at low glucose levels. An elevated C peptide could be of additional diagnostic benefit. The diagnostic criteria for HH are summarized in Table 2.

A rise in the blood glucose concentration of >1.5 mmol/l within 30 minutes following an intramuscular/intravenous injection of glucagon at the time of hypoglycemia provides

Table 2: Diagnostic criteria for patients with hyperinsulinaemic hypoglycaemia

Glucose infusion rate >8 mg/kg/min-Laboratory blood glucose <3 mmol/l with: -Detectable serum insulin/C-peptide* -Suppressed/low serum ketone bodies -Suppressed/low serum fatty acids Serum ammonia level may be raised (HI/HA syndrome) Raised plasma hydroxybutyrylcarnitine and urinary 3-hydroxyglutarate (HADH deficiency) Supportive evidence (when diagnosis is in doubt or difficult): Positive glycemic (>1.5 mmol/L) response to intramuscular/ intravenous glucagon Positive glycemic response to a subcutaneous/intravenous dose of octreotide Low serum levels of IGFBP1 [insulin negatively regulates the expression of IGFBP1] Suppressed branch chain (leucine, isoleucine and valine) amino acids Provocation tests (leucine loading or exercise testing) if indicated

*However, some assays (Delphia) will detect insulin even when the blood glucose is low making a cut off more difficult

supportive evidence in some patients.^[78] An improvement in hypoglycemia to a subcutaneous dose of octreotide may also aid diagnosis along with decreased serum levels of insulin growth factor binding protein 1 (IGFBP-1) as insulin suppresses the transcription of the IGFBP-1 gene.^[79] HI/HA syndrome is a possibility when elevated serum ammonia concentration is noted in a patient with CHI.^[25] Raised plasma hydroxybutyrylcarnitine and urinary 3-hydroxyglutarate might suggest HADH deficiency.^[30]

Diagnosis of certain types of CHI needs provocation testing. Patients with HI/HA syndrome have fasting as well as protein induced hypoglycemia. A protein/leucine load test will be necessary to demonstrate hypoglycemia in HI/HA patients.^[27] Similarly, patients with exercise induced CHI will require a formal exercise test and or a pyruvate load to demonstrate hypoglycemia.^[48,49]

As CHI is a rare disorder, it is important to network with the local tertiary centers where the expertise is available to guide on appropriate diagnosis and management.

Management of Congenital Hyperinsulinism

A prompt diagnosis and initiation of appropriate management will reduce the risk of brain injury in these patients. A higher threshold of blood glucose concentration is recommended to diagnose hypoglycemia and initiate treatment due to the metabolic effects of high insulin levels.^[2] It is important to maintain the blood glucose concentrations within the normal range (3.5-6 mmol/l),^[2] which may necessitate the use of high concentrations of glucose through a central venous catheter. A combination of intravenous fluids and oral feeds with a glucose polymer (such as Maxijul or Polycal) can be used to simulate orality and maintain normoglycaemia.

In situations where it is difficult to secure a venous access, intramuscular glucagon (0.5-1 mg) can be administered in order to temporarily improve blood glucose concentrations.^[1] Glucagon acts immediately by releasing glycogen stores from the liver and also has actions on gluconeogenesis, ketogenesis and lipolysis. It is important to follow this with intravenous glucose infusion to prevent rebound hypoglycemia (as glucagon in high doses causes paradoxical insulin secretion). Glucagon can also be administered as continuous intravenous or subcutaneous infusion to stabilize blood glucose concentrations in the acute management of infants with CHI.

Figure 2 gives an overview of the management of patients with CHI.

The mainstay of medical therapy is diazoxide, which is used

Senniappan, et al.: Hyperinsulinemic hypoglycemia

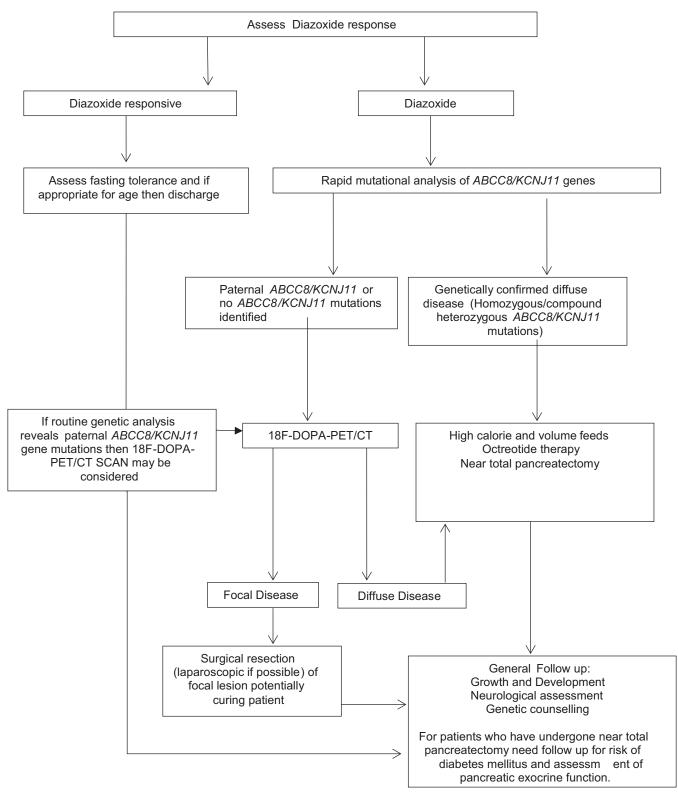


Figure 2: Outline of the suggested diagnostic and management cascade of patients presenting with CHI. The assessment of the response to diazoxide is critical in terms of planning further investigations

