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## Case

- A term male (birthweight 3.7kg, 80<sup>th</sup> percentile) with diazoxide-unresponsive congenital hyperinsulinism (CHI) was born to unaffected parents
  - natural conception
  - non-consanguineous and no family history
- He did **not** have the **cardinal** Beckwith-Wiedemann spectrum<sup>1</sup> features of
  - macroglossia, exomphalos or lateralised overgrowth
- He did **not** have the **suggestive** Beckwith-Wiedemann spectrum<sup>1</sup> features of
  - polyhydramnios, macrosomia
  - facial naevus simplex, ear creases or pits
  - umbilical hernia, diastasis recti
  - nephromegaly or hepatomegaly
- Placenta was not retained to assess for size or mesenchymal dysplasia<sup>1</sup>
- A targeted massively parallel sequencing (MPS) panel identified a heterozygous maternally inherited K<sub>ATP</sub> channel *ABCC8* variant (c.1332+4del) - minimal splicing effect predicted
  - classified as likely benign
- Intensive medical support was required and he could not be medically maintained with a trial of continuous subcutaneous octreotide
- [18F]-DOPA PET/CT imaging of the pancreas
  - unexpected finding of focal increased uptake in the pancreatic distal body/tail junction (**Figure 1A**)
- Histopathology of the subtotal pancreatectomy (day 22) showed (**Figure 2**)
  - focal adenomatous hyperplasia
    - trabeculae and islet nests composed of regular, oval or columnar cells
    - lacking atypia or conspicuous nuclear enlargement
  - adjacent lobules had a relatively normal distribution of islets and exocrine acini
  - chromogranin highlighted the islets
  - aberrant p57 expression in islet cytoplasm
    - nuclear in normal islets and in diffuse hyperinsulinism
    - would be negative in focal CHI due to a pathogenic paternal K<sub>ATP</sub> channel variant
- Within 2 weeks, medical support was again required with residual, increased [18F]-DOPA pancreatic uptake (**Figure 1B**)
- A second resection (5% left in-situ) (day 36) achieved normoglycaemia
- At 24 months of age
  - normoglycaemic with age-appropriate feeding (exocrine pancreatic supplements)
  - normal ultrasonographic appearance of liver and kidneys
  - normal neurodevelopmental progress

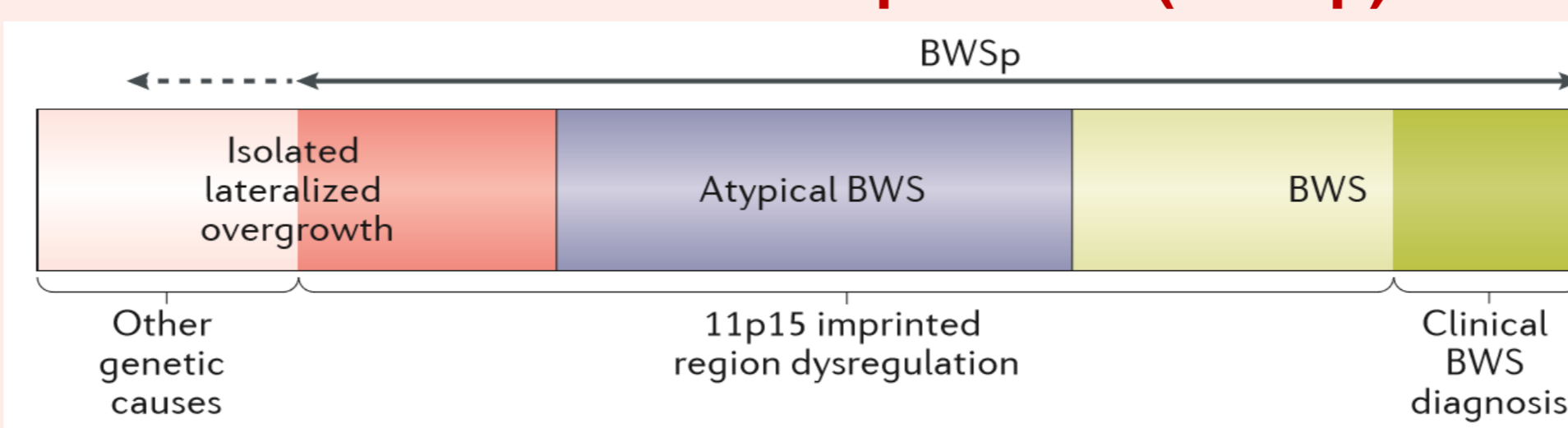
## Objectives

- Extended genetic analyses in the context of
  - Congenital Hyperinsulinism
  - focal increased [18F]-DOPA PET/CT pancreatic uptake and
  - atypical histology

