Cell Mass Increase Associated with Formation of Glucose-Controlling β-Cell Mass in Device-Encapsulated Implants of hiPS-Derived Pancreatic Endoderm

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ABSTRACT
Device-encapsulated human stem cell-derived pancreatic endoderm (PE) can generate functional β-cell implants in the subcutis of mice, which has led to the start of clinical studies in type 1 diabetes. Assessment of the formed functional β-cell mass (FBM) and its correlation with in vivo metabolic markers can guide clinical translation. We recently reported ex vivo characteristics of device-encapsulated human embryonic stem cell-derived (hES)-PE implants in mice that had established a metabolically adequate FBM during 50-week follow-up. Cell suspensions from retrieved implants indicated a correlation with the number of formed β cells and their maturation to a functional state comparable to human pancreatic β cells. Variability in metabolic outcome was attributed to differences in number of PE-generated β cells. This variability hinders studies on processes involved in FBM-formation. This study reports modifications that reduce variability. It is undertaken with device-encapsulated human induced pluripotent stem cell-derived-PE subcutaneously implanted in mice. Cell mass of each cell type was determined on intact tissue inside the device to obtain more precise data than following isolation and dispersion. Implants in a preformed pouch generated a glucose-controlling β-cell mass within 20 weeks in over 60% of recipients versus less than 20% in the absence of a pouch, whether the same or threefold higher cell dose had been inserted. In situ analysis of implants indicated a role for pancreatic progenitor cell expansion and endocrine differentiation in achieving the size of β- and α-cell mass that correlated with in vivo markers of metabolic control. STEM CELLS TRANSLATIONAL MEDICINE 2019;8:1296–1305

SIGNIFICANCE STATEMENT
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INTRODUCTION
Type 1 diabetes causes a depletion in pancreatic β cells and hence a complete loss in cell-regulated glucose control. β-Cell implants prepared from human donor pancreases can restore endogenous insulin production and glucose control, but shortage of human donor organs limits implementation of this form of cell therapy. Human pluripotent stem cells may overcome this limitation as they represent a large-scale cell source that can be differentiated into pancreatic cells in the laboratory, which were shown to generate insulin-releasing implants in mice [1]. Studies with human embryonic stem cell-derived pancreatic endoderm (hES-PE) have advanced this development to clinical translation [2, 3] (clinicalTrials.gov-ViaCyte).
The ability of hES-PE to generate functional β-cell implants in mice was demonstrated for nonencapsulated as well as for encapsulated grafts [1, 4, 5]. This potential was also observed with human induced pluripotent stem cell-derived pancreatic endoderm (hiPS-PE) and with further differentiation stages containing insulin-positive cells in the grafts [6–10].

We have assessed the functional β-cell mass (FBM) that is formed in device-encapsulated hES-PE implants in the subcutis of mice, using a combination of in vivo and ex vivo markers [4, 5, 11]. It was thus found that implants which resulted in plasma hu-C-peptide levels >6 ng/ml at post-transplant (PT)-week 20 achieved metabolic control from PT-week 20 to 50 while this was not the case for recipients with lower levels. These implants presented the following characteristics when analyzed following retrieval at PT-week 50: (a) more than 0.3 × 10^6 β cells at >50% endocrine purity and (b) a functional state of the β cells that was similar to that of isolated human pancreatic β cells in terms of secretory responses during perfusion, content in typical secretory vesicles, and nuclear PDX1 expression [5]. The processes leading to formation of this metabolically adequate FBM should now be investigated.

