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Pancreatic Pathology in Type 1 Diabetes Mellitus

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Dear Sir/ Madam,

Thank you for your recent correspondence confirming acceptance of our review for the Endocrine Pathology Jubilee Edition. Please find attached our revised review manuscript. I have checked the references and hope that these will now fit with the journal style. Please do let me know if you require any further alterations.

Yours sincerely

A handwritten signature in cursive script that reads "Sarah Richardson".

Dr Sarah Richardson

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Title: Pancreatic Pathology in Type 1 Diabetes Mellitus

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Introduction

Type 1 diabetes is a multifactorial disease resulting from a complex interplay between host genetics, the immune system and the environment, that culminates in the destruction of insulin-producing beta cells. The incidence of type 1 diabetes is increasing at an alarming rate, especially in children under the age of 5 [1-3]. Genetic predisposition, although clearly important, cannot explain this rise and so it has been proposed that changes in the ‘environment’ and/or changes in ‘how we respond to our environment’ must contribute to this rising incidence. In order to gain an improved understanding of the factors influencing the disease process, it is important, firstly, to focus on the organ at the centre of the illness; the pancreas. This review summarises our knowledge of the pathology of the endocrine pancreas in human type 1 diabetes and, in particular, explores the progression of this understanding over the past 25 years.

What was known 25 years ago?

While there had been numerous studies of the autopsy pancreas in young persons with diabetes before, the foundations of this subject were laid by the seminal article of Willy Gepts in 1965 [4]. Gepts noted the presence of insulinitis (a predominantly lymphocytic inflammatory infiltrate of islets) in 15 of 22 (68%) autopsies of persons, under the age of 30, dying within 6 months of a diagnosis of diabetes [4]. These findings were later confirmed in a larger collection of UK samples, where 47 out of 60 (78%) young patients (<20 years) with recent-onset type 1 diabetes were found to have evidence of insulinitis [5]. Prior to the development of immunohistochemistry Gepts also noted that the pancreas in persons who had had the disease for many years was characterized by an almost complete lack of insulin secreting beta cells. From these observations he raised two hypotheses: the islet inflammation could be secondary to a viral infection of beta cells or it could represent an immunologically driven process, akin to autoimmune thyroiditis.

Support for both these theories followed. Gamble, Taylor and Cumming [6] found that patients with recently diagnosed type 1 diabetes were more likely than controls to have antibodies to Cocksackie viruses, and Cocksackie B4 virus was cultured from the

1 pancreas of a 10 year old boy who had died at clinical presentation of type 1 diabetes
2 [7]. This case may have been exceptional in that the virus was also cultured from the
3 brain of the child, raising the possibility of an incidental viraemia. Many attempts
4 have since been made to culture viruses from the pancreas of recent onset diabetic
5 patients, largely without success.
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10 In 1974 two separate groups reported the presence of islet cell autoantibodies in
11 patients with type 1 diabetes [8-10], leading to the disease being later classified as an
12 organ specific autoimmune disease with the insulin secreting beta cells as the target.
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17 Following the advent of immunohistochemistry being applied to paraffin embedded
18 tissue, it was shown that within the pancreas of children dying at clinical presentation
19 of their disease that the majority of islets (70%) were small and had no beta cells [11].
20 They had a normal complement of glucagon secreting alpha cells, somatostatin
21 secreting delta cells and pancreatic polypeptide secreting PP cells and were termed
22 insulin deficient islets (IDI). The remainder of the islets had residual beta cells and
23 were called insulin containing islets (ICI). Insulinitis affected 18% of ICI but only 1%
24 of IDI. This was the first evidence that insulinitis affected ICI primarily and lent support
25 to the idea that it represented an immunologically driven destruction of beta cells [11].
26 Importantly the lobular distribution of the disease was also noted (Figure 1), where a
27 seemingly unaffected lobe, or one where ICIs with insulinitis could be observed, were
28 frequently surrounded by lobes containing only IDIs.
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42 It also became clear that the pancreas at clinical presentation could be studied to look
43 at the time course of events within the organ. Insulin deficient islets were islets where
44 beta cells had been destroyed in the past; ICI with insulinitis were islets where beta cells
45 were being destroyed at the time of the death of the patient, and ICI with no insulinitis
46 were islets where the beta cells were yet to be attacked and destroyed. Gepts [4] had
47 suggested that the destruction of beta cells must happen over years and support for
48 this was the finding of insulinitis affecting ICI up to 6 years after clinical onset of
49 diabetes [5].
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58 Prior to the development of antigen retrieval, many antigens could only be studied by
59 immunohistochemistry on tissues that had not been fixed in formalin.
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1 Characterization of the inflammatory cell infiltrate in insulinitis for many years relied
2 on one report where fresh frozen pancreas had been collected at the time of death
3 from a child with recent onset diabetes [12]. The majority of the inflammatory cells
4 were T lymphocytes and the majority of them expressed CD8, suggesting that they
5 were likely to be cytotoxic T cells. Macrophages were not abundant.
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10 Cytotoxic CD8 T cells can recognize an antigen when it is presented to them bound to
11 cellular class I major histocompatibility complex (MHC) molecules. This
12 presentation is enhanced if there is increased expression of class I MHC. Bottazzo *et*
13 *al* [12] noticed that endocrine cells in some islets in the single pancreas that they
14 studied hyperexpressed class I MHC but they did not fully characterize this
15 phenomenon and did not describe which cells were affected. This feature was later
16 elucidated in a study of 23 pancreases from persons dying of recent onset type 1
17 diabetes [13]. Hyper expression of class I MHC was found in all pancreases where
18 there were residual beta cells (an example is shown in Fig. 2). It affected 92% of ICI
19 but only 1% of IDI. The phenomenon was not seen in 95 control pancreases, which
20 included normal and diseased pancreases (graft versus host disease, Coxsackie viral
21 infection, type 2 diabetes, chronic pancreatitis, cystic fibrosis). It was thus as much a
22 characteristic of type 1 diabetes as insulinitis and insulin deficient islets.
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36 Within affected islets all endocrine cells (alpha, beta, delta and PP) hyperexpressed
37 class I MHC. Whilst islets complicated by insulinitis uniformly hyperexpressed class I
38 MHC if they had residual beta cells, many otherwise normal ICI, with no evidence of
39 insulinitis even on serial sectioning, also displayed this phenomenon [13]. This
40 suggested that hyper-expression of class I MHC must precede insulinitis in the disease
41 process.
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49 It was hypothesized that if alpha, delta and PP cells hyperexpressed class I MHC
50 when they were adjacent to beta cells in ICI, but ceased to do this when they were
51 physically divorced from beta cells in IDI, following the destruction of beta cells in
52 the insulinitis process, then perhaps the islet beta cells were releasing some paracrine
53 substance capable of causing hyper expression of class I MHC on adjacent islet
54 endocrine cells. Candidates for this role included type 1 interferons (interferon-alpha
55 (IFN α) and beta) and interferon-gamma, as they had been shown to cause hyper
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1 expression of class I MHC on islet endocrine cells *in vitro*) [14]. Interferon-gamma is
2 released only by T lymphocytes so attention was focused on type 1 interferons.
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5 The possibility that the secretion of IFN α by islet beta cells plays a role in the
6 induction of class I MHC hyper expression in ICI was then assessed [15]. In a study
7 of 34 patients with type 1 diabetes, insulin containing beta cells were shown to exhibit
8 IFN α in 93% of islets which hyperexpressed class I MHC but in only 0.4% of islets
9 which did not show this phenomenon. Among 80 control pancreases with a variety of
10 diseases, only in 4 neonates affected by Coxsackie B viral infection were beta cells
11 found to express IFN α [15].
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19 The following sequence of events leading to diabetes was then hypothesized. In a
20 given individual, pancreatic beta cells might harbour a non-cytopathic chronic viral
21 infection to which the body could react in a number of ways, perhaps partly
22 determined by their genetic susceptibility to diabetes. If there was no immune
23 response then no disease might result. By contrast if the infected cells expressed
24 interferon-alpha (as part of an innate immune response to the double stranded RNA of
25 the virus) this would result in hyper expression of class I MHC by the endocrine cells
26 of the islet. Such expression might provoke loss of immune tolerance and infiltration
27 of affected islets by inflammatory cells. Even a very weak autoimmune response to
28 beta cell antigens in the context of class I MHC hyper expression would eventually
29 (over years) lead to their destruction and the development of clinical diabetes (Figure
30 1; Foulis Oakley Lecture 1987 [16])
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42 **Insulinitis: 25 years on....**

