Congenital hyperinsulinism

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Wipple's triad

- Glucopenic symptoms
 - Drowsiness
 - Fainting
 - Seizures
 - Hallucinations
 - Any kind of neurological symptoms
- Glycemia <3,0 mmol/l (55mg/dl)
- Resolution of all symptoms after normalization of glycemia









Source: Expert Opinion on Orphan Drugs (2014) 2(8): 779-795

Biological diagnosis of HI

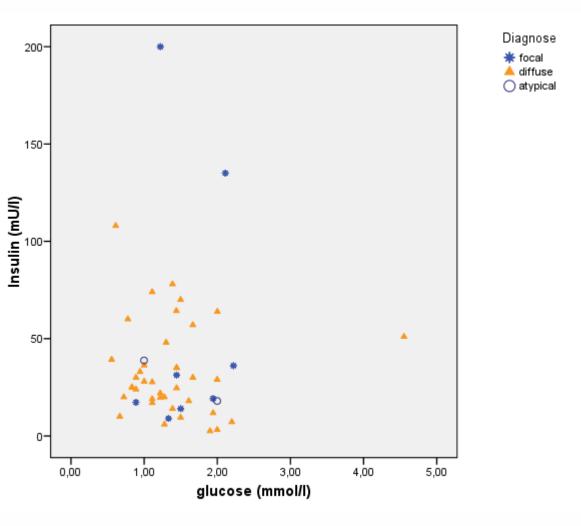
- Fasting and/or postpandrial hypoketotic hypoglycemia (<2,5 mmol/l)
- Inappropriate plasma insulin levels and c-peptide at time of hypoglycemia (in some cases, insulinemia can be seemingly null)
- Absent or low blood/urines ketone bodies and non-esterified fatty acids

JB

- An increase in blood glucose <1,7 mmol/l within 30 min after administration of 1 mg glucagon
- Need of high glucose infusion rate to keep blood glucose >3,0 mmol/l (>8 mg/kg/min)

Source: Expert Opinion on Orphan Drugs (2014) 2(8): 779-795

Insulin/ glucose at diagnosis





JB

Syndromic CHI

Overgrowth syndrome

Beckwith-Wiedemann AD or S11p15.5

Perlman AR ?

Sotos S NSD1

Developmental syndrome

Kabuki syndrome AD or SMLL2

CDG syndrome AR PMM2 / PMI

Costello AD or SHRAS

Turner S Monosomy X



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- Beckwith-Wiedemann syndrome
 - » Macrosomia, ear lobe creased, macroglossia, umbilical hernia, hemihypertrophia, heart defects...









 Congenital disorder of glycosylation (CDG) syndrome type la&b



» strabism, progressive cerebellar atrophy, inverted nipple, abnormal fat distribution, lipoatrophy areas, abnormal coagulation factors...



Kabuki make-up syndrome

• elongated palpebral fissures with eversion of the lateral third of the lower eyelid, skeletal anomalies (costal, vertebral or hips), fetal fingertip pads, failure to thrive, congenital heart defects, hormones deficiencies...



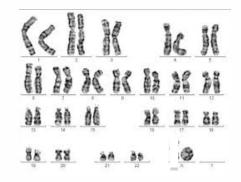




Turner syndrome

SGA, pterygium coli, heart, aortic, kidney malformation, growth delay...