To this end, the percent of recipients with adequate β-cell mass formation needs first to be increased. In our prior study, only 26% achieved hu-C-peptide >6 ng/ml at PT-week 20, the others exhibiting levels down to under 0.5 ng/ml, indicating a high variability in β-cell mass formation, confirmed by PT-week 50 counts of β-cell numbers in retrieved and dispersed implants. This variability hinders studies on mechanisms, timing, and duration of the growth in β-cell mass as it requires a higher number of recipients at start in order to establish a subgroup with similar rise to plasma hu-C-peptide levels that mark metabolic control. Moreover, it interferes with interpretation of in situ data during the first post-transplant weeks when plasma hu-C-peptide is not yet detected and thus cannot serve as in vivo marker for the degree of β-cell mass formation. To facilitate the investigation of early and late processes in β-cell formation we wished to identify conditions that help establish PE-generated β-cell implants with less variability than in the model studied so far. Our study has been undertaken with hiPS-PE instead of hES-PE in our previous work and also applied another device for its encapsulation [9, 12]. The similarity in 20-week plasma hu-C-peptide profiles of comparable cell numbers at start and in variability of individual values supports use of this hiPS-PE-device combination to address the aim. Two modifications were tested for their ability to help reach plasma hu-C-peptide >6 ng/ml at PT-week 20. The significance of this in vivo marker was evaluated by its correlation with glucose control and with the β-cell mass as measured in situ on intact implants, a more precise method than previous measurements on retrieved and dispersed implants. The first modification consisted in increasing the number of cells in the device at start, the second in implanting the device in a preformed pouch, a method that was reported to improve outcome of nonencapsulated islet cell and hES-PE implants [13, 14].

**MATERIALS AND METHODS**

**Transplantation of hiPS-Derived PE in Mice**

PE was derived from hiPS cells (produced by Fujifilm Cellular Dynamics, WI) by Nestlé Research (Lausanne, Switzerland) using a scalable and reproducible four-stage differentiation protocol [9]. Cryopreserved samples were shipped to Brussels. After thawing, aggregates were cultured for 72 hours in “stage 4” medium composed of Dulbecco’s modified Eagle’s medium high-glucose with Glutamax, 1% penicillin/streptomycin and 0.5× B27, supplemented by 50 ng/ml Noggin, epidermal growth factor, and keratinocyte growth factor (R&D Systems, MN) and 30 ng/ml Heregulin-β1 (tebu-bio, France) [9]. At end of culture, the preparation contained >90% living cells, the majority with nuclear PDX1 (Pancreatic and Duodenal Homeobox 1) positivity and 50% ± 5% being double-positive for PDX1 and NKX6.1 (NK6 Homebox 1) (average for three independent experiments; Supporting Information Fig. S1), similar to the characterization in [9]. An average 10% of cells were single- or double-hormone positive after staining for insulin, glucagon and somatostatin; they were negative for PDX1 and NKX6.1 (Supporting Information Fig. S1). Planar sheet macro-devices were also provided by Nestlé Research. They are composed of an outer 77 μm polyester mesh and a hydrophilic polytetrafluoroethylene 0.45 μm-porous (cell-impermeable) inner membrane held together by a polypropylene frame [12]. These devices have been shown to support long-term survival of allogeneic myoblasts implanted in the subcutaneous space [12]. They were loaded with 5 × 10^5 (Device 5M) or 15 × 10^5 (Device 15M) hiPS-PE cells before subcutaneous implantation in 8–12 weeks old male Fox Chase SCID/Beige mice (CB17.Cg-Prdke/sLy5<sup>+</sup>/Crl; Charles River, France). Devices with 5 × 10^6 cells were also tested after insertion in a preformed pouch (P-Device 5M; Fig. 1). Mice were followed for circulating markers of implant function over 20 weeks PT with blood taken under basal condition (2 hours-food removal) or following intraperitoneal glucose injection (3 g/kg body weight). Mice without implant were kept in parallel to collect control values on glycemia and plasma mouse C-peptide levels; they had not undergone surgery. Plasma proinsulin and hu-C-peptide were measured by trefoil-type time-resolved fluorescence immunoassay [15], plasma glucagon by radioimmunoassay (Merck-Millipore, Germany), mouse (m-)C-peptide by enzyme-linked immunosorbent assay (Crystal Chem, IL), and blood glucose using Glucocard (Menarini Diagnostics, Italy). These levels were also determined in a subgroup of recipient and control mice following alloxan injection (Sigma–Aldrich, MO; 50 mg/kg body weight intravenously) at PT-week 20, which is expected to destroy the mouse pancreatic β cells. Procedures were approved by the Ethical Committee of VUB and carried out according to European Community Council Directive (86/609/EEC). As in our previous studies we use the term “graft” for cells before transplantation and “implant” for the tissue after implantation. Device-encapsulated implants were retrieved for in situ (PT-weeks 2–20) and ex vivo analysis (PT-week 20).

