Presented antigen from damaged pancreatic β cells activates autoreactive T cells in virus-mediated autoimmune diabetes

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The induction of autoimmunity by viruses has been attributed to numerous mechanisms. In mice, coxsackievirus B4 (CB4) induces insulin-dependent diabetes mellitus (IDDM) resembling the final step of disease progression in humans. The immune response following the viral insult clearly precipitates IDDM. However, the molecular pathway between viral infection and the subsequent activation of T cells specific for islet antigen has not been elucidated. These T cells could become activated through exposure to sequestered antigens released by damaged β cells, or they could have responded to factors secreted by the inflammatory response itself. To distinguish between these possibilities, we treated mice harboring a diabetogenic T cell repertoire with either the islet-damaging agent streptozotocin (STZ) or poly I:C, which nonspecifically activates T cells. Significantly, only treatment of mice with STZ resulted in IDDM and mimicked the effects observed following CB4 infection. Furthermore, antigen-presenting cells from STZ-treated mice were shown to directly activate autoreactive T cells and induce diabetes. Therefore, the primary role of CB4 in the precipitation of IDDM is to damage tissue, causing release and presentation of sequestered islet antigen. These events stimulate autoreactive T cells and thereby initiate disease.


Introduction

Coxsackieviruses have long been associated with the development of insulin-dependent diabetes mellitus (IDDM) (1–4). Understanding how coxsackieviruses mediate diabetes has been complicated by this virus’s unique tropism for the pancreas as well as the apparent molecular similarity between the viral P2C gene product of the B4 strain (CB4) and the pancreatic islet antigen GAD65 (5). Recently, the mechanistic role of this virus was dissected by infecting mice representing different experimental models of diabetes (6), and the results favored a mechanism of T cell activation through bystander damage over induction through molecular mimicry. CB4 infection of NOD mice, which carry the NOD MHC allele to which presentation of the cross-reactive epitope is restricted, did not enhance or change the onset or progression of diabetes. Additionally, no proliferative response to either of the cross-reactive epitopes of P2C or GAD was observed following infection. However, CB4 infection of BDC2.5 transgenic (Tg) mice, which harbor a transgene encoding a diabetogenic T cell receptor (TCR) specific to an islet granule antigen other than GAD65 (7) and not cross-reactive with CB4, resulted in the rapid onset of diabetes (6). In our colony, BDC2.5 Tg mice develop peri-insulitis at an accelerated rate but do not progress to spontaneous diabetes. CB4 infection activated the Tg memory T cell population prior to disease onset, precipitating clinical diabetes in these mice.

Whether memory T cells can be activated due to the release of sequestered antigen from the damaged tissue or simply by cytokine induction following viral infection remained as an important question. To address this possibility, BDC mice were infected with lymphocytic choriomeningitis virus (LCMV), which can cross-activate nonspecific resting memory lymphocytes during infection but does not destroy islet cells. LCMV infection did not induce diabetes nor any increase in insulitis (6). Therefore, viral expansion of nonspecific T cell responses through cytokine induction is most likely not activating islet-specific memory T cells to mediate disease. Yet, the dynamics of the LCMV infection may somehow suppress (8, 9) or interfere with (10–12) disease development.

To more completely distinguish between the mechanisms that account for coxsackievirus-mediated IDDM, we treated BDC2.5 mice with two environmental agents that mimic aspects of viral infection. Treatment with poly I:C effectively induces type 1 cytokine activation similar to that observed following viral infection (13). Alternatively, administration of a single subdiabetogenic dose of streptozotocin (STZ) directly damages β cells in the pancreatic islets of Langerhans with only minimal immune cell involvement (14–17). If coxsackievirus induces IDDM through damage to β cells, antigen release, and activation of self-reactive T cells, then treatment of BDC2.5 Tg mice with STZ and not poly I:C should activate the resting anti-islet memory T cells,
leading to the rapid development of diabetes. If, however, viral infection merely activates the resting autoreactive T cells via a cytokine response, then poly I:C treatment alone should be sufficient to induce disease.