Transient CHI

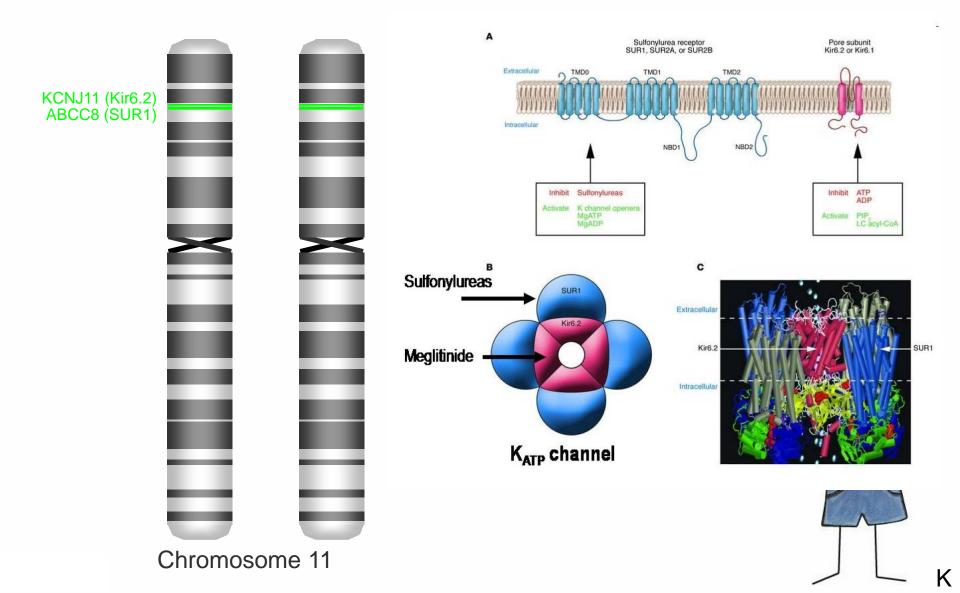
- Newborns from diabetic mothers
- Small for gestational age babies (SGA)
- Perinatal stress
 - Fetal distress
 - Asphyxia at birth

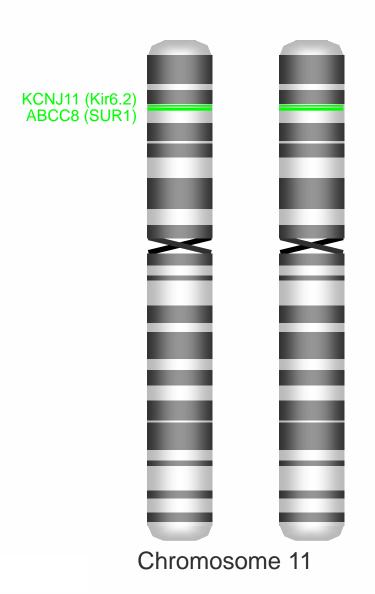
Isolated CHI

- Channelopathie: ABCC8, KCNJ11,
- Metabolopathie: GLUD1, GCK, UCP2, HADH, SLC16A1
- HNF1A, HNF4A



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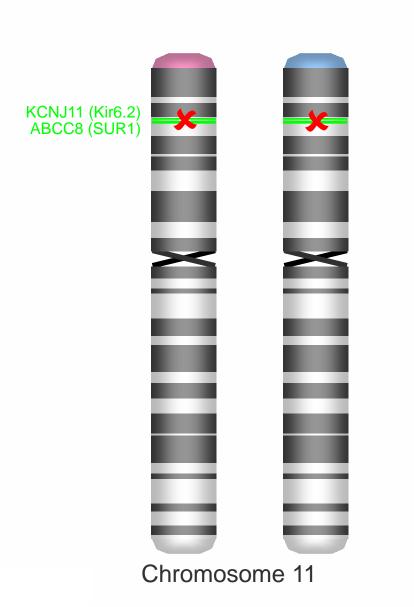




