

# Direct Effect of Cholesterol on Insulin Secretion

## A Novel Mechanism for Pancreatic $\beta$ -Cell Dysfunction

Mingming Hao, W. Steven Head, Subhadra C. Gunawardana, Alyssa H. Hasty, and David W. Piston

**OBJECTIVE**—Type 2 diabetes is often accompanied by abnormal blood lipid and lipoprotein levels, but most studies on the link between hyperlipidemia and diabetes have focused on free fatty acids (FFAs). In this study, we examined the relationship between cholesterol and insulin secretion from pancreatic  $\beta$ -cells that is independent of the effects of FFAs.

**RESEARCH DESIGN AND METHODS**—Several methods were used to modulate cholesterol levels in intact islets and cultured  $\beta$ -cells, including a recently developed mouse model that exhibits elevated cholesterol but normal FFA levels. Acute and metabolic alteration of cholesterol was done using pharmacological reagents.

**RESULTS**—We found a direct link between elevated serum cholesterol and reduced insulin secretion, with normal secretion restored by cholesterol depletion. We further demonstrate that excess cholesterol inhibits secretion by downregulation of metabolism through increased neuronal nitric oxide synthase dimerization.

**CONCLUSIONS**—This direct effect of cholesterol on  $\beta$ -cell metabolism opens a novel set of mechanisms that may contribute to  $\beta$ -cell dysfunction and the onset of diabetes in obese patients. *Diabetes* 56:2328–2338, 2007

**D** diabetes has become a global health problem, affecting >170 million individuals worldwide. The rapid rate of increase in the prevalence of diabetes has profound implications in terms of long-term complications and associated medical costs. Type 2 diabetes accounts for >90% of all cases of diabetes and is characterized by insulin resistance and a defect in insulin secretion from pancreatic  $\beta$ -cells (1). Whereas the relative importance of alterations in insulin sensitivity versus secretion is debatable, it is accepted that hyperglycemia, hence diabetes, does not develop without  $\beta$ -cell dysfunction (2). Alteration of pancreatic  $\beta$ -cell function

leading to an impaired insulin secretory response to glucose is hallmark of the transition from the pre-diabetic to the diabetic state (3). Among the many contributing factors, hyperlipidemia plays a critical role in the pathogenesis of  $\beta$ -cell dysfunction (4). The established link between obesity and diabetes, as well as observations that plasma levels of free fatty acids (FFAs) are elevated in most obese individuals, suggests that FFAs might induce hyperglycemia. Lipotoxicity, which refers to the diabetogenic effect of elevated circulating FFAs or cellular fat content, has been studied extensively (5). In addition to FFAs, plasma cholesterol is often elevated in obese patients; yet, the impact of cholesterol in glucose-stimulated insulin secretion (GSIS) from pancreatic  $\beta$ -cells has not been reported.

Cholesterol can regulate signal transduction through membrane microdomains and gene expression through cholesterol-activated transcription factors (6). For example, intracellular cholesterol regulates glucose metabolism and gene expression in adipocytes (7). GSIS is a complex process involving a cascade of regulatory factors. Misregulation of cholesterol could result in disruption of any one pathway and lead to partial or a near complete loss of secretory function. This study focuses on the function of cholesterol-rich membrane microdomains in GSIS.

The plasma membranes of mammalian cells contain 30–50% molar fraction of cholesterol (8). Cholesterol is also enriched in internal membranes (9). Cholesterol plays an essential role in membrane organization, dynamics, and function. It is thought to contribute to the tight packing of lipids by filling interstitial spaces between lipid molecules. Membrane microdomains are small membrane assemblies enriched in cholesterol and sphingolipids and coexist with more fluid domains enriched in phospholipids with unsaturated hydrocarbon chains (10). The formation of these membrane microdomains is seen only within critical concentrations of cholesterol (11). It is well known that both membrane fluidity and curvature are strongly modulated by the amount of cholesterol present in the membrane (12). Membrane microdomains can enhance cell signaling by locally concentrating or excluding selected protein components at specific sites on membranes. In this study, we examine GSIS in  $\beta$ -cells where membrane properties have been altered by depleting or overloading cholesterol.

