

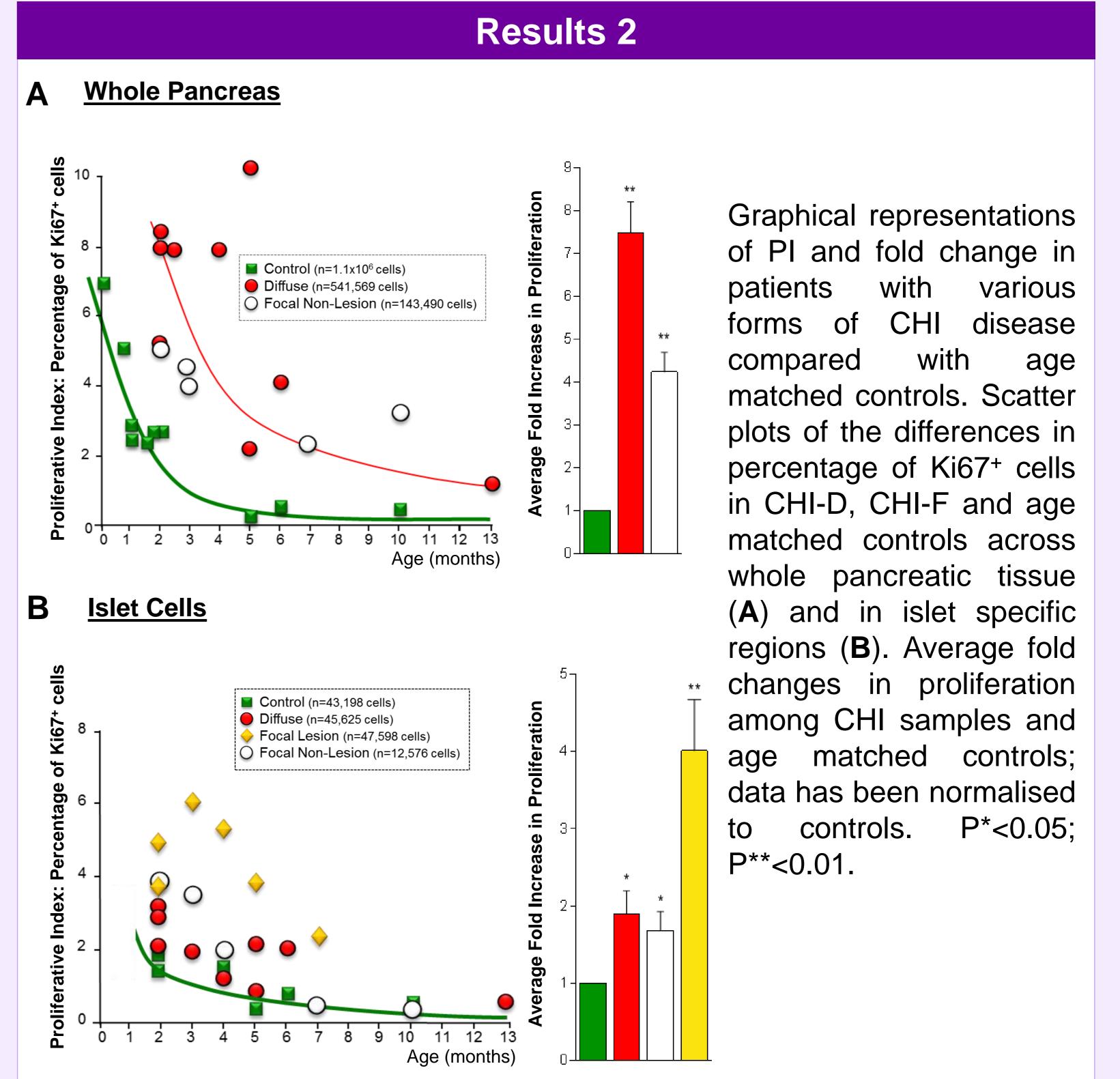
Islet Cell Proliferation is Inappropriately Maintained in the **Pancreas of Children with Congenital Hyperinsulinism in Infancy**

B Han¹, M Newbould², E Cheesman², G Batra², RJ Craigie², Z Mohamed^{1,2}, L Rigby², R Padidela², M Skae², KE Cosgrove^{*1}, MJ Dunne^{*1}, I Banerjee²

¹Faculty of Life Sciences, University of Manchester ²Central Manchester University Hospitals NHS Foundation Trust (CMFT) Oxford Road, Manchester, M13 9PT, UK

Background

Congenital Hyperinsulinism of Infancy (CHI) is a potentially lethal condition of profound hypoglycaemia caused by unregulated insulin release in the neonatal period and early infancy. CHI mainly arises due to mutations in ATPsensitive K-channel genes (ABCC8 and KCNJ11) which can manifest in all islets cells – diffuse CHI (CHI-D), or can be localised to a focal lesion, focal CHI (CHI-F). Increased rates of cell proliferation have been reported in the CHI-D and this may be liked to ABCC8 and KCNJ11 defects. Here, we examined the proliferative index (PI) of islet cells in CHI-D patients and compared this with focal CHI (CHI-F), which is caused by loss of cell cycle repression in β -cells specifically within the focal domain. We also examined islet PI in patient tissues with severe CHI unrelated to defects in ABCC8 and KCNJ11, atypical CHI (CHI-A).

