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## Review Articles

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# Extracellular Signals and Pancreatic $\beta$ -cell Development: A Brief Review

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### Abstract

Cell lineage development is a finely tuned process of proliferation and differentiation, survival and apoptosis, that is regulated by numerous extracellular signals. Here we review some of the extracellular signals—including insoluble cell–cell and extracellular matrix–cell interactions, as well as soluble factors—that appear

critical for pancreatic  $\beta$ -cell development. Knowledge of how these signals control the development of pancreatic endocrine stem/precursor cells into fully functional insulin-secreting  $\beta$  cells is a platform for the restoration of  $\beta$ -cell function and the cure therapy of type 1 diabetes.

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### Introduction

Type 1 (insulin-dependent) diabetes is due to insulin deficiency, following destruction of pancreatic islet  $\beta$  cells by autoreactive T lymphocytes. Restoration of  $\beta$ -cell function, the cure for type 1 diabetes, is far from a reality, partly because we understand so little about how extracellular signals (ECS) control precursor cell proliferation and differentiation into fully functional insulin-secreting  $\beta$  cells. Since the mid-1970s, it has been accepted that pancreas cell lineage development is independent of exogenous hormones, growth factors and innervation, and is controlled by autocrine and paracrine ECS, including extracellular matrix (ECM)–cell interactions. Thus the pancreas primordium dissected from an E11 fetal rat differentiates normally into exocrine and endocrine tissues after 9 days in culture in the absence of any exogenous factors (1). Recently, a great amount of information has appeared, witnessed by over 100 published reviews in last decade, on pancreas cell lineage development (2–4) and regulation at the cellular (5–10) and transcriptional (11–14) levels. Most transcription factor genes are activated by ECS (15). However, progress in determining the identity of ECS molecules and their mechanisms of action has been slow, because of early embryo lethality or functional redundancy after gene targeting and lack of simple and reliable in vitro assays.

There are, broadly, three types of ECS molecules: those that mediate cell–cell interactions such as Notch and cadherin receptors, insoluble ECM proteins such as laminins and collagens, and soluble factors such as growth factors, hormones, and vitamins. Here we review some ECS critical for pancreas development (Fig. 1).

### Cell–Cell Interactions

Molecules that mediate cell–cell interactions are illustrated by reference to Notch and cadherin families. Notch signaling is an evolutionarily conserved mechanism for development of multicellular organisms. Four isoforms of Notch have so far been identified. The Notch receptor is a single transmembrane protein composed of an extracellular domain, a transmembrane domain, and a cytoplasmic domain. The extracellular domain recognizes cell-bound ligands of the Delta/Serrate/Lag2 (DSL) family present on neighboring cells. Activation of the Notch pathway leads to proteolysis releasing the Notch intracellular domain (NICD), summarized in Figure 1 and reviewed elsewhere (16–19). By in situ hybridization, Notch 1 mRNA was detected uniformly in both the dorsal and ventral pancreatic buds of E10.5 mouse embryos, coexpressed with Hairy and Enhancer-of-split protein-1 (HES-1), a downstream molecule in Notch signaling (20). Mice deficient for *Delta-like gene 1* or *HES-1* showed premature differentiation of islet cells and exhaustion of the precursor population (21,22), demonstrating that Notch signaling is critical for maintaining the precursor cell pool and regulating its differentiation into islet cells. However,