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45 Since the seminal contributions of Gepts, the concept that type 1 diabetes arises from
46 a process of immune-mediated destruction of pancreatic beta-cells has continued to
47 develop and expand. However, an important question still remains – how strong is the
48 evidence that insulinitis is a characteristic feature of human type 1 diabetes? Certainly,
49 Gepts was persuaded of the concept when he noted in 1981 that insulinitis is “*a*
50 *common finding in the pancreas of recent onset juvenile diabetic subjects*”.
51 Unfortunately, however, firm corroboration has been achieved only slowly, to the
52 extent that In't Veld, in reviewing the weight of evidence, has recently defined
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insulinitis in the human pancreas as an “elusive lesion” [17]. In particular, he noted that the number of recorded cases of insulinitis worldwide has hardly increased over the last 25 years such that the total currently stands at only about 200. As a result, the essential features of the lesion are still debated.

An important issue arising from the study of insulinitis is that the number of infiltrating cells in any given islet is relatively few [18]. This stands in marked contrast to the situation in the NOD mouse (a model of spontaneous autoimmune diabetes) where, typically, inflamed islets are surrounded by an army of infiltrating cells which initially adopt a peripheral configuration before migrating deep within the islet structure to achieve close-contact with the endocrine cells [19]. In our experience, the equivalent situation is rare in human tissue. Rather, in the case of human islets, the inflammatory infiltrate often numbers only a few cells in any given islet cross-section and the majority of these reside around the islet perimeter (Figure 3: Peri-insulinitis and/or focal insulinitis). Very few immune cells actually cross the threshold to enter the margins of the islet structure and come into close contact with the endocrine cells. Thus, the process of beta-cell destruction appears rather inefficient and this may account for the relatively protracted time course preceding the onset of clinical symptoms in many patients.

This, in turn, prompts a further critical issue in that it becomes extremely important to establish a firm definition of “insulinitis” in human pancreas since it is probable that small numbers of immune cells are likely to be present within even “normal” (non-inflamed) islets. Distinguishing the “normal” from the “inflamed” situation then becomes a key priority. To help address this issue, a meeting was held in Florida in early 2013, at which a number of investigators who have studied human insulinitis, were present. Collectively, these workers developed a consensus statement to define insulinitis: In a pancreas containing insulin deficient islets the lesion should only be confirmed if at least 3 individual islet sections contain a minimum of 15 lymphocytes (located either at the islet periphery and/or within the islet structure) [20]. This figure was chosen as it represents at least double the number of such cells found in the islets of control tissue (after extensive analysis of several thousand islets across multiple pancreases). This consensus definition represents the state-of-the-art in 2013 and provides an important context in which the process can be studied [20].

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2 Definition of the numbers of infiltrating immune cells within islets represents an
3 important step forward in the study of insulinitis but it is not, itself, a fully informative
4 statistic. This is because the nature of the infiltrate will determine the cytokine profile
5 within the islet milieu and therefore, a second important objective is to establish the
6 phenotypes of the various immune cells that may be present. We undertook a
7 comprehensive analysis of the immune cell phenotypes present within insulitic lesions
8 in 29 separate individuals who had died shortly after being diagnosed with type 1
9 diabetes[18]. This revealed that both the absolute number and the profile of immune
10 cells present in inflamed islets is variable. Indeed, immune cells appear to enter and
11 leave islets as the process progresses, such that the profile of immune cell subsets
12 differs according to the extent of beta-cell loss. Despite this, some clear trends
13 emerged; notably that CD8+ T-cells comprise the majority of the population at all
14 stages of insulinitis. By contrast, CD4+ T-cells are relatively under-represented in the
15 immune cell population [18].
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29 It is self-evident that the dynamics of immune cell infiltration cannot be fully assessed
30 in fixed tissue preparations taken at a single time point but, nevertheless, it is still
31 possible to make reasonable inferences by studying islets at varying stages of beta cell
32 destruction. When the data were processed in this way, a dynamic model emerged in
33 which CD8+ T-cells featured as the most prominent cells leading the islet attack and
34 they were accompanied by reduced proportions of CD4+ T-cells and CD68+
35 macrophages. Importantly, the CD8+ cells increased in number as beta-cell
36 destruction proceeded, implying that they play a primary role in mediating this
37 demise. By contrast, CD4+ and CD68+ cell numbers remained much more constant.
38 More surprisingly, a further lymphocyte subset was found to mirror the profile of
39 CD8+ T-cells, in that CD20+ B-cells also increased in numbers markedly, to achieve
40 a status as the second most prevalent cell type in many of the patients[18]. The
41 functional activity of these cells remains to be determined but they do not appear to be
42 antibody-secreting plasma cells. By analogy with the situation in NOD mouse (which,
43 as indicated above, provides an exaggerated representation of human insulinitis) it is
44 possible that these B-cells are somehow trophic for the CD8+ T-cells, allowing them
45 to reach their full cytotoxic potential [21]. However, it has also become clear that
46 CD20+ cells are not absolutely required for CD8+ cells to promote beta-cell loss
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1 since, in a more recent study reported in abstract form, a subset of patients was
2 identified whose islets are not infiltrated by significant numbers of CD20+ cells but
3 who, nevertheless, progressed to clinical type 1 diabetes [22]. Conceivably, then, the
4 CD20+ cells may play a facilitating rather than a primary role in beta-cell
5 cytotoxicity.
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10 In an earlier study using samples recovered from a different (and much smaller) group
11 of patients, Dotta and colleagues had identified the presence of NK cells within some
12 insulitic lesions[23]. This suggests that such cells might also play a role in mediating
13 beta-cell loss under some circumstances, although in our series of patients we were
14 unable to detect significant infiltration of NK cells [18].
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21 Taken together, the weight of recent evidence supports the view that CD8+ T-cells are
22 likely to be primarily responsible for driving beta-cell loss and this conclusion is
23 supported by a more a recent development which represents an important step
24 forward. In particular, work by Coppieters and colleagues has attempted to define
25 whether the CD8+ T-cells present in inflamed human islets are antigen specific [24].
26 This critical goal has been achieved by exploiting the power of peptide-loaded,
27 engineered, tetrameric antibodies (“tetramers”) which allow the selective
28 identification of T-cells recognising the specific peptide in question. Using this
29 approach, it was demonstrated that a proportion (currently undetermined) of the T-
30 cells present in inflamed islets in humans are directed against specific beta-cell
31 antigens [24]. Thus, these T-cells may be homing directly to the islets, guided by the
32 presentation of antigens within the islet milieu. As such, these cells may be
33 responding to highly specific cues rather than simply to a generalised chemokine
34 gradient existing within the vicinity of the islet.
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49 In this context, chemokines are well known to play an important role in guiding
50 immune cells to sites of injury or infection and it has been surmised that the
51 generation of chemokine gradients may underlie the homing of immune cells to
52 inflamed islets. In support of this, a number of workers have shown that chemokines
53 can be elaborated from islet cells, including molecules such as CXCL10 which may
54 be important for T-cell recruitment [25,26]. However, the evidence base supporting a
55 chemokine-driven recruitment mechanism in inflamed islets in human type 1 diabetes
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remains relatively weak. Indeed, the morphology of inflamed islets in humans seems oddly inconsistent with such a mechanism. Thus, as discussed previously, the absolute numbers of immune cells that reach inflamed islets is relatively small, and it is difficult to understand why most immune cells appear to simply pass by if there are strongly chemotactic influences available to recruit them. While such observations do not exclude a role of chemokines, it must be acknowledged that important questions remain to be answered about immune cell targeting. The possibility of antigen-specific homing certainly represents a key step in this quest.

