

Stem Cell Therapy to Treat Diabetes Mellitus

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■ Abstract

Transplantation of pancreatic islets offers a direct treatment for type 1 diabetes and in some cases, insulin-dependent type 2 diabetes. However, its widespread use is hampered by a shortage of donor organs. Many extant studies have focused on deriving β -cell progenitors from pancreas and pluripotent stem cells. Efforts to generate β -cells *in vitro* will help elucidate the mechanisms of β -cell formation and thus provide a versatile *in vivo* system to evaluate the therapeutic potential of these cells to treat diabetes. Various successful experiments using β -cells in animal models have generated extensive interest in using human embryonic stem cells to

restore normoglycemia in diabetic patients. While new techniques are continually unveiled, the success of β -cell generation rests upon successful manipulation of culture conditions and the induction of key regulatory genes implicated in pancreas development. In this review, we compare successfully conducted protocols, highlight essential steps and identify some of the remarkable shortfalls common to these methods. In addition, we discuss recent advancements in the derivation of patient-specific pluripotent stem cells that may facilitate the use of autologous β -cells in stem cell therapy.

Keywords: type 1 diabetes · stem cell · beta-cell · transplantation · Pdx1 · Sox9 · NeuroD1 · ngn3

Diabetes and islet cell transplantation

Diabetes mellitus is a chronic metabolic disorder manifested by hyperglycemia due to a deficiency of insulin production by pancreatic β -cells. This can be a direct consequence of autoimmune destruction of β -cells, as seen in type 1 diabetes (T1D) [1-3]. Other types of diabetes, collectively known as type 2 diabetes (T2D), occur because of a combination of reduced insulin sensitivity (non-insulin dependent) and impaired β -cells function [4]. Diabetes can be inherited, such as that seen in maturity onset diabetes of the young adult (MODY), or caused by mutations in an autosomal dominant gene resulting in the disruption of insulin production. Gestational diabetes is another less known form of diabetes, but it

imposes an increasingly prevalent risk factor for the development of T2D after pregnancy [5].

Exogenous insulin supply is required to maintain normoglycemia in many diabetic patients. A promising way of treatment is using β -cell replacement therapy. For successful islet transplantation a sufficient amount of β -cells needs to be included in the transplant to control the blood glucose level without repeated insulin injections. In 2000, Shapiro and colleagues described the successful cure of T1D in a small number of patients using a procedure known as the 'Edmonton Protocol' [6]. The authors achieved independence of insulin injections in seven T1D patients by transplanting a large number of islet cells (obtained from two donor pancreases) combined with the use of glucocorticoid-free immunosuppressive regimen. Notably, both host-

versus-graft and autoimmune reactions were avoided. However, despite short term success, long term insulin independence is usually not sustainable. Their international clinical trial showed that only 13% of the patients (5 out of 36 subjects with T1D) maintained insulin independence at 2 years, and 28% had complete graft loss 1 year after the final transplantation [7].

One explanation for the poor long-term outcome is recurrent immune destruction of the transplanted islets despite immunosuppression. Indeed, late graft loss is commonly observed although graft function measured by C-peptide immunoreactivity was retained. This phenomenon of 'islet graft exhaustion' could be attributed to the toxicity of immunosuppression, autorejection or recurrence of autoimmunity [8]. Long term failure of the early transplantation therapies has prompted the search for more defined sources of β -cells. Intensive research is being conducted to look for alternative sources of β -cells because of the shortage of cadaver donors. An alternative resource for transplantable β -cells is the stem cell. This article reviews the current knowledge on the role of stem cells in pancreatic development and as a source for β -cell transplantation. In addition, it discusses recent advancements in the derivation of patient-specific pluripotent stem cells that may facilitate the use of autologous β -cells in stem cell therapy.

Formation of new β -cells: knowledge from developmental biology of the pancreas

Stem cells are cells that are able to proliferate while maintaining an undifferentiated status (self-renewal) and retaining a capacity to differentiate into specialized cell types under appropriate conditions. The sources of such stem cells for the purpose of generating β -cells were firstly identified through several studies on adult and fetal pancreas. It is generally assumed that islet neogenesis or replication occurs throughout life with a gradual decrease in capacity in elder humans. It is also assumed that the generation of β -cells can take place by two pathways: firstly via replication of already existing β -cells and secondly via neogenesis (differentiation) from pancreatic endocrine or ductal progenitors [9, 10]. During fetal life, the majority of new β -cells develop from precursors, but newly developed β -cells also proliferate. In the adult pancreas, the rate of both β -cell neogenesis and replication is more limited than in newborn, but it is thought to take place [11-13]. However, a study by Dor *et al.* (2004) has challenged the concept of neogenesis, because the authors did not observe new β -cells formed from non-insulin-

producing stem cells during adult life [14].

Early pattern of differentiation in the mouse and human pancreas

During early embryogenesis, the epiblast and primitive endoderm arise from the inner cell mass (ICM) of the blastocyst, from which pluripotent embryonic stem (ES) cells are derived. The primitive endoderm gives rise to extraembryonic tissue, whereas the epiblast differentiates to form the three primary germ layers during gastrulation. Cells from the epiblast are recruited into the primitive streak and migrate out to form the mesoderm and definitive endoderm (DE), from which the pancreas develops. Nodal and activin, members of transforming growth factor beta (TGF- β) superfamily, are essential for the initial endoderm specification, where they regulate the activation of many key regulatory genes, such as *foxa2*, *sox17* and *mixl1* in a strict temporal sequence. The primitive gut tube then arises from the DE, which later forms foregut, midgut and hindgut [15]. Signals from the notochord and mesenchyme specify the pancreatic domain at the endodermal region. The pancreas develops from the posterior foregut and the subsequent formation of the pancreatic anlagen relies on retinoid signaling and on inhibition of Indian and Sonic hedgehog (Shh) signaling. The developing pancreas is composed of pancreatic and duodenal homeobox 1 (Pdx1)-expressing epithelial precursor cells that give rise to the three differentiated compartments of endocrine, exocrine and ductal cells found in the adult pancreas. The endocrine cell mass aggregates in interstitial clusters adjacent to the ductal epithelial cells, to form the islets of Langerhans [16].

In mice, progression from foregut endoderm to insulin-producing cells is rapid (between E9.5 and E10). Nonetheless, in humans, although pancreatic differentiation from foregut endoderm is initiated at an equivalent time in human (26 days post conception, dpc), significant insulin expression is delayed. This observation supports the notion that embryological stages in the mouse and the human are not as closely equivalent as previously assumed [17]. In humans, by 35 dpc, the ventral pancreatic bud begins to migrate backwards and comes into contact and eventually fuses with the dorsal pancreatic bud during 6 weeks post conception (wpc) [18]. Both insulin and glucagon expression could only be detected at 8 wpc [19, 20], two weeks later than expected compared to the mouse [21].

Insulin is the most abundant hormone detected during the first trimester in human pancreas [22]. At 10 wpc, all four islet hormones can be detected. Islets of

Langerhans are formed at 12-13 wpc. At this stage, islets are comprehensively vascularized and contain cells independently immunoreactive for insulin, glucagon, somatostatin and pancreatic polypeptide (PP). This is accompanied by the expression of prohormone convertase 1/3 (PC1/3), islet amyloid polypeptide (IAPP) and β -cell specific glucose transporter-2 (GLUT2) [22]. The expression of these markers implies that fetal β -cells may be capable of processing and secreting insulin. However, the human fetal pancreas is not glucose-responsive, despite being able to secrete insulin upon stimulation with β -cell secretagogues [23]. These studies notwithstanding, the mechanisms that control endocrine cell formation from human neonatal pancreas are poorly understood because of the apparent scarcity of material and difficulties in obtaining human fetuses.