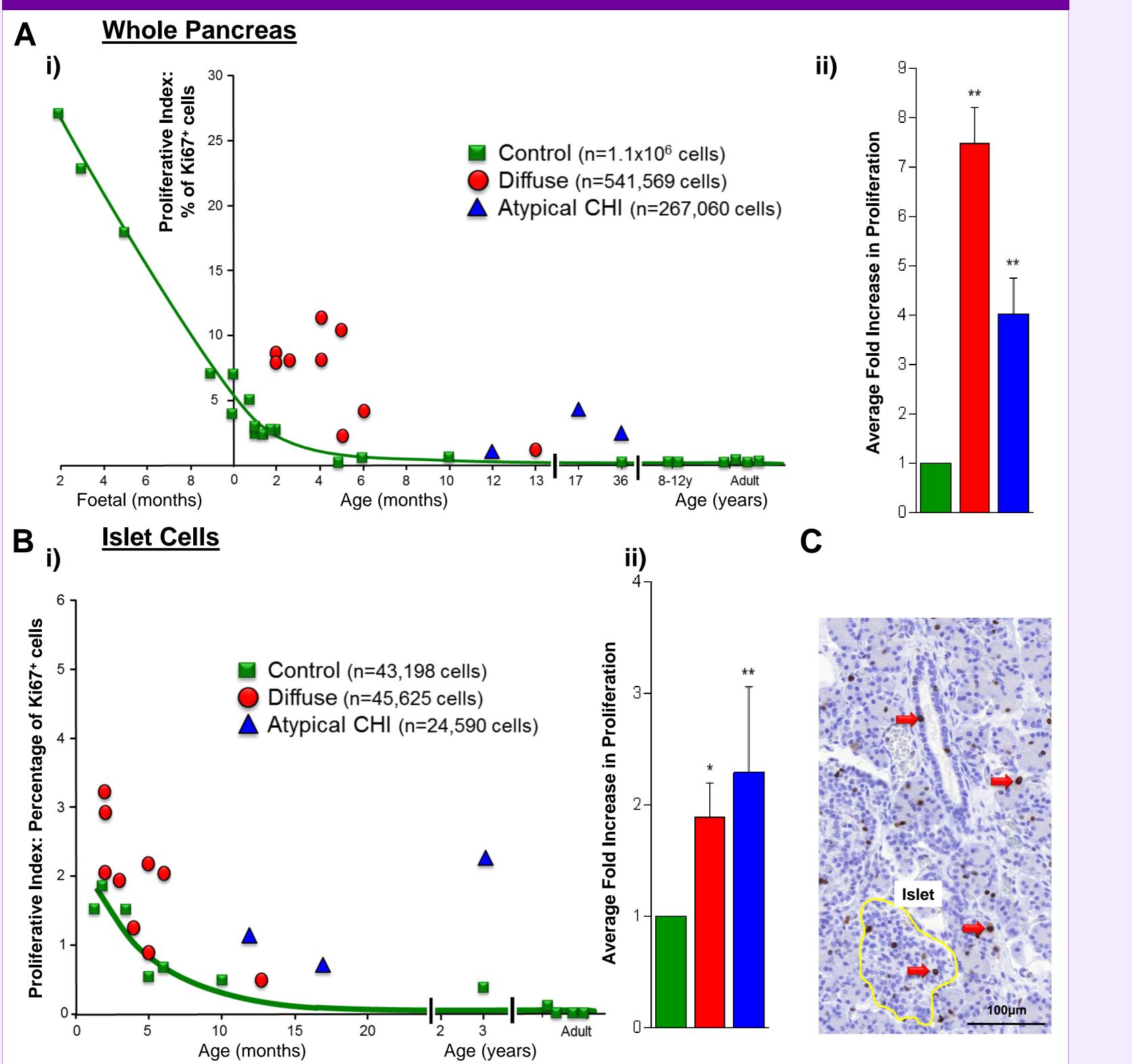


Results 3

Methods

Tissue was obtained following surgery from patients with CHI-D (n=10 patients) with ABCC8 gene defects), CHI-F (n=6 patients with ABCC8 gene defects), and CHI-A (n=3 patients with unknown genetic causes of disease). Neonatal control (n=12, 2-days to 36-months of age), foetal tissues (n=5, 10 to 35) weeks post conception) and adult tissues (n=4) were included as control groups. Immunohistochemistry staining with Ki67 on histological sections (5) µm) and high-content analysis were used to quantify the proliferation index (PI) and changes in PI expressed as average fold-changes across the various groups. Data were analysed with One-way ANOVA followed by Tukey's post hoc test.