To date, limited studies on the involvement of membrane microdomains in GSIS from pancreatic  $\beta$ -cells have focused only on proteins at the plasma membrane. It is shown that cholesterol depletion leads to redistribution of  $K^+$  channel  $K_v2.1$  and soluble *N*-ethylmaleimide-sensitive factor attachment protein receptor proteins from detergent-resistant to soluble domains in  $\beta$ -cells and in turn

From the Department of Molecular Physiology and Biophysics, Vanderbilt University Medical Center, Nashville, Tennessee.

Address correspondence and reprint requests to David W. Piston, Department of Molecular Physiology and Biophysics, Vanderbilt University Medical Center, 735 Light Hall, Nashville, TN 37232. E-mail: dave.piston@vanderbilt.edu.

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2-DG, 2-deoxyglucose; apoE, apolipoprotein E; FFA, free fatty acid; GK, glucokinase; GSIS, glucose-stimulated insulin secretion; M $\beta$ CD, methyl- $\beta$ -cyclodextrin; nNOS, neuronal nitric oxide synthase.

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TABLE 1  
Plasma metabolic parameters of mice used in this study

	C57BL/6J	<i>apoE</i> <sup>-/-</sup>	<i>ob/ob</i>	<i>ob/ob;apoE</i> <sup>-/-</sup>
<i>n</i>	5	5	5	4
TC (mg/dl)	121.0 ± 31.6	442 ± 65.0*	217 ± 66.6	728.3 ± 156.1†
TG (mg/dl)	51.3 ± 15.6	89.3 ± 14.7‡	107.0 ± 23.1	180.3 ± 36.0§
FFA (mmol/l)	0.49 ± 0.16	0.50 ± 0.11	0.78 ± 0.15	0.86 ± 0.23
Glucose (mg/dl)	125 ± 2.1	113 ± 6.7	118 ± 12	132 ± 25
Insulin (ng/ml)	0.89 ± 0.19	0.86 ± 0.22	17 ± 3.4	31 ± 8.5

Data are means ± SD. Six-month-old mice were kept on a standard chow diet. Plasma total cholesterol (TC), triglyceride (TG), and FFA levels were measured using enzymatic methods. Statistical analysis was performed using the Student's *t* test: \**P* < 0.01 vs. C57BL/6J, †*P* < 0.01 vs. *ob/ob*, ‡*P* < 0.05 vs. C57BL/6J, §*P* < 0.05 vs. *ob/ob*. ||Values were taken from a previously published study (16).

enhances GSIS (13,14). Membrane microdomains also play an essential role in interleukin-1 $\beta$ -induced nitric oxide (NO) release from  $\beta$ -cells (15). Our results in this study show that in addition to changes in proteins present on the plasma membrane, glucose metabolism involving glucokinase (GK) is affected by cholesterol alteration. Here, we show that excess cellular cholesterol is directly linked to reduced GSIS and that normal secretion could be restored by cholesterol depletion. We further demonstrate one potential mechanism for cholesterol regulation of GSIS that involves modification of neuronal NO synthase (nNOS) and GK activity through cholesterol-rich membrane microdomains on the insulin granules.

## RESEARCH DESIGN AND METHODS

**Mice, cells, and islets.** C57BL/6J mice (Harlan Laboratories, Indianapolis, IN) were used, unless otherwise noted. Apolipoprotein E (*apoE*)-deficient mice (*ApoE*<sup>-/-</sup>) (B6.129P2-*ApoE*<sup>tm1Unc/J</sup>), matching control mice (C57BL/6J), and nNOS knockout mice (B6.129S4-*Nos1*<sup>tm1Phh/J</sup>) were purchased from The Jackson Laboratory (Bar Harbor, ME). *ob/ob;apoE*<sup>-/-</sup> mice (on C57BL/6J background) were produced by intercrossing *ob/+* and *apoE*<sup>-/-</sup> mice (16). All mice were kept on a rodent chow (LabDiet 5001, 12% of calories from fat) and cared for according to the guidelines of the Vanderbilt Institutional Animal Care and Use Committee. 832/13 INS-1 cells were a kind gift from Dr. Newgard (Duke University, Durham, NC).  $\beta$ TC3 and 832/13 INS-1 insulinoma cells were cultured and prepared for experiments as previously described (17,18). Pancreatic islets were isolated and cultured using methods developed in our lab (19).