as a first line drug.^[1] Diazoxide binds and activates intact K_{ATP} channels, thereby reducing the insulin secretion. The side effects include fluid retention and hypertrichosis. The fluid retention is mostly observed in the neonatal period,

and may cause cardiac failure. A thiazide diuretic is usually added to prevent fluid retention. However, routine use of thiazide diuretic is not necessary in older children when there is no clinical evidence of fluid retention. Diazoxide responsiveness is noted when (a) feeding with normal frequency and volume (b) able to fast appropriately for age and maintain normal blood glucose levels (c) serum insulin level low or undetectable at the end of the fast (d) appropriate increase in serum fatty acids and ketone bodies at the end of the fast. However, diazoxide is generally ineffective in diffuse CHI due to inactivating mutations in *ABCC8* and *KCNJ11* and in most patients with focal CHI.

Rapid genetic analysis for mutations in ABCC8 and KCNJ11 allows identification of the majority of patients with diffuse disease (homozygous or compound heterozygous mutations in ABCC8 and KCN[11).[80] Patients with a paternal mutation in ABCC8 and KCNJ11 (or those with no mutations in these genes) potentially have focal disease. It is important to precisely locate the lesion in the pancreas preoperatively, as removal of this lesion can be curative in focal CHI. Fluorine-18-L dihydroxyphenylalanine positron emission tomography (18F-DOPA PET) offers precise pre-operative localization of the focal lesion, thus guiding the extent of surgical resection.^[8] The uptake of the positron emitting tracer 18F-DOPA is increased in β -cells with a high rate of insulin synthesis and secretion compared to unaffected areas allowing visualization of the focal lesion. The sensitivity for detecting focal lesions varies between 88 and 94% with a specificity of 100%.[81]

The treatment options for the diazoxide unresponsive infants with confirmed (genetically/by 18F-DOPA-PET

scanning) diffuse disease includes long term medical therapy with octreotide or surgery. Some of these patients respond to long term subcutaneous octreotide injections in combination with frequent feeding^[82] as the hypoglycemia gradually gets milder over time. A gastrostomy may be necessary in these patients to allow delivery of daytime bolus and overnight continuous feeds. The use of a long acting octreotide formulation has now been described in two studies.^[83,84] Tachyphylaxis has been observed on long term use of Octreotide.^[85] Recently, necrotising enterocolitis (NEC) has been described in infants treated with octreotide especially in the presence of other risk factors.^[86]

Table 3 summarizes the medical therapy for CHI.

In the severe forms of diffuse CHI patients, near total pancreatectomy may be required when all medical therapy has failed.^[87] Near-total pancreatectomy is associated with a high incidence of diabetes mellitus and pancreatic exocrine insufficiency. The surgery is traditionally carried out by open approach and is associated with peri and post-operative complications. The use of laparoscopy represents a new approach to the diagnosis and management of infants with both diffuse and focal CHI.^[88]

SUMMARY

CHI is caused by the unregulated secretion of insulin from pancreatic β -cells. The molecular basis of CHI involves

Medication	Route of administration	Dose	Mechanism of action	Side effects
Diazoxide	Oral	5-20 mg/kg/day divided into 3 doses	Opens a fully intact and functional K _{ATP} channel	Common: Fluid retention Hypertrichosis <i>Rare:</i> Hyperuricemia Eosinophilia Leukopenia Hypotension
Chlorothiazide	Oral	7-10 mg/kg/day divided into 2 doses	Used in conjunction with diazoxide for diuretic effect	Common: Hyponatremia Hypokalemia
Glucagon	SC/IV infusion for maintenance IM/IV injection for emergency	1-20 μg/kg/hour maintenance 0.5-1mg in emergency	Increases blood glucose levels by stimulating glycogenolysis and gluconeogenesis	Nausea, vomiting Paradoxical insulin secretion in high doses Skin rashes
Octreotide	SC/IV continuous infusion 6-8 hourly SC injections	5-35 µg/kg/day	Multiple actions: 1) Inhibition of insulin exocytosis 2) Stabilisation of K _{ATP} channels 3) ? Inhibiting entry of calcium into β-cells	Common: Cholelithiasis (not dose related) Tachyphylaxis Rare: Suppression of growth hormone, thyroid stimulating hormone and glucagon Diarrhoea Steatorrhoea Abdominal distension (Necrotising colitis) Growth suppression

defects in key genes (*ABCC8, KCNJ11, GLUD1, GCK, HADH, SLC16A1, HNF4A* and *UCP2*) which regulate insulin secretion. The advent of rapid genetic analysis, imaging with 18F-DOPA-PET/CT and new surgical techniques have changed the clinical approach to these complex patients.

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