Pancreas and islet-related transcription factors

In mice, embryonic pancreatic epithelial cells express nuclear Pdx1 and cytoplasmic cytokeratin-19 (CK19) [22]. *Pdx1* plays an important role in the transactivation of the *insulin* gene and so is required to maintain normal β -cell homeostasis [21]. Another more recently described pancreas-related transcription factor, Sox9, is predominantly expressed throughout the early developing pancreas (prior to 14 weeks of gestation). In contrast to Pdx1, the expression of Sox9 is down-regulated once endocrine cells are developed and is later restricted to ductal cells. A study using *sox9* heterozygous mouse mutants suggests that the role of *sox9* is as a determinant of multipotent pancreatic endocrine cells in the pancreas [24].

Pancreatic endocrine cell fate specification is also ensured by a lateral inhibition process mediated by Notch signaling pathways. Genetic studies that involved ectopic expression of *neurogenin3* (*ngn3*) and intracellular Notch in early pancreas progenitors collectively confirm the function of *ngn3* in controlling endocrine cell fate. Mice lacking *ngn3* function fail to generate pancreatic endocrine cells and die postnatally from diabetes [25]. Similarly, *neuroD1* knock-out mice fail to develop islets and develop severe diabetic ketoacidosis and perinatal death [26]. It has been shown that maturity onset diabetes of the young type 6 (MODY-6) in humans is also associated with mutations in *NEUROD1* and that the abnormality of islet morphogenesis is due in part to inadequate expression of the *INSULIN* gene [27].

Islet1 expression in pancreatic endodermal cells is required for the formation of dorsal mesenchyme and

generation of all endocrine islet cells. A number of genes control the differentiation of specific pancreatic endocrine cell subsets. *Pax4* is required for the initial commitment of early endocrine precursors to become β - and δ -cells, while *pax6* is required for the early differentiation of α -cells [28-33]. Fully differentiated β -cells first appear around E13 at the start of a massive wave of β -cell differentiation, which is known as "secondary transition" [34]. *Nkx6.1* expression is required by this second phase of β -cell neogenesis in the developing pancreas [35, 36]. A recently described transcription factor, MafA is induced at the final stage of β -cell differentiation and functions as a potent activator of *insulin* gene transcription [37].

Approximately 90% of β -cells and 15% of δ -cells in adult islets express Pdx1. Pdx1 regulates the expression of β -cell-specific genes such as *INSULIN*, *LAPP* (islet amyloid polypeptide), β -cell-specific glucose transporters glucokinase (*GCK*) and *GLUT2*. Hence, the expression of Pdx1, Gck and GLUT2 can be used as a key indicator of β -cell functionality [38]. The dual action of Pdx1, as a pancreas commitment factor during embryogenesis and as a regulator of islet cell physiology in mature islet cells, underscores the unique role of *PDX1* in maintaining the function of human pancreatic endocrine cells [39]. Apart from *PDX1*, mutations of which have been linked to MODY-4 [40, 41], several other genes have also been shown to be required in humans and in mice to assure β -cell functionality. These include the other five MODY genes: *GCK* (MODY-2), *hepatocyte nuclear factor 1a* (*HNF1A*; MODY-3), *HNF1B* (MODY 5), *HNF4A* (MODY-1) and *NEUROD1* (MODY-6) [27, 42].

Properties of a mature β -cell and insulin biosynthesis

The gold standard for defining β -cell function is glucose responsiveness. A functional β -cell exhibits an acute three-fold stimulatory insulin release in response to glucose. Zinc is required for packaging insulin, an integral part of insulin crystals for 2-Zn-insulin hexamer, as well as free ionized zinc in the extragranular space that acts as a reservoir for granular zinc pools [43-46]. The ability to regulate glucose uptake by the islet-specific glucose transporter GLUT2 is the first step necessary for the activation of the regulatory region of the *Insulin* gene to glucose [47]. In the absence of GLUT2, the endocrine pancreas shows the loss of first phase glucose-stimulated insulin secretion and an inverse α - to β -cell ratio [48]. Glucose signaling to secretion and insulin biosynthesis are also impaired [49]. The synthesis of the glucose-phosphorylating enzyme

glucokinase is a late event in β -cell maturation [50]. Patients with mutations of the *GCK* have mild fasting hyperglycemia throughout life [42].

Insulin mRNA is translated as a single chain precursor called preproinsulin, and the removal of its signal peptide during insertion into the endoplasmic reticulum generates proinsulin. Proinsulin consists of three domains: an amino-terminal B chain (30 amino acids), a carboxy-terminal A chain (21 amino acids) and a connecting peptide in the middle known as the C-peptide. Within the endoplasmic reticulum, proinsulin is exposed to several specific endopeptidases that excise the C-peptide, thereby generating the mature form of insulin. Insulin and free C-peptide are packaged in the Golgi into secretory granules, which accumulate in the cytoplasm. In pancreatic β -cells, C-peptide and mature insulin are present in equimolar ratios and co-localize in the secretory granules [51].

Sources of renewable β -cells

Pancreatic stem cells in pancreas ductal cells

Several studies have revealed the possibility that endocrine precursors lie within pancreatic ducts. In the adult pancreas, endocrine cell formation from ductal epithelial cells has been observed both in experimental models of pancreas injury [52] and in various clinical pathologies [53]. Islet neogenesis and replication may provide the source of pancreatic stem cells from which normal renewal of islets occurs throughout life [54].

Ramiya *et al.* (2000) generated pancreatic islets from adult murine ductal cells *in vitro* [55]. The pancreatic stem cells generated insulin-producing stem cells (IPSC) in a monolayer, from which islet progenitor cells (IPC) budded. The pancreatic islet mass significantly increased by growing IPC in culture; many of the cells expressed both glucagon and insulin, a phenotype reported for immature islet cells on their path to end-stage differentiation [56]. These IPSC-derived islets secreted insulin in response to glucose challenge, and upon addition of nicotinamide they further matured and differentiated into fully functional islets. After transplantation, the mice were able to regulate the levels of glucose in their blood within a week, and survive without further need for insulin.

Bonner-Weir *et al.* (2000) demonstrated that human pancreas duct tissue can also be expanded *in vitro* and then be directed to differentiate into glucose-responsive islet buds after being overlaid with matrigel [57]. The cells were grown in serum-free media with glucose, insulin, transferrin and selenium (ITS) sup-

plements, nicotinamide and keratinocyte growth factor (KGF). The epithelial cells formed three-dimensional cystic structures, characteristic of cultivated human islet buds. These cultivated human islet buds consisted of both CK19-positive ductal cells and hormone-positive islet cells. There was a significant increase in DNA and insulin content over 3-4 weeks of culture, suggesting that these human islet buds were immature and still in the process of differentiation.

Hui *et al.* (2001) demonstrated that overexpression of Pdx1 and treatment with glucagon-like peptide 1 (GLP-1) induce the differentiation of rat and human pancreas ducts into insulin-secreting cells [50]. GLP-1 is capable of restoring normal glucose tolerance in aging mice and increasing islet mass in adult animals previously subjected to subtotal pancreatectomy [9]. Additionally, endogenously *pdx1*-positive rat (ARIP) and stably *pdx1*-transfected human (PANC-1) cell lines attained a β -cell phenotype upon GLP-1 treatment. This study provides a significant basis for determining the minimum biological requirements for a non- β -cell to become a β -cell, whilst supporting the previous finding that pancreatic ductal cell lines are capable of giving rise to pancreatic endocrine cells.