The peri-islet capsule – a brake on beta cell destruction?

Questions surrounding the localisation and homing mechanisms employed by infiltrating immune cells in human islets are important and it remains unclear why the majority are localised around the islet periphery. A reasonable hypothesis would be to suggest that immune cells enter the islet vasculature from the wider circulation and are then subjected to homing signals (be they chemokines or other factors) which encourage the typical rolling adhesion as the cells transit the islet. Ultimately, strong adhesion and subsequent extravasation probably occur in post-capillary venules where shear forces are minimised. The post-capillary venules of the islet are located outside the peri-islet capsule [27] suggesting that extravasated immune cells may exit the vasculature at a point outside the islet and may then need to negotiate the capsule before they can gain direct access to the beta cells themselves. Important new advances have been made recently in understanding the components of the peri-islet capsule, which consists of an islet basement membrane (BM) and subjacent interstitial matrix (IM). The islet BM and IM enclose the islet, separating the endocrine cells from the surrounding exocrine acinar cells. The function of the BM is to act as a barrier to soluble molecules and to *cells* and the IM confers flexibility and elasticity [27-29]. The peri-islet capsule is made up of a complex mixture of extracellular matrix components [27-29]. Interestingly, it has been argued that immune cells cannot normally infiltrate the islet unless the structure of the capsule is destabilised. This may explain why the peri-insulinitis observed frequently in the islets of type 1 diabetes patients (Figure 3A, Figure 5) tends to occur in those with a near normal complement of beta cells [18].

1 The islet BM can be digested by a variety of enzymes, such as heparanases [28] and
2 cathepsins [29], which are probably released by peri-islet immune cells [28,29]. One
3 could therefore hypothesise, that as the number and type of recruited immune cells
4 increases around the islets, the damage inflicted on the capsule is enhanced (Figure 3:
5 Focal and progressive insulinitis, Figure 5) [20]. Once the peri-islet capsule has been
6 sufficiently degraded, the immune cells can infiltrate and destroy the beta cells
7 (Figure 3: Invasive insulinitis, Figure 5). Typically islets with heavy infiltration are
8 those in which the beta cells are most reduced in number. The need to penetrate the
9 peri-islet BM could therefore be one reason why the destructive process takes a
10 protracted period of time. Interestingly, following the complete destruction of the beta
11 cells, some islets are able to reform the BM (Figure 5) suggesting that it is not the beta
12 cells themselves that facilitate the production of this protective layer [29]. This may
13 have importance as a clinical target in the future. If the destruction of the islet BM can
14 be inhibited, the natural barrier to immune cell infiltration may prolong the survival of
15 the beta cells.
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27 *A role for enteroviruses in Type 1 diabetes*

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29 As noted previously, early studies on the human type 1 diabetic pancreas provided
30 indirect evidence that viral infection may play a role in disease pathogenesis. This,
31 coupled with mounting epidemiological evidence, has led investigators to consider
32 that direct infection of the beta cells could be driving the development of type 1
33 diabetes, at least in some patients. Many different viruses, including cytomegalovirus
34 (CMV), rubella, Epstein barr virus (EBV), rotavirus and in particular enteroviruses
35 have been nominated as candidates [30]. Among these, the most convincing evidence
36 implicates enteroviruses (single stranded, positive sense RNA viruses that belong to
37 the *Picornaviridae* family) [31]. Over the last 25 years, a plethora of epidemiological
38 studies have considered whether enteroviral infections are associated with the
39 development of islet autoimmunity or the onset of clinical type 1 diabetes. A recent
40 meta-analysis of 26 such studies concluded that there is a statistically significant
41 association between enteroviral infection and diabetes related autoimmunity/clinical
42 type 1 diabetes [31]. It was noted that the odds of having a detectable enterovirus
43 infection in people with type 1 diabetes are almost 10 times greater than in unaffected
44 individuals and 4 times greater in non-diabetic individuals with diabetes related
45 autoantibodies [31]. Many of these studies were however, performed on peripheral
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1 whole blood, preparations of PBMCs or serum samples, such that the authors also
2 remarked that firm evidence of viral infection within the pancreas is still lacking [31].
3 This is in part due to the limited amount of material available for analysis. Despite,
4 this the search for direct evidence of an infection within the pancreas (of those few
5 cases available) began over 34 years ago....
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10 Arguably the most convincing early evidence that enteroviruses can specifically target
11 cells within the pancreas and may play a role in the development of the diabetes, came
12 in the 1970s [32,7]. In these case reports, viral infection was associated with islet
13 inflammation and islet cell necrosis, indicative of an acute, lytic, infection [32,7].
14 Since then a range of studies have confirmed the tropism of enteroviruses for beta
15 cells both *in vitro* [33-35] and *in vivo* (particularly in fulminant type 1 diabetes,
16 described in more detail below and in Tanaka et al, [36]). Importantly, this is not
17 restricted to just one or two of the enterovirus family members, since Coxsackie virus
18 B (CVB), Coxsackie virus A (CVA) and several of the Echoviruses are capable of
19 infecting isolated human islets [33-35]. The tropism of enteroviruses for the pancreas
20 was further demonstrated *in vivo* in a series of neonates who had died from culture
21 proven Coxsackie viral myocarditis. In situ hybridisation (ISH), using a CVB-specific
22 probe [37,38] confirmed the presence of CVB RNA in the heart of all cases, but
23 interestingly also in the pancreas of five of the nine cases examined. The islets in
24 particular were targeted while only occasional acinar cell positivity was observed
25 [39,37].
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42 These studies suggested that detection of viral RNA in the pancreas might be a viable
43 means to verify whether enteroviral infection is associated with type 1 diabetes.
44 However, there are a number of important issues relating to the detection of viral
45 RNA in the pancreas which must be considered. Firstly, the pancreas is a particularly
46 noxious environment for RNA; it is rich in RNAses, meaning that free RNA is
47 susceptible to rapid degradation. Secondly, the majority of tissue studied to date have
48 undergone fixation with formalin, a process which has a dramatic impact on RNA
49 stability and integrity. Thirdly, the majority of samples tested so far are from
50 autopsies meaning that the pancreas may have been recovered many hours after death.
51 Therefore, despite the initial optimism arising from the demonstration of positive ISH
52 signals in Coxsackie-proven infections, the subsequent relative failure to reproduce
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this [37,40] in the pancreases of type 1 diabetes patients should be interpreted with caution.