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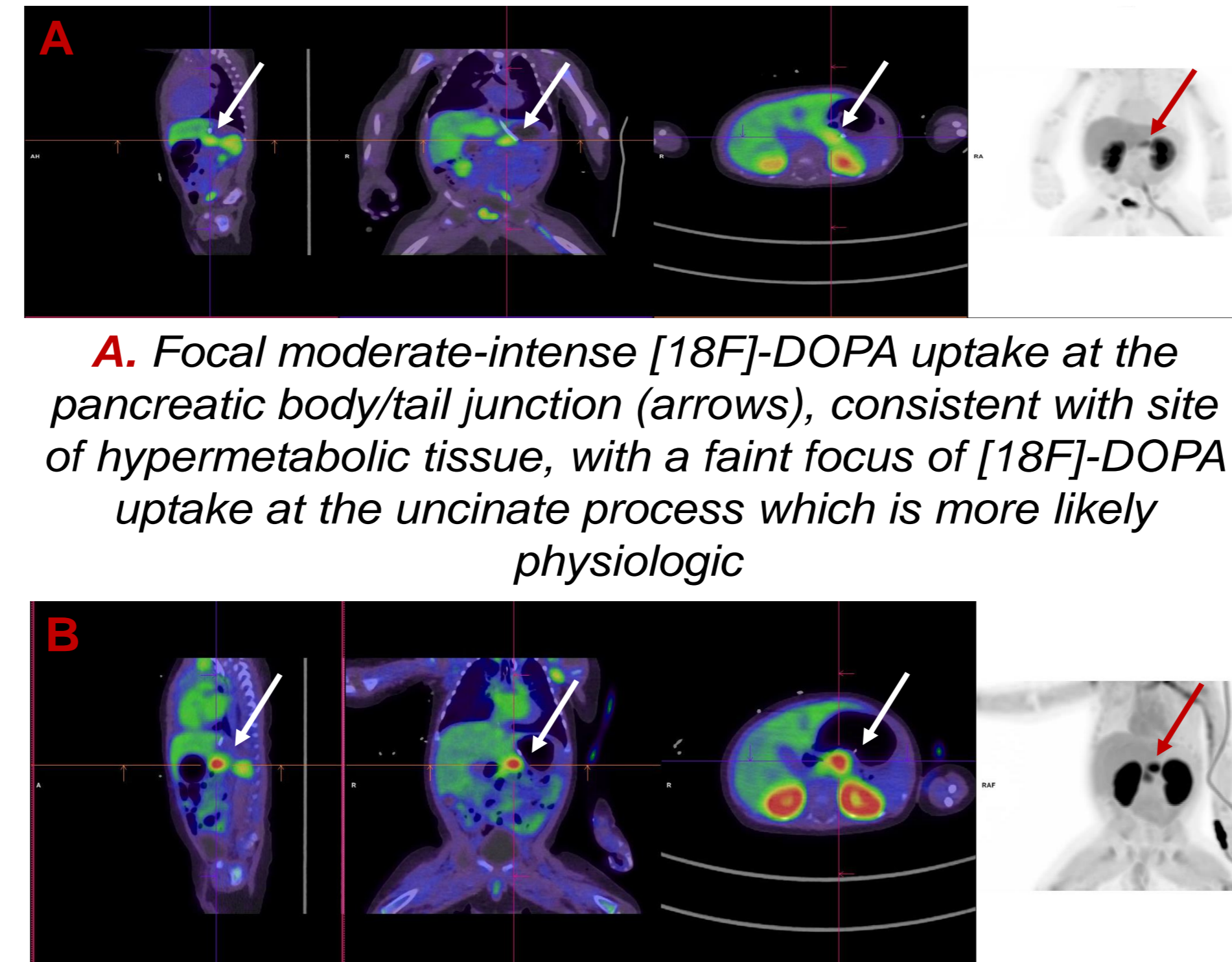
## Beckwith-Wiedemann syndrome (BWS)

