

# Enterovirus infection of human islets of Langerhans affects $\beta$ -cell function resulting in disintegrated islets, decreased glucose stimulated insulin secretion and loss of Golgi structure

M Hodik,<sup>1</sup> O Skog,<sup>1</sup> A Lukinius,<sup>1</sup> J M Isaza-Correa,<sup>2</sup> J Kuipers,<sup>2</sup> B N G Giepmans,<sup>2</sup> G Frisk<sup>1</sup>

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MH and OS contributed equally to the work.

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<sup>1</sup>Department of Immunology, Genetics and Pathology, The Rudbeck Laboratory, Uppsala University, Uppsala, Sweden  
<sup>2</sup>Department of Cell Biology, University Medical Center Groningen, University Groningen, Groningen, The Netherlands

**Correspondence to**  
Dr G Frisk;  
gun.ergida.frisk@gmail.com

## ABSTRACT

**Aims/hypothesis:** In type 1 diabetes (T1D), most insulin-producing  $\beta$  cells are destroyed, but the trigger is unknown. One of the possible triggers is a virus infection and the aim of this study was to test if enterovirus infection affects glucose stimulated insulin secretion and the effect of virus replication on cellular macromolecules and organelles involved in insulin secretion.

**Methods:** Isolated human islets were infected with different strains of coxsackievirus B (CVB) virus and the glucose-stimulated insulin release (GSIS) was measured in a dynamic perfusion system. Classical morphological electron microscopy, large-scale electron microscopy, so-called nanotomography, and immunohistochemistry were used to study to what extent virus-infected  $\beta$  cells contained insulin, and real-time PCR was used to analyze virus induced changes of islet specific genes.

**Results:** In islets infected with CVB, GSIS was reduced in correlation with the degree of virus-induced islet disintegration. The expression of the gene encoding insulin was decreased in infected islets, whereas the expression of glucagon was not affected. Also, in islets that were somewhat disintegrated, there were uninfected  $\beta$  cells. Ultrastructural analysis revealed that virus particles and virus replication complexes were only present in  $\beta$  cells. There was a significant number of insulin granules remaining in the virus-infected  $\beta$  cells, despite decreased expression of insulin mRNA. In addition, no typical Golgi apparatus was detected in these cells. Exposure of islets to synthetic dsRNA potentiated glucose-stimulated insulin secretion.

**Conclusions/interpretation:** Glucose-stimulated insulin secretion; organelles involved in insulin secretion and gene expression were all affected by CVB replication in  $\beta$  cells.

## INTRODUCTION

Type 1 diabetes (T1D) is generally regarded to be caused by a specific autoimmune

destruction of insulin-secreting  $\beta$ -cells. However, in addition to  $\beta$ -cell destruction and reduced  $\beta$ -cell mass,  $\beta$ -cell dysfunction has emerged as a possible contributor to the loss of insulin secretion, and subsequent  $\beta$ -cell death, which may enhance the immune response. Islets isolated from recent onset T1D organ donors have been found to have a decreased glucose-stimulated insulin secretion (GSIS), despite the presence of a large number of insulin-containing islets detected by immunohistochemistry (IHC).<sup>1–3</sup> Islets from T1D donors can sometimes regain their function after a time of culture *ex vivo*<sup>1</sup> and insulin-containing  $\beta$ -cells can be detected several decades postdiagnosis of T1D even in patients without detectable levels of C peptide.<sup>3–5</sup>

Development of T1D is to a large extent dependent on yet unknown factor(s). Human enterovirus (HEV) has for several decades been implicated in the pathogenesis of T1D,<sup>6</sup> both indirectly by frequent detection of immune responses against HEV in serum from patients with T1D,<sup>7–9</sup> and directly by demonstration of reactivity to EV-P1 by IHC<sup>10–11</sup> or viral genome<sup>12–13</sup> in islets of T1D close to onset. A recent meta-analysis of studies detecting HEV with molecular methods showed a significant association between HEV infection and T1D.<sup>14</sup> However, the causative relationship is still elusive and the mechanisms are unknown. Viruses with a lytic replication cycle and ability to replicate in  $\beta$  cells impair the oscillations of cytoplasmic  $Ca^{2+}$  induced by glucose,<sup>15</sup> and reduce the glucose-stimulated release of insulin from the  $\beta$  cells.<sup>16</sup>

Enterovirus replication results in the formation of a dsRNA intermediate in viral replication complexes formed in the cytoplasm by the virus.<sup>17–19</sup> A model for enteroviral

persistence has shown that the viral RNA can persist as dsRNA.<sup>20–21</sup> Synthetic dsRNA, poly(I:C), has been shown to affect  $\beta$ -cell function in rats<sup>22</sup> and mice<sup>23</sup> when the islets were exposed to interferon- $\gamma$ , and to induce apoptosis in primary rat  $\beta$  cells.<sup>24–26</sup> The sensing of dsRNA in the cytoplasm results in an innate immune response in human islets.<sup>27</sup> However, in contrast to poly(I:C), a virus infection leads to production of viral proteins that modulate the response of the infected cells to promote its own replication, resulting in a block of many of the pathways induced by poly(I:C).

The molecular mechanisms, by which host factors are involved in RNA-protein/protein-protein interaction, have been intensively studied; however, tissue-specific viral virulence remains unclear. Therefore, our aim was to study cellular function and ultrastructural changes in primary human cells in the islets of Langerhans.