**In Situ Histological Analysis of hiPS-PE Implants**

Retrieved devices were fixed in 4% neutral-buffered formaldehyde, processed, embedded in paraffin, and completely sectioned (5 μm). Sections were stained for cellular reactivity with guinea pig anti-insulin, rabbit anti-glucagon (each 1/1,000, in house-produced), rat anti-somatostatin (1/100, Abcam, U.K.), rabbit anti-vimentin (1/100, Dako, Denmark), mouse anti-CX19 (1/20, Dako), and rabbit anti-CD31 (1/100, Abcam). Cells were characterized for nuclear positivity after incubation with goat anti-PDX1 (1/100, R&D Systems, MN), mouse anti-NKX6.1 (1/100, Hybridoma Bank, NIH, IA), rabbit anti-MAFA (V-Maf...
β-cell mass formation in iPS-derived implants

Musculoaponeurotic Fibrosarcoma Oncogene Homolog A) (1/1,000, a kind gift of Dr. A. Rezania), and mouse anti-Ki67 (1/100, Dako) or rabbit anti-Ki67 (1/100, Acris, Germany). Secondary antibodies were Cyanine- or AlexaFluor-conjugated F(ab')2 fragments of affinity-purified antibodies allowing multiple labelling (1/500, Jackson ImmunoResearch, U.K.). Nuclei were stained by DAPI (Sigma–Aldrich). Digital images were analyzed with SmartCapture 3 software (DSUK, U.K.). Total volumes of cell types were determined semiautomatically according to an adaptation of a morphometric method originally established in rat pancreases [16] and described in Supporting Information Figure S2. For general histology, sections were stained with Hematoxylin & Eosin or Masson's Trichrome, images acquired with an Aperio CS2 (Leica Biosystems, Germany) and visualized with Pathomation software (Belgium).

Ex Vivo Analysis of hiPS-PE-Derived Cells Isolated from Retrieved Devices

At PT-week 20, implanted devices were retrieved and opened to isolate tissue content and disperse it to an aggregate suspension (<500 μm diameter) using type XI collagenase (1 mg/ml, Sigma–Aldrich). After washing, samples were taken for analysis of cell number (NucleoCounter YC-100, ChemoMetec, Denmark), composition and functions as previously described [4, 5]. Cell composition following dispersion was determined by manual count in immunocytochemistry on cells fixed in glutaraldehyde and embedded in araldite. Sections were stained using the same antibodies as described above. Insulin secretory responsiveness to glucose was determined in a perfusion assay. A sample with 500–700 × 10^3 cells was loaded on a P2-Bio-gel column (Bio-Rad, CA) and perfused for 60-minutes at 2.5 mmol/l glucose (basal), before pulsing with 5, 10, or 20 mmol/l glucose, with or without 10 mmol/l glucagon [4]. Perifusate was collected in 1 minute fractions and assayed for insulin [4]. Data were expressed per 10^3 β cells as determined in the sample that was added to the column.

Statistical Analysis

Results are expressed as means ± SD. Statistical analysis was performed using Prism 5.0 (GraphPad Software, CA) for comparison by one-way analysis of variance with Tukey’s post hoc test or Student’s t test (statistical significance at p < 0.05).

Results

Effect of Implantation in Preformed Pouch on Generation of Human β Cells in Device-Encapsulated hiPS-PE over 20 Weeks

Plasma hu-C-peptide was used as in vivo marker for the generation of β cells in hiPS-PE implants. The peptide was not detected in mice without device (<0.1 ng/ml detection limit). As in previous studies [4, 5], a glucose-induced level (>0.5 ng/ml) was set as criterium for the presence of functioning human β cells. This was the case for the average levels in the Device 5M group from PT-week 10 onward, progressively increasing up to PT-week 20 (Fig. 2A). They were similar for devices containing threefold more cells at start (Device 15M) but significantly higher when devices with 5M cells were implanted in a preformed pouch (P-Device 5M; Fig. 2A). As for hES-PE implants [5], individual values within each group varied widely. Increasing cell number from 5 to 15 million cells did not increase the percent recipients that reached the aimed in vivo marker (hu-C peptide >6 ng/ml; <20% in each group) but implants in a pouch achieved it in over 60% of recipients.