**Cholesterol assay in islets.** Cholesterol content was quantified in a 96-well plate by a fluorometric method using an enzyme-coupled reaction provided by the Amplex Red Cholesterol Assay kit (Molecular Probes). After isolation, islets were divided into two groups for cholesterol and insulin assays. For cholesterol measurement, material from each tube containing 10 islets was subjected to lipid extraction with chloroform/methanol (2:1; vol/vol), dried down to a thin film, and resuspended in 60- $\mu$ l 1 $\times$  working solution (Amplex Red Cholesterol Assay kit) supplemented with 0.1% Triton X-100. From that, 50  $\mu$ l was used for cholesterol assay and 10  $\mu$ l for protein assay (BCA Protein Assay; Pierce, Rockford, IL).

**Glucose-stimulated insulin secretion measurements.** Static incubations of pancreatic islets for insulin secretion measurements were done as previously described (19). Insulin radioimmunoassays were performed by the Diabetes Research and Training Center Hormone Core Resource at Vanderbilt University. Mouse serum total cholesterol, triglycerides, and free fatty acids were measured by the Vanderbilt Mouse Metabolic Phenotyping Center Lipid Core.

**Cholesterol manipulation.** For metabolic depletion, islets were grown for 3 days in metabolic depletion medium (RPMI-1640; similar to the growth medium but with 10% lipoprotein-deficient serum in place of FBS, supplemented with 200  $\mu$ mol/l mevalonate [required to maintain cell viability] and 10  $\mu$ mol/l mevastatin) to block cholesterol synthesis and deplete cholesterol stores (20). For depletion by methyl- $\beta$ -cyclodextrin (M $\beta$ CD) (Sigma), islets and  $\beta$ -cells were incubated with 10 mmol/l M $\beta$ CD at 37°C for 1 h, which removed cholesterol from both plasma membrane and intracellular stores because intracellular cholesterol effluxes efficiently to the plasma membrane where it is quickly removed by M $\beta$ CD as the acceptor (9). To cholesterol overload, cells were incubated with 10 mmol/l soluble cholesterol (Sigma; 1 g contains ~40 mg cholesterol) at 37°C for 1 h.

**Glucokinase activity assay.** Enzymatic activity of GK was measured by a continuous spectrophotometric rate determination method (21) using 60 mmol/l Tris, 20 mmol/l MgCl<sub>2</sub>, 2.5 mmol/l dithiothreitol, 100 mmol/l KCl, 4 mmol/l adenosine 5'-triphosphate, 0.9 mmol/l  $\beta$ -nicotinamide adenine dinucleotide phosphate, and 10 units glucose 6-phosphate dehydrogenase. One unit of GK activity is defined as phosphorylation of 1  $\mu$ mol D-glucose to D-glucose 6-phosphate per minute at pH 9.0 at room temperature. GK activity was measured by subtracting the activity measured at 0.5 mmol/l glucose from that at 100 mmol/l glucose to account for hexokinase activity in the cell lysates.