## MATERIALS AND METHODS

### Isolation and culture of human islets

Human pancreases were obtained from 17 organ donors (6 females and 11 males) free of any known pancreatic disease, from transplantation units in Sweden, Norway, and Finland. All samples were analyzed anonymously. The donors used for analyses of islet function (n=10) had an average age of 53±17 years (range 20–73) and an average body mass index (BMI) of 25.4±1.9 kg/m<sup>2</sup> (range 22.2–27.8). The donors of islets used for gene expression analyses (n=6) had an average age of 59±19 (range 22–74) and an average BMI of 25.8±2.0 kg/m<sup>2</sup> (range 23.3–28.9). The donors used for IHC analysis (n=1) and electron microscopic analysis (n=1) had an age of 55 years and a BMI of 22.5 kg/m<sup>2</sup>. Islets of Langerhans were isolated in Uppsala, Sweden, using a protocol approved by the Local Ethics Committee, as described previously.<sup>28</sup> Only islet preparations with an intact GISIS were included in the study and the islets were available for research only because the total islet volume was too low for clinical transplantation. The islets were kept in culture bags (Baxter Medical AB, Sweden) with 200 mL CMRL-1066 (ICN Biomedicals, Costa Mesa, California, USA) supplemented with 10 mM HEPES, 2 mM L-glutamin, 50  $\mu$ g/mL Gentamycin, 0.25  $\mu$ g/mL Fungizone (GIBCO BRL), 20  $\mu$ g/mL Ciprofloxacin (Bayer healthcare AG, Leverkusen, Germany), 10 mM nicotinamide, and 10% heat-inactivated human serum at 37°C in 5% carbondioxide and humidified air for 1–5 days. The culture medium was changed on day 1 and then every other day until the islet experiments were performed.

### Virus

Two strains of CVB4 (VD2921 and E2-Yoon) and two strains of CVB5 (Adr and V89-4557) were used. VD2921 (GenBank accession number AF328683) and V89-4557 were originally isolated from cerebrospinal fluid from patients with aseptic meningitis,<sup>16–29–31</sup> whereas Adr

(GenBank accession number FJ374273) and E2-Yoon (GenBank accession number S76772) originate from patients at onset of type 1 diabetes and were isolated from stool and pancreas, respectively.<sup>6–32–33</sup>

### Islet infection and observation of islet disintegration

The isolated human islets had an estimated purity ranging from 40% to 95% (mean 69%) and were purified further by hand-picking under a light microscope by islet experts. Forty hand-picked and size-matched islets per well were cultured in 2 mL RPMI-1640 containing 5.5 mM glucose (SVA, Uppsala, Sweden) supplemented with 5% fetal bovine serum and 2 mM L-glutamine, in 6-well plates for suspension cells (Sarstedt AB, Helsingborg, Sweden). Hand-picked islets were cultured for 24 hour before inoculation with CVB (CVB5/Adr, CVB5/V89-4557, CVB4/E2-Yoon, or CVB4/VD2921) to a final concentration of 100–1000 TCID<sub>50</sub>. Forty islets from each donor were exposed to 50  $\mu$ g/mL synthetic dsRNA (Poly(I:C), Sigma-Aldrich Sweden AB, Stockholm, Sweden). Samples of culture medium were collected from all wells, directly after inoculation and every day until days 4–7 postinfection, and stored at –20°C for subsequent TCID<sub>50</sub> titrations. Inoculated and uninoculated islets from the same donor were observed every day in a light microscope. The appearance of virus-induced islet disintegration was graded from 1+ to 4+ as previously described,<sup>29</sup> until the experiments were terminated on days 4–7 postinfection. Islets exposed to poly(I:C) and unexposed control islets were cultured for 24 hour before termination of the experiment. On termination, islets were washed twice with phosphate-buffered saline and challenged with high glucose in a dynamic perfusion assay or stored in RNA later (Qiagen AB, Sollentuna, Sweden) for subsequent analyses.

### Islet function test

GISIS of CVB-infected, poly(I:C)-exposed, and control islets was assessed in a dynamic perfusion system, Suprafusion 1000 (BRANDEL, Gaithersburg, Maryland, USA). Islets were perfused with two glucose concentrations (1.67 and 20 mM and then 1.67 mM). Fractions were collected at 6 min interval for 120 min and insulin concentrations were determined by ELISA (Mercodia, Uppsala, Sweden). Insulin secretion at each time point was normalized to control islets from the same donor, and is expressed as percent secretion compared to the first phase peak of the respective untreated control.

### Analysis of virus replication

Virus replication was determined by tissue culture infectious dose<sub>50</sub> (TCID<sub>50</sub>) titrations of samples of culture medium on an in-house cell line originating from Green Monkey Kidney cells, obtained from Gothenburg University, Sweden, as described previously.<sup>34</sup> The samples were collected from the infected islet cultures on the indicated days pi.

### RNA extraction and real-time RT-PCR

Total RNA from explanted islets stored in RNA later, was extracted with RNeasy Plus Mini kit (Qiagen AB, Sollentuna, Sweden). RNA quantity and quality were determined using spectrophotometry and gel electrophoresis. Up to 50 ng total RNA per sample were primed with Random Primers (Invitrogen) and reverse transcribed to cDNA with SuperScript II RT (Invitrogen) according to the manufacturer's instructions. Control reactions containing all reagents except the reverse transcriptase were included to confirm the absence of interfering genomic DNA. RNaseOUT (Invitrogen) was added to the reaction to avoid RNA degradation. Real-Time PCRs were run with Power SYBR Green master mix (Applied Biosystems, Sweden) on a StepOnePlus Real-Time PCR system (Applied biosystems, Sweden). Predesigned gene-specific primer sets (QuantiTect Primer Assays, Qiagen) were used for detection of insulin, glucagon, glut-1 (*SLC2A1*), glut-2 (*SLC2A2*), mafA, mafB, and PDX1 cDNA. The expression level of each gene was normalized against expression of the reference gene 18S ( $2^{-\Delta Ct}$ ), and is presented as fold induction compared to untreated islets from the same donor ( $2^{-\Delta\Delta Ct}$ ). PCR specificity was verified by melt curve analysis of all PCR products.