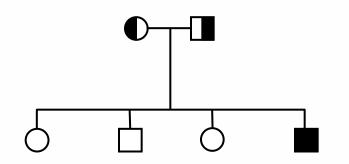
Wiedemann-Beckwith-Syndrome

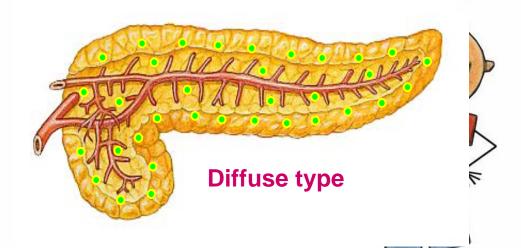


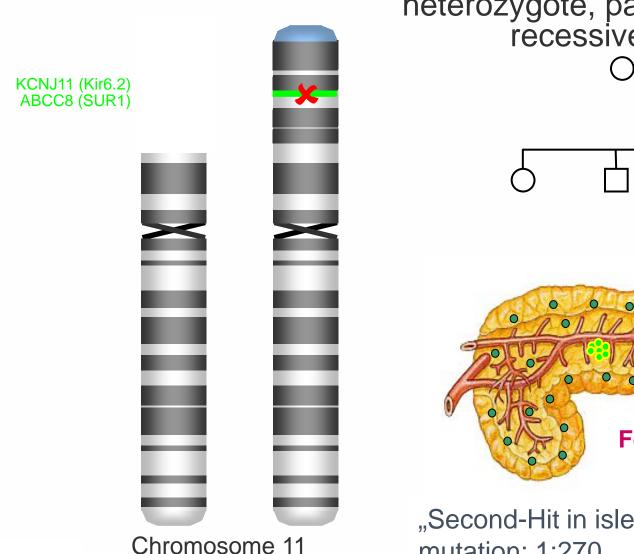
Silver-Russell-Syndrome



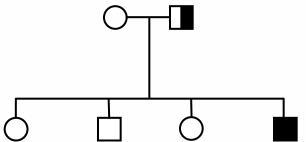
Autosomal-recessive

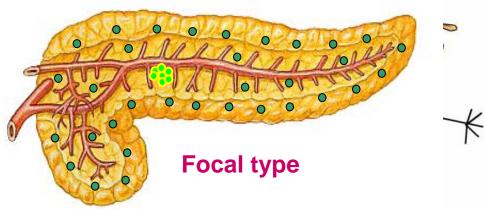






heterozygote, paternally inherited, recessive mutation





"Second-Hit in islets by paternally inherited mutation: 1:270

PET-scanning and CT-scan in one single device.

For simultaneous registration and fusion of 2 signals

- 1. tracer (e.g. F18-L-DOPA) by PET and
- 2. anatomical image by CT.





L-DOPA is a

transmitter substance in the nervous system.

precursor of catecholamines (= noradrenaline and adrenaline)

Neuroendocrine cells

take up and decarboxylate amine precursors,

e.g. L-DOPA and 5-hydroxytryptophan

store biogenic amines (= dopamine and serotonin)



L-DOPA

Pancreatic cells contain markers of neuroendocrine cells,

such as tyrosine hydroxylase, dopamine,

neuronal and vesicular dopamine transporter,

monoamine oxidases A and B.

Pancreatic islets take up L-DOPA and convert it to dopamine through

the Aromatic Amino acid Decarboxylase

AAAD.



- 1. No moving of patients necessary
- 2. No transmission radiation
- 3. Quick scanning times (< 4 min.)
- 4. Software allows:
 Alignment of PET and CT datasets measurement of distances
- 5. Data acquisition up to 5* repeated
- 6. multiple time point imaging: from 20 to 60 min p.i.



- 1. Between 2004 and 2015, a [18F]F-DOPA-PET/CT was performed in 200 patients.
- 2. Analyses of 150 patients (67 girls, 83 boys) (median age: 0.53y; range: 0.09-30.35y.).
- 3. Mutation analysis of ABCC8 and KCNJ11 were carried out in the index patient and their parents.
- 4. Pancreatic surgery was done in focal form