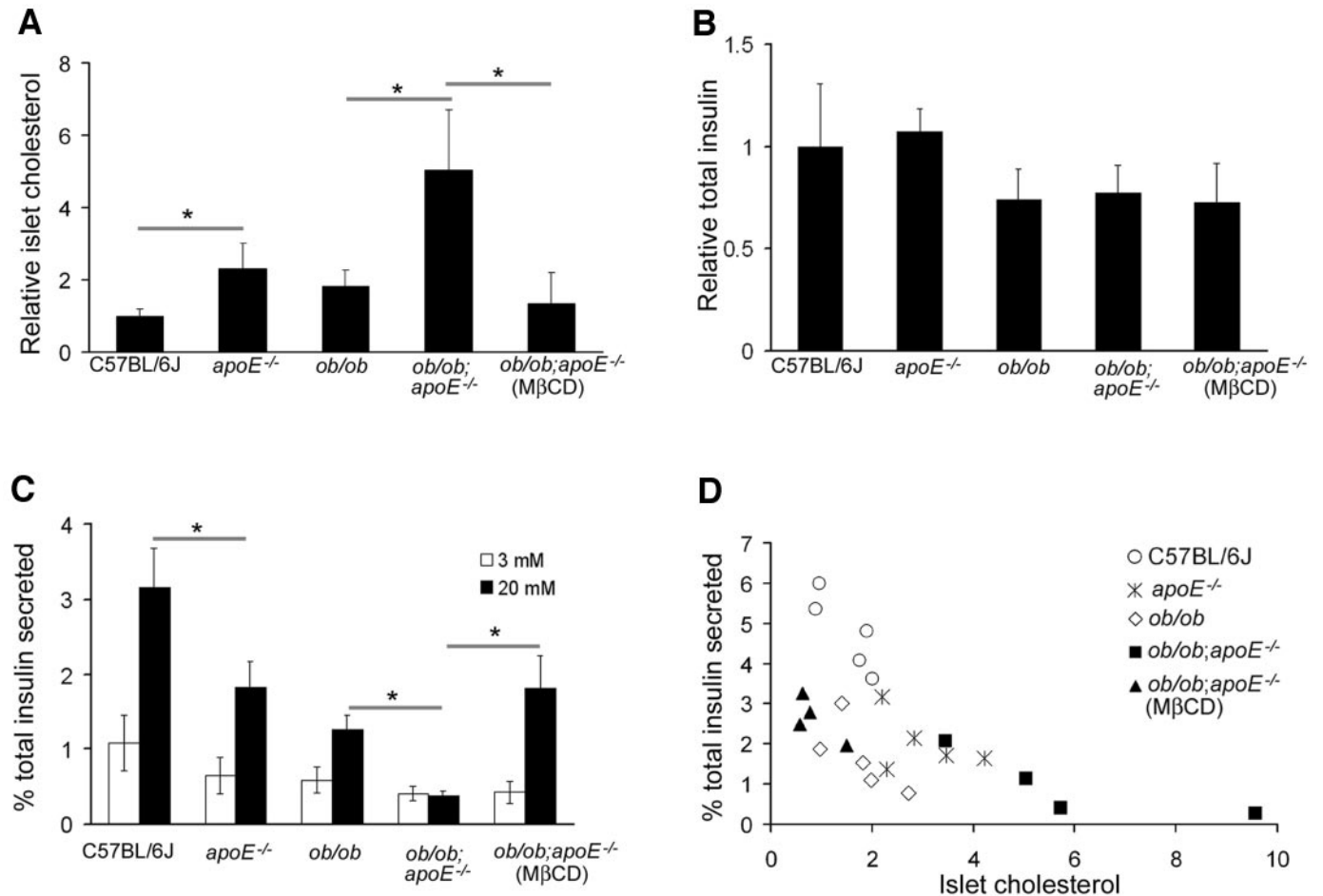
**Low-temperature SDS-PAGE.** Cells were collected in cold PBS and resuspended in cold lysis buffer (20 mmol/l HEPES, 100 mmol/l NaCl, 1 mmol/l EDTA, 1% cholate, 1% Triton X-100, and 1 $\times$  protease inhibitor cocktail for mammalian cells [Sigma]). Cold SDS loading buffer (0.05 mol/l Tris-HCl, pH 6.8, 2% SDS, 10% glycerol, 100 mmol/l dithiothreitol, and 0.1% bromophenol blue) was added to the cell lysate and incubated for 5 min at 0°C. Samples were loaded on 6% polyacrylamide gels, subjected to electrophoresis at 0°C, and visualized by immunoblotting with nNOS monoclonal antibody (BD Transduction Laboratories) and peroxidase-conjugated secondary antibody. To separate the cytoplasmic and membrane-bound fractions, digitonin permeabilization was performed as described previously (22), with the exception of using 10  $\mu$ g/ml instead of 20  $\mu$ g/ml digitonin.

**Statistical analysis.** Results are expressed as means ± SD. Unpaired two-tailed Student's *t* tests were performed for comparisons between treated and control groups. Differences were considered significant at *P* < 0.05. In cholesterol and insulin secretion assays, *n* denotes the number of mice used for each condition. Samples of 10 islets each were taken from each mouse. In all other experiments using cultured  $\beta$ -cells, *n* denotes the number of times each experiment was repeated with different plates of cells for each condition.