### Immunohistochemistry

Islets from a representative donor infected with CVB5/V89-4557 or CVB4/E2-Yoon resulting in moderate islet disintegration (graded 2+ to 3+) and mock-infected control islets were fixed in 4% PFA and paraffin-embedded day six postinoculation. Sections (5  $\mu$ m) were dried on SuperFrost slides (Menzel Gläser, Braunschweig, Germany), followed by deparaffination and rehydration in xylene and decreasing concentrations of ethanol, respectively. Endogenous peroxidase was blocked for 10 min followed by incubation of primary antibodies (guinea-pig anti-insulin 1:200) in a humidified chamber at 37°C. Washing steps were performed in TBS with 0.05% Tween 20. Visualization was achieved by Envision using DAB as a substrate chromogen. All reagents and antibodies were purchased at DAKO, Glostrup, Denmark.

### Electron microscopy

Isolated pancreatic islets from one donor were either inoculated with CVB5/Adr to a final concentration of 5.5 TCID<sub>50</sub> or left uninoculated. On day three postinfection, all islets were fixed in 2% glutaraldehyde in 0.1M cacodylate buffer supplemented with 0.1M sucrose, followed by 1.5 hour postfixation in 1% OsO<sub>4</sub>, dehydration, and embedding in epoxy plastic Agar 100 (Agar Aids, Stansted, England). Ultrathin sections (50 nm) were placed on Formvar-coated Cu-grids and contrasted with uranyl acetate and lead citrate. The ultrastructural morphology was analyzed in an electron microscope (Tecnai G<sup>2</sup> Spirit BioTWIN). Subsequently, representative islets were analyzed by large-scale EM<sup>35</sup> on a Zeiss Supra55 (Germany) with an external large-scan generator (ATLAS,

Fibics, Canada) at 29kv; 2nm pixel-size-using the STEM detector.

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Quantification of the insulin granule-, lipofuscin- and mitochondrial cytoplasmic area (%) was made on the outer layer of cells on cross-sections of both CVB-5/Adr infected and uninoculated control islets. Lipofuscins were defined as secondary lysosomes with electron-lucent (ie, lipid-like material) vesicles<sup>36</sup> and insulin granules after the irregular crystalline electron-dense core in an electron-transparent halo.<sup>37</sup> The organelles were manually color-coded in Photoshop CS4 and analyzed with Fuji32/ImageJ using the color deconvolution tool. Lipofuscin, insulin, and mitochondrial analyses were performed on virus-particle-containing  $\beta$  cells (n=11),  $\beta$  cells with no signs of infection (n=7), and non-inoculated  $\beta$  cells in the periphery of the islet (n=11). The density of insulin granules was analyzed in the same cells.

### Statistical analysis

Differences in insulin secretion and gene expression levels were analyzed with Wilcoxon matched-pairs signed rank test. Ultrastructural quantitative data were analyzed with Mann-Whitney two-tailed tests. p Values <0.05 were considered statistically significant. Data values are presented as mean $\pm$ SEM.

### Ethics statement

The work performed on human tissue was in accordance with the principles expressed in the Declaration of Helsinki and the European Council's Convention on Human Rights and Biomedicine. The study was approved by the Regional Ethics Committee in Uppsala, Sweden according to the Act concerning the Ethical Review of Research Involving Human (2003:460), Permit Number: Dnr 2009/043, 2009/371 and Ups 02-577. The pancreas was obtained from the Nordic Network of Islets transplantation.

## RESULTS

### Virus-induced islet disintegration decreases GSIS in human islets

The level of replication and degree of islet disintegration varied between the different virus strains and islet donors (table 1). In six cases (n=6), the virus had replicated (mean titer increase 3.1 logTCID<sub>50</sub>/200  $\mu$ L) without causing any observable islet disintegration (table 1). Despite viral replication, GSIS in these islets was not significantly different compared to that of the non-inoculated control islets (figure 1A). When a low but clear (graded 1+ to 2+) virus-induced islet disintegration was observed along with a mean titer increase of 3.7 logTCID<sub>50</sub> (n=6; table 1), the rapid first as well as sustained second phase of GSIS was slightly decreased compared to non-inoculated islets (p<0.05). The basal insulin release at low (1.67 mM) glucose was not altered

**Table 1** Virus replication and virus-induced islet disintegration in different donors

Donor number	Strain of CVB (day of termination)	Islet disintegration* (graded from 0 to 4)	Titer change† from day 0
1	Adr (6 dpi)	0	+4
	V89-4557 (6 dpi)	3–4	+4
2	VD2921 (4 dpi)	0	+3
	V89-4557 (4 dpi)	0	+2.5
	V89-4557 (7 dpi)	3	+4.5
3	Adr (7 dpi)	0	+3.5
	VD2921 (7 dpi)	0	+3.5
4	E2-Yoon (7 dpi)	3	+3
	VD2921 (7 dpi)	2	+6
5	E2-Yoon (4 dpi)	2	+1.5
	E2-Yoon (7 dpi)	3–4	+2
	V89-4557 (7 dpi)	2	+2.5
6	E2-Yoon (6 dpi)	3–4	+2.5
7	E2-Yoon (6 dpi)	0	+2
8	E2-Yoon (6 dpi)	3	+2.5
9	VD2921 (6 dpi)	2	+4.5
	V89-4557 (6 dpi)	3–4	+4
	E2-Yoon (6 dpi)	3–4	+3
	VD2921 (6 dpi)	2	+5
10	V89-4557 (6 dpi)	3–4	+6
	E2-Yoon (6 dpi)	2	+2.5

\*Virus-induced islet disintegration evaluated under a light microscope.