Guided by the expression of *ngn3*, β -cell precursors were identified in an injury model of adult pancreas [58]. Following partial duct ligation, cells located in the ductal lining were found to reactivate *ngn3*. These Ngn3-positive cells express ductal cytokeratins, but did not stain for insulin, suggesting they are of islet progenitor phenotype. However, they developed into cells that subsequently proliferated, both *in situ* and *in vitro*, and were found to also express Pdx1. Ngn3-positive cells were then isolated and injected into *ex vivo* embryonic mouse pancreas, and were found to autonomously (not by fusion) increase glucose-responsive β -cell mass in the explants.

Dor *et al.* (2004) have challenged the concept of neogenesis despite the aforementioned evidence of the existence of β -cell precursors [14]. To prove this, they labeled β -cells in transgenic mice with *insulin* promoter driving an inducible Cre/lox system. Following tamoxifen induction, a human placental alkaline phosphatase (HPAP) reporter gene was expressed ('pulse'). Thus, during turnover ('chase'), β -cells could be identified with HPAP dye. Surprisingly, all islets analyzed in adult pancreas in the 'pulse and chase' experiments contained numerous HPAP-immunoreactive β -cells. Nevertheless, no insulin-immunoreactive and HPAP-negative cells were observed. Notably, partial pancreatectomy resulted in the same observation but with increased bromodeoxyuridine (BrdU) incorporation, in-

dicating that these β -cells retained their full proliferative capacity following injury. Thus, their observation indicates that terminally differentiated β -cells are capable of self-renewal and differentiation into new β -cells. To date, this study remains controversial because it does not prove the absence of stem cells during neonatal life or after pancreas injury, and the possible transcriptional activity of *insulin* promoter in such overlooked stem cells.

Pancreatic stem cells in islets of Langerhans

The endocrine cells of the adult rat pancreatic islets of Langerhans, including β -cells, turn over every 40-50 days by a process of apoptosis, and are replaced by neogenesis from progenitor epithelial cells located in the pancreatic ducts [55, 57]. However, the administration of glucose or GLP-1 to rats for 48 hours resulted in a doubling of islet cell mass, suggesting that islet progenitor cells were able to proliferate and that the precursors may reside within the islets themselves [59].

In another study, rat and human pancreatic islets were isolated *post-mortem* and cultivated in medium supplemented by growth factors. Focal regions of nestin-positive cells were identified in large, small, and centrolobular ducts of the rat pancreas. These nestin-expressing cells were not ductal epithelial cells as they were CK19-negative. When these cells were grown in medium containing basic fibroblast growth factor (bFGF) and epidermal growth factor (EGF), they were able to proliferate. When the medium was switched to low glucose medium containing hepatocyte growth factor (HGF), betacellulin, activin-A, exendin-4 or nicotinamide, they displayed a ductal/endocrine phenotype with expression of *CK19*, neural-specific cell adhesion molecule (*NCAM*), *insulin*, *glucagon*, *glut2* and *pdx1*, besides the expression of liver and exocrine pancreas markers. These findings provide evidence that pancreatic islets themselves, apart from the ducts contain multipotential progenitor cells that are able to differentiate to β -cells.

However, controversy exists regarding the significance of nestin-expressing cells in islet neogenesis. Selander and Edlund did not find nestin expression in the pancreatic ductal epithelium [60], where the potential progenitors reside. Instead, nestin was detected in mesenchymal cells, called pancreatic stellate cells [61, 62] and was not detected during pancreas development [21]. Nonetheless, many groups have tested nestin selection protocols to derive β -like cells from mouse embryonic stem cells (MESC) later.

Pancreas and β -cell precursors from exocrine tissue

Human cells derived *ex vivo* from pancreatic exocrine tissue isolated from healthy donors dedifferentiated into a ductal phenotype (CK19- and CK7-positive) after adherence to culture surfaces [63]. After 2 days in culture, these cells reactivated *PDX1* expression, suggesting their pancreatic precursor potential; *PDX1* re-expression in ductal cells has been considered to be a prerequisite for their differentiation. This phenomenon of 'transdifferentiation' is questionable, since these *in vitro* studies are subject to ductal cell contamination [64]. However, a recent report has demonstrated that pancreatic exocrine cells can rapidly adopt a β -cell phenotype when the genome of these cells was altered and reprogrammed [65].

Zhou *et al.* (2008) developed a strategy to re-express multiple embryonic genes using adenoviruses in pancreatic exocrine cells of adult mice [65]. Nine genes that exhibit β -cell phenotypes when mutated were selected for initial reprogramming experiments. Of these, the combination of Ngn3, Pdx1 and MafA transcription factors could efficiently reprogram 20% of differentiated pancreatic exocrine cells into insulin-positive β -cells *in vivo*. Although emergent extra-islet insulin-positive cells did not appear to self-cluster as islets, angiogenesis was remarkably increased. Importantly, exocrine cells that adopted a β -cell phenotype improved hyperglycemia in STZ-induced diabetic mice. Taken together, those studies suggest that fully differentiated or even somatic cells could serve as an important source of tissue for generating functional insulin-producing cells in diabetic patients. Zhou's study was largely based on earlier studies that have provided the evidence that these lineage-committed cells can further be reprogrammed back to an ES cell fate, a characteristic termed induced pluripotency, or more commonly referred to as reprogramming.

β -cells from other adult tissues

The ability of adult stem cells to produce differentiated cells from embryologically unrelated tissues is an example of metaplasia and shows that embryological commitments can be changed or reversed under certain circumstances [66]. For example, both rodent and human bone-marrow stromal cells have been demonstrated to differentiate into other mesodermally derived tissues such as cardiac muscle cells [67], liver or neurons [68, 69], which are ectodermal in origin.

Although no significant *in vivo* differentiation of bone marrow into β -cells was observed in a study of adult mice [70], transdifferentiation from liver to pan-

creas has been seen. Ferber *et al.* (2000) demonstrated that transdifferentiation of liver to pancreas could be induced by ectopic expression of *pdx1*; transient expression of *pdx1* in mouse liver cells *in vivo* induced expression of the otherwise silent endogenous *insulin* gene [71]. A follow-up study indicated that insulin- as well as glucagon-producing cells were found mainly located in the proximity of central veins [72]. This indicates that *pdx1* plays an important instructive role in pancreas differentiation, not only from primitive gut endoderm but also to reprogram extra-pancreatic tissue towards a β -cell phenotype [72]. It was also sufficient to direct the production and secretion of mature, biologically active insulin from a restricted population of cells in liver *in vivo* [71]. Insulin secreted from the liver of *pdx1*-transfected mice ameliorated streptozotocin (STZ)-induced diabetes.

β -cell precursors from fetal pancreas

Islet-like cell clusters (ICC) cultured from human fetal pancreatic tissue (18 to 24 weeks gestation) were able to mature functionally and morphologically when grafted to kidney or pancreas in normoglycemic mice [73]. Additionally, exendin-4, used to treat those human fetal pancreas-derived ICC *in vitro*, upregulated *PDX1* expression [74]. Although insulin content did not increase in culture, ICC transplanted under the kidney capsule were found to give rise to grafts that had significantly higher levels of insulin compared to adult islets and completely reversed diabetes in STZ-treated nude rats.

