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β-Cell Growth and Regeneration: Replication Is Only Part of the Story

Susan Bonner-Weir, Wan-Chun Li, Limor Ouziel-Yahalom, Lili Guo, Gordon C. Weir, and Arun Sharma

“I am accordingly of the opinion that the normal regulation of islet content in the pancreas is by interstitial growth of pre-existing islets and by the formation of new islets from the duct epithelium, and not at all by the formation of new islets out of acini.”

R.R. Bensley
Am J Anatomy 1911;12:297–388

For almost a century (for historical review, see 1) both β-cell replication and neogenesis (the differentiation of new islet cells from progenitors or stem cells) have been thought to be responsible for postnatal growth of the endocrine pancreas. Even though doubts have been raised in recent years about the existence and importance of neogenesis, this skepticism may be subsiding. Replication and neogenesis are not mutually exclusive. Both processes often occur simultaneously, as seen during the regeneration that follows pancreatic injury. However, there are important differences in the balance of these two pathways that depend upon species and age. Replication of β-cells is an important mechanism particularly in adult rodents, but there are compelling data that after-birth progenitors also have a role in renewal and growth of islets. Eventually one or both of these pathways may be manipulated for therapeutic treatment of diabetes. Since we and others have extensively reviewed the regulation of β-cell mass (2,3), this Perspective will specifically address the contributions of the neogenic pathway to new β-cell formation, considering whether postnatal neogenesis occurs, to what extent, possible differences between mammalian species; and whether it might be exploited therapeutically.

β-Cell expansion in normal growth. The concept that β-cell mass is dynamic and increases and decreases both in function and mass to maintain the glycemic level within a very narrow physiological range (4) is now generally accepted. In both normal and pathophysiological states, the mechanisms responsible are changes in replication and neogenesis, changes in individual cell volume, and changes in cell loss or death rates. In rodents, β-cell replication is significantly higher during late gestation and the neonatal period than after weaning and changes little beyond 30–40 days of age except in response to physiologic/pathological changes (2,3,5). Other than during the neonatal remodeling of the endocrine pancreas that occurs before weaning (6), the frequency of apoptosis is low and unchanging (7). In the initial months of life, β-cell mass increases with body weight through increases in both β-cell number and size, but in old (15–20 months) rats the increment in mass is largely accounted for by increased β-cell size (7). While replication and apoptosis of the β-cell have been the focus of many studies in the last decade, changes in cell volume are usually not measured; however, changes have been well documented in a number of other models (8–10). In the case of neogenesis, there is strong evidence that the process occurs normally and can be activated in mouse, rat, pig, and human.

The concept of neogenesis or the formation of new islet cells from pancreatic progenitor/stem cells after birth has been built upon many observations from different models and species over many years. An appealing assumption has been that the islet hormone-positive cells within the ductal epithelium or that appear to be “budding” from the ducts represent neogenesis (Figs. 1 and 2). However, it can be argued that they are static with no active “budding” or neogenesis (11). Other than lineage tracing, no direct means of showing a dynamic process of neogenesis exists, but with the limited labeling efficiency of current lineage-tracing approaches, detection of marked cells rather than the lack of such cells represents stronger evidence. Many studies have quantified neogenesis as increased hormone-positive cells within ducts or small clusters of hormone-positive cells in the parenchyma that accompany growth whether normal growth, after surgical perturbation or addition of various growth factors and/or cytokines. These data have strengthened the hypothesis that this budding appearance actually represents the dynamic process of neogenesis, such that hormone-positive cells within ducts may be used as quantifiable markers. Table 1 shows more quantitative studies over the past several decades; more detail is provided below about some of these models. It should be noted that not all rodent models of β-cell growth show neogenesis: no increases in islet number were seen in pregnancy in mice (12) or in adult ob/ob mice compared with ob/+ controls (13); both of these studies showed increased β-cell mass with larger islets, supporting replication as the mechanism of expansion. One must be aware of possible species differences, as illustrated by neogenesis not being reported in adult mice fed a high-fat diet but reported in adult pigs that became obese after ad libitum feeding for 1–2 years (14). Additionally, in contrast to the data from rodents, the adaptive increase in β-cells during pregnancy in humans is accompanied by decreased mean islet size, increased number of β-cells in “apparently new small islets” with no change in β-cell size, replication, or apoptosis frequency (15).