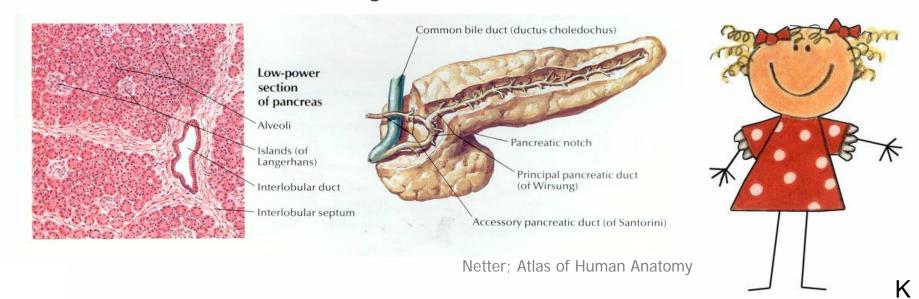


Pancreas

located behind the stomach between the spleen and duodenum

Islets of Langerhans

- α cells glucagon
- β cells insulin
- ∆ cells somatostatin/gastrin



Focal form



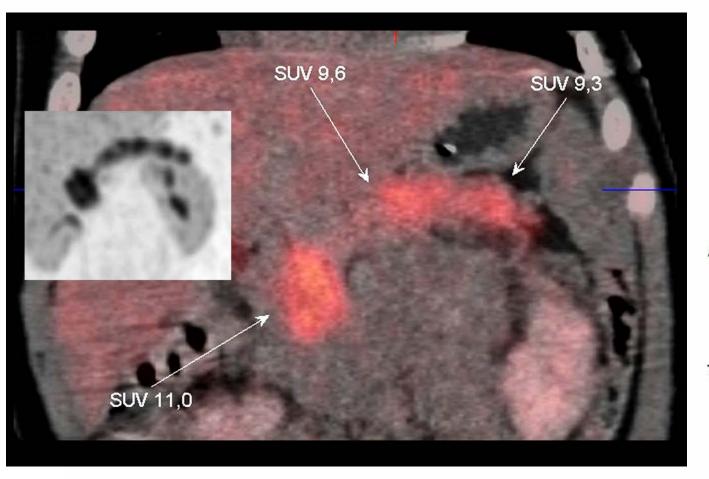


Focal form



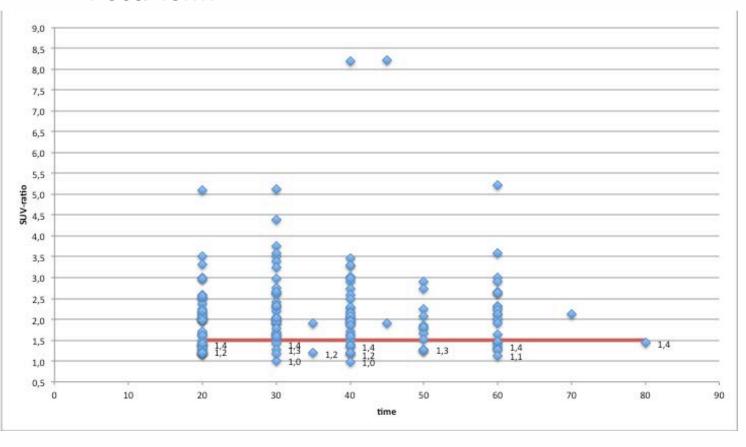


Diffuse form



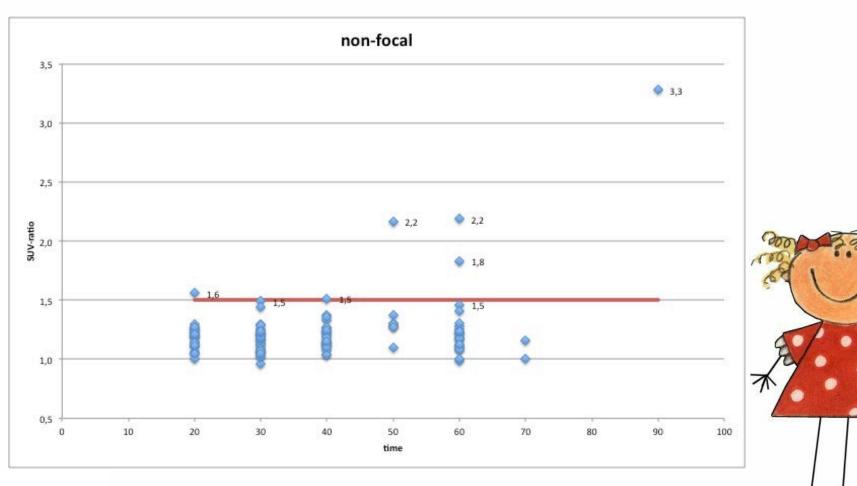