In another study, nicotinamide treatment induced maturation of human fetal pancreatic islet cells, but not adult β -cells, and resulted in otherwise unresponsive glucose-stimulated insulin release [75]. Although fetal β -cells failed to proceed through the differentiation process to achieve a terminal phenotype in culture, this finding suggests that they are able to go through the maturation process *in vivo*. Taken together, these tissue culture models have provided groundwork for further studies to dissect the molecular mechanisms of islet cell differentiation from developmentally earlier multipotent or pluripotent stem cells such as ES cells.

Embryonic stem cells and β -cells

Derivation of embryonic stem (ES) cells and induced pluripotent stem (iPS) cells

Apart from the derivation of β -cells from various pancreatic stem cells in adult tissues and pancreas, pluripotent ES cells are another source to search for renewable β -cells. Mouse and human ES cells are de-

rived from the inner cell mass of pre-implantation blastocysts [76, 77], but they require different signaling pathways to maintain pluripotency. Recently, cells from epiblast of post-implantation mouse and rat embryos have also been shown to give rise to pluripotent stem cells. These epiblast stem cells share many features with human embryonic stem cells (HESC), and differ from mouse embryonic stem cells (MESC), suggesting that the differences between mouse and human ES cells are due to their relationship to different developmental stages, rather than distinct species differences in the control of pluripotency [78].

Assuming that techniques were available for the efficient production of β -cells from HESC, there are still other factors restricting its widespread use for application in therapy. Besides ethical concerns, because of their derivation from human embryos, difficulties remain due to tissue rejection following transplantation. Much effort has been spent in the development of alternative methods of generating patient-specific stem cells. One way to circumvent the problem is to reprogram adult cells back to pluripotent ES-like cells. Takahashi *et al.* (2006 and 2007) derived these so-called induced pluripotent stem cells (iPSC) by transducing first mouse, and then human adult dermal fibroblasts using retroviruses carrying four stem cell factors, namely *OCT4*, *SOX2*, *KLF4* and *C-MYC* [79, 80]. They later improved the protocol by eliminating *C-MYC* due to its function as an oncogene [81]. This potentially allows the generation of iPSC from patients with genetic disorders, providing a platform in which human diseases including diabetes mellitus may be studied *in vitro* [82], apart from the prospect of patient-specific cells that would obviate immune rejection in transplantation regimens.

β -cells differentiated from MESC by cell-trapping

In the first successful attempt to induce pancreatic differentiation from MESC, Soria *et al.* (2001) employed a cell-trapping system, followed by cell-lineage selection and maturation protocols to obtain insulin-producing clones [83]. The cells were transfected with a plasmid containing a neomycin resistance gene under the control of the *insulin* promoter, and hygromycin selection cassette driven by a constitutively active phosphoglycerate kinase (PGK) promoter. The hygromycin-resistant clones were cultured in suspension to induce embryoid body (EB) differentiation. Neomycin antibiotic pressure was applied later to select for cells expressing factors that activate the *insulin* promoter. In one clone, the cells released insulin in response to glucose, although the insulin content was

low. These insulin-positive cells normalized hyperglycemia when transplanted into the spleen of STZ-induced diabetic mice. In these mice, blood glucose levels remained normal and insulin-positive cells were present in the liver and in the spleen even after 42 weeks of transplantation [84].

Several genes, such as *nkx6.1*, bind to the *insulin* promoter and regulate *insulin* transcription in β -cells. It was reasoned that insulin-producing cells could be enriched if cells that activated its upstream promoters were isolated and expanded. Hence, the aforementioned approach was refined by replacing the *insulin* promoter with the *nkx6.1* promoter to allow the selection of cells expressing pro-endocrine genes that have an activated *nkx6.1* gene [85]. The hygromycin-resistant clones were first differentiated as EB in the presence of several exogenous agents. Treatment with nicotinamide, anti-Shh, and co-culture EB in the presence of pancreatic rudiments of E17.5 mice resulted in optimal *nkx6.1* expression. An almost pure population of insulin-positive cells was produced, of which 20% were Pdx1 co-expressing cells from several clones. Subsequent transplantation of these cells into STZ-induced diabetic mice normalized their glycemia. The same group later demonstrated that substitution of insulin for pancreas-conditioned medium did not result in differentiated β -cells, suggesting that the differentiation effect was mostly due to soluble factors released by the pancreatic rudiment [86].

β -cells differentiated through neural progenitors from MESC

Pancreas organogenesis shares similarities with the development of the nervous system, despite pancreas and CNS originating from different germ layers [16, 87, 88]. Pancreatic endocrine cells also share several characteristics with neurons, for instance, islet cells are electrically excitable [38, 87]. Additionally, insulin-producing cells have also been observed in the invertebrate nervous system [89-91] and in primary cultures of mammalian fetal brain [92]. Hence, many protocols for deriving β -cells from MESC were designed first to produce or select for neural progenitors defined by nestin expression [93] and then to direct pancreatic islet differentiation in subsequent steps. As described in the previous section, nestin has been detected in a small population of pancreatic cells that have been proposed as possible islet precursors [59, 87].

Lumelsky *et al.* (2001) demonstrated that a small proportion of insulin-producing cells could be derived from MESC with five sequential *in vitro* differentiation steps, during which cultures were highly enriched in

cells expressing nestin [94]. Nestin-positive cells derived from EB were expanded in B27/N2 neurobasal medium supplemented first with bFGF and subsequently cultured in nicotinamide. The outcome of their protocol was the formation of cell clusters connected by a web of neurons, with 10-30% insulin-expressing cells localized inside these clusters. Although the differentiated cells were immunoreactive to insulin, RT-PCR revealed that *insulin2*, *glut2* and *pdx1* mRNA levels were low and β -cell-specific *insulin1* mRNA was never detected. These insulin-positive cells contained 50 times less insulin than normal islet cells. Not surprisingly, with such small amounts of insulin, the cells failed to correct hyperglycemia when grafted into STZ-treated diabetic mice. Nonetheless, the transplanted cells underwent rapid vascularization. The grafted mice were able to maintain their body weight and survived for longer periods of time than hyperglycemic sham-grafted controls.

With minor modifications in Lumelsky's differentiation protocols, populations of nestin-expressing MESC derivatives differentiated into insulin-producing cells showing increased levels of β -cell transcripts and insulin production. Hori *et al.* (2003) substituted the phosphoinositide 3-kinase (PI3K) inhibitor, LY294002 for B27 at the later stage of differentiation [95]. LY294002 has been shown to increase total endocrine cell number and insulin content from the human fetal pancreas [96], as well as to prevent neurite outgrowth from neuroendocrine cells [97]. This modification yielded insulin-producing cell clusters (IPCC) with morphological similarities to pancreatic islets. Immunocytochemistry staining showed that Pdx1, GLUT2 and glucokinase were co-expressed by insulin-positive cells. However, the cells were also distinct from islets in numerous ways; while 95-97% of the cells were insulin-positive, and 2-3% were glucagon-positive, no staining for somatostatin and PP was detected. Nonetheless, these IPCC increased circulating insulin levels, reduced weight loss, improved glycemic control and completely rescued the survival of diabetic mice. Intriguingly, when LY294002 was not used, transplanted cells formed tumors that resembled teratomas, suggesting that undifferentiated cells were still present.