However, largely based on Dor et al. in 2004 (11), the occurrence of neogenesis has been questioned (16,17).
The Dor et al. study, in which genetic marking of β-cells in rat insulin promoter:CreER mice was not diluted over several months of chase in adult mice, showed that replication as a major mechanism of β-cell expansion in adult mice. However, with possible low levels of leakiness of the Cre-lox system over time with the strong insulin promoter (18), the small proportion of the islets counted, and only 30% labeling of the β-cells, neogenesis in the adult cannot be ruled out by negative data of a lack of dilution of labeled β-cells. Furthermore, the common extrapolation of these data that neogenesis does not occur after birth is flawed since neither the neonatal period nor the new lobes after pancreatectomy were examined.

In building a case for a given hypothesis, it has become fashionable to rely more heavily upon genetic methodology than correlative observations. However, each approach has its strengths and limitations. In the case of neogenesis, years of strong observational support had now been supplemented by molecular tracing experiments, with two strong studies (19,20) in 2008 that complement past evidence for postnatal neogenesis.

**Neogenesis in normal growth during the neonatal period.** During the fetal stage, differentiation is the major mechanism for forming new β-cells, but replication or self-duplication is enhanced during the perinatal period. During the neonatal period, β-cell replication continues and significant neogenesis occurs under these normal physiological conditions, as has been demonstrated using various methods, including Cre-lox lineage tracing experiments.

Bouwens et al. (21) studying neonatal rat pancreas in the first 2 weeks after birth suggested that cytokeratin 20+ (a marker of rat duct epithelium) cells at the periphery of islets served as islet progenitors. Between 2 and 5 days of age, the β-cell mass more than doubled, but the cross-sectional area of the β-cells was unchanged, indicating that new cells rather than larger cells accounted for the increased mass. The BrdU incorporation in β-cells of 2.4% at day 2 could account for only 12% of the observed growth by day 5. It was concluded that most of the new β-cells originated from neogenesis. Their observation that CK20+ cells had higher BrdU incorporation than hormone-positive cells is consistent with the finding that duct cell replication precedes neogenesis (22).

Our mathematical modeling study (5) predicted two waves of neogenesis: one immediately after birth and the other 2–3 weeks after birth. This model used existing data on β-cell mass and its determinants (cell volume, replication, and apoptosis frequency) to estimate turnover of β-cells. In our subsequent longitudinal study of β-cell mass and its determinants over the first month after birth (6), we documented the increased appearance of islets budding from the ducts at the same times (shortly after birth and just before weaning), confirming the predicted waves of neogenesis. Using these data to estimate the number of β-cells at each time point, we estimated that ~70% of the β-cells seen at day 31 could be accounted for by replication of preexisting β-cells, while the remaining 30% were from neogenesis (22). This estimate is consistent with the
findings from our duct-specific lineage tracing experiments described next (20).

Carbonic anhydrase II (CAII), which only starts to be expressed in ducts at the very end of gestation (23), has been considered a marker for ducts to distinguish the mature ductal phenotype from the embryonic tubular structures often called embryonic ducts. Thus, bigenic CAIICre:ROSA26R mice can provide duct-specific lineage tracing with only genetically marked ducts expressing the reporter at birth with no CAII or Cre expressed in /H9252 cells (20). We showed that in the 4 weeks after birth, both islets and acini were formed from cells that once expressed CAII: 38% of the islets were marked (17% of all insulin cells) as well as a number of acinar cells; some lobes were marked and others not. Since the pancreatic weight increases fourfold between day 17 and day 31 (6), we interpret the lobular pattern of marked islets and acini as evidence of new lobe formation in the neonatal period. It should be noted that two recent lineage-tracing studies using inducible Cre-ER driven by either the hepatocyte nuclear factor (Hnf) 1β (24) or mucin1 promoter (25) found no marked islet nor acinar tissue in neonates when tamoxifen was administered at the end of gestation (24) or birth (25). These studies found only marked ducts during this time of rapid pancreatic expansion, but their negative data may be due to 1) their low efficiency of marking ducts (20 and 7.6%, respectively) and only ~1,000 insulin+ cells counted per animal and 2) a marked heterogeneity within the ducts of the expressed product of their driver gene.