As RNA stability is an issue when examining pancreatic tissue, researchers have also taken an alternative approach by attempting to detect viral proteins in this tissue, since these are likely to be better preserved in autopsy pancreas. In the first such study [39], two rabbit polyclonal antibodies raised either against a fusion protein containing sequences from CVB3 capsid proteins (VP4, VP2 and VP3) or against the CVB3 VP1 structural protein, were employed [41]. Both detected the presence of enterovirus in the heart and pancreas of confirmed Coxsackie-infected neonates, suggesting that they were suitable for detection of widespread, systemic, enteroviral infection in formalin-fixed autopsy tissues [39]. In addition, these antibodies were shown to identify various CVBs (CVB2, CVB4 and CVB5) despite being raised against CVB3 [39]. However, when applied to sections of autopsy pancreases from 88 patients who had died at, or shortly after, clinical presentation of type 1 diabetes, no positive signals were obtained [39]. This might be because viruses were not present in the tissue or alternatively it might also be because these antisera failed to bind to proteins expressed by the serotype of virus present. A third possibility is that the sensitivity of the antisera was insufficient to detect the very modest levels of protein expression achieved during a sub-lytic infection. Fourthly, because antigen retrieval was not in use at the time of this work, other technical issues might also have militated against virus detection.

Why is it easier to find the culprit in Coxsackie-infected neonates & fulminant type 1 diabetes?

As hinted above, direct evidence of an enteroviral infection has been relatively easy to find in the pancreases of individuals presenting with an acute Coxsackie infection or with fulminant type 1 diabetes. The latter is characterised by the acute-onset of clinical symptoms and differs pathologically from “classical” type 1 diabetes in that there is evidence of *lysis* of both beta and alpha cells [42,43]. Fulminant diabetes accounts for approximately 25% of type 1 diabetes cases in Japan, but is much rarer in European populations. By contrast with fulminant diabetes, in typical autoimmune type 1 diabetes there is *no* evidence of cell lysis and the cell loss is selective for beta cells. Islet inflammation is present in both forms of diabetes, but the complement of

1 immune cells in fulminant disease differs from that observed in typical type 1 diabetes
2 (reviewed in [44]). This has led some to suggest that fulminant type 1 diabetes
3 represents a non-autoimmune form of the disease [45], characterised by the absence
4 of diabetes-related autoantibodies, normal expression of class I MHC, lymphocytic
5 infiltration of the exocrine and endocrine tissue, elevated serum pancreatic enzyme
6 levels and a remarkably aggressive disease progression [45]. It appears therefore that
7 the ease of identification of enterovirus in these pancreas samples may be due, in part,
8 to the development of an acute lytic infection where abundant amounts of viral
9 protein are present in multiple cell types. In contrast, evidence of viral protein
10 production has been much harder to find in the pancreases of patients with
11 autoimmune type 1 diabetes. *Why is this?*

21 In autoimmune type 1 diabetes, diabetes-related autoantibodies appear in most
22 individuals years before clinical onset of the disease and the appearance of these islet
23 specific autoantibodies is believed to be an indicator of a beta cell stress.
24 Accumulating evidence from birth cohort studies has shown that the appearance of the
25 first autoantibody correlates with evidence of an enteroviral infection in the preceding
26 6 months [46,47]. This had led some to hypothesise that the initial insult to the beta
27 cell is a viral infection. This might lead to secretion of interferon-alpha by beta cells
28 and hyperexpression of class I MHC in infected islets, leading to the activation of
29 auto-reactive cells in genetically susceptible individuals [16].

40 *Increasing the sensitivity of detection of viral proteins in the pancreas*

41 The development of antigen retrieval techniques (often termed “Heat Induced Epitope
42 Retrieval” (HIER)) in the 1990s was game changing in the field of pathology. This
43 technique allowed certain antigens that had previously been inaccessible to antibodies
44 in formalin-fixed paraffin-embedded tissues, to be unmasked. HIER also allows the
45 use of reduced primary antibody incubation times (and lower dilutions) such that
46 staining is revealed in formalin-fixed tissues that fail to stain by conventional methods
47 [48]. In summary, the application of this technique increases the sensitivity of antigen
48 detection by individual antisera. A second major development in the enterovirus field
49 was the production of more sensitive, broad spectrum antisera directed against
50 enteroviral capsid proteins. One such antibody (clone 5D8/1) is marketed
51 commercially by Dako. The antibody is monoclonal in origin, was raised against the

1 VP1 protein of Coxsackievirus B5 and has provided particularly sensitive and specific
2 detection opportunities such that it has now been used extensively to detect
3 enterovirus in formalin fixed tissues [49-55]. In combination, therefore, the use of
4 HIER and the availability of new antisera have enabled an increase in both the
5 spectrum of enteroviral serotypes which can be detected in fixed tissues and an
6 improvement in the sensitivity of their detection [49-57] (and unpublished results
7 Richardson *et al*).