The discrepancy in *insulin1* mRNA and insulin protein expression, and the lack of C-peptide release in these two studies leads us towards the hypothesis that cells differentiated with the nestin protocol take up exogenous insulin from the media. By adding fluorescein-labeled insulin in the media, cells were found to absorb these insulin molecules, hence were immunoreactive to anti-insulin antibody. An explanation for

this active insulin uptake is the prevalence of apoptosis and necrosis due to suboptimal culture condition. Although these cells release insulin in response to glucose, C-peptide release was never detected, suggesting that *de novo* insulin synthesis was not a common outcome of cells differentiated from nestin-expressing neuroendocrine progenitors [98, 99].

Later, Blyszczuk *et al.* (2004) reported that MESC can be induced to differentiate into insulin-producing cells without employing the 'nestin-selection' step [100]. In this study, the use of ITSF (insulin, transferrin, selenium and fibronectin) and bFGF were avoided after replating EB on monolayer, so that selection or enrichment of specific cell types before induction of

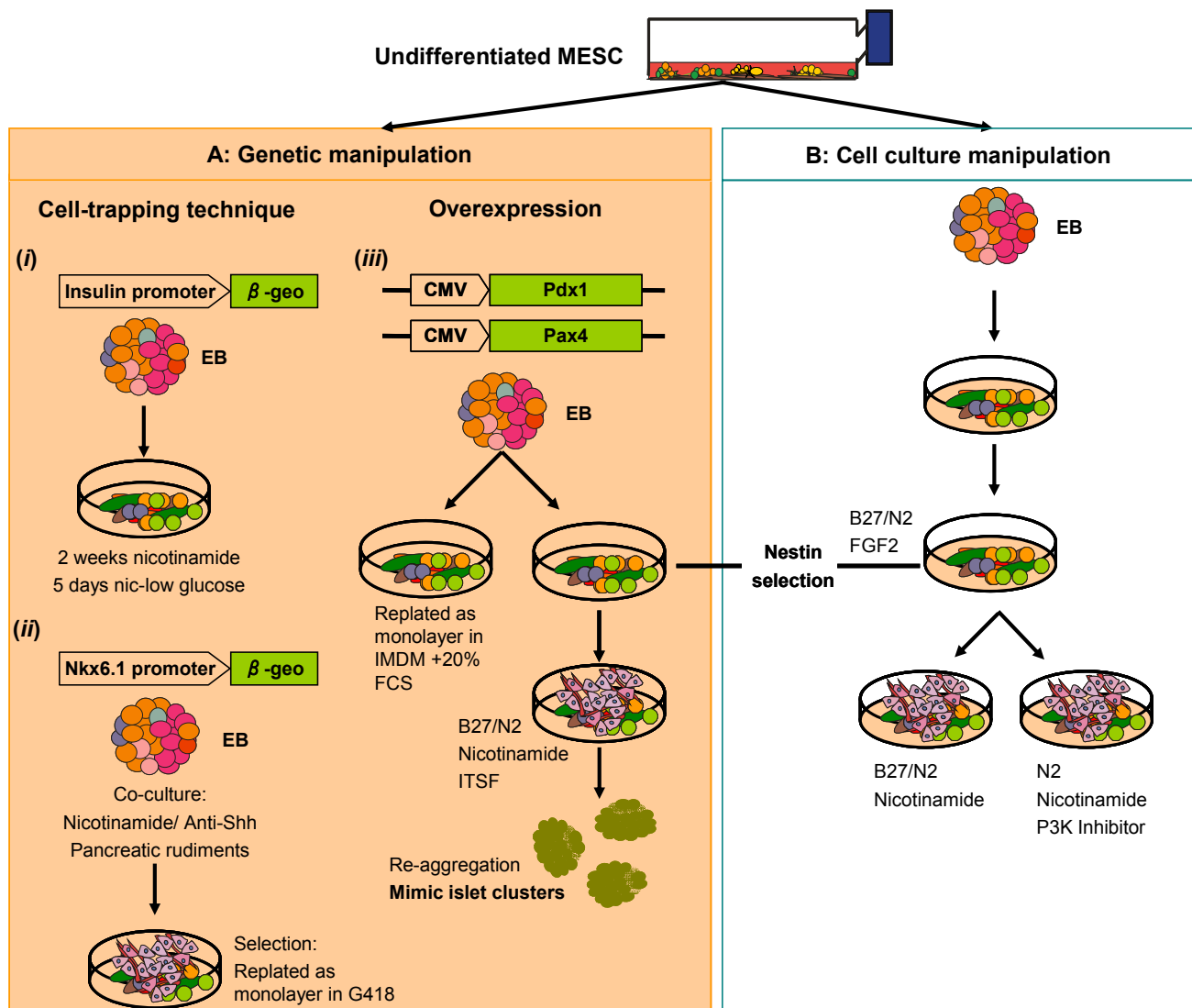


Figure 1. *In vitro* differentiation of putative β -cells from mouse embryonic stem cells (MESC). A: Genetic manipulation. *i*: Soria *et al.* (2001) used a 'cell-trapping' protocol to select for insulin-producing cells expressing the β -geo gene under the control of the human *Insulin* gene promoter [84]. This strategy was later refined by placing the β -geo gene under the control of the promoter of *Nkx6.1* (*ii*). *iii*: Enforced expression of Pax4 or Pdx1 increased the frequency with which insulin-producing cells were isolated from differentiating MESC. **B:** Cell culture manipulation of ES cells. Lumelsky *et al.* (2001) developed a five-step protocol based on methods known to promote the generation of neural cell types from MESC [94]. Nestin-expressing cells were cultured in B27/N2 neurobasal medium went on to form cell clusters. Although the nature of insulin-staining cells derived by this method remains controversial, other groups have successfully used variations on this procedure to generate similar cell types. EB: embryoid bodies.

pancreatic differentiation would not occur. Functional β -like cells were generated following pancreatic differentiation by serum-free medium containing nicotinamide and laminin. However, ITSF were still added in the later stage of the differentiation protocol to compensate for serum-free culture conditions. Although it is not clear whether the addition of ITSF to the medium would select for nestin-positive cells, they concluded that β -cell precursors transiently expressed nestin during differentiation *in vitro*.

Exogenous expression of β -cell transcription factors

Constitutive expression of transcription factors with a role in pancreatic development such as *pdx1* or *pax4* has also been used to enhance differentiation of insulin-producing β -cells from MESC [101]. During *in vitro* differentiation, important changes in expression levels of pancreatic genes were detected in Pax4-positive cells, and to a lesser extent, in Pdx1-positive cells. *LAPP*, *glut2* and *insulin* transcripts were upregulated in Pax4-positive cells at a later stage of differentiation, although *ngn3* expression remained constant [101]. There was also a significant increase in the number of nestin-expressing cells in Pax4-positive cells, and this increase was corroborated by an increased level of intracellular insulin. As expected, no significant differences in glucagon-expressing cells were observed. Pax4-positive cells were then cultivated in Spinner rotation cultures to promote their histotypic maturation into spheroid islet-like clusters grown in suspension. Electron microscopy detected insulin-labeled secretory granules in Pax4-derived cells, and notably, these cells were able to normalize blood glucose levels in STZ-treated diabetic mice.