Methods

Mice and virus. NOD/Shi and NODSCID mice were obtained from the rodent breeding colony at The Scripps Research Institute. BDC2.5 Tg mice were a gift from Diane Mathis (Joslin Diabetes Center, Harvard University, Boston, Massachusetts, USA; formerly of Institut de Génétique et de Biologie Moleculaire, Strasbourg, France) and have been maintained and bred to NOD/Shi in our mouse colony. Virus stocks of coxsackievirus group B type 4 Edwards strain 2 (CB4 strain E2) were obtained from Charles Gauntt (U. of Texas–San Antonio, San Antonio, Texas, USA) and were derived from stocks obtained from Roger Loria (Medical College of Virginia, Commonwealth University of Virginia, Richmond, Virginia, USA; ref. 3). CB4 was originally isolated from patients with IDDM, passed through murine pancreatic β cells, and subsequently found to induce a diabetes-like disease in mice (3, 4). Virus stocks of CB4 strain E2 were prepared and titered as previously described (18). Mice were infected at 12 weeks of age intraperitoneally with 100 plaque-forming units (pfu) of CB4. Mice were treated with 100 µg per mouse of poly I:C (Sigma Chemical Co., St. Louis, Missouri, USA) in PBS with an intraperitoneal injection. Mice were treated with STZ (Sigma Chemical Co.) at various concentrations as indicated in the text. STZ was prepared in citrate buffer immediately prior to intraperitoneal injection. Blood glucose was measured in tail vein or eye bleeds from nonfasting mice at various times from 1–28 days postinfection or posttreatment on a standard glucometer with a range of 20–400 mg/dl. Mice were considered diabetic when two or more consecutive readings (24 hours apart) were greater than 300 mg/d.

Immunohistochemical staining. Organs were removed and placed in 10% formaldehyde, then processed for paraffin sectioning. Alternatively, organs were snap-frozen in OCT on dry ice. Immunohistochemistry was performed on 4-µm-thick paraffin sections for insulin staining or 7-µm cryocut sections for the other antibodies that were prepared and blocked with avidin and biotin (Vector Laboratories Inc., Burlingame, California, USA). The primary antibodies were to insulin (DAKO Corp., Carpinteria, California, USA), F4/80 (Serotec Inc., Washington, DC, USA), CD4 and CD25 (PharMingen, La Jolla, California, USA), and the secondary antibody was a biotinylated anti–guinea pig IgG (for insulin staining) or anti-rat IgG (for the other antibodies) used in conjunction with Vectastain ABC (peroxidase) kits (Vector Laboratories Inc., Carpinteria, CA). Staining was detected using diaminobenzidine as a chromagen. Sections were counterstained in Mayer’s hematoxylin (Sigma Chemical Co.) and mounted in Permount (Fisher Scientific Co., Pittsburgh, Pennsylvania, USA). Quantitation of the degree of pancreatic insulitis was evaluated using both anti-insulin–stained and multiple hematoxylin and eosin–stained pancreatic sections from infected and uninfected, age-matched NOD and BDC2.5 Tg mice, and the mice were scored for peri-insulitis and insulitis. Statistical analysis using χ-square tests were done on paired groups.

FACS analysis. Peripheral blood was obtained from mice before and 1 week after treatment with STZ or poly I:C. Erythrocytes were lysed by NH4Cl treatment and lymphocytes incubated with phycocerythrin-labeled (PE-labeled) antibody to the TCR variable chain Vβ4 and Cy-chrome-labeled CD4 (PharMingen), and FITC-labeled antibody to CD25 (Sigma Chemical Co.), CD44, or CD69 (PharMingen). Cells were fixed in 1% paraformaldehyde and analyzed with a Becton Dickinson FACS. A minimum of 200,000 events were analyzed for each sample.