†Determined by TCID<sub>50</sub> titration and presented as logTCID<sub>50</sub>/200 µL.

CVB, coxsackievirus B; Dpi, days postinfection; TCID, tissue culture infectious dose<sub>50</sub>.

(figure 1B). Pronouncedly disintegrated islets (graded 3–4) and mean titer increase 3.6 logTCID<sub>50</sub> (table 1) showed poor or none GSIS (n=9, p<0.05) (figure 1C).

There were no difference in GSIS between the virus strains, but that might be due to the limited number of analyses, due to the limited number of islets available.

### Synthetic dsRNA potentiates GSIS in human islets

Islets exposed to poly(I:C) showed a potentiated response to high glucose (figure 2A; p<0.05); the rapid first phase of insulin release was increased 2.5-fold compared to that of untreated control islets, and the sustained second phase was about doubled. At low (1.6 mM) glucose, the insulin release was similar in islets exposed to poly(I:C) and untreated control islets.

A dose–response relationship could be observed when GSIS was tested with dynamic glucose perfusion 24 hour after poly(I:C) exposure, where 10 µg/mL had no effect, 50 µg/mL had moderate effect and 250 µg/mL had the strongest effect (figure 2B; n=1). Four days after exposure to 250 or 50 µg/mL poly(I:C), the islets were still strongly potentiated to release insulin in response to high glucose compared to control islets, but no difference could be seen between these two concentrations (figure 2C; n=1).

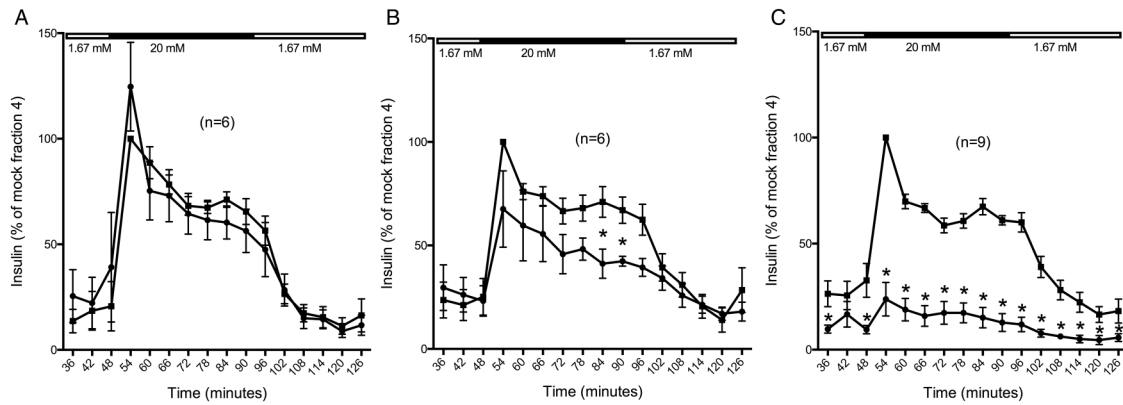
### Virus-induced islet disintegration correlates specifically to reduced transcription of islet genes

Six days postinfection, islets infected with CVB4/VD2921 showed moderate islet disintegration (graded 2+), which

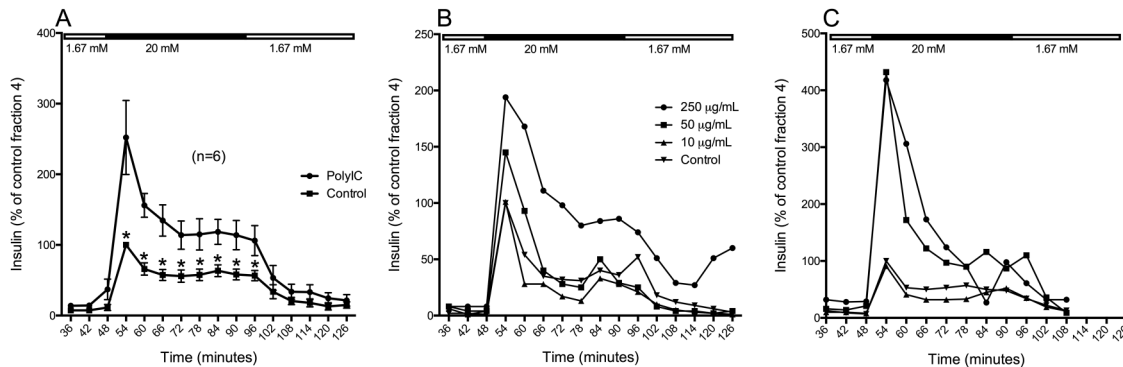
was more pronounced (graded 3+ to 4+) in islets infected with CVB4/E2-Yoon or CVB5/V89-4557. The relative islet expression of insulin mRNA was reduced in all islets with virus-induced disintegration (figure 3A), whereas the relative expression of glucagon mRNA was unaffected by infection with CVB4/E2-Yoon or CVB4/VD2921 (figure 3B). The third virus strain, CVB5/V89-4557, reduced the relative islet expression of both insulin and glucagon mRNA (figure 3A, B). Expression of the two transcription factors *pdx1* and *mafa* was also reduced to a similar extent by infection with the three viruses, whereas *mafB* (mainly expressed in  $\alpha$  cells<sup>38</sup>) was only reduced by CVB5/V89-4557 infection (figure 3C–E). The expression of mRNA encoding the glucose transporter protein *glut2* was downregulated by the three viruses to a similar extent as the expression of *ins*, *mafa*, and *pdx1* (figure 3G). The expression of the gene encoding *glut1* was not affected by virus infection, except in islets infected with CVB5/V89-4557 in which it was upregulated significantly (figure 3F). Exposure of islets to poly(I:C) did not affect the expression of any of these genes (figure 3A–G).