- A multisystem human genomic imprinting disorder with variable clinical expression and complex molecular aetiology<sup>1</sup>
- An international consensus statement has introduced the concept of **Beckwith-Wiedemann spectrum (BWSp)**<sup>1</sup>



- Hyperinsulinaemic hypoglycaemia is common (30-60%) and usually resolves within a few days
  - persistent, severe cases refractory to medical management are usually associated with the paternal uniparental disomy (pUPD11) molecular defect
    - majority do not have a paternal inactivating K<sub>ATP</sub> channel variant but those that do have even more refractory hypoglycaemia

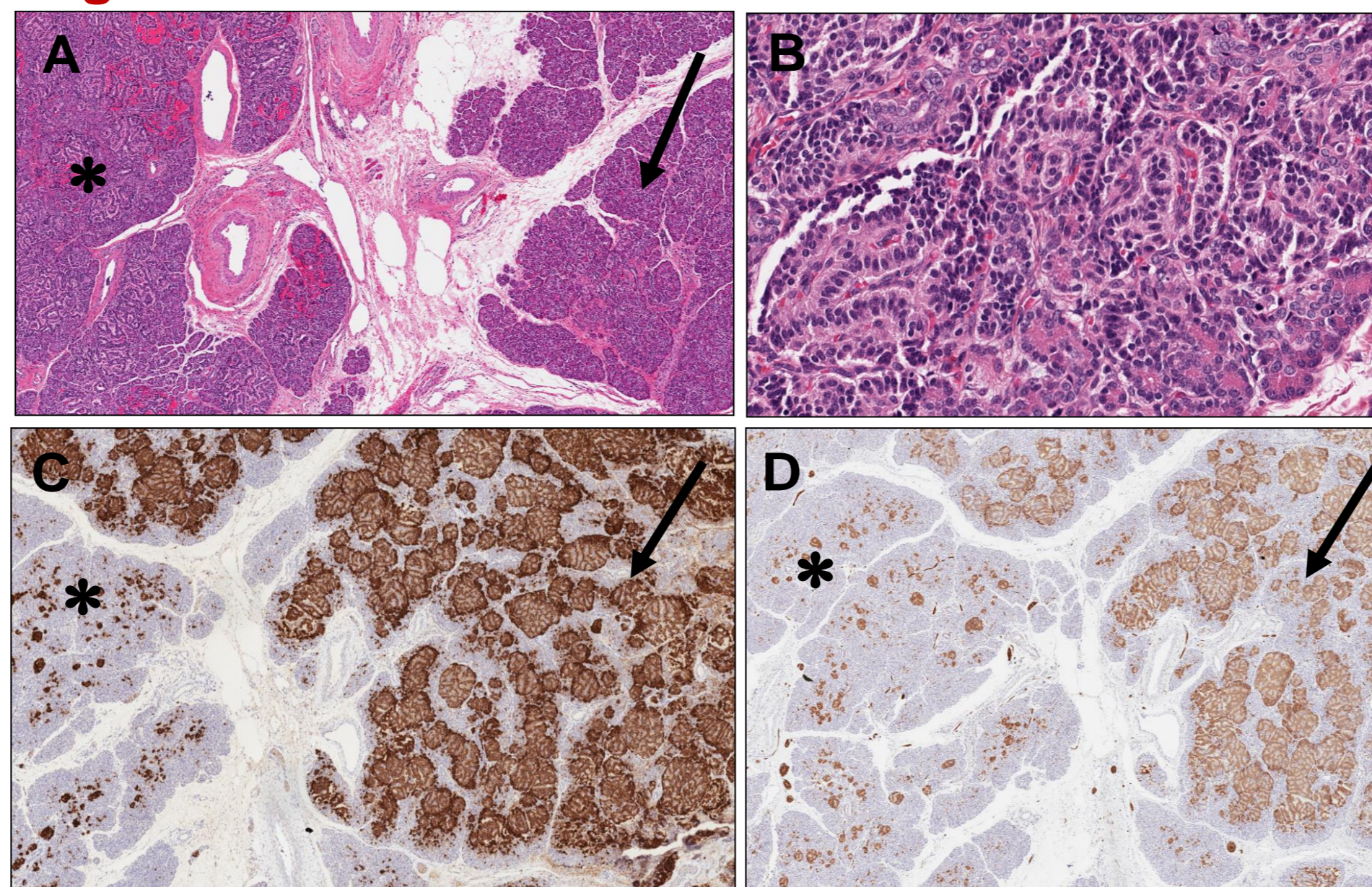
**Figure 1**



**A.** Focal moderate-intense [18F]-DOPA uptake at the pancreatic body/tail junction (arrows), consistent with site of hypermetabolic tissue, with a faint focus of [18F]-DOPA uptake at the uncinate process which is more likely physiologic

**B.** Focal moderate-intense [18F]-DOPA uptake at the subtotal pancreatectomy resection margin (arrows), suggesting that significant residual hypermetabolic tissue had not been resected

**Figure 2**



**A.** Focal adenomatous hyperplasia (arrow), normal pancreas (\*) H&E, 10x

**B.** Islet hyperplasia, H&E, 20x

**C.** Chromogranin expression in hyperplastic islets (arrow), normal islets (\*), 4x

**D.** p57 expressed in hyperplastic islets (arrow) and normal islets (\*), 4x

### References

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## Methods

- Pancreas (region of islet hyperplasia)
  - Targeted MPS hyperinsulinism panel with mosaic variant calling programme on the sequence data (detects variants to level of 1%)
    - KCNJ11, ABCC8, AKT2, GLUD1, GCK, GPC3, HADH, HNF4A, INSR, KDM6A, KMT2D, SLC16A1, CACNA1D, PMM2, TRMT10A, HNF1A*
  - Single-nucleotide polymorphisms (SNP) array analysis (Affymetrix Cytoscan 750K)
- Peripheral blood and buccal cells
  - SNP array analysis