Proliferation assays. Antigen-presenting cells (APCs) were enriched from the spleens of untreated and STZ-treated NOD and NODSCID mice by plastic adherence as previously described (19). Briefly, spleens were harvested and forced through mesh screens to generate single cell suspensions. Erythrocytes were lysed by NH4Cl treatment, and the remaining cells were resuspended in RPMI with 10% FCS at 10 million cells per milliliter. Cells were incubated on Petri dishes for 3 hours (37°C, 5% CO2). Afterwards, nonadherent cells were removed by multiple washes with RPMI. Adherent cells were removed by scraping, counted, and placed in appropriate media for use in either T cell proliferation assays or adoptive transfer. For proliferation assays, the APC-enriched population was irradiated (15 Gy).

For T cell proliferation assays, spleens from untreated or STZ-treated BDC2.5 TCR Tg mice were harvested, and erythrocytes were removed and cultured in triplicate on 96-well plates at 2 × 105 cells per well in HL-1 media. To the BDC splenocytes, in triplicate, varying amounts of adherent cells were added (0, 1.25 × 104, 2.5 × 104, 5 × 104, or 10 × 104 cells). Additionally, triplicate controls were loaded with added APCs and peptide (20) (7 µM) were performed. Cultures were incubated for 2 days at 37°C and pulsed for 18 hours before harvest with 1 µCi 3H-thymidine. Incorporated 3H was measured using a liquid scintillation counter. The stimulation index was calculated for each splenocyte/APC sample, as compared with control wells incubated on the same plate but lacking any additional adherent cells or antigen.

Adoptive transfer. Donors were either 12-week-old untreated BDC2.5 TCR Tg mice or 6- to 8-week-old untreated or STZ-treated NODSCID mice. From BDC Tg mice, splenocytes were harvested, single cell suspensions generated, and erythrocytes removed, counted, and transferred at 2 × 105 to 5 × 106 cells per mouse. Recipient mice for the BDC splenocytes were NODSCID mice that had been treated with either STZ or PBS 7 days before. Alternatively, splenocytes from treated NODSCID mice were harvested and enriched for APCs by adherence to plastic as described above. APCs were trans-
ferred into untreated 12-week-old BDC mice at $5 \times 10^5$ to $10 \times 10^5$ cells per mouse. Mice were monitored weekly for changes in blood glucose.

**Results and Discussion**

Prior work has established the ability of poly I:C, a double-stranded nucleic acid, to invoke a type 1 IFN antiviral response without the accompanying damage of a viral infection (21). Additional work by Tough et al. (13) had shown that poly I:C treatment could act similarly to LCMV infection and stimulate nonspecific memory T cells. Whether these activated T cells could then participate in an autoimmune or pathologic response has been controversial. To examine whether the upregulation of the type 1 IFN pathway could induce IDDM through stimulation of memory T cells, we treated BDC2.5 Tg mice with poly I:C (100 µg per mouse) and monitored for disease onset. While CB4 induces diabetes in 67% of the infected BDC2.5 Tg mice within 4 weeks, poly I:C treatment was unable to induce clinical disease (Figure 1a). The IFN response to poly I:C was comparable to that observed following LCMV infection, as IFN-γ levels were similar in mice 3 days after treatment or infection as measured by ELISA (350 ng/ml). To establish whether treatment was able to increase the level of insulitis without inducing diabetes, we stained pancreatic tissue sections with insulin-specific antibody to demonstrate any loss of insulin-containing β cells. However, no increase in the level of peri-insulitis or insulitis was observed following treatment (Table 1). No histological changes were observed in the pancreas between age-matched treated and untreated BDC2.5 Tg mice (Figure 2a). To determine whether poly I:C treatment was able to activate lymphocytes in the BDC2.5 Tg mouse, lymphocytes were isolated from treated and untreated mice and stained with antibodies to cell surface molecules for analysis by flow cytometry. Although none of the CD4+ Vβ4+ (Tg TCR) lymphocytes showed changes in CD25, CD44, or CD69, indicating a lack of activation, the CD4– Vβ4+ T cell subset, representing primarily CD8+ T lymphocytes, demonstrated upregulation of these markers, denoting their activation (Figure 3a). Therefore, treatment of BDC2.5 Tg mice with poly I:C was comparable to infection with the noncytopathic virus, LCMV, and in neither case were the resting memory Tg CD4+ cells activated or disease induced.