Recipients of device-encapsulated hiPS-PE implants exhibited higher fasting plasma glucagon levels (Fig. 2A), as

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was also the case for device-encapsulated hES-PE implants [5]. The increase was noticed from PT-week 3 onward and remained present until the end of the study. Levels were comparable for implants with $5 \times 10^6$ cells whether placed in a preformed pouch or not, but significantly higher with $15 \times 10^6$ cells at start. For this higher cell dose, the majority of recipients presented levels $>200$ pg/ml at PT-week 3–7, which was never achieved in the other groups or in mice without implant (Fig. 2A). The higher plasma glucagon levels were not correlated with higher plasma hu-C-peptide levels at PT-week 20; they also occurred in recipients with hu-C-peptide $<0.5$ ng/ml throughout follow-up.

**Basal and Stimulated Glucose-Responsiveness of Device-Encapsulated hiPS-PE Implants Associated to Glucose Control in Recipients**

Recipients exhibiting functioning human $\beta$ cells at PT-week 20 (hu-C-peptide $>0.5$ ng/ml at minute 60 postglucose injection)
Table 1. Metabolic parameters at PT-weeks 17–20 in recipients of functioning hiPS-PE implants

| Device-Encapsulated hiPS-PE Implants | Plasma hu-C-peptide (ng/ml) postglucose load | Glycemia (mg/dl) postglucose load | Plasma glucagon (pg/ml) 2 hours-fasting |
|--------------------------------------|---------------------------------------------|----------------------------------|----------------------------------------|
| Device SC 5 x 10⁶ cells              | 0 minutes: 0.7 ± 0.7 15 minutes: 1.9 ± 2.4 60 minutes: 4.0 ± 4.5††† | 0 minutes: 161 ± 15 15 minutes: 501 ± 62††† 60 minutes: 311 ± 107††† | 0 minutes: 144 ± 50‡‡‡ |
| Device SC 15 x 10⁶ cells             | 0.7 ± 0.5 18 ± 1.8 3.8 ± 2.5†††† | 145 ± 18** 495 ± 80††† 302 ± 106††† | 195 ± 57* ‡‡‡ |
| P-device SC 5 x 10⁶ cells            | 2.4 ± 1.0*** 5.9 ± 3.0*** 1.9 ± 5.3*** ††† | 149 ± 30 419 ± 165*** 165 ± 76*** ††† | 153 ± 88 |
| Controls                             | NA NA NA | 147 ± 19 473 ± 83** 287 ± 103††† | 70 ± 23 |

Glycemia, plasma human C-peptide, and glucagon levels measured at PT-weeks 17 (glucagon) and 20 (glycemia, C-peptide). Measurements were performed before (0 minutes), 15 minutes, or 60 minutes postglucose load (3 g/kg body weight, intraperitoneal) in 2 hours-fasted animals.

Statistical differences at a given time point by Student’s t-test:
Versus Device SC 5 x 10⁶ cells: *p < .05. **p < .01. ***p < .001.
Versus Controls: ‘p < .05. †p < .05. ††p < .01. †††p < .001.
Statistical differences between values at 0 minutes, 15 minutes, and 60 minutes within a given condition, by one-way analysis of variance with Tukey’s post hoc test:
15 minutes and 60 minutes versus 0 minutes: ′p < .05. †p < .05. ††p < .01.
15 minutes versus 60 minutes: ′′p < .05. ††p < .05. †††p < .01.
Abbreviations: hiPS-PE, human induced pluripotent stem cell-derived pancreatic endoderm; PT, post-transplant; SC, subcutaneous.