Recently, Peng et al. (26) published studies that showed an increasing number of islets from 1 week to 2 months of age. The obvious caveat for enumeration of islets over time in that small clusters of islet cells, which initially may have been below the measurable limit, may increase in size due to proliferation and be counted as newly formed. Interestingly, they also report that Ki67 mRNA and telomere length were significantly correlated in single islets at either 2 weeks or 4 months but differed for islets within the same animal, suggesting islets of differing ages in adult mice.

If new islets were generated from preexisting ductal tissue, one might expect expression of residual ductal markers for a short period of time after hormone expression. Indeed, transient expression of markers of duct cells has been demonstrated in β-cells of newborn rats (27) as well as in regenerates islets after partial pancreatectomy (22,28), in grafts of purified human duct cells (29), and in islets of mice conditionally expressing Pax4 in glucagon-producing cells (POE::GluCre) (30), suggesting their recent passage through the ductal phenotype.

Models of regeneration associated with neogenesis. Both partial pancreatectomy (31) and partial duct ligation (32), classic models of regeneration in rodents, are followed by both β-cell replication and neogenesis. Adding to the weight of evidence, newly formed or “budding” islets have been found in a number of other experimental conditions (Table 1), including dietary treatment with soybean trypsin inhibitor (33), after glucagon-like peptide (GLP)-1/exendin-4 treatment (8), treatment with betacellulin (34), and after cellophane wrapping of the head of the pancreas (a partial duct obstruction) (35). Additional strong evidence of formation of new β-cells from differentiation of progenitors comes from a number of transgenic models, including overexpression of interferon-γ in the β-cells of transgenic mice (36), overexpression of transforming growth factor (TGF)-α in pancreatic ducts (37), and Pax4 ectopic expression in glucagon-positive cells (30). These regeneration models may not be normal physiology but may show pathways that can be exploited, as much as transgenic and knockout mice are hardly physiological but quite informative.
Partial pancreatectomized rodents. A 90% pancreatectomy in rats demonstrates the substantial regenerative capacity of the adult pancreas (31,38,39). By 8 weeks after surgery, the 10% remnant had regenerated such that the pancreatic weight was 27% and the β-cell mass was 45% of the sham-operated pancreas. This was accounted for by both increased β-cell proliferation and formation of whole-new lobes of pancreas that contained newly formed islets (31,40). Expansion of the ductal tree by replication could be demonstrated with BrdU incorporation and subsequent appearance of branching ductules in well-defined areas (foci of regeneration) that correspond to new lobes. The pancreatic duct cells after replication transiently express pancreatic duodenal homeobox (PDX)-1 protein (41) and lose expression of key markers of the mature duct phenotype (40). The appearance of these foci regions and their disappearance with maturation coincide with the marked growth of the pancreatic remnant. Within a single pancreas, there are multiple focal areas of varying stages of maturation, as assessed by their histological appearance and gene expression profiles. Ductules in these foci express many markers of embryonic pancreatic progenitors, including transient expression of the
endocrine lineage-specific transcription factor neurogenin 3 (40). Islets in these foci resemble embryonic islets with higher proportion of insulin+-cells lacking MafA expression; the proportion of MafA+/insulin+ cells increases as foci differentiate further, but even in mature foci the proportion of MafA-expressing β-cells was lower than in the remnant pancreas of the same animals (40). We conclude that in response to pancreatectomy, pancreatic duct cell clusters recapitulate aspects of embryonic pancreas differentiation and contribute to the regenerating pancreas.

After partial pancreatectomy, regeneration is not limited to young animals. Both replication and neogenesis could be demonstrated in retired breeder (500 g) rats subjected to the same surgery (42). Whole-new lobes of pancreas may even be formed spontaneously in normal adult animals; occasional pancreatic lobes with high BrdU incorporation were seen in 6-month-old rats when most of the pancreas had almost none (5,22). When the extent of resection of pancreas is less, there is less regeneration (43). Even so, Jetton et al. (44) have utilized the 60% pancreatocex in adult rats extensively and reported enhanced β-cell proliferation and neogenesis.