Treatment with a single subdiabetogenic dose of STZ damages β cells directly with only minimal immune cell involvement (14–17). While a multiple high- or low-dose regime of STZ leaves the mouse without functional β cells and induces autoimmune diabetes, a single low-dose treatment causes significantly less damage and does not induce clinical disease (14, 15, 17). During coxsackievirus infection of most mouse strains, limited damage to the β cells is observed. The loss of a few β cells is most likely counterbalanced by the limited ability of β cells to regenerate. To mimic the minimal level of viral damage to the islet, mice would need to be treated with a single low dose of STZ that would damage only a few β cells, leaving the majority of insulin-producing cells intact without altering clinical disease incidence on its own.
Local increases of activated macrophages in pancreatic islets accompany the progression of insulinitis in mice developing diabetes (22, 23). This sequence is clearly visible in both NOD mice developing spontaneous diabetes and CB4-infected BDC Tg mice. Following single-dose STZ treatment of NOD mice, insulin production decreased without significant insulinitis (Figure 2a). Consequently, fewer activated macrophages were present in the affected islets. To determine whether the islets of the STZ-treated BDC mice were phenotypically similar to the islets from virus-infected mice, pancreatic tissue sections were immunostained with F4/80 antibody to locate activated macrophages. As depicted in Figure 2c, the presence of macrophages in the islets of STZ-treated BDC mice closely resembled the situation in the islets of CB4-infected BDC mice. In contrast, BDC mice treated with poly I:C and NOD mice given STZ did not exhibit pronounced insulinitis, nor was there any significant increase in activated macrophages.

To correlate the shift in state from resting to activation in the transgenic Vβ4+ lymphocytes with the increase in insulinitis and onset of diabetes, lymphocytes were isolated from the spleens and pancreatic lymph nodes of treated and untreated BDC2.5 mice and stained with antibodies to cell surface molecules including Vβ4 (Tg TCR), IL-2 receptor (CD25), CD44, CD69, and L-selectin (CD62L) for analysis by flow cytometry. As previously reported, the predominant phenotype of the transgenic CD4+ Vβ4+ population of T cells signified resting memory as demonstrated by high expression of CD44 and absence of expression of CD25, CD69 (Figure 3b), and CD62L (data not shown). Strikingly, five days after STZ treatment, transgenic lymphocytes were activated, as confirmed by the increased numbers of Vβ4+ lymphocytes staining for the activation markers CD25 and CD69, as well as the memory and activation marker CD44 (Figure 3). However, NOD mice had no obvious increase in activation in CD4+ T cells following STZ treatment. Additionally, similar STZ treatments of OT-II TCR Tg mice that express an MHC class II restricted TCR specific for ovalbumin (24) did not activate T cells when measured by a similar FACS analysis for upregulation of the activation markers CD25, CD44, or CD69 (data not shown). Thus, the single low-dose treatment of STZ is acting solely as a β cell toxin and did not directly activate T cells. The shift in T cell activation by a significant portion of Vβ4+ cells correlated with the rapid and significant increase in insulinitis composed predominantly of Vβ4+ (Tg) cells, β cell loss, and observed diabetes similar to that observed following CB4 infection. However, in this case, a β cell toxin replaced the virus, further underscoring the requirement for β cell damage in the development of IDDM.