Results 1

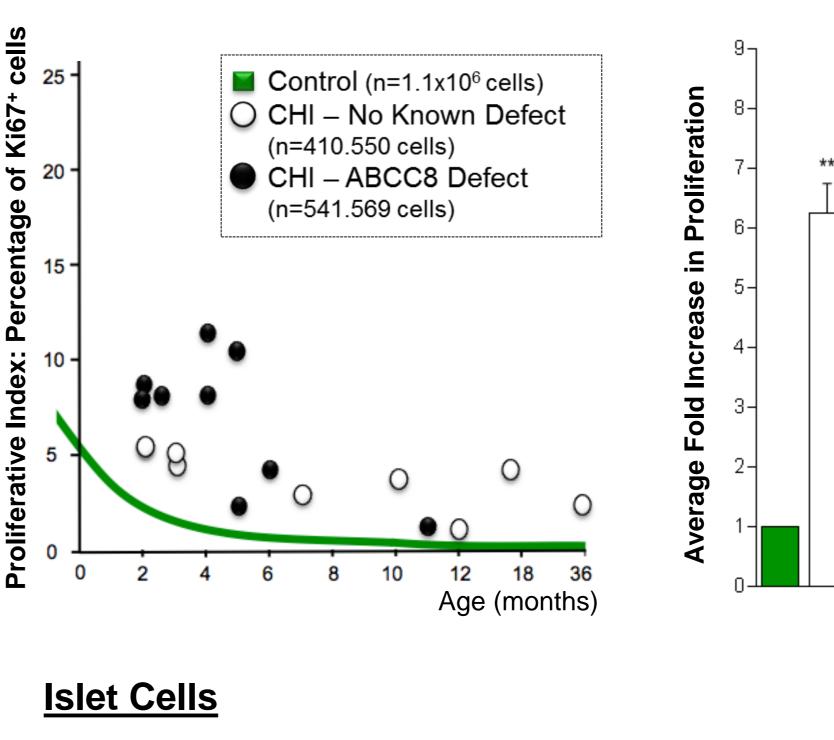


Whole Pancreas Α

B

Ki67

itage



Control (n=43,198 cells)

CHI – ABCC8 Defect

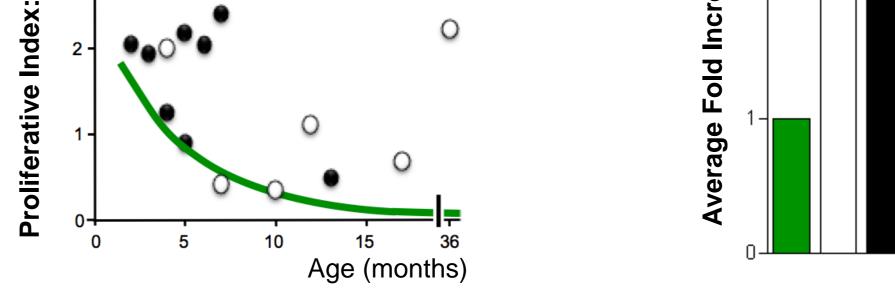
(n=12,576 cells)

(n=93,223 cells)

O CHI – No Known Defect

Graphical representations of PI and fold change in tissues with *ABCC8*⁺ gene defect known VS. no mutations compared with matched controls. age Scatter plots of the differences in percentage of Ki67⁺ cells in CHI tissues and age matched controls across whole pancreas (A) within islet specific and regions (B). Average fold proliferation changes in between CHI samples and age matched controls; data been normalised to has P*<0.05; controls. P**<0.01

Graphical representations of proliferation index (PI) and fold changes in patients groups compared with age matched controls, foetal and adult control samples. Scatter plots of the differences in percentage of Ki67⁺ cells in CHI-D, CHI-A and age matched controls across whole pancreatic tissue (shown in Ai) and in islet specific regions (shown in **B-i**). Average fold changes in proliferation among CHI samples and age matched controls; data has been normalised to controls (A-ii and B-ii). Immunohistochemistry staining of Ki67 in a 10-week post-natal control pancreatic tissue. A single islet has been annotated in yellow and Ki67⁺ cells have been indicated by red arrows (**C**). P*<0.05; P**<0.01.



Conclusions

ation

olife

1. There is a negative correlation between age and cell proliferation in control pancreas. 2. All three forms of CHI are associated with increased rates of cell proliferation. 3. Although there is a negative correlation between age and cell proliferation in CHI tissue (islets and pancreas), it is quantitatively different and not directly associated with ABCC8 mutations.