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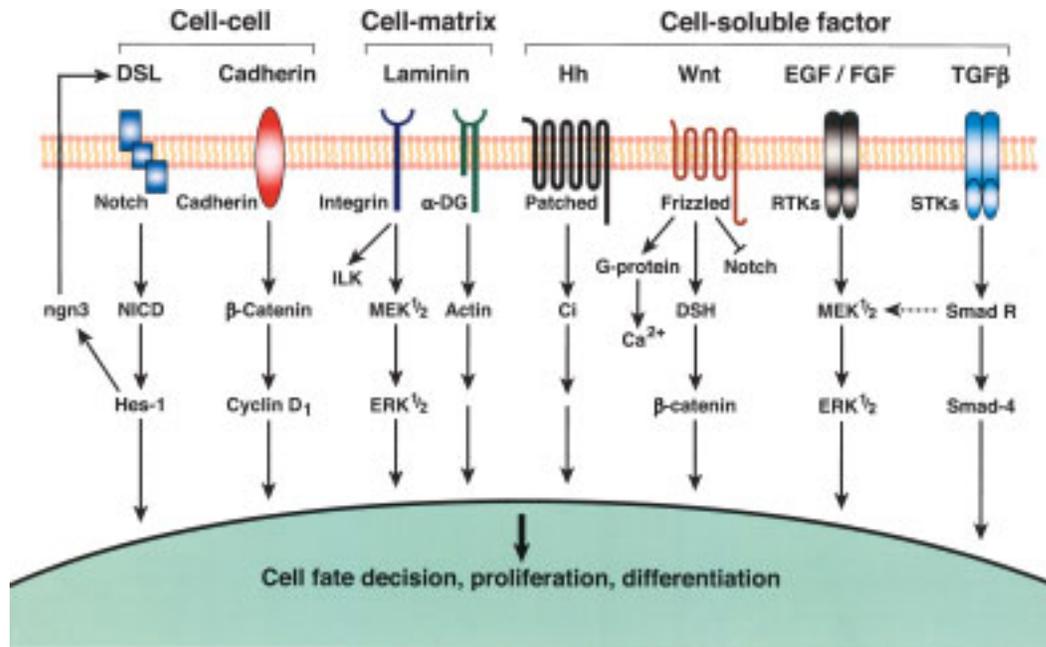


Fig. 1. Schema depicting three major classes of extracellular factors involved in pancreatic cell lineage development. See main text for details.

studies on other Notch isoforms are required to determine their physiologic roles.

Cadherins comprise 80 members of a superfamily of homophilic cell-cell adhesion molecules (CAMs) (23,24). The cytoplasmic domains of cadherins bind to the submembranal proteins,  $\beta$ -catenins, which link to the actin cytoskeleton via  $\alpha$ -catenin.  $\beta$ -catenin-mediated transcription is activated by the Wnt signaling pathway (see below).  $\beta$ -catenin activates genes involved in cell proliferation, such as *Cyclin D1* (25). Data from other cell systems and the pancreas demonstrate that cadherins play an important role during development. For example, N-cadherin interactions regulate cell proliferation in the sensory epithelia of the inner ear in response to changing cell density (26). Ovarian granulosa cell proliferation in vitro is significantly inhibited in the presence of monoclonal E-cadherin antibody (27). Blockage of epithelial CAM by a monoclonal antibody induces the differentiation of insulin and glucagon-positive cells in fetal human pancreatic cells (28). Though not affecting normal cell differentiation, abnormal islet architectural development was observed in N-CAM deficient mice (29) and in dominant-negative mutant E-cadherin mice (30). The expression and function of other cadherins during pancreas development remains to be investigated. Availability of more reagents for studies on cadherins could facilitate in vitro manipulation of islet cell development.