The ability of STZ to elicit diabetes with kinetics and phenotype similar to those of diabetes elicited by CB4 infection demonstrates that virus-mediated diabetes is the result of direct damage to the β cells followed by

| Table 1 |
| Summary of histological analysis of pancreatic sections from treated mice |
| Peri-insulitis | Insulitis | Total islets |
| Poly I:C - BDCs | 43% | 42% | 124 |
| STZ - BDCs | 7% | 85% | 114 |
| Untreated - BDCs | 42% | 56% | 113 |

Mice were sacrificed 4 weeks after treatment with poly I:C or STZ and examined for peri-insulitis and insulinitis. Islets were scored for inflammatory cells as clean, peri-insulitis, or insulitis. Mice were matched for age and sex. Over 100 islets were scored per group. Scored groups were compared for statistical significance by standard χ-square tests. Of the relevant comparisons, significant differences were observed for both peri-insulitis (P < 0.001) and insulinitis (P < 0.001) between STZ-treated BDC mice and untreated BDC mice or poly I:C-treated BDC mice. Additionally, no significant difference was observed between poly I:C-treated and untreated BDC mice.

For this purpose, a pilot study was performed with a single treatment of STZ. One of three concentrations of STZ, 40, 80, or 160 mg/kg, was administrated and mice were monitored for clinical disease. Initial treatments with STZ clearly indicated that induction of diabetes was dose-dependent (Figure 1b), and a single dose of 80 mg/kg was chosen for further study. After administration of 80 mg/kg of STZ, BDC2.5 Tg mice developed IDDM in a manner and time frame similar to that induced by CB4 infection of BDC2.5 Tg mice (Figure 1b). That is, 67% of the mice became diabetic within 2–4 weeks of treatment. To ensure that this was not an effect of the NOD MHC background, since this murine strain spontaneously develops diabetes, a large cohort of age-matched (12-week-old) female NOD mice were treated simultaneously (Figure 1c). The NOD mice had a slight increase in the incidence of diabetes compared with that of spontaneous disease in our colony, but this incidence of spontaneous disease in our colony, but this incidence of the NOD MHC background, since this murine strain spontaneously develops diabetes, a large cohort of age-matched (12-week-old) female NOD mice were treated simultaneously (Figure 1c). The NOD mice had a slight increase in the incidence of diabetes compared with that of spontaneous disease in our colony, but this incidence of STZ-induced NOD disease was significantly lower than that in the BDC2.5 Tg treated mice (P < 0.001).

STZ-treated BDC2.5 Tg mice had a marked loss of insulin-producing β cells, which was accompanied by a dramatic rise in insulinitis. The quantitated insulinitis index showed a significant increase (P < 0.001) that coincided with the onset of diabetes (Table 1). To characterize the nature of these infiltrating lymphocytes, tissue sections were immunostained with antibody to CD4, CD25 (Figure 2b), and Vβ4 (data not shown). The infiltrating cells were predominantly CD4+ lymphocytes and a sizable percentage of the infiltrating population was indeed CD25+ or IL-2 receptor–positive, reflecting an activated state (Figure 2b). As we had observed previously in the islets of CB4-infected mice (6), the predominant cell type was indeed the transgenic T cell and therefore both CD4+ and Vβ4+. Additionally, while loss of insulin staining in β cells without strong insulinitis is a common event in STZ-treated NOD mice (Figure 2a), this was not the case in the STZ-treated BDC mice since loss of insulin production by β cells was closely associated with immune cell infiltration and insulinitis. This outcome strongly suggests that resting memory T cells become activated due to the release of sequestered antigen during the environmental insult.
activation of an autoreactive memory response that attacks and destroys the islets. The direct damage of \( \beta \) cells associated with either the single low-dose STZ (at 80 mg/kg) treatment or CB4 infection is, by itself, not sufficient to cause diabetes, since nontransgenic mice show little enhancement in disease. However, in both instances, sufficient \( \beta \) cell damage occurs to allow for release of sequestered \( \beta \) cell antigen and stimulation of the resting memory T cell population. This result, most likely, reflects the ability of pancreatic APCs to process and present \( \beta \) cell antigen and thereby stimulate the BDC T cell.