Inducible expression of Pdx1 in MESC also resulted in increased expression of pancreatic endocrine transcription factors during EB differentiation [102]. A MESC line, in which exoge-

nous Pdx1 expression was regulated by a tet-off system integrated into the ROSA26 locus, was employed to examine the effect of Pdx1 expression during *in vitro* differentiation using a protocol similar to Lumelsky's protocol. For differentiation, cells were grown in the absence of doxycycline to induce the expression of Pdx1 and incubated with the addition of B27/N2, nicotinamide, KGF and EGF. The results showed that Pdx1 induction enhanced the expression of the *insulin*, *somatostatin*, *Kir6.2*, *glucokinase*, *ngn3*, *p48* and *pax6* genes. An overview of the methods used to generate β -cells from MESC is illustrated in Figure 1.

Early progress with human ES cells (HESC)

Expression of genes characteristically involved in the establishment of the mouse embryonic pancreas, including *NGN3* and *PDX1* transcription factors as well as *INSULIN*, *GCK* and *GLUT2* were detected by RT-PCR analysis in HESC cultures during EB and monolayer culture differentiation [103]. Immunohistochemistry analysis revealed that 60-70% EB contained insulin-expressing cells, although only up to 2% of cells in an EB stained for insulin. Not surprisingly, with such low numbers of insulin-positive cells, further characterizations were impossible. Furthermore, glucose-responsiveness of these insulin-positive EB could not be determined.

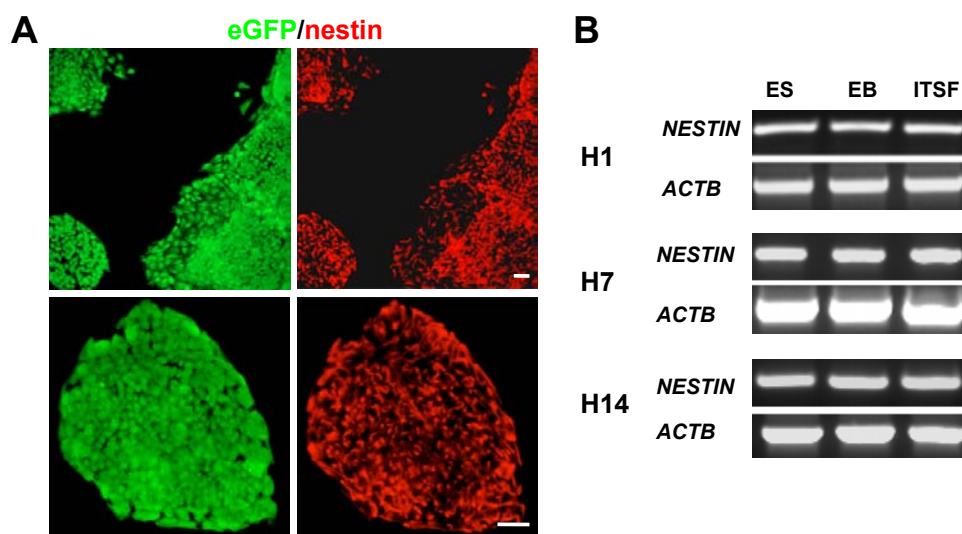


Figure 2. Nestin expression in human embryonic stem cells (HESC). **A:** Fluorescent microscopy showing that HESC colonies were immunoreactive to nestin. **B:** RT-PCR analysis of *NESTIN* expression in H1, H7 and H14 HESC lines during *in vitro* differentiation. *NESTIN* transcripts were detected in undifferentiated stem cells and embryoid bodies (EB). There was no significant difference in the levels of *NESTIN* expression when the cells were grown in ITSF serum-free medium, indicating that ITSF medium did not increase the expression of *NESTIN* in HESC. Scale bars: 50 μ m.

Three years after the first report with MESC was published, it was shown that HESC could be induced to form insulin-producing islet-like clusters similar to immature pancreatic β -cells [104]. The differentiation strategy in this study was a modification of the 'nestin-selection' method. Similarly, HESC were differentiated in EB and replated on monolayer to allow the growth of nestin-positive neuroendocrine progenitors. Similar to MESC, differentiated cells expressed low insulin. However, by reducing the glucose concentration in later stages of differentiation and by reaggregating these clusters in suspension, insulin expression and secretion was increased. Insulin-producing clusters produced by this method resemble immature pancreatic embryonic cells. A substantial number of cluster cells were co-stained for insulin and glucagon or somatostatin. RT-PCR detects the expression of immature islet genes such as *NGN3* but low expression of *ISLET1*, *PDX1* and *GCK*. Although they showed a response to the different β -cell agonists and antagonists, their responsiveness to glucose was minimal. According to our experience, nestin is expressed by undifferentiated HESC and its expression remained constant during differentiation (Figure 2). Thus, the protocol to differentiate β -cells using nestin selection should be reevaluated considerably.

Co-culture of HESC with mouse embryonic pancreas

Given the inefficiency and controversies for deriving β -cells from nestin-expressing neuroendocrine progenitors, a more plausible approach is to direct ES cells through a process that mimics normal pancreas development. HESC grown to confluency gave rise to spontaneously differentiated cells that yielded FOXA2 and PDX1 co-expressing definitive endoderm (DE) cells at the periphery of the colonies. These early pancreatic progenitors were manually picked and co-transplanted with E11.5 mouse embryonic dorsal pancreas under the kidney capsule of SCID mouse. Eight weeks after transplantation, clusters consisted mainly of proinsulin-expressing β -like cells and a few glucagon and amylase-secreting cells were detected in the grafted section. Intriguingly, transplanting DE cells alone did not result in the appearance of β -like cells. This finding suggests that the stimulatory cues from co-grafted pancreas and an *in vivo* milieu were necessary to coax HESC to differentiate into β -cells.

β -cells from HESC-derived definitive endoderm cells

Studies with MESC revealed that by restricting serum in the presence of activin A in media induced the

formation of Brachyury-expressing mesendoderm cells in EB, from which endoderm cells may develop later [15]. D'Armour modified this protocol by inducing HESC to differentiate in monolayer in the presence of 100ng/ml activin A and low serum [105]. 80% of cells in activin A-treated cultures were immunoreactive to FOXA2 and SOX17, suggesting the enrichment of DE cells. High activin/nodal, but reduced insulin/insulin-like growth factor (IGF) signaling and suppression of PI3K-dependent pathway, are critical for the development of HESC to DE cells [106].

These DE cells can further differentiate into pancreatic endocrine cells in the presence of Hedgehog and Notch-signaling inhibitors [107]. In brief, KAAD-cyclopamine and FGF10 were used to promote endoderm patterning and proliferation from DE cells. The outcome of this stage was the generation of PDX1-expressing gut-tube endodermal like cells. These cells were subsequently cultured in retinoic acid (RA), α -secretase inhibitor and exendin-4 to promote the transition towards hormone-expressing pancreatic endocrine cells. This protocol generated 7-12% of insulin-positive cells, and similar to fetal β -cells, they released C-peptide in response to depolarizing reagents such as potassium chloride (KCl) and tolbutamide. Nevertheless, there was only minimal glucose-induced C-peptide release. With a slight modification to this protocol, Jiang *et al.* (2007) generated islet-like clusters from feeder free HESC-derived DE cells in low adherent cultures to form aggregates in the presence of noggins and nicotinamide [107]. Similarly, these aggregates produced cells secreting various pancreatic endocrine hormones, and notably, released C-peptide in response to glucose. The use of xeno-free HESC will be prerequisite for utilizing these cells in stem cell therapies [108, 109].