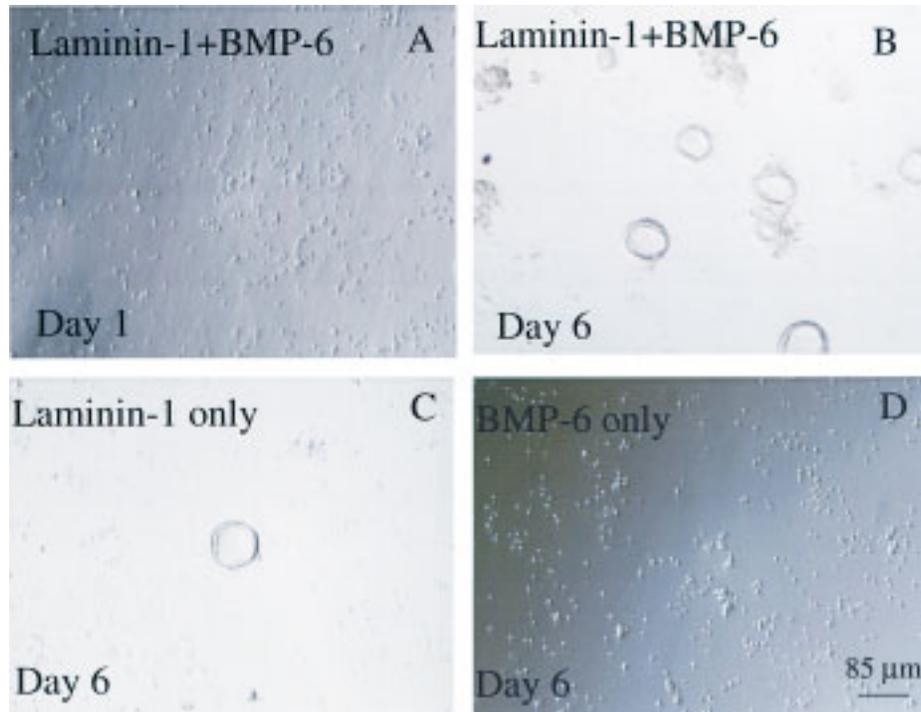
To avoid any complicating effects of cell-cell interactions, for example, via cadherins and Notch receptors, and paracrine effects from high cell density, we developed a low cell density cell culture system for pancreas precursor cells to investigate the

effects of added ECS (Figure 2A) (31). A similar system was also established for multipotent adult bone marrow mesodermal progenitor cells, because cell-cell interaction at a high cell density was shown to induce their differentiation (32,33).

### Cell-Matrix Interactions

The ECM is an organized network composed of numerous glycoproteins, typically including laminins, fibronectin, collagens, proteoglycans, and glycosaminoglycans. The ECM is not a static structure, but is continually produced and remodeled. ECM-cell interactions operate through receptor-mediated signaling and directly or indirectly modulate the cell response to growth factors (Fig. 1) (34–38).

The pancreas is an endoderm-derived, branching epithelial organ. At the interface between epithelial and mesenchymal tissues, there is a dense, sheet-like specialized ECM, the basement membrane (BM), composed mainly of laminins (80%) and collagen IV (39,40). Laminin is biochemically a heterotrimeric glycoprotein ( $M_r = 850,000$ ) composed of disulfide bonded  $\alpha$  (400 kDa),  $\beta$  (210 kDa), and  $\gamma$  (200 kDa) chains, whose structure and function has been well-reviewed recently (41,42). Laminin-1, the earliest described prototype and most extensively studied member of this family, plays an important role in proliferation and differentiation of many cell types during development (43). We detected laminin  $\alpha_1$  chain, specific for laminin-1, in the BM of the developing mouse pancreas from E13.5 to E17.5 (31), but it is not present in adult pancreas (44–46). Thus laminin  $\alpha_1$  is expressed in the epithelial BM during early



**Fig. 2.** Phase contrast images showing dissociated E15.5 mouse pancreas cells cultured with and without BMP-6 (10 ng/ml) and laminin-1 (160  $\mu$ g/ml). (A) Day 1 in the presence of both laminin-1 and BMP-6. (B, C, D) Day 6: BMP-6 induced pancreatic cell colony formation in the presence of laminin-1. Without BMP-6, fewer colonies formed in the presence of laminin-1 only. BMP-6 alone failed to promote colony formation in the absence of laminin-1.

development but is subsequently down-regulated (45,47). We found that laminin-1 was required for differentiation of isolated E13.5 pancreas cells to insulin-positive  $\beta$  cells in vitro (31). Expression of various collagen isoforms during pancreas development has been documented (48,49). Collagen IV was shown to inhibit fetal pancreatic cell survival, whereas the non-BM ECM molecule, fibronectin, had no effect (31). Our findings demonstrate that laminin-1 and collagen IV have opposing effects on  $\beta$ -cell development.