The partial pancreatectomy model has been transferred to mice using less extensive resection, often only 50%. In several reports, only enhanced replication was reported (11,16,45), but two groups using 60% pancreatectomy reported neogenesis in addition to enhanced replication (46,47). These latter studies have even defined some of the regulatory pathways involved in neogenesis. Using 70–80% pancreatectomy in C57BL/6 mice, we obtain mild to moderate hyperglycemia and have found the same regeneration pattern as in 90% pancreatectomy rats (Fig. 1F) (48).

Partial duct ligation in rodents. The partial duct ligation has been used for several decades in rats (32,49–52). Wang et al. (32) did a comprehensive study of this model and showed that the β-cell population distal to the ligation doubled in 1 week and that the labeling index with BrdU pulse labeling in β-cells could not account for this increase. With increased number of small islets and islet cell clusters in the distal ligated portion, they concluded that there was islet neogenesis. A value of this model is that regeneration is limited to the portion distal to the ligation, there was islet neogenesis. A value of this model is that regeneration is limited to the portion distal to the ligation, allowing an internal control. In later experiments, this model was used to show that gastrin would stimulate β-cell neogenesis in the distal portion but not in the pancreas proximal to the ligation (52).

Importantly, this model transferred to mice has been useful in lineage-tracing studies. For example, with duct ligation on acinar-specific elastase 1 promoter:ROSA 26R bigenic mice, Desai et al. (53) found no marked islets after ductal ligation and concluded that acinar cells did not contribute to in vivo new islet formation after pancreatic ligation. Then, in a series of elegant experiments with various neurogenin 3 reporter mice, Heimberg and colleagues (19) showed that neurogenin 3 was induced in cells in or adjacent to the pancreatic ducts after partial duct ligation. By isolating these cells with flow cytometry and subsequent transplanting them into explants of embryonic pancreas from neurogenin 3–null mice, they showed that these cells gave rise to islet cells including β-cells. While lineage tracing was not performed, these data provide strong support for the concept that multipotent progenitor cells can be activated to increase the islet mass. Complementing these experiments, we tested the hypothesis that duct cells could give rise to β-cells by doing partial duct ligation in inducible duct-specific carbonic anhydrase II CreERT2:Rosa26R bigenic mice (20). In the distal, regenerated pancreas, 42% of the islets (24% β-cells) were marked compared with 12% (5.5% β-cells) in the nonligated pancreas of the same animals. Thus, these studies using duct ligation in adult mice have clearly shown that new islet cells can be formed from progenitors activated in the ducts.

Transgenic mice overexpressing cytokines/growth factors/transcription factors. A number of transgenic mice have reported increased neogenesis, but only three will be described herein. The most studied transgenic model of neogenesis is the human insulin promoter, interferon-γ mouse (54). These mice were found to have continual inflammatory destruction of islets associated with continued proliferation of ductal epithelium and “budding” of new islets from the ducts. Originally the insulin signal in this model was thought to trigger the neogenesis, but similar results were found with this transgene in immunocompromised mice. In these proliferating ducts, the expression of PDX-1 protein and other transcription factors seen in embryonic pancreatic development suggested that this regeneration recapitulates the embryonic process (55,56).

A second transgenic model, the metallothionein1–TGF-α mouse, also provides evidence of neogenesis from ducts. The induction of TGF-α in the exocrine pancreas by zinc in the drinking water (57) resulted in sustained proliferation of the ductal epithelium and metaplastic ducts. In these metaplastic ducts, 6% of the cells were immunostained for insulin (37) (Fig. 1A). Moreover, enhanced PDX-1 expression and focal expression of Pax6 suggested the initiation of islet neogenesis (58). When these mice were crossed with mice in which gastrin expression was driven by the insulin promoter, the number of metaplastic ducts was reduced and islet mass increased, leading to the hypothesis that gastrin could drive the differentiation of progenitors to islets (37).