Focal form



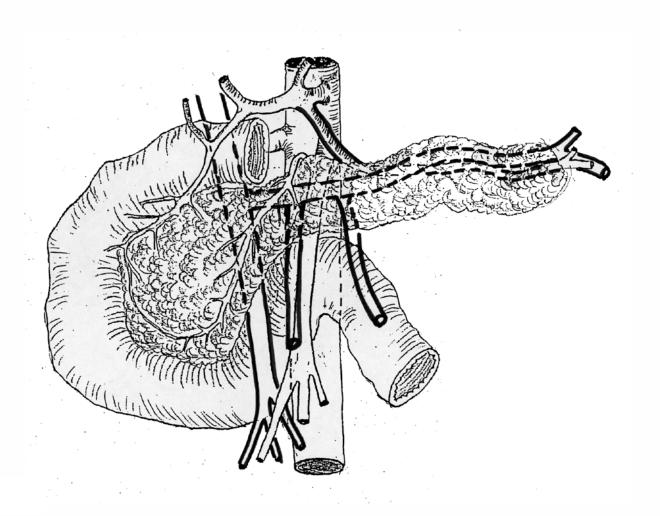


non-focal form





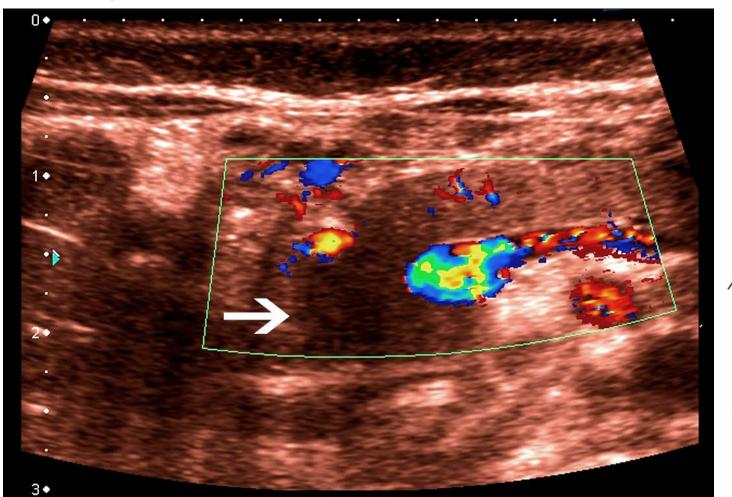
Localisation of important landmarks





Intra operative diagnosis

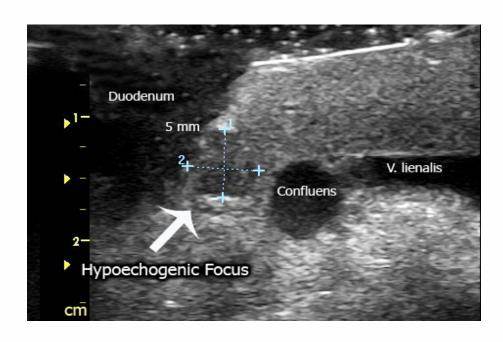
Pre-operative ultra-sound

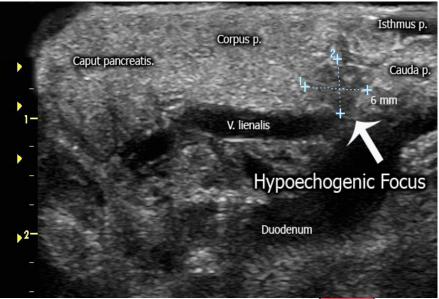




Intra operative diagnosis

Intra operative ultra-sound

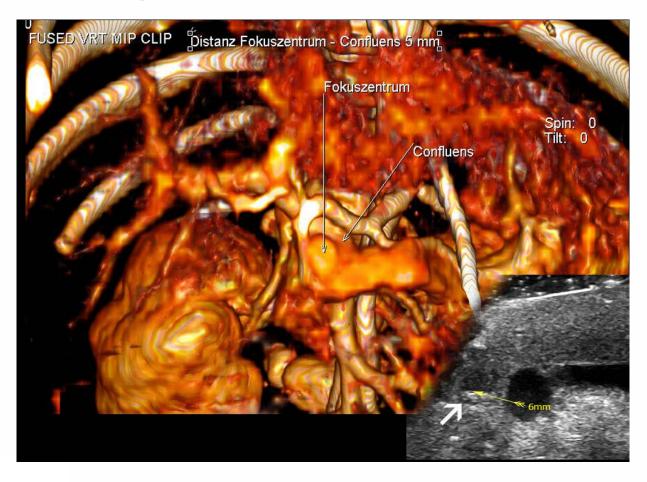