The major laminin receptors, the integrins, comprise at least 22 isoforms. They are a well-characterized family of heterodimeric transmembrane glycoprotein molecules composed of noncovalently bound  $\alpha$  (120–180 kDa) and  $\beta$  (90–110 kDa) subunits, of which there are 16 and 8 isoforms, respectively. Integrins mediate both cell–matrix and cell–cell interactions in multicellular organisms (50). In addition to their ability to link cells to their extracellular microenvironment, integrins play a role in cellular signaling via association of their cytoplasmic domains with signal transduction cascades (51). All known epithelial integrin receptors for laminin are among the isoforms composed of  $\alpha_6$ ,  $\alpha_3$ ,  $\beta_1$ , and  $\beta_4$  subunits (52). However,  $\alpha_v\beta_3$  and  $\alpha_v\beta_5$  integrins are also detected in the developing human pancreas (53). Disruption of  $\beta_1$  subunit, a partner of at least 12 integrins, causes peri-implantation lethality (54). Because the  $\alpha_6$  subunit shares significant (40%) identity with the  $\alpha_3$  subunit (55), the function of  $\alpha_6$  integrins

can be compensated by  $\alpha_3$  integrins (56). This probably explains why knockout of the  $\alpha_6$  integrin gene failed to result in any abnormality of  $\beta$ -cell development in vivo (57). Although a comprehensive profile of integrin expression during pancreas development is still lacking, blocking laminin-1 binding to  $\alpha_6$  integrin or  $\alpha_6$  integrin downstream signaling pathways, such as the extracellular activated receptor kinase 1 pathway, by monoclonal antibody or specific inhibitors, respectively, promotes  $\beta$ -cell differentiation (Fig. 1) (57), illustrating that the role of integrins not only to physically support cells but also to transduce signals for development.

The nonintegrin receptor for laminin,  $\alpha$ -dystroglycan ( $\alpha$ -DG), is a glycoprotein initially identified in muscle (58) and subsequently in other tissues including pancreas (59). It is a highly glycosylated peripheral membrane protein associated with a membrane-spanning protein  $\beta$ -dystroglycan ( $\beta$ -DG), comprising the dystrophin–glycoprotein complex (DGC). In muscle, DGC is structurally organized into three distinct subcomplexes: the dystroglycans ( $\alpha$ -DG and  $\beta$ -DG), the sarcoglycans (SGs,  $\alpha$ ,  $\beta$ ,  $\gamma$ ,  $\delta$ , and  $\epsilon$  subunits) and the cytoskeletal proteins, dystrophin (Dp), syntrophin and dystrobrevin (60,61).  $\alpha$ -DG is associated with the F-actin cytoskeleton through dystrophin (62,63). In epithelial cells, however, neither  $\alpha$ -,  $\beta$ -,  $\gamma$ -, nor  $\delta$ -SG is expressed (59). The intracellular binding partners of  $\beta$ -DG have not been identified but may include utrophin and/or the shorter dystrophin isoforms

Dp71 and Dp140 (64). To study the role  $\alpha$ -DG in laminin-1-induced  $\beta$ -cell differentiation, dispersed E13.5 fetal mouse pancreatic cells were cultured for 4 days with IIH6, a mouse monoclonal anti- $\alpha$ -DG antibody that blocks laminin-1 binding to  $\alpha$ -DG (62,65,66), and heparin, a known inhibitor of laminin-1 binding to  $\alpha$ -DG. Both IIH6 and heparin significantly decreased the number of both total cells and  $\beta$  cells in a dose-dependent manner, indicating that  $\alpha$ -DG was essential for survival and differentiation of the pancreatic precursor cells in vitro (57). Further studies are required to examine the role of  $\alpha$ -DG in islet cell development in vivo.

