

CONGENITAL HYPERINSULINEMIC HYPOGLYCEMIA IN INFANTS: GENOTYPE AND PHENOTYPE OF 102 CASES

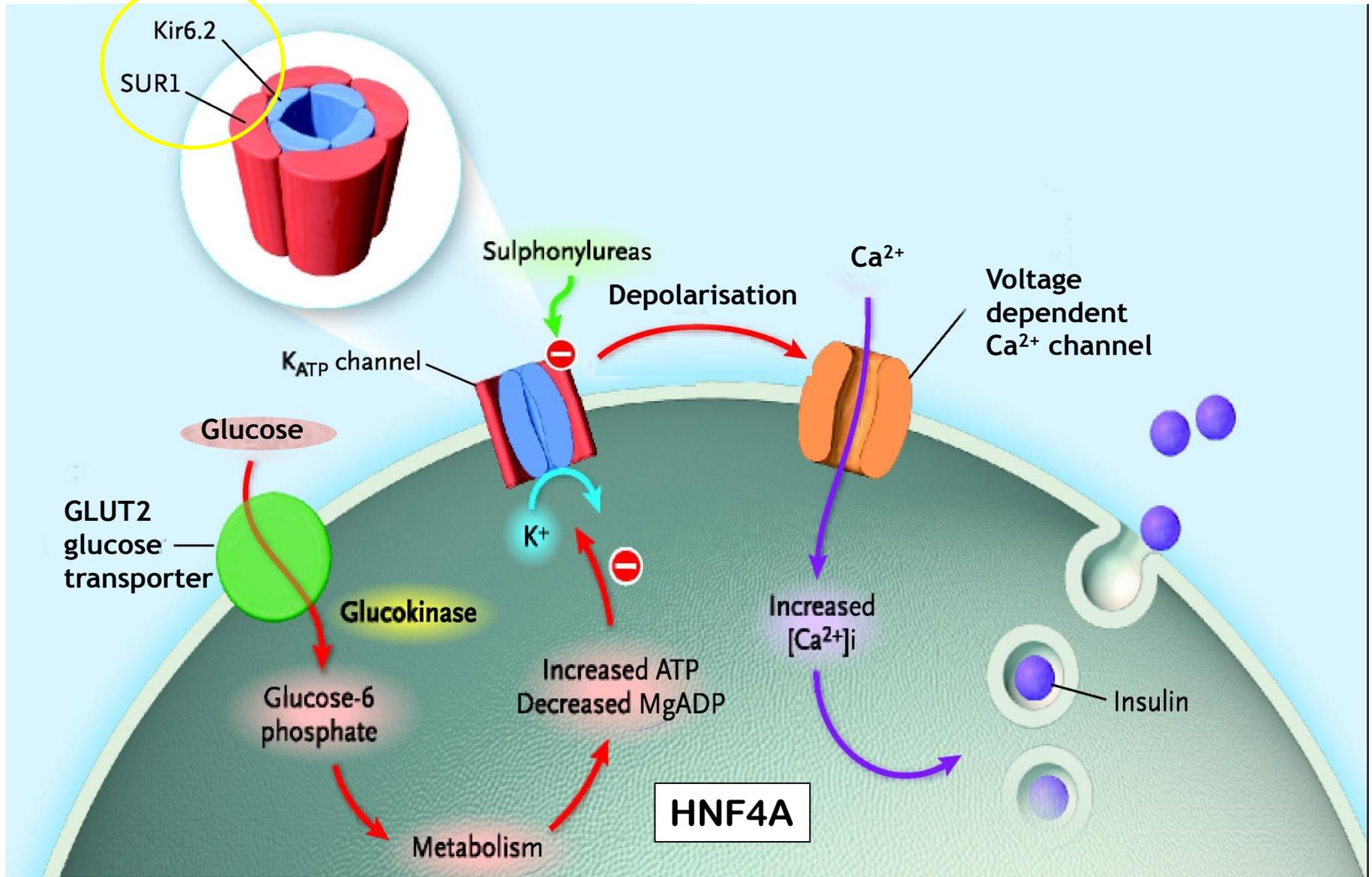
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Introduction

- Congenital hyperinsulinism (CHI): inappropriate of insulin secretion despite low blood glucose levels
- Absence of treatment → irreversible brain damage
- Incidence $1/50,000 \rightarrow 1/2,500$ live births