To demonstrate that APCs take up islet antigen and stimulate the BDC T cells, we performed in vitro proliferation assays. APCs were enriched using plastic adherence from mice either treated or not treated with STZ; BDC T cells from untreated mice were then cocultivated with these APCs. As a control, APCs from untreated mice were cultured with BDC T cells and antigenic peptide (20). Previous work has demonstrated that enriched APCs can harbor antigen following viral infection or during local pathogenic inflammation and can present this antigen in in vitro proliferation assays (19, 25–27). In this situation, the pancreat-

**Figure 2**

(a) Histological analysis of pancreata of treated BDC Tg and NOD mice. Representative sections of the pancreas from STZ-treated BDC Tg mice and poly I:C-treated BDC Tg mice reveal islet cell integrity and function. The function of \( \beta \) cells was confirmed by immunohistochemistry using antibody to insulin (INS) (×400), present as brown staining. Mice were sacrificed at 2 weeks after treatment. (b) Immunohistochemical analysis of the islets from treated BDC Tg mice for activated T cells. Representative immunostained sections of islets from the pancreas of either STZ- or poly I:C-treated BDC Tg mice. Immunohistochromic staining for either CD4 or CD25 was performed and revealed CD4+ lymphocytes in both mice, but only STZ-treated mice have activated CD4+ cells as demonstrated by CD25 staining. Immunostained cells are brown (×400). The mice were sacrificed at 14 days after treatment. (c) Immunohistological analysis of pancreata from treated BDC Tg and NOD mice for macrophages. Representative immunostained sections of the pancreas from CB4-infected and STZ- and poly I:C-treated BDC Tg and NOD mice reveal differences in the number and activation state of resident macrophages. Antibody to the macrophage activation marker F4/80 was used to identify activated macrophages, which, following CB4 infection and STZ treatment of BDC Tg mice, appear to become activated in the peri-insular inflammation and to mobilize into the islets of Langerhans during insulitis (×400). Brown antimacrophage staining is present over macrophages. Mice were sacrificed 2 weeks after infection or treatment.
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**Figure 3**
Analysis of the activation state of lymphocytes from BDC2.5 mice following treatment with either poly I:C (a) or STZ (b). Cells were stained with antibody to Vβ4 TCR, CD4, and antibody to either CD25, CD44, or CD69. Cells were analyzed by FACS and compared to isotype controls. Shown are histograms of mean fluorescent intensities generated by gating on 10,000–20,000 CD4+ Vβ4+ cells, or Vβ4–, CD4+ cells. (a). For both a and b, the plots are representative of one of six mice analyzed, and all six had similar results. (a) Analysis of peripheral blood lymphocytes before (narrow) and three days following poly I:C treatment (wide) from identical mouse are displayed. No shift of activation is observed in the Vβ4+, CD4+ T cell population while activation is observed in the Vβ4+, CD4– (CD8+) T cell population posttreatment (PT). Changes in cell populations and gated markers CD4+, Vβ4+: CD25, 5.7% to 7.2%; CD44-high, 28.6% to 25.2%; CD69, 6.7% to 6.9% and CD4–, Vβ4+; CD25, 2.9% to 27.8%; CD 44-high, 22.9% to 65.0%; CD69, 10.7% to 41.8%. (b) Analysis of lymphocytes from the spleen and pancreatic lymph node before (narrow) and seven days following STZ treatment (wide) are displayed. Mice were analyzed at three, five, and seven days PT. Similar results were observed at five days, but not at three days. A dramatic shift in all three activation markers was observed in the spleen PT. A less dramatic shift was observed in the pancreatic lymph node. Changes in cell populations and gated markers in spleen CD4+ Vβ4+: CD25, 10.4% to 32.9%; CD44-high, 19.8% to 56.5%; CD69, 9.0% to 32.3%; and in pancreatic lymph nodes, CD4+ Vβ4+: CD25, 5.4% to 25.3%; CD44-high, 7.2% to 18.9%; CD69, 8.7% to 27.0%.
TCR could activate BDC T cells, the results convincingly demonstrated that this could not occur in vivo to induce diabetes (32). One experimental drawback of this system is the lack of knowledge of the true BDC-specific antigen; however, recently, antigenic peptides have been identified that stimulate the BDC2.5 T (20). Although the nature of the natural islet autoantigen is not known, it is clear that an active response from F4/80-positive cells is required within the islet itself before disease develops. Activated macrophages or dendritic cells are most likely picking up released antigen and presenting it to resting memory Tg T cells. Conversely, STZ-treated NOD mice do not have the same level of activated F4/80 staining cells and likewise exhibit less insulitis. Further, while APCs isolated from STZ-treated NOD or NODSCID mice successfully activated the BDC T cell and induced diabetes (Figure 4), they did not induce disease in NOD mice (data not shown). Therefore, mere stimulation of macrophages by STZ or β cell damage does not necessarily activate T cells to induce disease. A critical level of autoreactive precursor resting memory T cells is most likely required (33, 34). Clearly, once initiated, the responses of both T cells and APCs amplify each other and heighten pathologic effects. The requirements of costimulation for this initiation are probably reduced, since the memory state of the T cells lessens the priming requirements of these cells.