### Soluble Factor–Cell Interactions

A plethora of soluble factors, mainly polypeptides signal via binding to cell surface receptors. In development, they include hedgehog signaling molecules, Wnt proteins and several families of growth factors.

The hedgehog family of secreted signaling molecules comprise Sonic hedgehog (Shh), Indian hedgehog (Ihh) and Desert hedgehog, which regulate growth and differentiation of many organs during development. Hedgehog signaling inhibits the proteolytic cleavage of the transcriptional factor cubitus interruptus (Ci) and has been recently reviewed (67,68). Although Shh expression is not detectable during pancreas development, *Shh*<sup>-/-</sup> and *Shh*<sup>-/-</sup> *Ihh*<sup>+/-</sup> mice had a 3-fold increase in pancreas mass and a 4-fold increase in islet cell number (69), demonstrating the importance of hedgehog signaling for pancreas development. Ectopic expression of *Shh* under the control of the homeodomain pancreas duodenum transcription factor-1 (PDX1) promoter induced myoid cell formation around pancreatic cells, but islet cell development still occurred (70).

The *Wnt* genes encode a large family of secreted glycoprotein growth factors (19 in humans) (71). The *Wnt* genes are highly conserved between vertebrates, sharing overall sequence identity. Once secreted, Wnt proteins associate with glycosaminoglycans in the ECM and are tightly linked to the cell surface. Wnt signals are transduced by binding to two distinct families of cell surface receptors: members of the Frizzled (FZD) gene family (Fig. 1) and members of low-density-lipoprotein receptor-related family, and activated FZD receptors signal through Disheveled (DSH) (71–74). Before binding to their receptors, Wnt proteins are also modulated extracellularly by various secreted proteins including Fzd-related proteins (Frzbs), Wnt-inhibitory factor-1, and Cerberus. Wnts-2b and -11 mRNAs are expressed in many human fetal organs including pancreas (75,76). A human Frzb homolog termed HFZD-1b was exclusively expressed in the human pancreas (77). Similarly, HFZD-5 was highly expressed in fetal liver and pancreas (78). Recently, Wnt6 protein was showed to induce neural crest cell differentiation (79). All these data suggest that Wnt

signals play important roles during pancreas lineage development. Therefore these signals deserve closer attention with respect to pancreas development. In addition, together with the hedgehog family, Wnt signaling regulates stem cell numbers in epithelia such as those of the skin and intestine, which undergo constant renewal (80). Wnt proteins can bind to the Notch extracellular domain and the Notch signaling is inhibited through binding to the Wnt downstream molecule, DSH (19).

Growth factor signals can be transduced by receptor tyrosine kinases, for example, for the epidermal growth factor (EGF) family, fibroblast GF (FGF) family, hepatocyte GF (HGF) and nerve GF (NGF), or by serine/threonine kinases, for example, for the transforming GF [TGF]- $\beta$  superfamily members such as TGF- $\beta$  itself, bone morphogenetic proteins (BMPs) and activin.

Serum contains over 1000 components, including various kinds of growth factors and poorly defined growth modifying molecules. Without fetal bovine serum (FBS), for example, islet cell differentiation in chicken dorsal pancreas buds embedded in Matrigel, a purified BM component, was 5-fold more than in 1% FBS (81). Similarly, proliferation of bone marrow mesodermal progenitor cells required a serum-free environment (32,33). We chose serum-free complete medium for culture of dispersed fetal pancreas cells (31,57,82).