Insulin secretion in the pancreatic beta-cell



BACKGROUND

Summary of genetic causes of isolated HI

	Gene	Protein	Inheritance	Diazoxide-Resp.	Histology	Comment
K_{ATP} Channel	ABCC8	SUR1	AR	No	F or D	
			AD	Usually	D	
	KCNJ11	Kir6.2	AR	No	F or D	
Enzymes/Transporters	GLUD1	GDH	AD or DN	Yes	D	HIHA syndrome
	GCK	GCK	AD or DN	Usually	D	MODY 2
	HADH	SCHAD	AR	Yes	D	
	SLC16A1	MCT1	AD	Usually	D	EIHI
	UCP2	UCP2	AD	Yes	D	
Transcription Factor	HNF4A	HNF4A	AD or DN	Yes	D	MODY 1

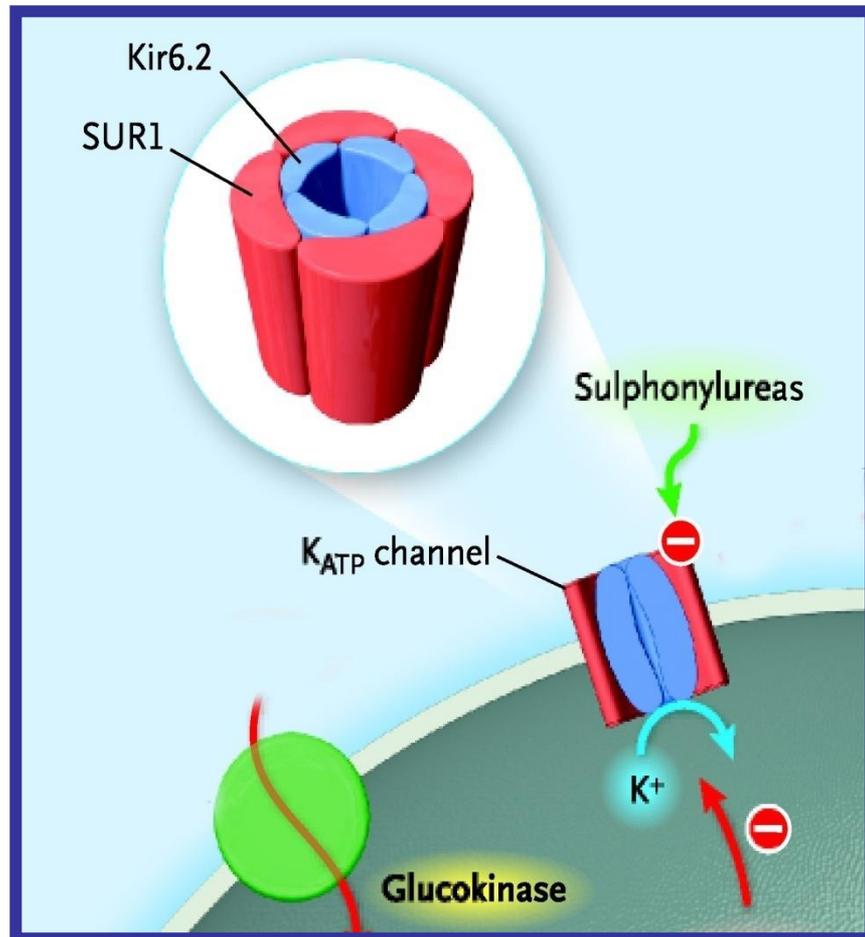
AR: autosomal recessive; **AD:** autosomal dominant; **DN:** De Novo; **F:** Focal Form; **D:** Diffuse Form; **HI/HA:** hyperammonemia/hyperinsulinism syndrome; **EIHI:** Exercise-induced hyperinsulinism; **GDH:** Glutamate Dehydrogenase; **GCK:** Glucokinase; **HADH:** Hydroxy-Acyl-CoA Dehydrogenase; **MCT1:** Monocarboxylate transporter 1; **MODY:** Maturity-onset diabetes of the young; **UCP2:** Uncoupling protein 2.