were analyzed for their plasma hu-C-peptide levels at basal glycemia and following glucose injection, an index for their glucose responsiveness (Table 1). Those in Device 5M group exhibited a response, but this was only statistically significant at minute 60, at which time glycemia was still elevated as in mice without implant. This was also the case in the Device 15M group, with similar values. In the P-Device 5M, they presented higher hu-C-peptide levels than those in the Device 5M group, basal as well as after glucose injection. Both minute 15 and minute 60 values were statistically higher than basal with a reduction in glycemia to fasting values at minute 60 (Table 1). These observations indicate that the β-cell mass in pouch-implants is secretory active at basal glycemia in mice (>3 ng/ml in mice without implant (black curves in Fig. 2B). The β-cell mass in pouch-implants maintained glucose control after injection of alloxan, without decline in hu-C-peptide levels and rise in glycemia (red curves in Fig. 2B, 2C). In mice without implant, alloxan abolished mouse C-peptide release and caused hyperglycemia and body weight loss (black curves in Fig. 2B, 2C).

Correlation Between Glucose-Induced Plasma Hu-C-Peptide Levels and Size of Formed β-Cell and α-Cell Mass in Devices with hiPS-PE Implants

At PT-week 20, glucose-induced plasma hu-C-peptide levels correlated with total β-cell volume as measured in the device (Fig. 3A). They were also strongly correlated with α-cell volume, and weakly with that of nonendocrine CK19-positive cells (Fig. 3A). When total cell volume measured in the device was compared with that in the initial grafts, a wide variability was noticed. For hu-C-peptide levels <0.5 ng/ml, it was lower than at start (Fig. 3A) and/or mainly consisted of hormone-negative cells (Fig. 4) that were vimentin (VIM)- or CK19-positive. For

Identification of Cells in Proliferative Activity in Device-Encapsulated hiPS-PE Implants

The observed growth in cell mass between PT-weeks 2 and 20 led us to search for cells in proliferative activity. At PT-week
2, all analyzed implants—whether with or without pouch—were mainly composed of PDX1+/NKX6.1+/SOX9+ positive cells, with <10% hormone-positive cells, similar to grafts at start. However, their PDX1+/NKX6.1+ cells were strongly CK19-positive while this staining was weak and spotty in the initial aggregates (Fig. 5). A large proportion of these CK19+/PDX1+/NKX6.1+ were now organized as an epithelial cell layer around a lumen, with VIM expression at their basal pole and CK19 at their apical pole (Fig. 5). A smaller fraction with weaker CK19+ staining occurred clustered between the duct-like structures and was found associated with hormone-positive cells (Fig. 5), some double-positive, and mostly glucagon-positive and NKX6.1-negative.

At PT-week 20, all implants—whether with or without pouch—still presented CK19+ cells in epithelial layers around a lumen but these were now NKX6.1−/VIM−; they represented the major fraction of cells in proliferative activity but with a lower and variable percentage (0.5%–9%). Between these duct-like structures, clusters of single-hormone positive cells were observed, including PDX1+/NKX6.1+/insulin (INS)+ cells with low and variable KI67-positivity (0%–3%).

**DISCUSSION**

This study shows that subcutaneous implants of device-encapsulated hiPSC-PE can generate a FBM that controls glycemia in mice. It thus reproduces the outcome that we previously reported for hES-PE using another device [5]. Glucose control by the hiPS-PE-generated β cells was demonstrated by

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**Figure 3.** Correlation between glucose-induced plasma hu-C-peptide levels and pancreatic endocrine cell mass in device-encapsulated human induced pluripotent stem cell-derivated pancreatic endoderm (hiPS-PE) implants at post-transplant (PT)-week 20. (A): In situ cell mass measurements (as volume) in devices at PT-week 20 with initially 5 × 10⁶ hiPS-PE cells are plotted against the corresponding glucose-induced plasma hu-C-peptide levels. Linear regression analysis indicates high correlations for the volumes of insulin- (μl) and of glucagon- (μl) positive cells and lower correlations for the volume of CK19-positive cells (μl) and total cell volume (expressed as percent of initial volume). (B): In situ total and hormone-positive cell mass in devices of pouch-implants with plasma hu-C-peptide >3 ng/ml at PT-week 20. Comparison with values at PT-week 2 (no hu-C-peptide detected). (C): ex vivo β-cell secretory response of pouch-implants retrieved from mice with plasma hu-C-peptide >3 ng/ml. Glucose concentrations are indicated on top. Curve shows insulin release at baseline (2.5 mmol/l glucose) and at higher glucose concentration in absence and presence of glucagon (10 nmol/l). Data are expressed as pg × 10⁻³ β cells × minute⁻¹. Statistical differences in first-phase insulin release peaks at 5–10–20 mmol/l glucose versus baseline, by one-way analysis of variance with Tukey’s post hoc test: **, p < .01. Student’s t test of second-phase release at 20 mmol/l glucose with or without glucagon: †, p < .05.
C-peptide levels within this follow-up period. In addition, quantification of β-cell mass on intact tissue inside the device is more precise provided that it is based on representative sampling as validated in the present study. This in situ approach will also help identify processes in the generation of the β-cell mass and possible causes of failure.