### Insulin is detected immunohistochemically and ultrastructurally in disintegrated infected islets with decreased or lost GSIS

Insulin was detected by IHC in CVB-infected islets with a degree of virus-induced disintegration (graded 2+ to 3+) and unresponsiveness to 20 mM glucose (figure 4A). Ultrastructural analysis confirmed the presence of



**Figure 1** Glucose-stimulated insulin secretion in virus-infected human islets. Forty hand-picked and size-matched human islets from each donor were inoculated with serotypes/strains of CVB (black circles) or uninoculated (black boxes; medium only). On days four to seven postinoculation, the islets were perfused with low glucose (1.67 mM) for 42 min, high glucose (20 mM) for 48 min, and then low glucose again. Fractions were collected at 6 min intervals and the secreted insulin was measured by ELISA. Islets were grouped based on the virus-induced islet disintegration in three groups: no visible disintegration (A), moderate disintegration (B), and pronounced disintegration (C). \*  $p < 0.05$ . CVB, coxsackievirus.



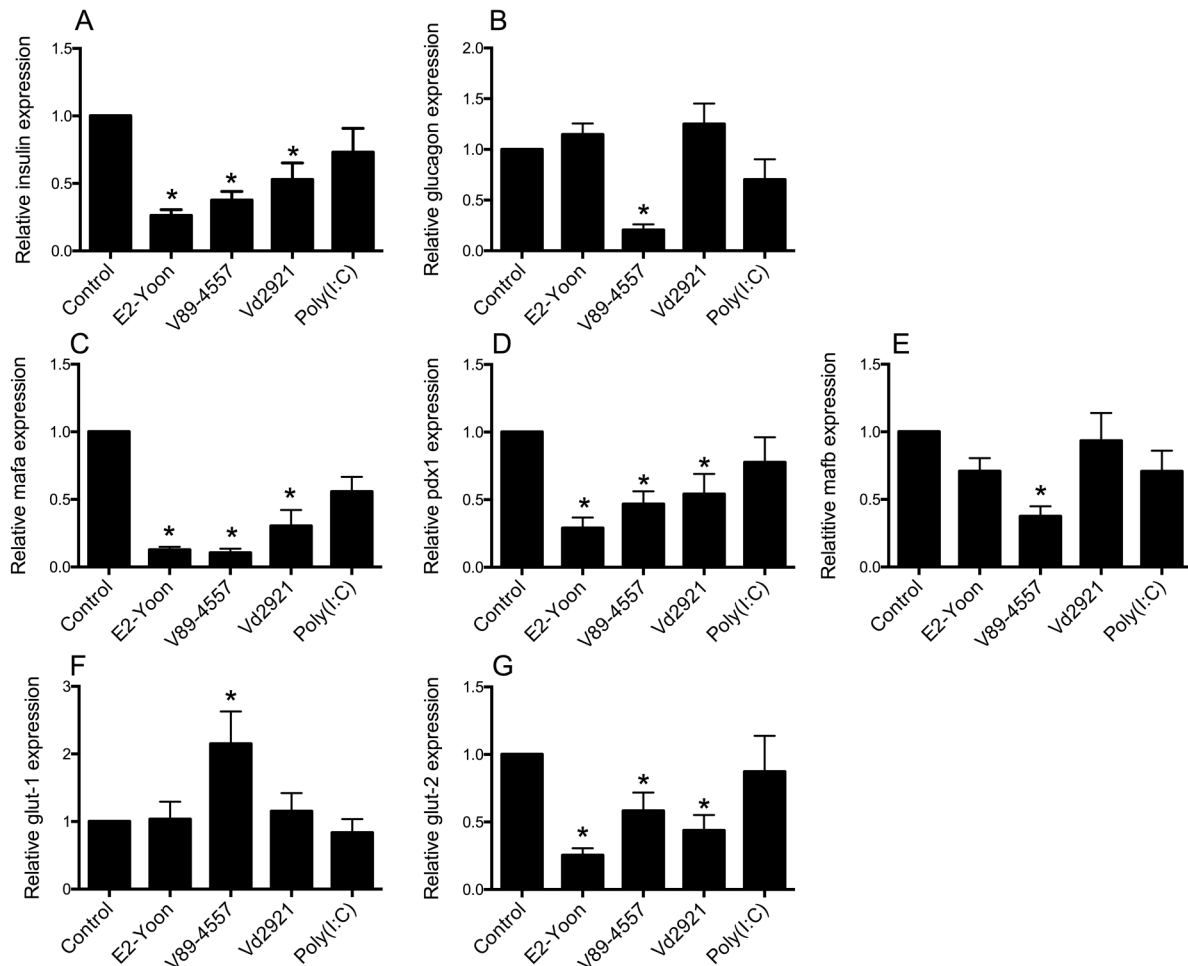
**Figure 2** Glucose-stimulated insulin secretion in human islets exposed to poly(I:C). Forty hand-picked and size-matched human islets were exposed to poly(I:C) or left untreated. 24 hour (A, B) or 96 hour (C) postexposure, the islets were perfused with low glucose (1.67 mM) for 42 min, high glucose (20 mM) for 48 min, and then low glucose again. Fractions were collected at 6 min intervals and the secreted insulin was measured by ELISA. Mean insulin secretion in islets exposed to 50 µg/mL poly(I:C) was measured after 24 hour (A) (n=6). Insulin secretion in islets exposed to 250, 50, or 10 µg/mL poly(I:C) was measured after 24 hour (B) or 96 hour (C) (n=1). \*  $p < 0.05$ .

insulin granules in CVB5/Adr-infected  $\beta$  cells in various necrotic stages (figure 4B).