The EGF family consists of at least 10 members including EGF, amphiregulin (AR), betacellulin, epiregulin, heparin-binding EGF (HB-EGF), neuregulins 1–4, and transforming GF- $\alpha$  (TGF- $\alpha$ ). The EGF receptor (EGFR) family members, EGFR/ErbB1/HER1, ErbB2/neu/HER2, ErbB3/HER3, and ErbB4/HER4, are tyrosine kinases. They have some specificity for particular ligands; for example EGF, AR, and TGF- $\alpha$  preferentially bind EGFR/ErbB1 (83). Each EGFR family member can form a homodimer or heterodimer with another, except ErbB2, which is the critical partner for several heterodimers (84). Structures and signaling of the EGF family have been reviewed (83–87). By immunocytochemistry, ErbBs 2–4 were detected in the mouse pancreatic epithelium around E14.5 (88). Similarly, HB-EGF was observed in the rat pancreatic epithelial precursors, with a pattern similar to that of PDX1 (89). Mutant ErbB3/HER3 impaired development of the mouse pancreas (90). In vitro, EGF was shown to promote proliferation of mesenchyme-free rat pancreatic epithelium at E13 (91). All these data implicate the EGF family in pancreas development. However, in *EGFR/ErbB1* gene knockout (-/-) mice, lineage development of the pancreas was normal, except for somewhat impaired migration and delayed differentiation of the islet cells (92). This perhaps illustrates the shortcomings of gene targeting to examine the function of molecules with redundant, overlapping actions. The effect of EGF family members on pancreas development in vitro deserves further investigation.

To date, the FGF superfamily consists of 23 members. All contain a conserved 120 amino acid core region and act extracellularly through four tyrosine kinase FGF receptors (FGFRs 1–4) with various affinities (93). The extracellular region consists of two or three immunoglobulin-like domains called loops I, II, and III, the last determining ligand specificity. In isolated chick endoderm, FGF2 can repress Shh expression and activate PDX1 and insulin expression (94), while in the mouse ventral foregut endoderm FGF2 or FGF8b induces Shh expression and leads to liver development (95). RT-PCR analysis detected FGFs-1, -7, and -10 and FGFR2b in the developing rat pancreas from E12–E18 (96). In mesenchyme-free cultures of embryonic pancreatic epithelium, FGF-1, -7, and -10 stimulated growth, morphogenesis, and differentiation of pancreatic exocrine cells (96). mRNA in situ hybridization revealed only FGF-10 before E12.5, and exogenous FGF-10 restored proliferation of the pancreatic precursor cell population in FGF-10 mutant ( $-/-$ ) mice (97). Overexpression of FGF10, but not FGF8, disturbed  $\beta$ -cell differentiation (98). FGFR1 and FGFR4 mRNAs were detected throughout pancreas development in rats (99). Expression of dominant-negative FGFR2IIIb, specific for FGF-7 (also called keratinocyte GF), caused pancreatic aplasia (100). Similarly, replacing FGF2IIIb translational stop codons with an IRES-LacZ cassette resulted in dysgenesis of many organs including the pancreas (101). This further supports the idea that FGF7 promotes pancreas precursor cell proliferation. Furthermore withdrawal of FGF7 led to the differentiation of endocrine cells in vitro (102). FGF2IIIb signaling also induces FGF4 expression and the latter stimulates Shh expression (101). However, perturbing FGF signaling, by expressing dominant negative forms of FGFR1b and FGFR1c (dnFGFR1c) in the developing pancreas on the PDX1 promoter, did not interfere with normal lineage development, although the adult mice with a dnFGFR1c developed diabetes (103).