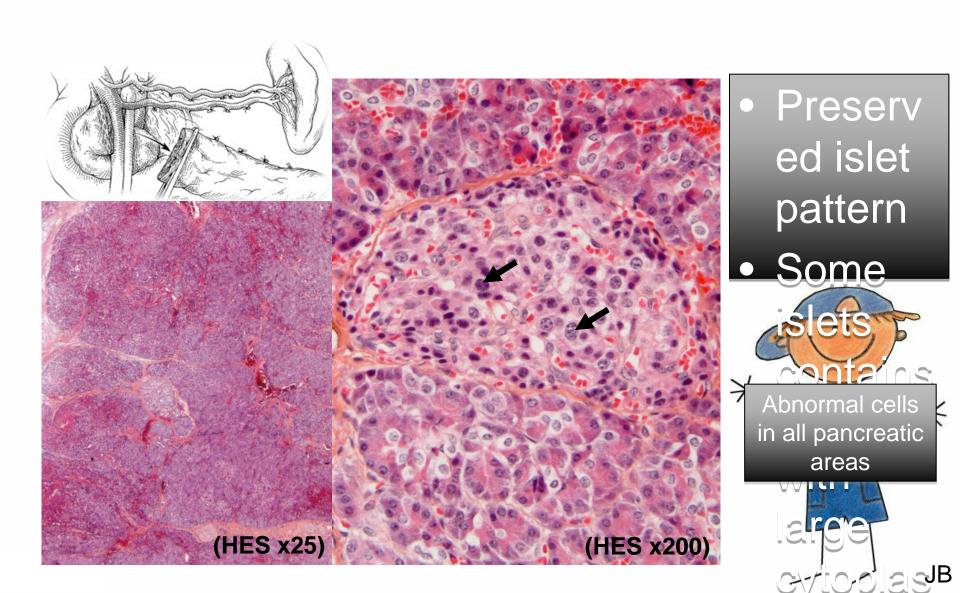
Intra operative diagnosis

Intra operative ultra-sound



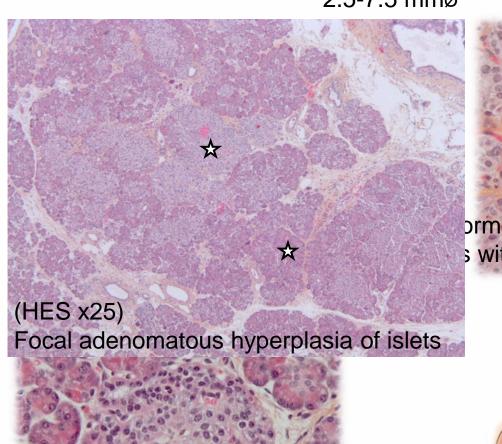


Pathology / CHI diffuse

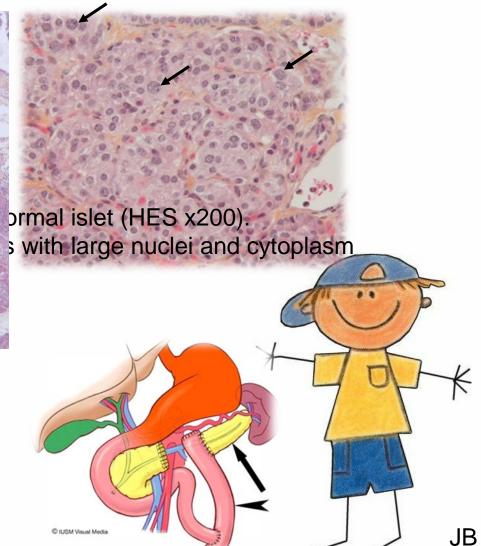


Pathology / CHI focal

2.5-7.5 mmø

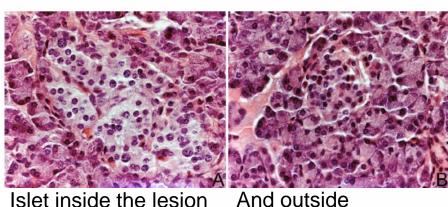


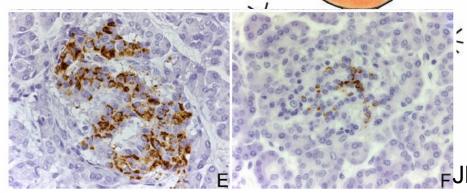
Normal islet outside the focal lesion. Shrunken ß cells (HES x200)



Pathology / CHI atypical

- Mosaicism of the pancreatic islets (+/- 5% of severe CHI patients)
 - Described in two series 7/282 [Snider JCEM 2013] and 16/217 [Sempoux JCEM 2011]
- Pathology:
 - Normal pancreatic architecture
 - Large ß-cells & islets in several adjacent lobules
 - Outside of the lesion:
 ß-cells normal or at rest.