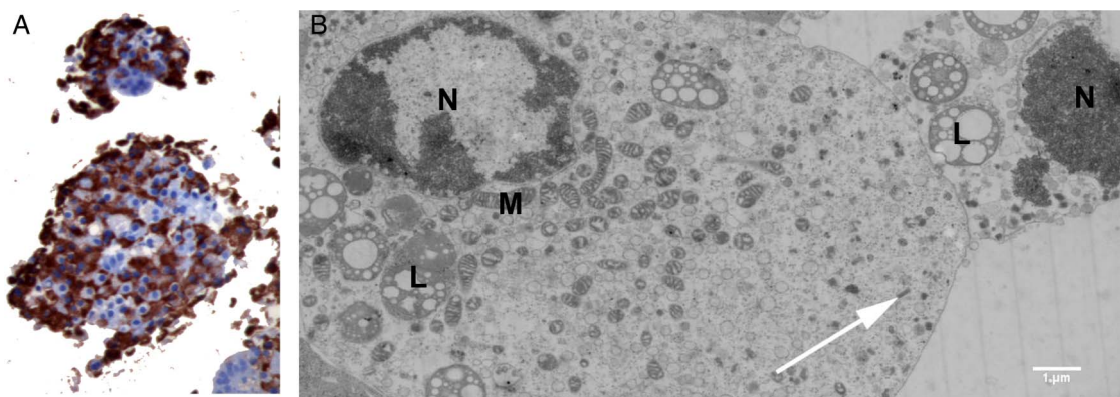
### Virus-induced ultrastructural changes can be seen in $\beta$ cells in moderately disintegrated islets

Islets from five donors were inoculated and analyzed for the presence of virus particles, viral replication complexes, and virus-induced ultrastructural changes. Two representative islets from one donor were selected for large-scale electron microscopy (EM) to evaluate if this could be used for quantitation. Both infected and uninoculated islets lacked basal lamina, due to the isolation process, although the plasma membranes and the different cell to cell contacts were normal. Infected islets had an irregular outer form, in line with the findings in the light microscopy studies, due to detachment of dead cells. Virus particles were found in  $\beta$  cells in the

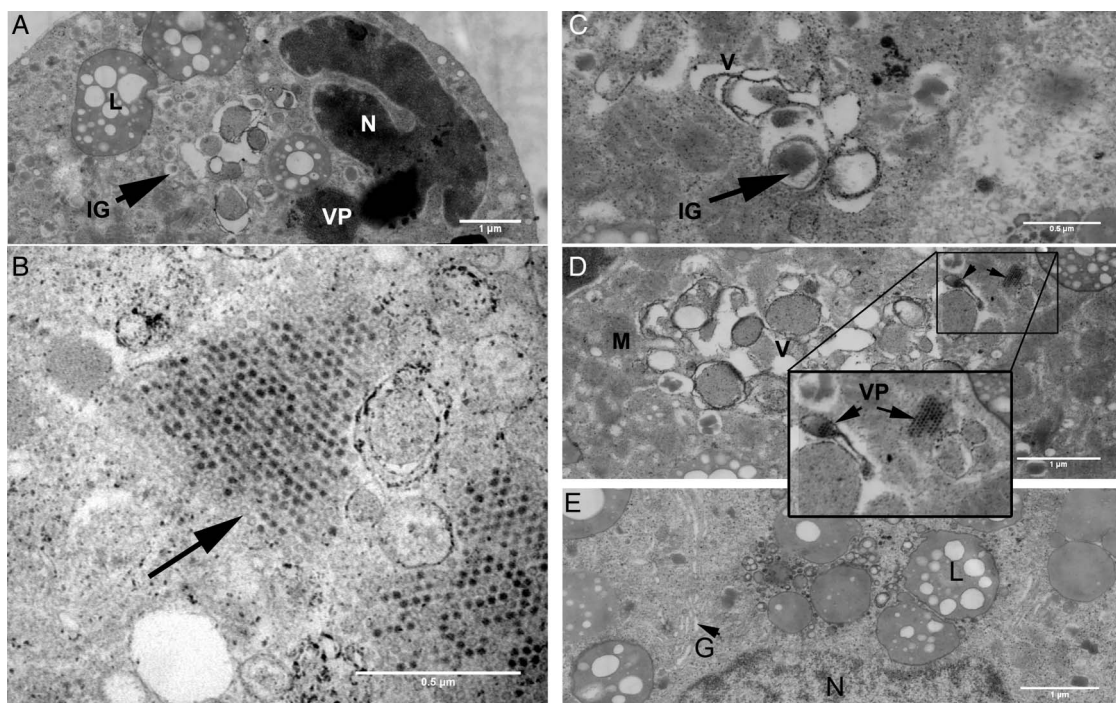
outermost cell layers of the islet, assembled in one or several crystal rafts scattered in the cytoplasm (figure 5A). The finding of virus particles only in  $\beta$  cells of the islets, 3 days postinfection, can be explained by the lack of basal lamina resulting in virus infecting the first layer of cells and the released progeny virus that will bind to and infect the next layer of  $\beta$  cells. Occasionally, empty viral capsids were observed in the viral rafts (figure 5B, black arrow). In infected  $\beta$  cells, nuclei were condensed with invaginations and marginated condensed chromatin. Endoplasmic reticulum (ER) was dilated with dropped off ribosomes and often broken up into small vesicles and vacuoles. Golgi apparatuses were also decomposed into vesicles and vacuoles of various electron densities (figure 5C, D). In virus-infected  $\beta$  cells (n=11), no recognizable Golgi apparatus was observed compared to apparently non-infected  $\beta$  cells (n=7) in