TGF- $\beta$  superfamily members, including the TGF- $\beta$ , the BMPs and the activins, bind to two types (I and II) of serine/threonine kinase receptors on the cell surface, both receptors being necessary for signal transduction (104,105). The TGF- $\beta$  family consists of at least three members. Its signal transduction, regulation and function have been recently reviewed (106–108). Transgenic mice expressing a dominant negative TGF- $\beta$  receptor II controlled by the mouse metallothionein 1 promoter display increased proliferation and impaired differentiation of pancreatic acinar cells (109). Transgenic mice expressing a dominant negative activin receptor controlled by the human insulin promoter have hypoplasia of pancreatic islets (110). Activin B represses endodermal expression of Shh, a prerequisite for expression of PDX1, required for pancreas development (111,112). Follistatin, an inhibitor of activins, was expressed by E12 in rat pancreas mesenchyme and its addition in vitro induced the development of exocrine tissues and

repressed the differentiation of islet cells (113). TGF- $\beta$ 1 and activin A were detected in the developing pancreas and, as in other epithelial cells, shown to inhibit pancreatic cell proliferation in vitro (82).

For the BMPs, there are three type I receptors, activin-like kinase (ALK)-3 (also called BMPR-IA), ALK-6 (BMPR-IB), and ALK-2 (or activin receptor type IA, ActR-IA) and three type II receptors, BMP type II receptor and activin type II receptors (ActR-IIA and ActR-IIB). ALK-2, ActR-IIA, and ActR-IIB are also activin receptors (114,115). Unlike TGF- $\beta$ , BMPs appear to bind cooperatively to both type I and II receptors (104). The type II receptor transphosphorylates the type I receptor at a specific region of its cytoplasmic domain resulting in its activation. The activated type I receptor then recruits and phosphorylates downstream signaling molecules belonging to the Smad family, via Smad1, Smad5, and Smad8. The activated Smads then form a complex with Smad4, translocate into the nucleus and regulate the transcription of various target genes. Smads2 and 3 are phosphorylated by TGF- $\beta$  type 1 and activin type IB receptors, respectively (116,117). Smads6 and 7 are inhibitory Smads, serving as transcriptional repressors to silence the transcription of target genes.

The BMPs have been shown to be important in development of kidney tubule, lung and other organ epithelia (118,119), including the pancreas. BMP-7 was detected immunocytochemically in human fetal pancreas duct epithelium (120) and by mRNA in situ hybridization in mouse pancreas epithelium between E12.5 and E14.5 (121). In our RT-PCR analysis of mRNAs from E13.5, E15.5, and E17.5 fetal mouse pancreas, BMPs-4, -6, and -7 were detected at each age, whereas BMP-5 and BMP-2 were only detected later at E17.5 (82), consistent with other reports (120–122). In the presence of laminin-1, we found that BMPs-4, -5, and -6 promoted the development of E15.5 isolated pancreas cells into cystic epithelial colonies (Fig. 2) containing insulin-positive cells (82). At this time, one can presumably harvest a maximal number of precursor cells, because expression of neurogenin-3, a marker of islet precursors (123,124), peaks at E15.5 (125). In the absence of laminin-1, BMPs-4, -5 or -6 failed to promote colony formation (Fig. 2), indicating that both laminin-1 and BMP signaling act in concert. BMP-6-induced colony formation was completely abolished by TGF- $\beta$ 1 or activin A (82), indicating that BMPs and TGF- $\beta$  and activins may have opposing roles on islet development.

Other growth factors are implicated in pancreas development. Overexpression of vascular endothelial GF under the control of the PDX1 promoter, and hence hypervascularization of the pancreas, is associated with islet hyperplasia (3-fold increase in islet number and area) (126). HGF and platelet-derived GF were also shown to increase proliferation of immature human pancreas  $\beta$  cells in vitro (127–129). High-affinity NGF receptor gp140<sup>Trk-A</sup> (tyrosine receptor kinase A [TrK-A]) was detected by immunofluorescence in the

fetal rat pancreas ductal epithelium (130), implying that NGF may influence pancreas development.

In summary, understanding and manipulating pancreas lineage development will require investigation of many ECS and their complex interactions. Together with genetic analysis, low cell density, serum-free culture of pancreatic precursor cells facilitates the identification of ECS molecules required for pancreas development. An outcome of these studies will be the generation of insulin-secreting  $\beta$  cells in vitro for cell replacement therapy in type 1 diabetes.

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