## RESULTS

### GSIS is directly affected by islet cholesterol levels.

Traditionally, a high-fat diet is used to increase serum cholesterol levels in mice. However, this is also accompanied by elevations in FFA levels. To circumvent this issue, we used *apoE*-deficient (*apoE*<sup>-/-</sup>) mice fed a standard diet. Compared with control C57BL/6J animals, *apoE*<sup>-/-</sup> mice have elevated plasma cholesterol levels without changes in plasma FFA. Furthermore, when the *apoE*<sup>-/-</sup> mice are made obese by placing them on a leptin-deficient *ob/ob* background, plasma cholesterol levels are further elevated, again without concomitant increases in FFA compared with *ob/ob* controls (Table 1). Because *apoE* is not present in the pancreatic islets of wild-type mice (23), its absence does not directly affect islet function when studied *in vitro*, and we observed similar islet morphology between *apoE*<sup>-/-</sup> and control mice (*apoE*<sup>-/-</sup> vs. C57BL/6J and *ob/ob;apoE*<sup>-/-</sup> vs. *ob/ob*). Cholesterol measurements in pancreatic islets show that an increase in serum cholesterol (Table 1) leads to increased islet cholesterol (Fig. 1A). Although total insulin content does not vary significantly among different mouse models (Fig. 1B), the increase in islet cholesterol causes a dramatic reduction in the islets' ability to secrete insulin in response to basal and high glucose concentrations (Fig. 1C). This result indicates



**FIG. 1.** Reduced GSIS from pancreatic islets in mice with elevated cholesterol levels. Islets were isolated from four mouse strains, as indicated, and the number of mice used from each strain is listed in Table 1. Each sample consisted of 10 islets, and the results were taken from an average of triplicate samples. For results, labeled *ob/ob; apoE*<sup>-/-</sup> (MβCD) islets from *ob/ob; apoE*<sup>-/-</sup> mice were incubated with 10 mmol/l MβCD for 1 h shaking at 37°C before taking measurements. GSIS was measured using static incubations of isolated islets: islets were first incubated in 2 mmol/l glucose for 1 h before transferred to different concentrations of glucose. At the end of the 1-h stimulation period, samples were collected for insulin measurement followed by the addition of 1% Triton X-100 to determine total insulin content. **A:** Islet cholesterol levels were determined from lipids extracted from each sample and normalized to the protein content in the same sample, \**P* < 0.05. **B:** Total islet insulin levels were measured from Triton X-100-extracted samples and normalized to the protein content in the same sample. **C:** Insulin secretion in response to 3 and 20 mmol/l glucose from each sample, expressed as fractional release, \**P* < 0.05. **D:** GSIS (20 mmol/l glucose) plotted against islet cholesterol levels. Each data point represents an average of triplicate samples from one mouse.