More recently, another variation of neogenesis of islet cells from ducts has been shown in bigenic Pax4:glucagon Cre (POE::gluCre) transgenic mice (30), in which Pax4 is expressed in glucagon-expressing α-cells. This ectopic expression of Pax4 reprogrammed the α-cells to insulin-producing β-cells. However, even with continuing conversion to β-cells, there were still α-cells, suggesting a renewal of the α-cell population. The authors concluded that α-cell neogenesis occurred since 1) BrdU incorporation was mainly in cells within or adjacent to the duct epithelium rather than in hormone-expressing cells; 2) using a lentiviral reporter injected into the ductal tree, and both glucagon and insulin-positive cells were labeled in the induced transgenic mice but not in noninduced controls, suggesting that the hormone+ reporter + cells were derived from the labeled ducts; and 3) knockdown of neurogenin 3 with shRNA blocked the replenishment of the α-cells.

β-Cell compensation in human pancreas. Our main interest in studying rodent pancreas is to understand what happens in the human pancreas with which we cannot do the same experiments. We know that the pancreatic β-cell mass is increased in human obesity albeit less than in rodents. Whereas mice can have a 30-fold increase in β-cell mass with insulin resistance or obesity (59), the estimate for the increase in humans is more like 30–40% (60,61). Yet, β-cell proliferation in adult humans is ex-
tremely low, and greatly enlarged islets are rarely found. In autopsied human pancreas, β-cell replication (Ki67+ β-cells) drops to <0.2% already by 5 years of age (62) and can be almost negligible in adults (63,64). However, this low level of detection may result from the tissue being only retrieved after death. In a recent study on human organ donor pancreases, all pancreases had Ki67+ β-cells, with a stable, albeit low, percentage in those donors 40–65 years of age (65). An intriguing report documents increased Ki67+ β-cells (0.69 ± 0.15%) in a surgically resected pancreas from an 89-year-old with recent-onset type 1 diabetes (66), suggesting that β-cell replication could still be stimulated.

A major difference between mice and humans is that in humans telomere shortening limits replication and leads to senescence, a process described as replicative aging (67). In contrast, mice have long telomerases and do not show impairment of replication for several generations after ablation of telomerase (68,69). Similarly rats do not show replicative aging (70,71). This difference in replicative aging could account for the differential response (proliferation versus differentiation from stem cells/progenitors) in β-cell compensation (72).

**Evidence of neogenesis in human pancreas.** Due to the limits of experimentation possible on human pancreas, our knowledge is based mainly on observations made from pancreases obtained at autopsy, pancreas donation, and surgical resection. A number of pathological reports cite increased hormone-positive cells within the ducts in some human diseases, such as recent-onset type 1 diabetes and severe liver disease (73). In a recent study on pancreata harvested from organ donors (65), insulin+ cells in pancreatic ducts were usually present at low levels (0.4%) across the age span of 7–70 years. Similarly, in autopsied pancreata after the age of 12, 0.5–1.2% of the duct cells were insulin+, although in some pancreata none could be found (62). In other studies by Meier and colleagues (74,75) in lean controls, 0.6 ± 0.2% of duct cells were insulin+ but in pancreata from obese patients, 1.2%. In 10 pancreata from patients with chronic pancreatitis, there were significant increases in the percentage of duct cells positive for glucagon, amylase, Ki67, Pdx1, or insulin compared with control pancreata (76). It is of interest that in a small series of patients with 50% pancreactectomy, there was no evidence of pancreatic or β-cell regeneration and no change in Ki67+ or insulin+ cells in ducts (77), but this is not surprising because replication would only be expected shortly after the surgery. Further evidence that these hormone-positive cells within the ducts were dynamic comes from the comparison of 16 donor pancreata and biopsied pancreata from eight simultaneous pancreases and kidney transplantations (78). In the donor pancreata, the frequency of 0.45% insulin+ duct cells was similar to the other above-mentioned studies and in three transplants without autoimmune recurrence (0.5%). However, in five pancreatic transplant recipients with recurrent autoimmunity, 57.5% of the duct cells expressed insulin protein. This unexpectedly high value needs to be confirmed. Still more evidence suggesting neogenesis in humans comes from a small group patients who develop hypoglycemia after gastric bypass surgery (79). They have been found to have increased β-cell mass and impressive numbers of islet cells within the ducts (Fig. 2A), accompanied by high circulating levels GLP-1, which has been shown to stimulate neogenesis in rodents (8). From these accumulating circumstantial data, one can be more confident that neogenesis is an important process in humans.