BACKGROUND

Beta-cell potassium ATP (K_{ATP}) channel genes

- *ABCC8* gene: 39 exons, 100 kb, encoding a 1582-amino acids protein (SUR1)
- *KCNJ11* gene: single exon encoding a 390-amino acid protein (Kir6.2)
- Interestingly, location of *KCNJ11* only 4.5 kb from *ABCC8* gene on 11p15.1
- *GLUD1*: 45 kb; 13 exons on 10q23.2
- *HNF4A*: ~74 kb; 10 exon on 20q13.12

Hyperinsulinism results from loss-of-function K_{ATP} channel mutations



Protein does not reach cell surface/or channels do not function



Membrane Depolarised



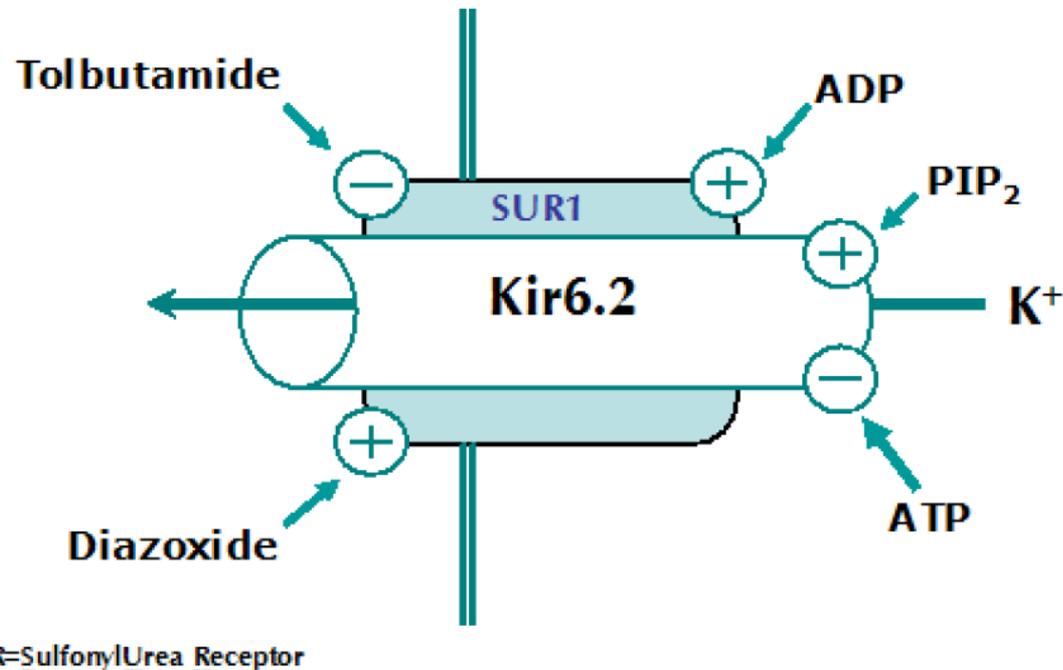
Calcium influx



Unregulated insulin secretion

= Hyperinsulinism

Control Elements for the K_{ATP} Channel in Pancreatic β -Cells, 2006



- Diazoxide blocks insulin secretion by activating (opening) SUR1
- Sulfonylureas (tolbutamide) stimulate insulin secretion by closing SUR1

SPECIFIC AIMS

- *To identify mutations in the ABCC8 and KCNJ11, HNF4A and GLUD genes*
- *To describe genotype and phenotype correlations of Vietnamese children with congenital hyperinsulinism*

PATIENTS

- Patients

102 cases with CHI at NHP (male: 60; female:42)

Diagnosis age: 1 - 30 days of age

- From Jan.2010 to Dec. 2016 at the National Children's Hospital

PATIENTS

Diagnostic criteria (Hussain K. 2008)

1. Fasting & post-prandial hypoglycemia ($< 2.5\text{--}3$ mmol/l) with unsuppressed insulin secretion & c-peptide levels (plasma insulin concentrations > 1 mU/l).
2. Positive response to subcutaneous or intramuscular administration of glucagon (plasma glucose concentration increase by 2 to 3 mmol/l following a 0.5 mg glucagon subcutaneous injection)
3. Negative ketone bodies in urine or blood
4. Prolonged dependence on treatment to prevent hypoglycemia throughout first months/years of life

PATIENTS

Excluded criteria

- Syndromic: e.g.
 - ✓ Beckwith-Wiedemann
 - ✓ Trisomy 13
 - ✓ Mosaic Turner
- Metabolic conditions
- Secondary to (usually transient)
 - ✓ Maternal diabetes mellitus (gestational & insulin dependent)
 - ✓ Intra-uterine growth retardation
 - ✓ Perinatal asphyxia

METHODS

- Genomic DNA was extracted from peripheral leukocytes using standard procedures.
- Single exon of *KCNJ11*; 39 exons of *ABCC8*; 10 exons of *HNF4A* & 13 exons of *GLUD1* were amplified & sequenced.
- Sequencing reactions were analyzed on an ABI 3730 capillary sequencer & were compared to published sequences using Mutation Surveyor version 3.24.