NOD mice treated with STZ did not exhibit an increase in diabetes development despite the release of islet antigen and the prevalence of autoreactive lymphocytes. The major difference between the BDC mice, in which diabetes is STZ-inducible, and the NOD mice is the frequency of autoreactive precursor resting memory T cells. The transgenic population consists primarily of CD4+, CD44-high, CD62L-low islet-specific lymphocytes with few requirements for activation. In the NOD mouse, these autoreactive lymphocytes are present at a much lower frequency. Despite the predominance of a single specific T cell population, the BDC Tg mouse had enough flexibility to combat viral infections and counter-regulate its large number of autoreactive cells. This abnormal T cell repertoire of the BDC mouse is obviously an exaggerated version of a human preclinical situation.

In patients, genetic predisposition and environmental exposures could lead to a similar expansion of autoreactive T cells, leading to peri-insulitis. This repertoire could undergo further distortion in response to viral infection, as has been reported by others (35–37). Ultimately, release of sequestered antigen by viral infection or environmental insult would be the triggering event in disease initiation, as observed in the STZ-treated or CB4-infected BDC mouse. It was previously proposed that, in a situation with a distorted lymphocyte repertoire, infection could shift the balance and result in a loss of regulatory T cells, allowing disease to occur. Similarly, viral interference of the host immune regulatory process has been observed in mice with normal T cell repertoires, and the result was an increased survival rate of nonviral-specific T cells (11, 38, 39). However, a single low dose of STZ (acting in place of CB4) did not directly activate T cells or interfere with the turnover or regulation of autoreactive cells. STZ can