Glucose-controlling hiPS-PE implants exhibited a β-cell volume of 1.1 μl ± 0.3 μl at PT-week 20, which is in the range of the β-cell volume measured in the normal mouse pancreas (0.8–1.1 μl). Their β cells were single-hormone positive, with nuclear positivity for PDX1, NKX6.1, and MAFA, markers of differentiation. When perfused ex vivo, they were shown to rapidly adjust their insulin release to increased or decreased glucose concentrations, a sign of functional maturation and a cellular basis for the observed in vivo hu-C-peptide responses. The amplitude of the in vivo secretory responses appears determined by the corresponding β-cell volume in the implants as indicated by their close correlation.

Implants of device-encapsulated hiPS-PE in a preformed pouch reduced the number of recipients with low β-cell mass at PT-week 20. Pepper et al. showed that such site pre-treatment facilitates survival and function of nonencapsulated pancreatic islet cells under the skin of mice, possibly by prevascularizing their microenvironment [13]; they also found it to support development of in vivo β-cell functions in non-encapsulated hES-PE implants, but also of large cysts lined by PE-generated epithelial cells [14, 18]. In implants with metabolic control at PT-week 20, we only noticed small cell lined cavities in the devices, with endocrine cells composing the major tissue fraction. Total cell volume was threefold higher than at start. This increase was not yet detected at PT-week 2; on the contrary, total cell mass was lower than initially, possibly the result of cell loss during engraftment. Between PT-weeks 2 and 20, it increased sevenfold comprising appearance of single-hormone positive cells representing 64% of total cell volume at PT-week 20.

The PDX1+/NKX6.1−/hormone-negative pancreatic progenitor cells are considered as precursor for hormone-positive cells and as drivers for cell expansion. In the start preparation they contained the majority of KI67-positive cells and this was also the case in PT-week 2 implants. At the latter time point, the PDX1+/NKX6.1− population was however strongly CK19− and organized as epithelial cell layer around small cavities. It also exhibited a VIM-positivity at the basal pole of the cells, similar to that reported in proliferating duct cells of the mouse pancreas, and suggested to precede budding of newly formed β cells [19, 20]. Our protocol was not designed to investigate such process but we did notice that the PDX1+/NKX6.1−/CK19+/VIM− (Kl67+) cell lining of the cavities had juxtaposed cell clusters containing weakly CK19+ cells and a small fraction of hormone-positive cells, most expressing glucagon, with very few Kl67+ cells. In the subsequent 18 weeks, both compartments increased in cell mass whereby the PDX1+/CK19+ epithelial cell layer became NKX6.1+/VIM− with less Kl67 positive cells, and the neighboring cell clusters presented single-hormone positive cells, including PDX1+/NKX6.1+/INS− cells with few Kl67+ cells. The model in which these data have been collected allows to investigate the processes between these two time points and the possible transition from epithelial cell layer to endocrine cell clusters.