Ellard S et al. Am J Hum Genet 2007: 81: 375-382.

Flanagan SE, et al. Diabetologia 2006: 49: 1190-1197.

Congenital hyperinsulinism

Table 1

Infusion of glucose.

Peripheral catheter: glucose 10%
2 ml/kg/h (= 3.3 mg/kg/min)
4 ml/kg/h (= 6.7 mg/kg/min)
6 ml/kg/h (= 10 mg/kg/min)
8 ml/kg/h (= 13.3 mg/kg/min)
Central catheter: glucose 10%, 20%, 30% or 50%
e.g. Glucose 30%
0.5 ml/kg/h (= 2.5 mg/kg/min)
1 ml/kg/h (= 5 mg/kg/min)
2 ml/kg/h (= 10 mg/kg/min)
3 ml/kg/h (= 15 mg/kg/min)

A specialized team approach to diagnosis and medical versus surgical treatment of infants with congenital hyperinsulinism

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Table 1 Timeline for diagnosing HI, initiating medical therapy, and referring to specialized center

Day 1	Establish diagnosis of HI (see Table 3)
	Begin 5-d trial of diazoxide If HI is severe begin at max dose (15 mg/kg/d) If HI less severe/perinatal-stress, start diazoxide at 5-10 mg/kg/d*
Day 2-5	Consider starting a diuretic with diazoxide, especially if on high GIR Determine minimum GIR required to maintain blood glucose between 70 and 100 If HI is severe or GIR is >10 mg/kg/min, send mutation analysis on HI genes for infant and parents
Day 6	Determine fasting tolerance on diazoxide Failure to fast >12 h with BS >70 mg/dL indicates diazoxide unresponsiveness Diazoxide failure suggests a K_{ATP} channel HI and potential surgical candidate Begin arrangements for transfer to a specialized HI center with ^{18}F -DOPA PET scan capability
Day 7	Discontinue diazoxide; consider octreotide, 5 $\mu\text{g kg}^{-1} \text{d}^{-1}$ divided every 6-8 h Desensitization to octreotide is common after 2-3 doses If required, octreotide can be increased to maximum of 15 $\mu\text{g/kg/d}$
Day 8-14	Evaluate effectiveness of octreotide with fasting test while awaiting transfer of patient

Abbreviations: GIR, glucose infusion rate (mg/kg/min); HI, hyperinsulinism.

*See text for further discussion of tachyphylaxis.

METHODS

- Definition of diazoxide efficiency: normalization of glycemia > 3 mmol/l measured before & after each meal in patients fed normally with a physiological overnight fast, after stopping intravenous glucose & any other medications for at least five consecutive days

Arnoux JB et al. Early Human Development 2010;86:287–294

- Non responsive with diazoxide
 - Surgery
 - Octreotide

RESULTS

CLINICAL SYMPTOMS

- ❖ Weight of birth: 4.1 ± 0.9 (2.3 – 5.6) kg
- ❖ Age at presentation: < 24 hours: 47/102 (46.1%)
- ❖ Symptoms:
 - ✓ Poor feeding, lethargy: 89/102 (87.3%)
 - ✓ Seizures 14/102 (13.7%)
 - ✓ Apnea, cyanosis 9/102 (8.8%)
- ❖ Glucose infusion rate: 12 – 28 mg/kg/mn

RESULTS

Distribution of mutations in different genes

Gene	Number of patients	%
<i>ABCC8</i>	47	46.1
<i>KCNJ11</i>	5	4.9
<i>HNF4A</i>	1	0.9
<i>GLUD1</i>	0	0
Total	53	51.9

HNF4A: **c.659T>C** (p.L220P): novel mutation
& mother inheritance

RESULTS

Mutations in *ABCC8*

- 25 different mutations: 13 novel; 12 reported one in *ABCC8*
- Homozygous/compound heterozygous mutations in *ABCC8*
27/47 (57.4%)
- Hemizygous mutations in *ABCC8* from father or mother
20/27 (42,6%)