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14 These important methodological advances have provided better tools for use in the
15 search for enteroviral proteins in tissue from recent-onset type 1 diabetes patients.
16 Accordingly, in a landmark paper published in 2007, Dotta *et al* reported that
17 enteroviral VP1 protein was present in the islets of 2 of 5 recent-onset type 1 diabetes
18 cases and in a whole pancreas graft recovered from a 26 year old recipient [23].
19 Importantly, this work revealed the presence of small numbers of intensely-stained
20 endocrine cells, shown to be beta cells, within the islets. We now term these “Dotta”
21 cells to recognise the importance of this contribution. The immunohistochemical
22 evidence was supported by electron microscopic studies revealing the presence of
23 virus particles within islet cells and by the isolation of a serotype of CVB4 (now
24 referred to as the “Tuscany” isolate) from one of the cases [23]. Subsequent to this,
25 we conducted a more comprehensive analysis of a cohort of recent-onset type 1
26 diabetes cases collected within the UK and were able to demonstrate the presence of
27 VP1+ “Dotta” cells, again proven to be beta cells, in 44 of the 72 (61%) cases
28 examined. In contrast, only 4 “Dotta” cells were identified in 3 islets from the 50
29 (7.7%) neonatal and paediatric control cases examined [58]. We further demonstrated
30 that HIER allowed the antiserum used in the original study that had failed to detect
31 viral proteins [39], to now reveal staining in specific islets. Strikingly, these islets
32 were also stained positively by clone 5D8/1 in serial sections. A further pan-
33 enterovirus antibody (Clone 9D5; Millipore) was also found to label individual
34 endocrine cells in the islets of patients with type 1 diabetes [58]. A comparison of the
35 staining patterns achieved with these three antisera in heart tissue from CVB-infected
36 neonates revealed that the Dako 5D8/1 clone was both more sensitive and produced
37 more consistent staining in the face of a variety of different fixatives when compared
38 to the other two [58,57] (and unpublished results Richardson *et al*). Hence, when
39 considered together, these studies provided the first firm evidence that enteroviral
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1 infection, confined to beta cells, can be detected in the islet cells of a significant
2 number of people with a recent diagnosis of type 1 diabetes.
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5 The UK cohort used in our work represents the largest single collection of recent-
6 onset type 1 diabetes pancreases in the world and has proved invaluable as a means to
7 study the underlying pathology of this illness. However, despite this there are several
8 limitations (notably the historical nature of the collection; its specific geographical
9 location and the non-uniform fixation methods employed). To address these, we have
10 recently had the opportunity to examine a second cohort of pancreases from patients
11 with type 1 diabetes, collected under the auspices of the Juvenile Diabetes Research
12 Foundation's network of Pancreatic Organ Donors (nPOD) with Diabetes programme
13 (<http://www.jdrfnpod.org/index.php>) [59]. This important and growing resource
14 promises much, since it should facilitate future collaborative studies to better
15 understand the pathogenesis of type 1 diabetes. However, the number of recent-onset
16 cases (<1 year) in this collection is still very small. Indeed, initially we were only able
17 to access 1 case of short clinical duration among 10 cases which retained insulin-
18 containing islets[60]. We also examined a further 7 cases in which only insulin-
19 deficient islets were present. Importantly, the mean time since diagnosis in the 17 type
20 1 diabetes cases studied was 11.9 ± 2.3 years; which compared to only 8.2 ± 4.1 months
21 (i.e. 0.68 years) in the UK cohort. However, despite this, we noted that many of the
22 pathological features highlighted in the UK cohort could still be observed in the
23 American cases. We were therefore intrigued to investigate whether islet enteroviral
24 infection could be detected in this cohort.
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44 In short, the firm answer to this question is “yes”. In 8 of the 10 cases where residual
45 beta cells were present, intensely stained VP1+ (“Dotta”) cells were seen [60]. By
46 contrast, no evidence of staining was observed in the 7 cases with only insulin-
47 deficient islets. Serial sections of a representative islet from an nPOD case are shown
48 in Fig2 demonstrating VP1 expression in an insulin-containing islet with hyper
49 expression of class I MHC. Importantly, VP1+ cells were seen in only 1 of 12 age-
50 matched non-diabetic controls [60]. Within this (and the UK) cohort, it is important
51 to emphasise the absolute numbers of VP1+ cells within any given patient is
52 vanishingly small. Indeed, even in those cases with the highest proportion of “Dotta”
53 cells, we calculate that less than 0.005-0.01% of the cells within the entire pancreas
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1 section stain positively for VP1 suggesting that viral infection of beta-cells does not
2 proceed with a typical, acute, lytic course.
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5 One further consideration is important since it has been proposed that clone 5D8/1
6 can, under certain conditions, cross-react with two additional proteins in the pancreas,
7 creatine kinase B (CKB) and ATP5B [61]. To address this issue, we have examined
8 this cross-reactivity in greater detail and show that, under optimised conditions, the
9 immunostaining achieved with clone 5D8/1 in formalin-fixed paraffin embedded
10 tissue or cells, retains its specificity for VP1 [62].
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18 Additional support for the hypothesis that the VP1 staining of the beta cells in the
19 islets of type 1 diabetes patients represents a *bona fide* infection is the finding that the
20 pathogen recognition receptor, protein kinase R (PKR) is selectively up-regulated in
21 VP1+ islet cells. This has been confirmed in both the UK and nPOD cohorts [60]. It
22 has been shown by others that PKR is both induced and activated following an
23 enteroviral infection and that this leads to phosphorylation of an elongation factor,
24 eIF2alpha, involved in protein synthesis [63,64]. This causes translational arrest and,
25 as such, will result in the selective depletion of the more labile proteins. Therefore, we
26 took advantage of this situation to study the expression of a labile, anti-apoptotic
27 protein, Mcl-1, in VP1+ (PKR+) islet cells. Mcl-1 is constitutively expressed at high
28 levels in most beta cells [60] and from in vitro studies, the protein is understood to be
29 a critical determinant of beta cell fate in response to stressors (such as pro-
30 inflammatory cytokines and viral infection) [65]. Importantly, we observed that Mcl-1
31 was selectively depleted from beta-cells expressing VP1 and PKR suggesting that
32 these individual cells might then be rendered more sensitive to the detrimental effects
33 of the pro-inflammatory milieu existing within inflamed islets [60].
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49 *Is islet VP1 immunopositivity the tip of an iceberg?*

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51 The presence of interferon-alpha within most residual beta cells patients at clinical
52 onset of type 1 diabetes contrasts with the relative scarcity of “Dotta” cells expressing
53 enteroviral VP1 protein. *How can this be explained?* In an acute lytic enteroviral
54 infection the positive strand RNA of the virus is transcribed within the infected cell to
55 produce a negative strand RNA template. This in turn is transcribed to produce
56 hundreds of copies of positive strand RNA viral molecules which are then translated,
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1 with synthesis of complete viral particles. Upon reaching a critical mass of these
2 particles the cells bursts and releases virus. By contrast, in a chronic non-lytic
3 enteroviral infection there are equal numbers of positive and negative viral RNA
4 molecules and there is little synthesis of complete viral particles expressing viral
5 capsid proteins [66-68]. Thus the “Dotta” cells seen in type 1 diabetes may represent
6 the ‘tip of the iceberg’ of viral infection within the islet. The beta cells synthesizing
7 interferon-alpha may well be chronically infected by enterovirus existing, not as
8 complete viral particles, but as double stranded RNA, a well-recognised stimulant of
9 interferon synthesis (reviewed in [69]). In our study [58], we found that among young
10 children, enteroviral infection of beta cells was much more frequent in patients with
11 type 1 diabetes than those without. However, we suspect that it is not just the presence
12 or absence of virus *per se* within a beta cell which is of most significance for the onset
13 of diabetes. Rather, it may be the response of the cell to an on-going viral infection
14 that matters. Conceivably, it is those children who respond most vigorously to an
15 early beta-cell enteroviral infection, by mounting a strong interferon response, who
16 are at greatest risk of triggering islet autoimmunity. Interferon secretion will cause
17 hyper expression of class I MHC within the islet endocrine cells, which in turn
18 probably initiates insulinitis. Those children who respond minimally (or not at all) to an
19 enteroviral infection of beta cells may be relatively protected from developing islet
20 autoimmunity.
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38 *Are the beta cells trying to fight back?*