and outside the lesion ProInsulin staining in

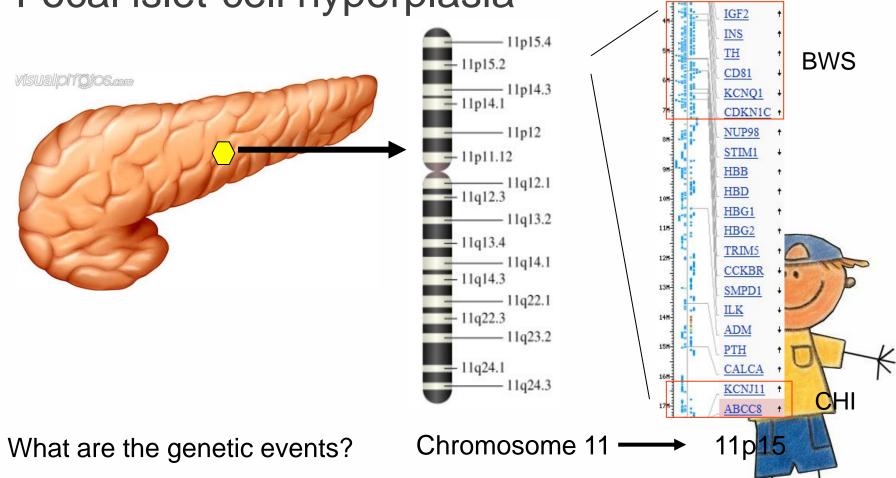
Pathology / CHI atypical

- Distinct clinical course compared to DZX-unresponsive K_{ATP} CHI
 - Normal birth weight
 - Late median age at presentation (150-165 days)
 - Can be cured by partial pancreatecomy
 - Increased incretin secretion after oral glucose load in some pts? [Shy J Pediatr 2013]
 - Undue expression of HK1 in the lesion (5 pts) [Henquin Diabetes 2013]
- Cause:
 - Somatic GCK mutation (1 pt/6 [Henquin Diabetes 2013])
 - Unknown



Post-operative diagnosis

Focal islet-cell hyperplasia



Post-operative diagnosis

Patient no 6:

Age at surgery: 6 months

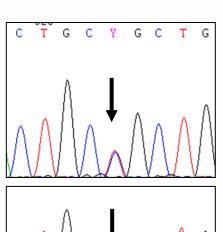
ABCC8 mutation:

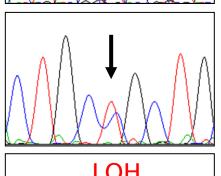
c.4241C>T, p.P1414L

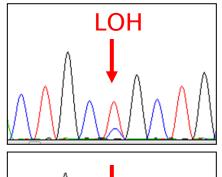
Bi-allelic expression of both parental alleles

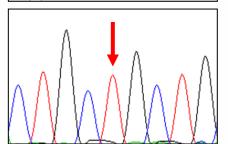
Loss of heterozygosity

Monoallelic expression of paternal mutant allele









gDNA normal pancreatic tissue

cDNA normal pancreatic tissue

gDNA focal lesion

cDNA focal lesion



Post-operative diagnosis

Summary: pUPD and expression of focal CHI

| Patient. No. | Exon | Mutation | | Observed | Age at | mRNA | LOH | Paternal |
|--------------|------|-----------------------|------------------------|-------------------------------|---------------------|-------------------------------|-----|----------|
| | | Nucleotide | Protein | freq.* [Ref.] | surgery (months) | expression | | UPD11p15 |
| ABCC8 | | | | | | | | |
| 1 | 1 | c.50T>C | p.V17A | 2* [11] | 10 | monoallelic mutant | ++ | ++ |
| 2 | 10 | c.1530G>T | p.K510N | 1 [11] | 10 | monoallelic mutant | ++ | ++ |
| 3 | 12 | c.1792C>T | p.(R598*) | Multiple [CM050968] | 7 | no (NMD) | ++ | ++ |
| 4 | 22 | c.2560- ?_2697+? | p.(D854_W8 99del46) | 2 [CG107114] | 8 | monoallelic r.2560_2697del | ROH | n.d. |
| 5 | 34 | c.4162_4164d eITTC | p.F1388del | Multiple [CD962164] | 9 | monoallelic mutant | ++ | ++ |
| 6 | 35 | c.4241C>T | p.P1414L | Multiple [CM068331] | 6 | monoallelic mutant | ++ | ++ |
| 7 | 35 | c.4259C>T | p.S1420L | 1 [Barthlen et al, submitted] | 2 | monallelic mutant | + | + |
| KCNJ11 | | | | | | | | 1 |
| 8 | 1 | c.286G>A | p.A96T | 1* [11] | 2 | mutant/wt 75%/25% | + | + |
| 9 | 1 | c.612C>A | p.D204E | 2 [CM083531] | 2 | monoallelic mutant | ++ | ++ |
| 10 | 1 | c.844G>A | p.E282K | 3 [CM071810] | 17 | monoallelic mutant | ++ | ++ |
| 11 | 1 | c.901C>G | p.R301G | Multiple [CM088147] | 6 | monoallelic mutant | (+) | + |
| | | | | | | | | |

Questions?