Table 2
Transfer of autoimmune diabetes with APCs from mice with damaged β cells

<table>
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<tr>
<th>Donor cells</th>
<th>Recipient mice</th>
<th>Weeks post-transfer (% diabetic)</th>
<th>1</th>
<th>2</th>
<th>3</th>
<th>4</th>
<th>5</th>
<th>6</th>
</tr>
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<tbody>
<tr>
<td>APCs from untreated NODSCID mouse</td>
<td>BDC2.5 TCR Tg (n = 9)</td>
<td>0%</td>
<td>0%</td>
<td>0%</td>
<td>0%</td>
<td>0%</td>
<td>0%</td>
<td>0%</td>
</tr>
<tr>
<td>APCs from STZ-treated NODSCID mice</td>
<td>BDC2.5 TCR Tg (n = 9)</td>
<td>11%</td>
<td>33%</td>
<td>33%</td>
<td>33%</td>
<td>33%</td>
<td>33%</td>
<td>33%</td>
</tr>
<tr>
<td>BDC2.5 TCR Tg splenocytes</td>
<td>Untreated NODSCID mice (n = 9)</td>
<td>0%</td>
<td>0%</td>
<td>0%</td>
<td>44%</td>
<td>44%</td>
<td>55%</td>
<td>55%</td>
</tr>
<tr>
<td>BDC2.5 TCR Tg splenocytes</td>
<td>STZ-treated NODSCID mice (7 days PT, n = 10)</td>
<td>60%</td>
<td>70%</td>
<td>70%</td>
<td>80%</td>
<td>90%</td>
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Autoimmune diabetes was induced following either adoptive transfer of APCs from mice with damaged β cells or transfer of autoreactive T cells into immune-deficient mice with damaged β cells. Adherent mononuclear cells from NODSCID mice treated with STZ or not treated were adoptively transferred into BDC2.5 Tg mice at a concentration of 5 × 10^6 to 10 × 10^6 cells per mouse. Splenocytes from BDC2.5 TCR Tg mice were adoptively transferred into NODSCID mice that had been STZ-treated or not 7 days previously at a concentration of 2 × 10^6 to 5 × 10^6 cells per mouse. Mice with consecutive BG measurements of greater than 300 mg/dl were considered diabetic. *Age-matched. PT, post-transfer.
fully replace the virus in IDDM development; thus the role of CB4 is clearly to damage β cells and not to act as an immune modulator.

The inability of poly I:C to upregulate the BDC Tg T cell and initiate disease argues against a role for bystander T cell amplification in the development of virus-mediated autoimmune disease. Although viruses have the ability to upregulate nonspecific responses and to enhance disease, this bystander amplification is subject to limitations in the scope of its involvement. Results from other investigators (40, 41) imply that an additional factor is required to reactivate lymphocytes and drive autoimmunity. Merely activating potential autoimmune T cells is not enough to cause disease. Yet, in some instances, viral cross-activation has increased the incidence of disease (42, 43). Cross-activation during infection is predominately associated with CD8+ T cells through the type I IFN pathway and involves IL-15 and the use of the IL-2β receptor that is specifically found on CD8+, but not CD4+, T cells. Conversely, CD4+ T cells are not similarly reactivated. This activation of CD8+ T cells by IFN was induced by IL-15 and appeared to utilize the IL-2 receptor β that is found on CD8+ T cells but not CD4+ T cells (44). Although CD8+ T cells participate in nonspecific cross-activation, systems involving CD4+ lymphocytes, as in coxsackievirus-induced IDDM of BDC2.5 Tg mice (6), herpes simplex virus 1-induced herpetic stromal keratitis (45), and pathogen-induced experimental autoimmune encephalomyelitis (46), show little cross-activation. In systems dependent on CD8+ lymphocytes, such as the LCVM transgenic models (43, 47), systemic cross-activation may be the only requirement for activating memory CD8+ T cells, inducing pathogenic T cells and producing disease. Additionally, in general, CD8+ T cell effectors have fewer counter-regulatory mechanisms than do CD4+ T cells. In the case of human autoimmune disease, the actions of CD8+ T cells may, in turn, be more important during relapses and be critical to the development of both IDDM and multiple sclerosis. Certainly CD4+ and CD8+ T cells are both important participants in human autoimmune disease. Their reactivation, apparently via separate mechanisms, is clearly mutually dependent. Differences in the inducibility of diabetes between the BDC2.5 Tg mouse and its NOD littermate further elucidate not only the crucial importance of breakdown and release of sequestered antigen, but also the requirement for a sizable preexisting memory population of autoreactive lymphocytes in the development of overt clinical disease.

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