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40 Studies in the UK recent-onset paediatric cohort of type 1 diabetes patients have
41 provided evidence that endocrine cells (both beta and alpha) display a 10-fold
42 increase in proliferation when compared to controls [70]. This was strongly associated
43 with the presence of insulinitis suggesting that factors released from the immune cells
44 could be driving an attempt by the beta cells to increase their numbers in the face of
45 the attack [70]. The presence of proliferating endocrine cells was also frequently
46 observed in islets with VP1 staining, implying that a viral infection could be inducing
47 the recruitment of the immune cells, which in turn release factors that promote
48 endocrine cell proliferation [71]. The finding that endocrine cell proliferation was
49 increased in some islet autoantibody positive organ donors with evidence of insulinitis
50 [72] and that proliferation was not observed in adult, longer duration type 1 diabetes
51 patients [73] who rarely have evidence of insulinitis, lends support to this hypothesis.
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2 *Proposed sequence of events - 2014*
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4 In individuals at-risk of developing type 1 diabetes, a non-cytopathic chronic viral
5 infection of pancreatic beta cells is sensed by host pathogen recognition receptors.
6 Activation of these receptors induces the expression of interferon-alpha (as part of an
7 innate immune response to the double stranded RNA of the virus) which results in
8 hyper-expression of class I MHC by the endocrine cells of the islet and release of
9 factors that promote the recruitment of immune cells. However the islet has a
10 protective basement membrane that can prevent direct contact between the immune
11 cells and the beta cells. As the recruitment of more immune cells to the affected islets
12 occurs a slow progression of insulitic destruction may occur (demonstrated by peri-
13 insulitis, focal insulitis, progressive insulitis and finally invasive insulitis), invasive
14 insulitis occurring once the basement membrane has been sufficiently degraded by
15 enzymes released by the infiltrating cells. At this point the invasive immune cells
16 actively target and kill the beta cells leading to the formation of insulin-deficient
17 islets, where under certain circumstances the basement membrane can reform. This
18 progressive destruction of islets (over years) leads to the development of clinical
19 diabetes (Fig5).
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34 *Summary*
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36 We have come a long way in 25 years - but we still have a very long way to go. Many
37 questions remain unanswered and the limited amount and type of material available in
38 which to address these questions restricts the speed at which they can be answered.
39 However, the establishment of large international collaborative networks such as
40 nPOD and EU Framework 7 funded PEVNET (Persistent virus infection as a cause of
41 pathogenic inflammation in type 1 diabetes – an innovative research program of
42 biobanks and expertise) are enabling expert researchers to combine their skills,
43 question hypotheses and push forward our knowledge in this area. In particular the
44 nPOD (organ donor pancreases) and the Norwegian DiViD (distal pancreatectomy
45 specimens in patients with recent onset type 1 diabetes) collections are likely to prove
46 critical to finally answering the virus question as these tissues are optimally
47 preserved, can be used to perform islet dissection and are suitable for next-generation
48 sequencing analysis. We are sure that the next 25 years will prove to be even more
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1 fruitful than the last, as this collective effort really starts to help us understand the
2 changes that beset the pancreas of type 1 diabetes patients.
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5 Twenty-five years ago findings in the diabetic pancreas suggested that both
6 hypotheses raised by Gepts might be right: viral infection of islet beta cells might
7 lead to a destructive autoimmune response directed against them. Twenty-five years
8 further on we still think he is right!
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28 1 diabetes research project sponsored by the Juvenile Diabetes Research Foundation
29 International (JDRF) and with a JDRF research grant awarded to the nPOD-V
30 consortium. Organ Procurement Organizations (OPO) partnering with nPOD to
31 provide research resources are listed at www.jdrfnpod.org/our-partners.php.
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FIGURE LEGENDS

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4 Fig1: The lobular nature of type 1 diabetes. A photomicrograph of a pancreas from a
5 type 1 diabetes patient (from the nPOD collection) stained via immunohistochemistry
6 for the presence of glucagon (A) and a serial section stained for insulin (B). The lobe
7 containing ICIs can be seen on the right (hashed line), whereas the neighbouring lobe
8 contains only IDIs.
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14 Fig2: A photomicrograph of a non-diabetic control (A-C) and a type 1 diabetes patient
15 (D-F) immunostained for insulin (red) and glucagon (brown – A and D), class I MHC
16 (B and E) and enteroviral VP1 (C and F).
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22 Fig3: The proposed sequence of events in islets in type 1 diabetes 25 years ago (Alan
23 Foulis, Pathology Society 1987 Oakley Lecture).
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27 Fig4: Insulitic lesions in type 1 diabetes. Representative islets from nPOD cases
28 immunostained for insulin (blue), the Pan lymphocyte marker CD45 (green) and
29 TOPRO (Red) demonstrating the proposed progression of the insulitic lesion. A. Peri-
30 insulitis, B. Focal insulitis, C. Progressive insulitis and D. Invasive insulitis.
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36 Fig 5: Proposed sequence of events in type 1 diabetes – 2014.
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Figures

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Fig1: Lobular loss of insulin-containing islets

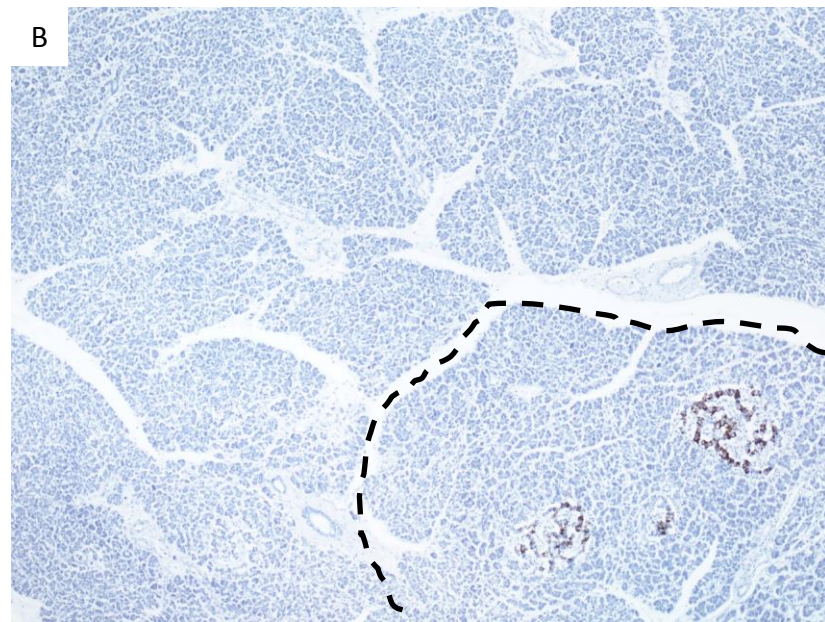
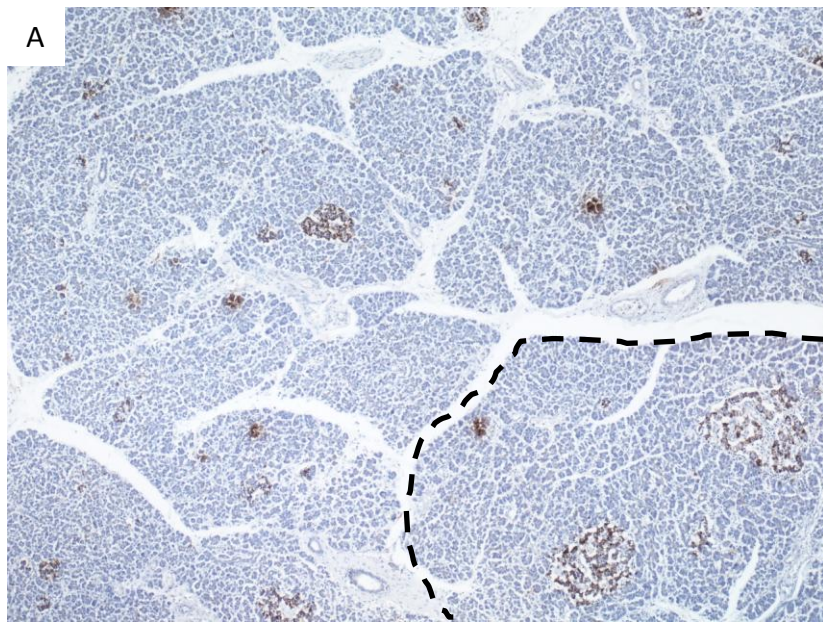