## Results

- Pancreas
  - Targeted MPS hyperinsulinism panel
    - maternal *ABCC8* variant that was identified in blood was again detected, but only in a small number of reads with skewed allelic frequency → suggesting mosaicism
  - SNP array analysis
    - mosaic loss of heterozygosity (LOH) was observed for chromosome 11
    - observed pattern suggested high-level mosaicism for a cell line with whole-chromosome isodisomic UPD for chromosome 11, as well as a normal biparental cell line
    - Trio analysis suggested the UPD to be paternal in origin (isodisomic UPD11 pat)
    - a mosaic gain of one copy of chromosome 12 was also detected, consistent with mosaic trisomy 12 (mosaicism level 50%)
- Peripheral blood and buccal cells
  - SNP array analysis
    - no mosaic paternal uniparental disomy (pUPD) or trisomy 12 identified (cannot exclude low-level mosaicism of <10%)

## Conclusions

### Pancreatic mosaicism for pUPD11

- Most likely cause of CHI
- With 2 cardinal BWSp features
  - hyperinsulinism >1 week duration, escalating treatment and pancreatic adenomatosis
  - the BWS clinical diagnosis is met<sup>1</sup>
- In BWSp, pUPD11 predicts a high risk for Wilms tumour and hepatoblastoma, with 3-monthly ultrasound recommended for 7 years<sup>1</sup>
- α-fetoprotein screening is debated<sup>1,2</sup>
- Even in the absence of overt 11p overgrowth features, BWSp due to pUPD11 should be considered if**
  - persistent, severe CHI without an identified pathogenic K<sub>ATP</sub>-channel mutation(s)
  - large focal pancreatic lesions (with/without a K<sub>ATP</sub> mutation) or
  - atypical histology<sup>3,4</sup>

### Pancreatic mosaicism for trisomy 12

- Unreported previously → significance unknown
- Embryonic lethal when not in mosaic form
- A patient with trisomy 12 in 25% of peripheral blood cells has been reported
  - mild dysmorphic features at birth
  - normal development at 6 months of age<sup>5</sup>