RESULTS

Mutations in *ABCC8* and genotype

Genotype with <i>ABCC8</i> mutations	Number of families
c.3403-1G>A	13
c.3403-1G>A/c.3403-1G>A	1
c.3403-1G>A/c.2995C>T	1
c.2057T>C	2
c.2057T>C/c.2057T>C	1
c.2417G>A/c.2995C>T	1
c.4160_4162del	2
c.1467+5G>A/c.2800C>T	1
c.2041-21G>A	1

RESULTS

Mutations in *ABCC8* and genotype

Genotype with <i>ABCC8</i> mutations	Number of families
c.2041-21G>A/c.3978del	1
c.2041-21G>A/c.2041-21G>A	1
c.2056T>A/c.2057T>C	1
c.2057T>C/c.3403-1G>A	2
c.2057T>C/c.2995C>T	1
c.2995C>T	3
c.3293A>G	1
c.3403-1G>A/c.4462C>T	1
c.4415-13G>A	1

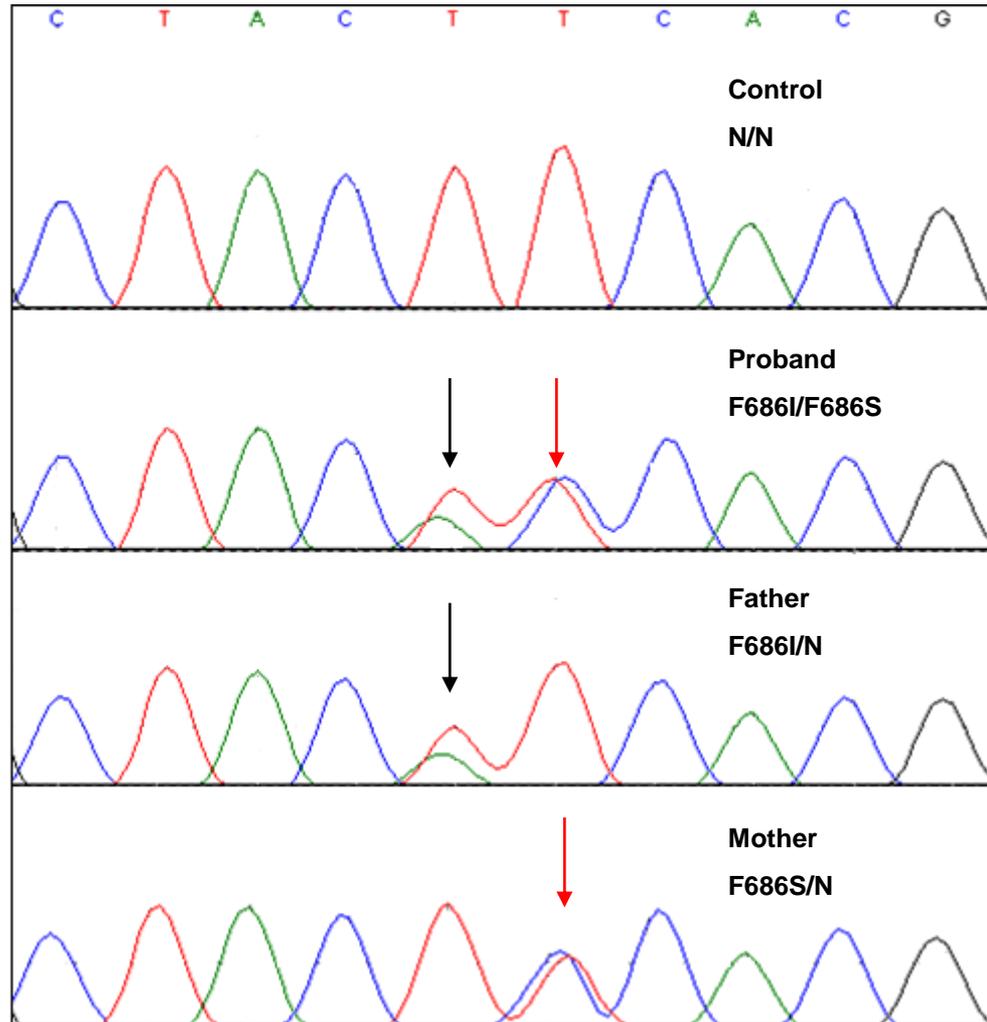
RESULTS

Mutations in *ABCC8* and genotype

Genotype with <i>ABCC8</i> mutations	Number of families
c.4610C>T	1
c.655C>A/c.892C>T	2
c.1106A>G/ c.4611G>A	1
c.1183A>T	1
c.2056T>A/c.2057T>A	1
c.3293A>G	1
c.4061A>G *	1
c.4135G>A	1
Deletion of exons 22-23	1