Fig2: Hyper expression of Class I MHC in type 1 diabetes

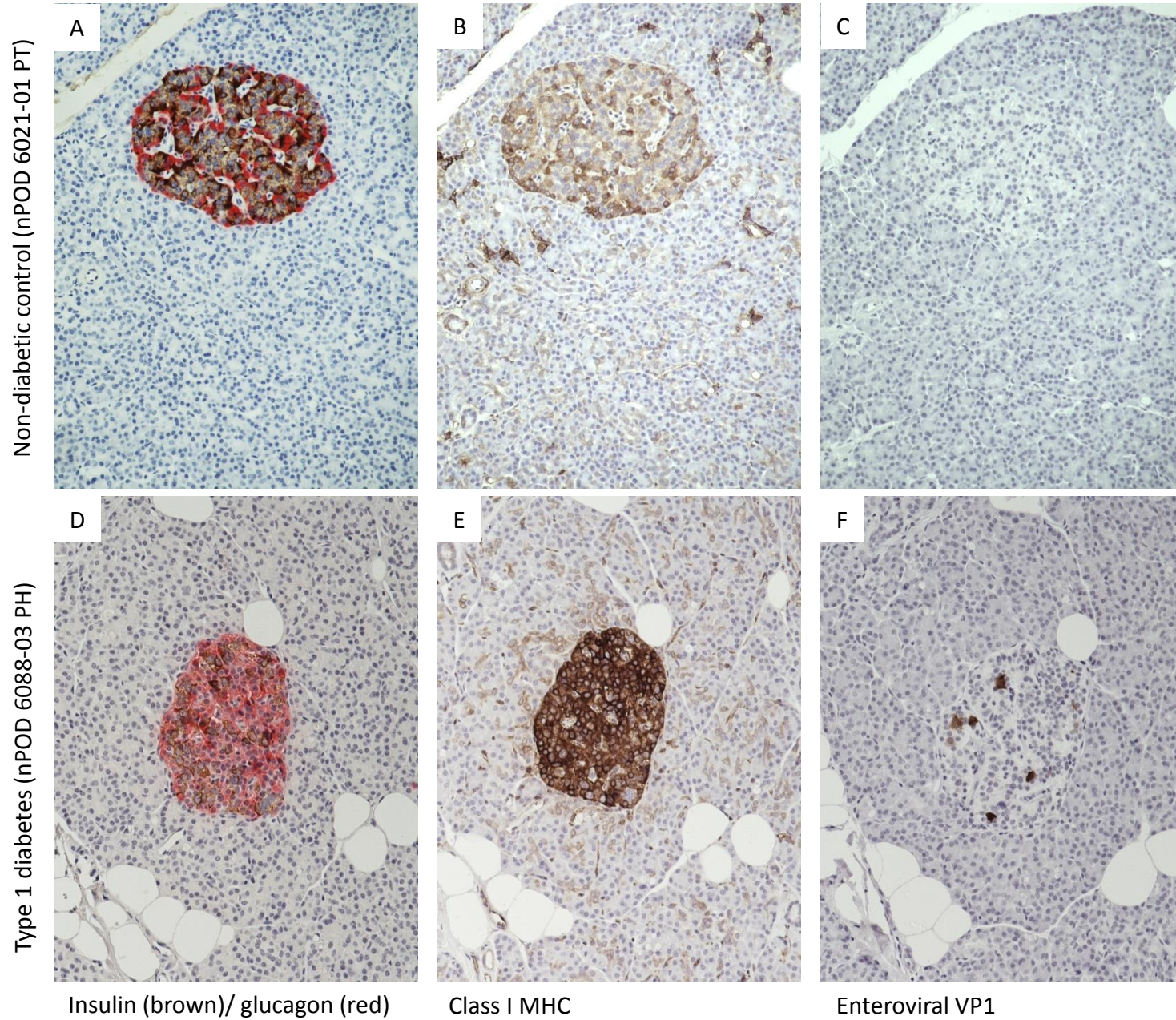


Fig3: Possible sequence of events in islets in type 1 diabetes (1987 Oakley Lecture)