It is reasoned that *in vivo* maturation might be necessary for the functional maturation of HESC-derived pancreatic endocrine or β -cells. Pancreatic endocrine cells, owing to their precursor status, may be more potent than β -cells in undergoing maturation should they respond to *in vivo* milieu. Harvested pancreatic endocrine cells went on to form aggregates on gelatin foam sponges and matrigel overlays. These cells were then harvested and grafted into STZ-treated immunocompromised mice. Although no significant level of human C-peptide was detected within the first month, long term engraftment data revealed that glucose-stimulated serum levels of human C-peptide increased rapidly during the next sixty days [110]. Nonetheless, because purification steps were not used, teratomas containing various other tissues of ectodermal and

mesodermal were formed in most of the animals. Thus, future improvement will require purification of pancreatic cell types and depletion of inappropriate cell types prior to transplantation.

β-cells isolated from Pax4-expressing HESC

Our own laboratory has described an efficient method to enhance β-cell differentiation from HESC [111]. We first overexpressed Pax4 in HESC from a

constitutively active CAG (human cytomegalovirus major immediate early enhancer- chicken β-actin- rabbit β-globin) hybrid promoter and induced differentiation in EB model. We reasoned that a proportion of DE cells spontaneously differentiated from EB, substantiated by the detection of *FOXA2* and *NEUROD1* transcripts. Various key genes expressed in mature β-cells such as prohormone convertase 1/3 (*PC1/3*), *GCK* and *GLUT2* were upregulated in Pax4-expressing

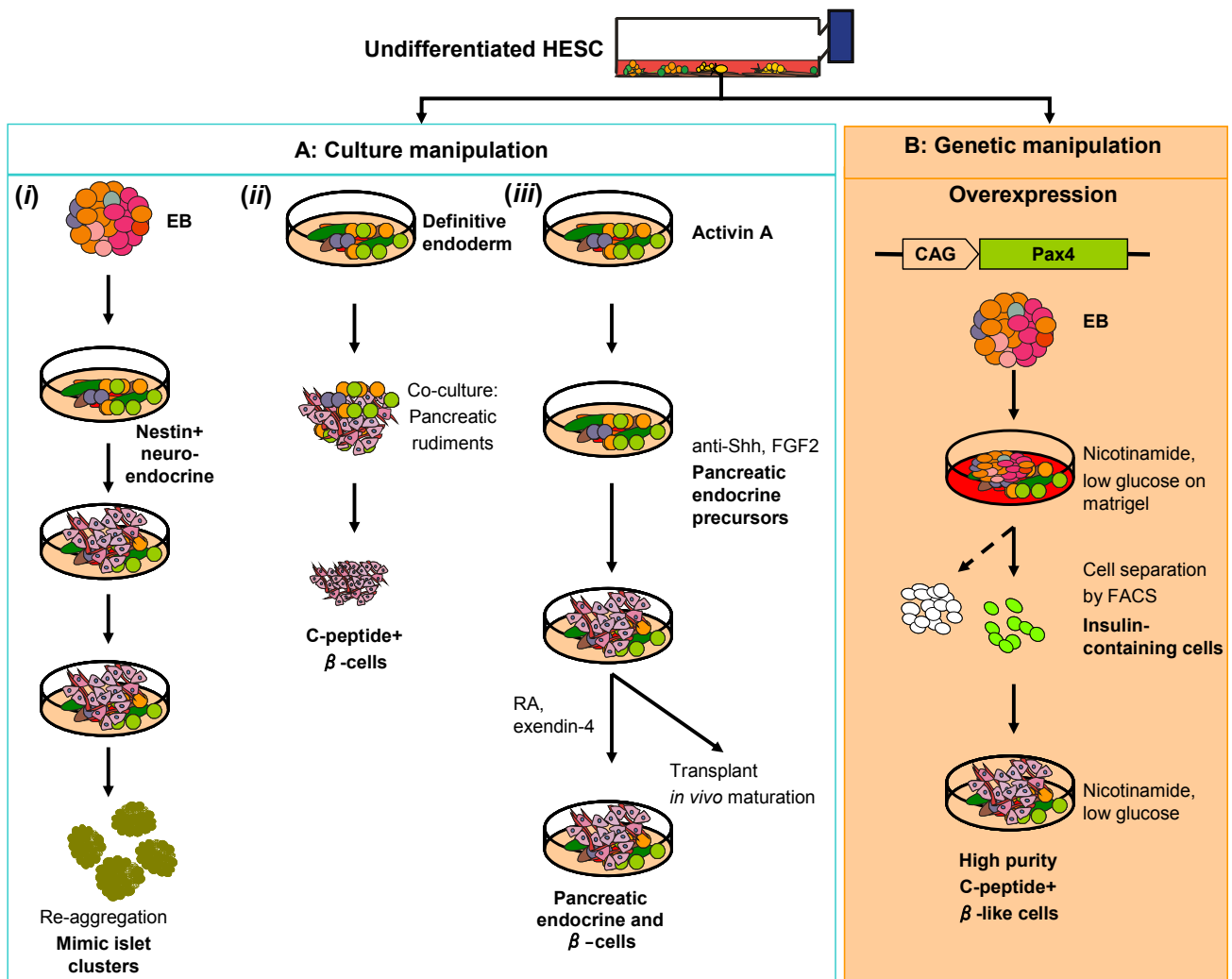


Figure 3. Overview of studies on generation of putative β-cells from HESC. **A:** Cell culture manipulation. *i:* Nestin-expressing cells from EB, similar to those differentiated from MESC gave rise to insulin-immunoreactive cells. Cells were re-aggregated in suspension cultures to yield islet-like clusters consisted of cells expressing pancreatic endocrine hormones. *ii:* Definitive endoderm (DE) cells differentiated at the edge of HESC colonies were manually picked, co-transplanted with mouse embryonic pancreas further differentiated to β-cells *in vivo*. *iii:* A 2D culture generated DE cells by activin A treatment. DE cells responded to hedgehog and Notch inhibitors, giving rise to cells that express pancreas endocrine hormones. Cells transplanted in STZ-induced mice ameliorated hyperglycemia. **B:** β-like cells were enriched from Pax4-expressing EB and replated on monolayer with nicotinamide and low glucose. A zinc chelating dye was used to isolate insulin-containing cells. These cells went on to form C-peptide-positive cells that responded to β-cell secretagogues. It remains to be determined whether these isolated β-cells rescue hyperglycemia in a mouse model.

EB. These cells also expressed markers of voltage-gated Ca^{2+} -channels (VGCC) at comparable levels to human fetal pancreas. In addition, the increase in VGCC genes correlated with an enhanced responsiveness of EB to muscarinic receptor agonists such as KCl. When replated on monolayer, EB went on to form putative β -cells. These cells could be isolated by fluorescence activated cell sorting (FACS) using Newport Green dye based on the high zinc content in insulin-containing cells. Purified β -like cells were enriched in *PDX1* and *INSULIN* transcripts but depleted in *OCT4* expression, suggesting the absence of non-pancreatic cell types and residual undifferentiated cells. Isolated cells also synthesized proinsulin, expressed C-peptide and responded to tolbutamide. Whereas this study shows the possibility to increase the differentiation of β -cells from HESC, further work is required to characterize *in vivo* maturation and function of these cells. A summary of the methods used to generate β -cells from HESC is illustrated in Figure 3.