**Evidence of neogenesis from human tissue in vitro or transplanted into mice.** With the caveat that tissue in vitro may be released from a number of regulatory controls and thereby appear more plastic, there is strong evidence that neogenesis can occur in human pancreatic tissue. Starting with the islet-depleted pancreatic digest remaining after islet isolation, several groups were able to show increased islet cells and increased insulin mRNA (80–83). Indeed, in our own in vitro studies, collections of new islet cells could be seen budding from cysts of duct epithelium (80). Since all of these studies were confined by the possible inclusion of β-cells in the starting material, we purified human ductal cells, using immunomagnetic beads and an antibody against the cell surface antigen CA19–9, and transplanted aggregates of expanded duct cells into immunocompromised mice (29). Here, too, we could show the induction of insulin+ cells and insulin mRNA. Some insulin+ cells in the grafts coexpressed duct markers (CK19 and CA19-9) and HSP27, a marker of nonislet cells, suggesting their transition from duct.

Suarez-Pinzon and colleagues (84,85) have provided further support to activation of neogenesis in human pancreatic ducts. First, using human pancreatic preparations that included islets, ducts and acinar cells treated 4 weeks in vitro with the combination of EGF and gastrin were found to have a doubling of β-cells with similar increases in the number of cells expressing the duct marker cytokeratin 19 and the transcription factor IPF-1/PDX-1 (84). Then, in a later study (85), similar low-purity human islet preparations were dispersed and transplanted under the kidney capsule of streptozotocin-induced diabetic NOD-scid mice, which then were treated with GLP-1 and gastrin for 5 weeks. In those treated with this combination, there was a fourfold increase in insulin+ cells compared with vehicle-treated controls from the same donor pancreas, with a high proportion of insulin+ cells also expressing cytokeratin 19.

**What is the cell of origin of the neogeneic islet cells?** A major question about neogenesis under debate is, What is the cell of origin? Most reports have suggested a cell of origin in or adjacent to the pancreatic duct epithelium, but is it a true stem cell, a differentiated duct epithelial cell capable of multipotent redifferentiation, or some other cell type? If a duct cell, can any duct cell have the potential or just a subpopulation? Xu et al. (19) clearly showed neurogenin 3 activation in cells within or adjacent to the ducts, and our report (20) showed that carbonic anhydrase I was expressed in the cell of origin. However, the recent Solar et al. (24) study using Hnf1β promoter as a driver for the Cre recombinase excision did not find any marked β-cells after partial duct ligation. Their interpretation was that differentiated duct cells do not contribute to new β-cell formation but a more precise interpretation would be that duct cells with high enough levels of Hnf1β transcription to cause excision do not have this potential (86).

In light of the report of Solar et al. (24), a redefined question should be, Are there identifiable subpopulations within the ductal tree and which ones have progenitor potential? We asked whether there is a heterogeneous expression of progenitor markers throughout the ductal tree (87) by immunostaining with titration of antibodies against Hnf1β and Sox9. Both proteins can be found throughout the pancreatic ductal tree, but the intensity and proportion of the expressing population varies within
the various categories of ducts, and only some cells express both. Better understanding of this heterogeneity may define a progenitor population. The recent identification of a population of terminal ductule/centroacinar cells that express high activity of aldehyde dehydrogenase 1 and have the ability to self-renew and to differentiate into both endocrine and acinar cells (88) adds new exciting directions for study. Moreover, expansion of these cells was found in chronic caerulein-induced pancreatitis. Further characterization of this population of cells in various other models is now a clear priority.

In summary, considering the overall body of evidence, we can only conclude that both replication of preexisting β-cells and neogenesis are pathways that contribute to maintaining an adequate β-cell mass after birth. Data show that both pathways are functional in all mammalian species studied, but different contributions, which are quantitative rather than qualitative, are made in different species and under different conditions. Eventually, one or both of these pathways may be manipulated for therapeutic treatment of diabetes. Our next efforts should address important questions such as, What is the cell of origin? what are the pathways that drive the activation of the progenitors? can we selectively stimulate neogenesis and to what extent? how applicable will these treatments be for human therapy?

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S.B.-W. wrote the manuscript. W.-C.L., L.O.-Y., and L.G. researched data. G.C.W. and A.S. contributed to the discussion and reviewed/edited the manuscript.

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