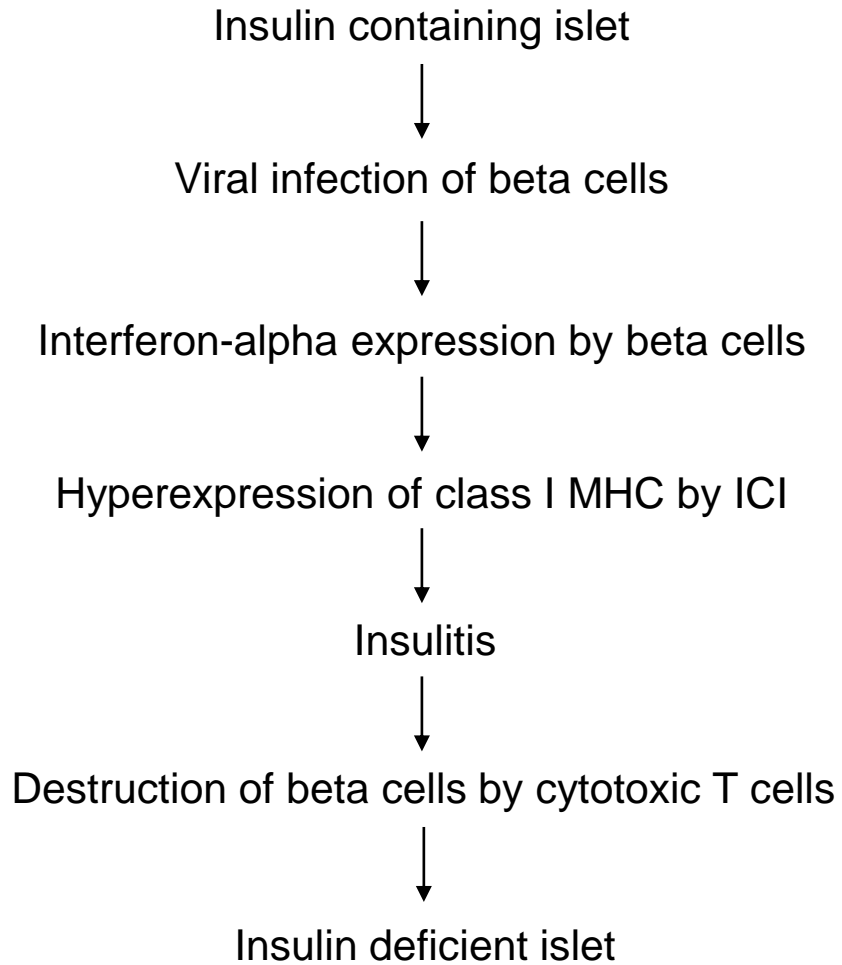
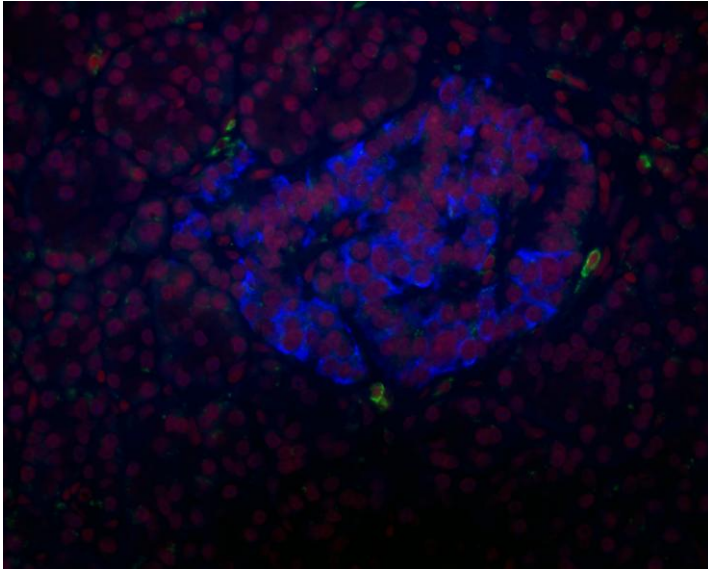
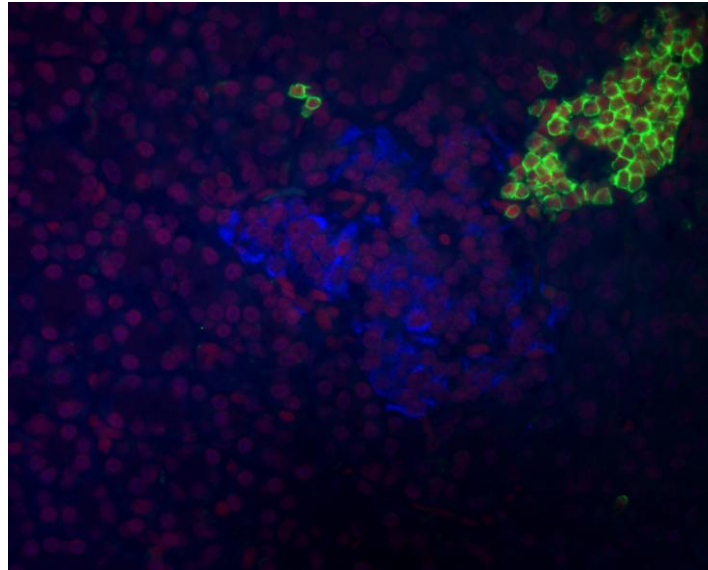


Fig4: Heterogeneity of insulinitis; is it related to the stage of destruction (peri-islet, polar, progressive and invasive)?

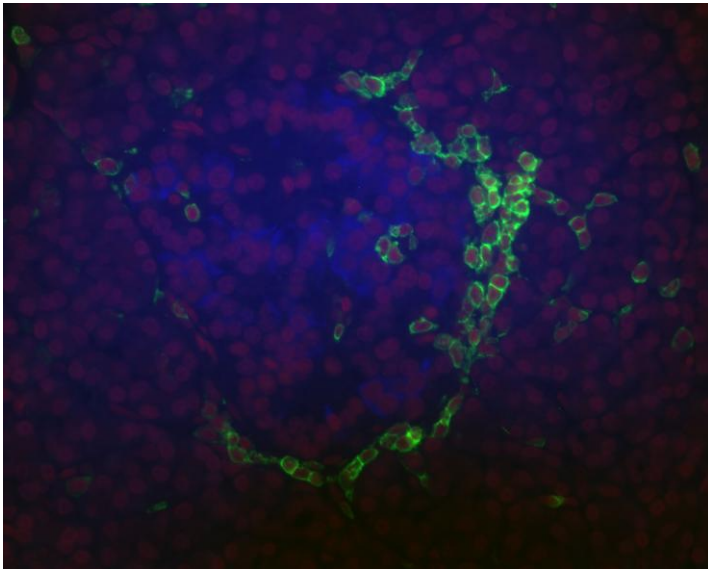
A. nPOD 6070 – Peri-islet insulinitis



B. nPOD 6070 – Polar insulitis



C. nPOD 6052 – Progressive insulitis



D. nPOD 6052 – Invasive insulitis

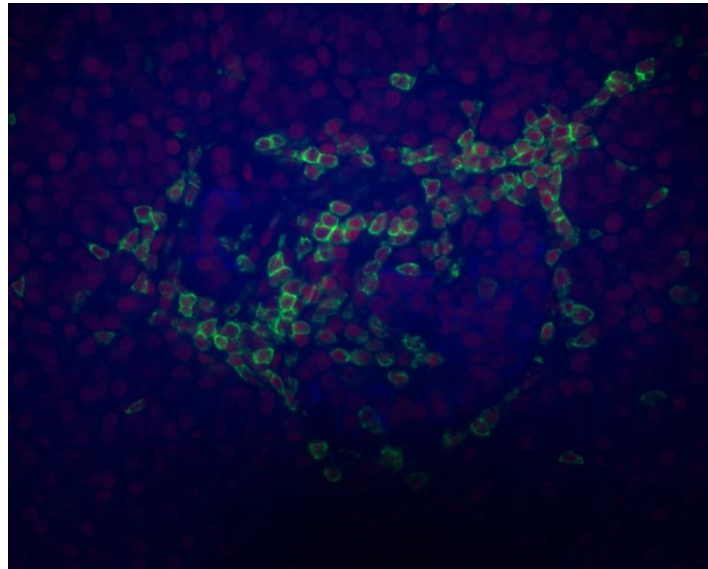


Fig5: Progression in type 1 diabetes