Generally, the formation of a β -cell from a pluripotent stem cell represents a part of the complex choreography of embryonic development. If the formation of a β -cell *in vivo* is to be strictly followed, expression of key regulatory genes for this sequential commitment should be regulated in a time-dependent manner. A major challenge is that ES cells support the early stages of endoderm and pancreatic endocrine cell formation *in vitro* but do not robustly generate glucose-responsive β -cells, making diabetes treatment using stem cell therapy untenable. Inefficiencies in deriving a scalable source of β -cells from HESC *in vitro* likely reflect the inability of undifferentiated stem cells to mimic these normally intricate *in vivo* gene

expression patterns. Furthermore, published studies so far revealed that HESC-derived β -cells were not glucose-responsive, but acquired further maturation once transplanted into mice. These findings add support to the contention that these cells mimic β -cells in the human fetal pancreas. In order to recapitulate pancreas organogenesis *in vitro*, an improved three-dimensional EB differentiation model and manipulation of extrinsic signals may be employed to support the formation of pancreatic mesenchyme, acini, islets and β -cells in a spatially and temporally controlled pattern.

Future prospects of stem cell therapy using HESC- or iPSC-derived β -cells

Several findings have underscored the necessity for deriving more HESC for research purposes. As HESC

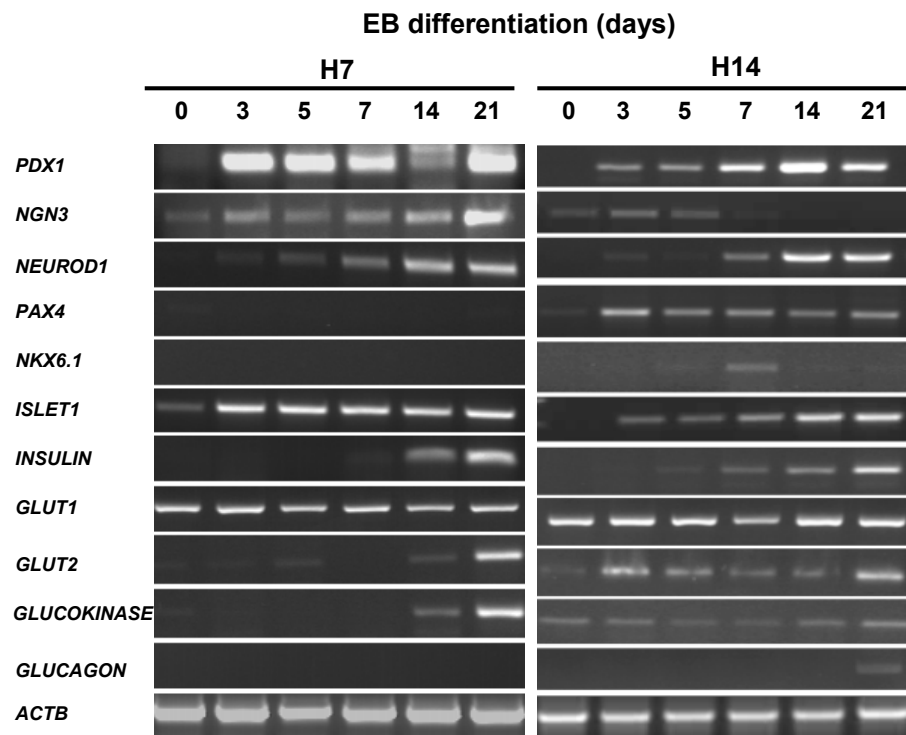


Figure 4. RT-PCR analysis of β -cell gene expression in two different HESC lines (H7 and H14) during *in vitro* differentiation. 0-day cells are undifferentiated stem cells. Embryoid bodies (EB) of different stages were collected for RNA extraction. There were transient expressions of *NGN3*, *PAX4* and *NKX6.1* transcription factors at early stages of H14 EB differentiation. However, the expressions of *PAX4* and *NKX6.1* in H7 EB were minimal. Up-regulation of *NEUROD1* and *ISLET1* transcripts was detected in the course of EB differentiation. Differentiated EB expressed *INSULIN*, *GLUT2* and *GLUCOKINASE* β -cell markers at a later stage of EB differentiation, suggesting that H7 and H14 cells were able to produce β -cells in the course of spontaneous differentiation. There was no change in *GLUT1* transcript levels during EB differentiation. *GLUCAGON* transcripts were only detected in H14 EB.

lines are derived from a genetically heterogeneous population, their biological variations, heterogeneity, genetic and epigenetic differences contribute to the variations in developmental potential. Various studies have reported that HESC lines exhibited marked differences in differentiation propensity into specific lineages [112, 113]. Notably, genes that play the key role in maintaining pluripotency, such as *OCT4*, *NANOG* and *CRIP1* showed little variation; however, marker expression of multiple tissue-specific lineages became variable after the cells began to differentiate [114]. Some lines exhibited a marked potential to differentiate into one lineage, often with more than 100-fold differences in lineage-specific gene expression. When gene expression levels of the same HESC lines with different passage numbers were compared, consistent patterns within the same line but significant variations between lines remained. This observation argues against the possibility that variability of culture technique or senescence of HESC contributed to marked differences in differentiation propensity in different lines. We have also observed variations of pancreatic lineage differentiation potential from different HESC lines (Figure 4). Thus, it would be advantageous to work with HESC that are inclined toward pancreatic differentiation.

The derivation of iPSC opens new possibilities to reprogram pluripotent stem cell lines from individual patients to avoid immunological rejections in cell therapy. However, time and costs necessary for derivation, differentiation and safety testing of good manufacturing practice (GMP)- or clinical-grade iPSC from each individual may exceed practical and economical limits. Another concern is the age of the patient; iPSC obtained from young adults are healthier than those derived from mature adults.

An attractive alternative to eliminate the age concern and to reduce the likelihood of graft rejection in stem cell therapy is through the use of human leukocyte antigen (HLA)-haplotype cell banking, from which a best match could be selected. The results of pancreas transplantation will also be improved by minimizing HLA mismatches [115]. Recently, Taylor *et*

al. (2005) estimated the required number of HESC lines for beneficial HLA-matching in the UK. They proposed that about 150 HESC lines would be needed for most of the UK population, and that as few as ten might be sufficient if one were to prospectively identify cell lines that could serve a larger number of patients, such as lines homozygous for common HLA types [116, 117]. Based on estimates using Japanese population, Nakatsuji *et al.* (2006) also estimated that with a bank of HESC lines from 170 randomly selected donated embryos, 80% of the patients could be expected to find at least one HESC line with only a single mismatch at one HLA locus or even a better match [118]. Thus, the production of HLA-mapped clinically-safe iPSC would greatly minimize the ethical concern and enable the creation of more efficient HLA-haplotype banks. Therefore, in cases when stem cells cannot be obtained from the patient themselves (e.g., in an aged patient), iPSC lines derived from an unrelated individual may eventually become alternative sources of various cell types for transplantation therapy.

However, whether cell transplantation will ultimately be the approach of choice for regenerative medicine is a moot point. Many technical challenges remain to be overcome, and it is possible that a better understanding of the basic biology of the pancreas, from the point of view of both its development and subsequent homeostasis, may offer insight that will provide alternative approaches to curing diabetes and related diseases. Studies of ES cells and other stem cells may well play a substantial role in acquiring the necessary knowledge, and the development of patient-specific iPSC certainly offers a new approach to advancing disease models as well as tools for screening new drugs that could play a role in novel treatments in the future.

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