RESULTS

Sequencing of *ABCC8*



RESULTS

Mutations in *KCNJ11*

- 3 novel mutations from father (**c.482C>T**, **c.512C>A**, **c.820G>C**) in 2 unrelated families
- Homozygous **c.185delC** of *KCNJ11* in two sibling of 1 family.

RESULTS

Correlation of genotype - phenotype

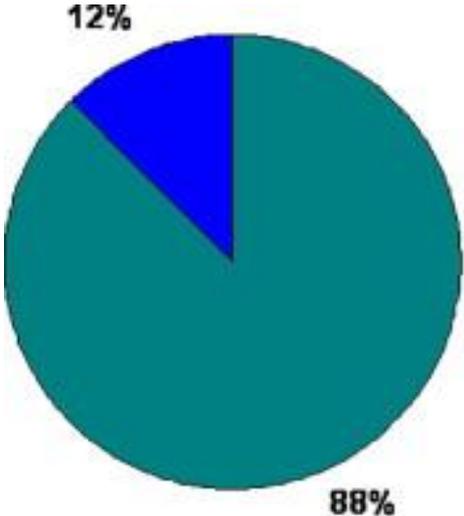
- ❖ Responsive with diazoxide: 52 cases:
 - 49 without mutations
 - 1 case with maternal mutation in *ABCC8*
 - 1 case with mutation in *HNF4A*
 - 1 case with mutation in *KCNJ11*

Kết quả

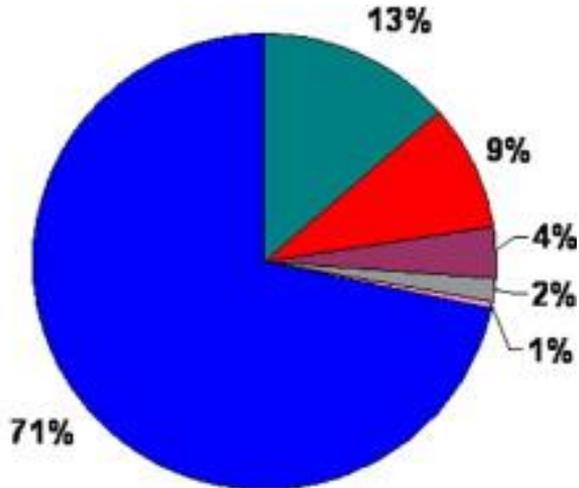
Correlation of genotype - phenotype

- ❖ **Non responsive with diazoxide** (surgery and/or octreotide): 48 cases
 - 4 cases with mutations in *KCNJ11*
 - 44 cases with homozygous/compound heterozygous or paternal mutations in *ABCC8*

DISCUSSION



Diazoxide Unresponsive cohort (n= 105)



Diazoxide Responsive cohort (n= 183)