**Figure 3** Islet mRNA expression of insulin (A), glucagon (B), mafa (C), pdx1 (D), mafb (E), glut-1 (F), and glut-2 (G). Forty hand-picked islets were exposed to 50  $\mu$ g/mL poly(I:C), inoculated with CVB4/E2-Yoon, CVB5/V89-4557, or CVB4/VD2921 or left untreated. Six days postinfection, or 1 day postexposure to poly(I:C), the intracellular mRNA content was measured by qPCR. The gene specific expression relative to expression of the reference gene 18S was normalized to the expression in mock-infected control islets from the same donor ( $2^{-\Delta\Delta Ct}$ ). Data are expressed as mean  $\pm$  SEM from experiments on six different islet donors (n=6). \*p<0.05. CVB, coxsackievirus.



**Figure 4** Representative images of human islets infected with strains of CVB. (A) Immunohistochemical staining of insulin in islets infection with CVB5/V89-4557 6 days postinfection. Original magnifications  $\times 20$ . (B) Electron micrograph of  $\beta$ -cells in different stages of necrosis in CVB5/Adr inoculated islet, showing presence of insulin (white arrows). CVB, coxsackievirus; M, mitochondria, N, nuclei, L, lipofuscin.



**Figure 5** Ultrastructural analysis of islet infected with a strain of CVB. (A) Infected  $\beta$ -cell with virus particles and a condensed and invaginated nucleus. CVB, coxsackievirus; IG, insulin granule; L, lipofuscin; N, nucleus, VP, virus particles. (B) Empty-virus capsids (arrow) (C and D) Virus-induced vesicular and vacuolar structures of varying complexity. CVB, coxsackievirus; IG, insulin granule; M, mitochondria; V, vesicles and vacuoles. Inset: VP at higher magnification. (E) Recognizable Golgi complexes (G) in  $\beta$ -cell with no signs of virus infection and normal nuclei (N) in an infected islet. L, lipofuscin. (F) Upper row from left: Cytoplasmic insulin area (%), insulin granule density (insulin granules/ $\mu\text{m}^2$ ), and observed insulin granule diameter (nm). Lower row from the left: Mitochondrial area (%) and lipofuscin area (%). Bars correspond to  $\pm$ SEM. In figure 5F, quantification of insulin granules, area of lipofuscin and mitochondria. Number of insulin granules (insulin granules/ $\mu\text{m}^2$ ) and cytoplasmic insulin, lipofuscin and mitochondrial area (%) in CVB infected (n=11), no signs of virus infection (n=7) and uninfected (n=11)  $\beta$ -cells. Bars correspond to  $\pm$ SEM. CVB, coxsackievirus B.

the same islet in which well-developed Golgi apparatuses were observed in  $\sim 71\%$  of cells (figure 5E). In infected  $\beta$  cells, the ultrastructure of mitochondria were apparently normal with well-preserved cristae. In severely decomposed  $\beta$  cells also, the mitochondria started to condense (figure 4B).

The cross-section of a whole non-inoculated islet and a CVB5/Adr-infected islet displayed  $\sim 76\%$  and  $\sim 40\%$   $\beta$  cells, respectively. Virus particles were found in  $\sim 16\%$  of the  $\beta$  cells (n=11) in the CVB5/Adr-infected islet periphery, and signs of virus infection were observed in  $\sim 80\%$  of the  $\beta$  cells. The insulin granule-, lipofuscin- and mitochondrial cell area and insulin granule density quantification comparison between infected islet and uninfected control islet was performed on the outer cell layers. Other endocrine cell types present in the same islet were unaffected by the virus infection even when surrounded by virus-infected  $\beta$  cells. Cytoplasmic insulin granule (IG) area (%) and IG density (IG/ $\mu\text{m}^2$ ) were significantly higher in CVB5/Adr-infected  $\beta$  cells with virus particles ( $10.7 \pm 1.17\%$  and  $3.2 \pm 0.35$  IG/ $\mu\text{m}^2$ ; n=11) compared to peripheral  $\beta$  cells in a non-inoculated islet ( $4.4 \pm 0.63\%$  and  $1.5 \pm 0.22$  IG/ $\mu\text{m}^2$ ; n=11) (figure 5F). The IG area, but not the IG density, was also significantly higher in CVB5/Adr-infected  $\beta$  cells with virus particles

than in  $\beta$  cells without visible virus particles in the same islet ( $6.1 \pm 0.8\%$ ; and  $2.6 \pm 0.36$  IG/ $\mu\text{m}^2$ ; n=7) (figure 5F). The observed mean IG diameter was significantly higher in the islet peripheries both in virus-infected  $\beta$  cells ( $209 \pm 8.7$  nm; n=11) and non-inoculated  $\beta$  cells ( $194 \pm 8.6$ ; n=11) compared to apparently uninfected  $\beta$  cells in the infected islet ( $175 \pm 3.8$  nm; n=7) (figure 5G). There was no statistical difference in cytoplasmic mitochondrial and lipofuscin cytoplasmic area between infected and non-inoculated cells (figure 5H).

## DISCUSSION

Here, we demonstrate that virus-induced disintegration of human islets leads to reduced GSIS and decreased expression of the  $\beta$  cell specific genes encoding insulin, *mafA*, *pdxq*, and *glut2*. These results correlate well with the sign of virus infection only in  $\beta$  cells revealed by ultrastructural analysis. In infected  $\beta$  cells, the Golgi apparatuses are resolved, and nuclei with deep invaginations and condensed chromatin, indicating various stages of virus-induced cell death, can be seen.