On basis of the measured volumes of β-cell and α-cell population at PT week 20, and of the averages of individual β-cell and

**Figure 4.** Pancreatic endocrine cells in device-encapsulated human induced pluripotent stem cell-derived pancreatic endoderm (hiPS-PE) implants at post-transplant (PT)-week 20. (A): In situ histological analysis of device-implants at PT-week 20 and stained with Masson’s trichrome, illustrating cell recovery and phenotype in implants from recipients without or with glucose-stimulated plasma hu-C-peptide (<0.5 ng/ml, top; >3 ng/ml, bottom). Scale bar: 100 μm. (B): In situ analysis of device-implants in pouch with plasma hu-C-peptide levels >3 ng/ml at PT-week 20. Presence of single-hormone positive cells staining for insulin, glucagon, or somatostatin (top). Most insulin-positive cells exhibit strong nuclear expression of transcription factors PDX1, NKX6.1 and MAFA (bottom). Scale bars: 100 μm.
α-cell surface areas, we calculated that $10^6$ β cells and $2 \times 10^6$ α cells had been formed since PT-week 2; the number of somatostatin-positive cells was not determined but their total mass suggests that it is not lower than that of the insulin-positive cells. This endocrine cell formation occurs during a sevenfold expansion of total cell mass. It is at present unknown to which extent it depends on this cell amplification, and whether this is expressed as budding of endocrine progenitor cells from epithelia with replicating pancreatic progenitor cells, and/or whether it involves replication of newly formed endocrine cells. Formation of more α cells than β cells was also observed in device-encapsulated hES-PE implants at PT-week 50 [5]; it may be induced by local influences such as hypoxia. The PE-generated α cells have been shown to release glucagon under glucose-dependence [4]. They may thus exert positive paracrine effects on formation, maturation, and function of β cells [11, 21–23]. They are also likely responsible for the higher plasma glucagon levels in recipients of hIPS-PE or hES-PE implants [5], occurring at an early stage, more than 5 weeks before hu-C-peptide was detected; implants with threefold higher cell number at start indeed resulted in a more pronounced elevation of plasma glucagon. It is so far unknown whether this elevation marks a local or systemic effect.

**CONCLUSION**

This study reports modifications that reduce variability in the formation of a FBM and thus facilitate studies on underlying mechanisms. Implants of device-encapsulated hIPS-PE cells in a preformed subcutaneous pouch were shown to generate, within 20 weeks, a glucose-controlling β-cell mass in the majority of recipient mice. This outcome was defined by in vivo markers that were applicable in normoglycemic mice. They correlated with β-cell volume as measured in situ within intact implants. A dose of $5 \times 10^6$ cells at start with less than 10% mostly double-hormone-positive cells increased threefold over this period with more than 60%

![Figure 5.](image-url)
single hormone-positive cells. In situ measurements of the mass of composing cell types indicated that the formed α-cell mass was twofold larger than the formed β-cell mass, raising the possibility of positive paracrine influences. Comparison with in situ analysis at PT-week 2 indicated a role for pancreatic progenitor cell expansion and endocrine differentiation in achieving in vivo markers of metabolic control at PT-week 20. Further studies using this model and markers can clarify their respective contribution and interactions.

**ACKNOWLEDGMENTS**

We thank Dr. Rezania (when at BetaLogics, Venture, Janssen Research and Development, Raritan, NJ) for providing the anti-MAFA antibody, and the Developmental Studies Hybridoma Bank (NIH, IA) for providing the anti-NKX6.1 antibody. We are grateful to their collaborators at VUB-Diabetes Research Center for their support, and the team of the Clinical Chemistry department of UZB for the peptide assays. This work was supported by grants from the European Commission (H2020 681070), Juvenile Diabetes Research Foundation (17-2013-296), and the Flemish Government (IWT130138).

**AUTHOR CONTRIBUTIONS**

T.R.: conception and design, collection and/or assembly of data, data analysis and interpretation, manuscript writing, final approval of manuscript; I.D.M., F.O.V.H., G.M.S., K.G.S., Z.L.: collection and/or assembly of data, data analysis and interpretation, final approval of manuscript; B.K.: final approval of manuscript; M.R.C.K., C.H., N.B.: provision of study material, final approval of manuscript; D.G.P.: conception and design, financial support, data analysis and interpretation, manuscript writing, final approval of manuscript.

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T.R. is Ph.D. fellow of Research Foundation Flanders-FWO. The other authors indicated no potential conflicts of interest.

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