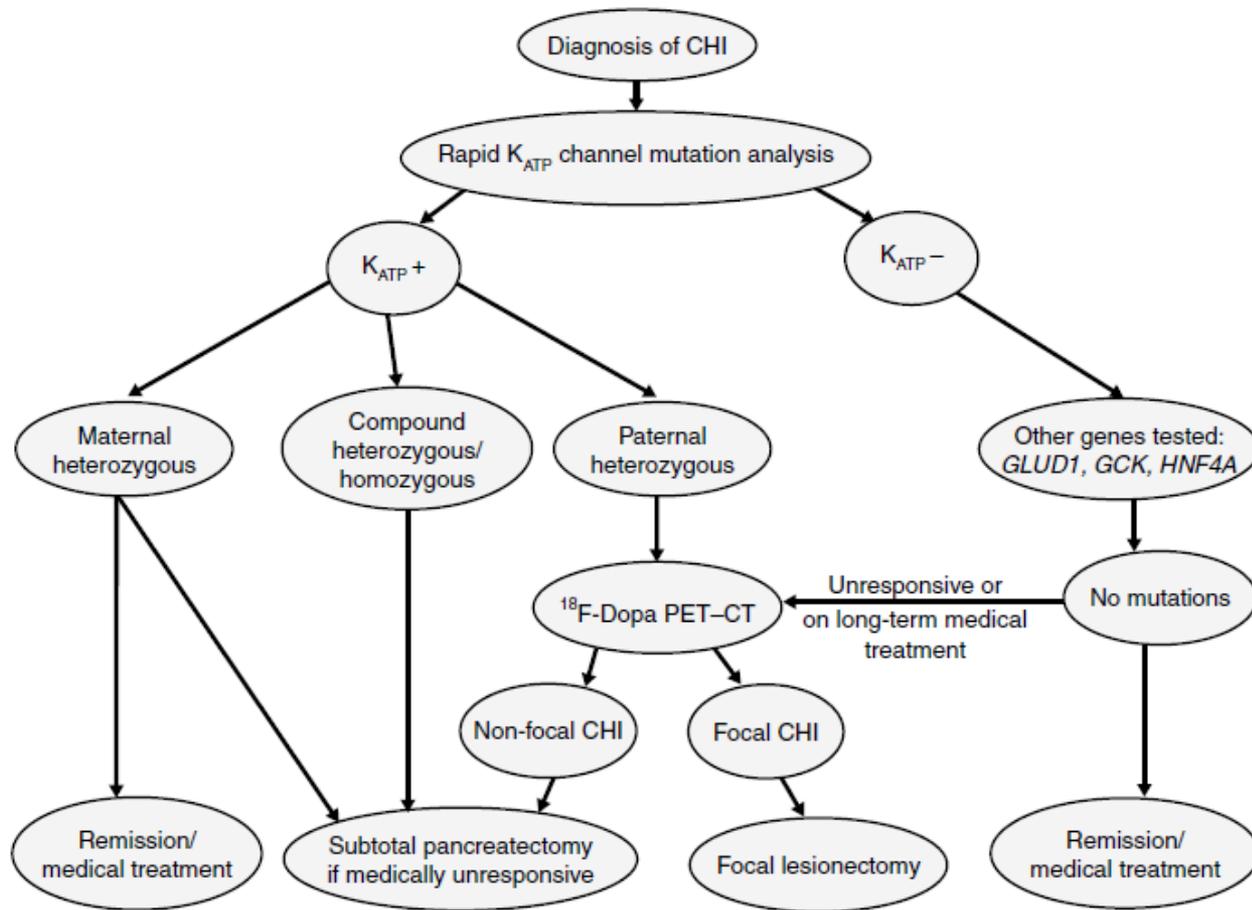
DISCUSSION

- Mutation in ABCC8 (SUR1): most common cause of CHI and were first to be described
- Approximately 45% of affected individuals have mutations in *ABCC8* [[Nestorowicz et al 1998](#), [Aguilar-Bryan & Bryan 1999](#), [Meissner et al 1999](#), [Fournet & Junien 2003](#), [Tornovsky et al 2004](#)].
- Almost 20 years after discovery of first mutation
- Over 200 mutations identified
- Distribution of mutations throughout the gene

DISCUSSION

- Diazoxide is effective in virtually all forms of CHI except in inactivating recessive mutations in *ABCC8*
- Rapid genetic analysis for mutations in *ABCC8* & *KCNJ11* → identification of majority of patients with diffuse disease (homozygous or compound heterozygous mutations)

Kapoor RR et al. Arch Dis Child 2009;94:450-457



Flowchart of investigation and management of children with CHI.

CONCLUSIONS

- Understanding genetic basis of CHI provide novel insights into β -cell physiology
- Prediction phenotype, management & genetic counseling
 - Genetic analysis for mutation in CHI can help in genetic diagnosis → treatment
 - Prenatal diagnosis of CHI → immediate medical management at the time of birth



**Lê Thiện N, WOB 5 kg (39 weeks)
responsive with medical treatment**



**Nguyen Thi Diem H. Responsive with medical treatment. WOB
3.5 kg (37 weeks). Two sibling died at Provincial Hospital at 3
days of age (cyanosis)**



Vũ Hải Y. WOB 5.4 kg, responsive with medical treatment



Vuong Ha M; WOB 3.8 kg
Unresponsive with medical treatment
Mutation of ABCC8: (F686I/F686S)





**Cao Bao N. WOB 5 kg;
Unresponsive with medical treatment
Mutation of ABCC8 F686S/IVS27-1G>A**

Thank you very much!