The degree and kinetics of virus-induced islet disintegration varied between the different strains of CVB, as well as between islets from different donors (table 1).

Non-disintegrated islets or islets with low-grade disintegration with a high viral titer increase and no effect on GSIS could contain more  $\beta$  cells or the time from infection to cell death might be longer compared to that in highly disintegrated islets. In islets that were only moderately disintegrated (grade 2+ to 3+), it was shown that ~80% of the  $\beta$  cells had signs of infection, though the effect on islets function was less pronounced, suggesting that the remaining and non-infected  $\beta$  cells to some extent compensate for the loss.

The finding that infected  $\beta$  cells contain somewhat larger insulin granules compared to non-infected  $\beta$  cells from the same islet might be due to accumulation of cholesterol to replication sites by viral proteins.<sup>39</sup> Cholesterol accumulation has been shown to increase insulin granule size and also to impair membrane trafficking.<sup>40</sup>

Accumulation of the viral protein 2B or 2BC on the Golgi apparatus changes the permeability of these membranes<sup>41</sup> and causes disassembly of the Golgi apparatus<sup>42</sup> and ultimately lysing cells, halting insulin secretion. In addition, in virus infected cells, the membrane that integrates the 2B/2BC complex also reduces the  $\text{Ca}^{2+}$  levels in the ER,<sup>43–45</sup> this is also related to antiapoptotic properties.<sup>46</sup> The effect of virus infection on cholesterol accumulation and on  $\text{Ca}^{2+}$  concentrations will affect GSIS.

It is also known that these viruses cleave the translation initiation factor eIF4GI, which leads to block of cap-dependent translation;<sup>46</sup> this will result in transcription of IRES containing sequences such as the viral genome and cellular IRES containing genes such as bcl-2 and bip. Recently, it was shown that the major part of the synthesis of insulin is cap-dependent,<sup>47</sup> suggesting that in an enterovirus infected  $\beta$  cells this would eventually lead to hampered insulin secretion.

In support of our findings, it has been shown that an increase in intracellular insulin content has been implicated in HEV-infected islets, suggesting that a block of secretion might precede the block of translation.<sup>16</sup>

Our finding that virus particles only can be detected in  $\beta$  cells might be explained by the fact that HEVs have been shown to translocate key proteins in the secretory pathway, including adenosine diphosphate (ADP) ribosylation factor 1 (ARF1) and its activator, Golgi-specific brefeldin A-resistance guanine nucleotide exchange factor 1 (GBF1) to virus-induced replication membranes. Interestingly, the GBF1 protein, which is crucial for virus replication, is expressed specifically in  $\beta$  cells, and that could be one explanation for the predominant virus replication in these cells.<sup>48</sup>

The mitochondria in severely decomposed cells have condensed but well-discernible cristae, which are signs of an actively respiratory state and this have been observed in apoptosis and autophagy;<sup>49 50</sup> both these cell death pathways have previously been implicated in enterovirus-infected cells.<sup>25 51</sup>

The finding that the expression of insulin but not glucagon mRNA was lower in islets infected with E2-yoon or VD2921 than in uninfected control islets supports

previous studies arguing for a selective tropism of many HEV for  $\beta$  cells.<sup>1 10 12</sup> Ultrastructural analysis supported this notion; virus particles were only seen in  $\beta$  cells. In islets infected with the V89-4557 strain, both the transcription of glucagon and insulin was reduced in contrast to in islets infected with other CVB strains. In contrast to most other CVB serotypes/strains, this virus has previously been shown to infect cells lacking both CAR and DAF,<sup>30</sup> giving it a broader tropism. Perhaps this strain also has tropism for  $\alpha$  cells. Interestingly, the relative expression of *glut1* was increased significantly in islets infected with this virus strain, perhaps reflecting high expression of *glut1* in a non-permissive islet cell-type, but this needs to be investigated further.

The ideal thing would have been to infect islets with all strains and study them with an external large-scan generator (ATLAS, Fibics, Canada); however, this was not possible to do due to practical and logistical reasons. It should be stressed that all strains were studied with EM (some in previous publications), but the purpose of this study was to see if the ultrastructural changes that could be seen in islet cells containing virus particles could be quantitated and that on a whole islet basis to where the virus could be seen, and if similar ultrastructural changes could be seen in islet cells with no visible virus particles. We have shown that the method works and it can now be used to study the effect of virus infections, toxins, etc.

Synthetic dsRNA (poly(I:C)) potentiated GSIS in human islets. This is in contrast to findings in murine islets where synthetic dsRNA alone has no effect on GSIS. poly(I:C) exposure of interferon- $\gamma$ -treated murine islets, induced apoptosis and reduced GSIS.<sup>22–26</sup> The viral dsRNA could not increase GSIS, probably because the virus infection causes reduced  $\text{Ca}^{2+}$  levels, indicating that poly(I:C) is not a good model for viral infections.

In summary, the data presented here show that CVB infection can impair  $\beta$  cell function and  $\beta$  cell specific gene expression. The decreased GSIS in the virus-infected and disintegrated islets is most likely due to a functional loss of virus infected and dispersed cells, whereas non-infected  $\beta$  cells in islets with low-grade disintegration probably still can secrete insulin in response to glucose stimulation. The functional loss of the infected  $\beta$  cells is supported by the loss of the Golgi apparatus and the condensed chromatin. The  $\beta$  cell specific expression of GBF1, an essential protein for CVB replication, might explain why virus particles are only detected in these cells.

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