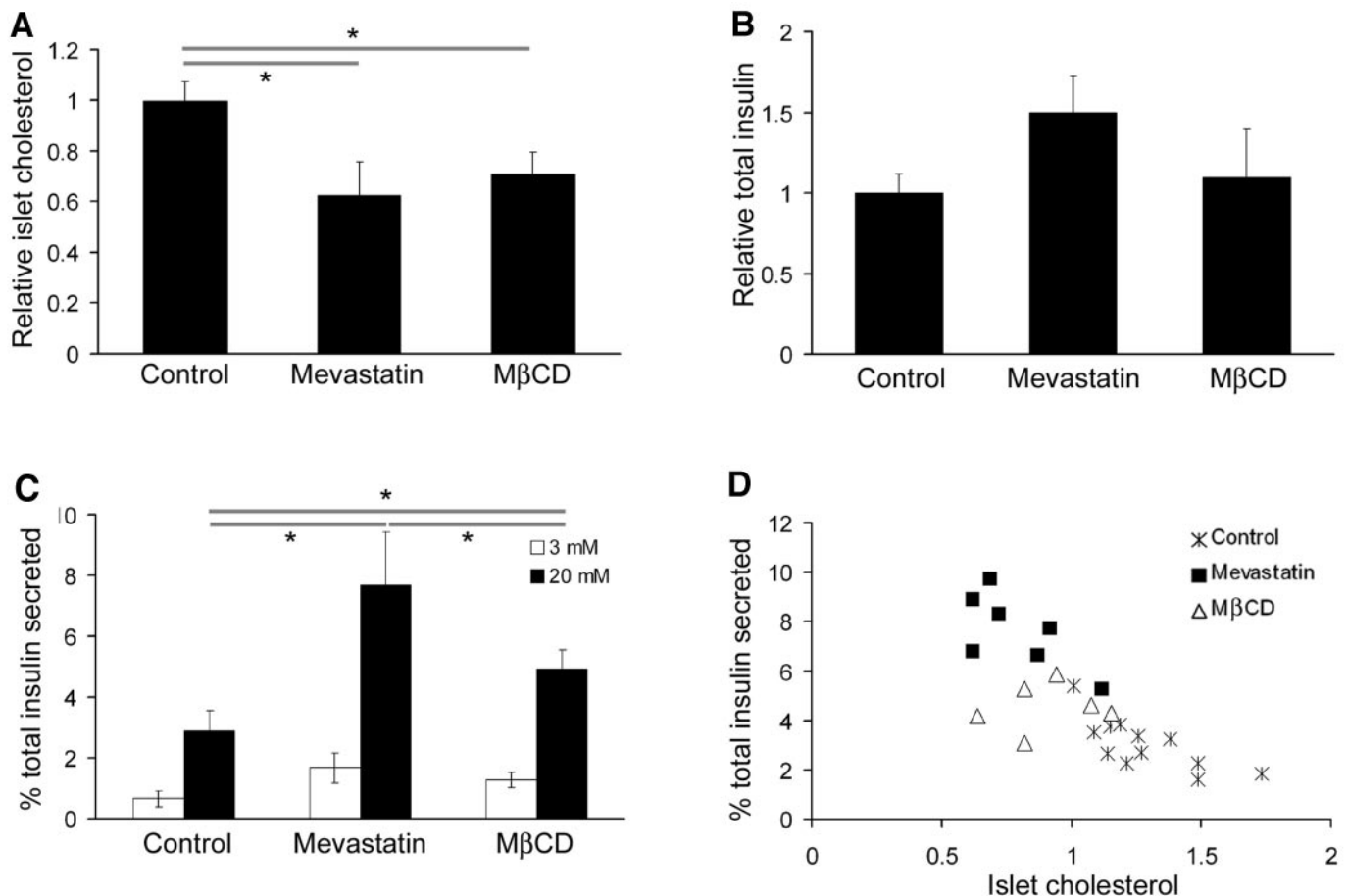
that cholesterol directly impacts GSIS independent of the cellular FFA content.

Because plasma triglyceride levels are also elevated in our mouse models relative to their respective controls (Table 1), we studied isolated islets in which cholesterol can be specifically manipulated. To determine whether direct cholesterol depletion could enhance GSIS from isolated C57BL/6J islets, MβCD was used for 1 h to cause acute depletion of cellular cholesterol (9) and mevastatin (a cholesterol biosynthesis inhibitor) to cause metabolic depletion over several days in culture (20). With these treatments, islet cholesterol was reduced by 30–40% (Fig. 2A), whereas total insulin content remained unaffected (Fig. 2B). Direct cholesterol depletion from isolated islets significantly enhances insulin secretion under both basal and high glucose concentrations (Fig. 2C). Cholesterol depletion of the *ob/ob; apoE*<sup>-/-</sup> islets using MβCD also partially restores GSIS (Fig. 1C). GSIS (20 mmol/l glucose) is shown as a function of islet cholesterol for the genetically modified mouse models (Fig. 1D) or isolated wild-type islets with different cholesterol levels (Fig. 2D). Although many factors affect insulin secretion, in all of

these cases, the islet cholesterol level is inversely correlated to β-cell GSIS.

**Glucose metabolism is involved in cholesterol-regulated GSIS from β-cells.** To investigate the molecular mechanisms underlying cholesterol-dependent GSIS, we turned to cultured pancreatic β-cells. Both 832/13 INS-1 and βTC3 cells secrete insulin in a regulated manner (18,24). INS-1 cells are used for the functional experiments because of their robust responsiveness to glucose (18), and βTC3 cells are used for the imaging experiments because of their low autofluorescence. The capability to manipulate cholesterol in these insulinoma cells is greatly improved relative to that in islets, so that cholesterol overloading using soluble cholesterol (cholesterol loaded on MβCD) is possible (Fig. 3A). In INS-1 cells, using MβCD to deplete cholesterol before glucose stimulation increases insulin secretion at all tested glucose concentrations (Fig. 3B). In contrast, cholesterol overload suppresses GSIS (Fig. 3B). Similar results are observed in βTC3 cells for changes in intracellular Ca<sup>2+</sup>, a critical signaling molecule in GSIS (Fig. 3C). The increase in intracellular Ca<sup>2+</sup> on glucose stimulation is reduced in





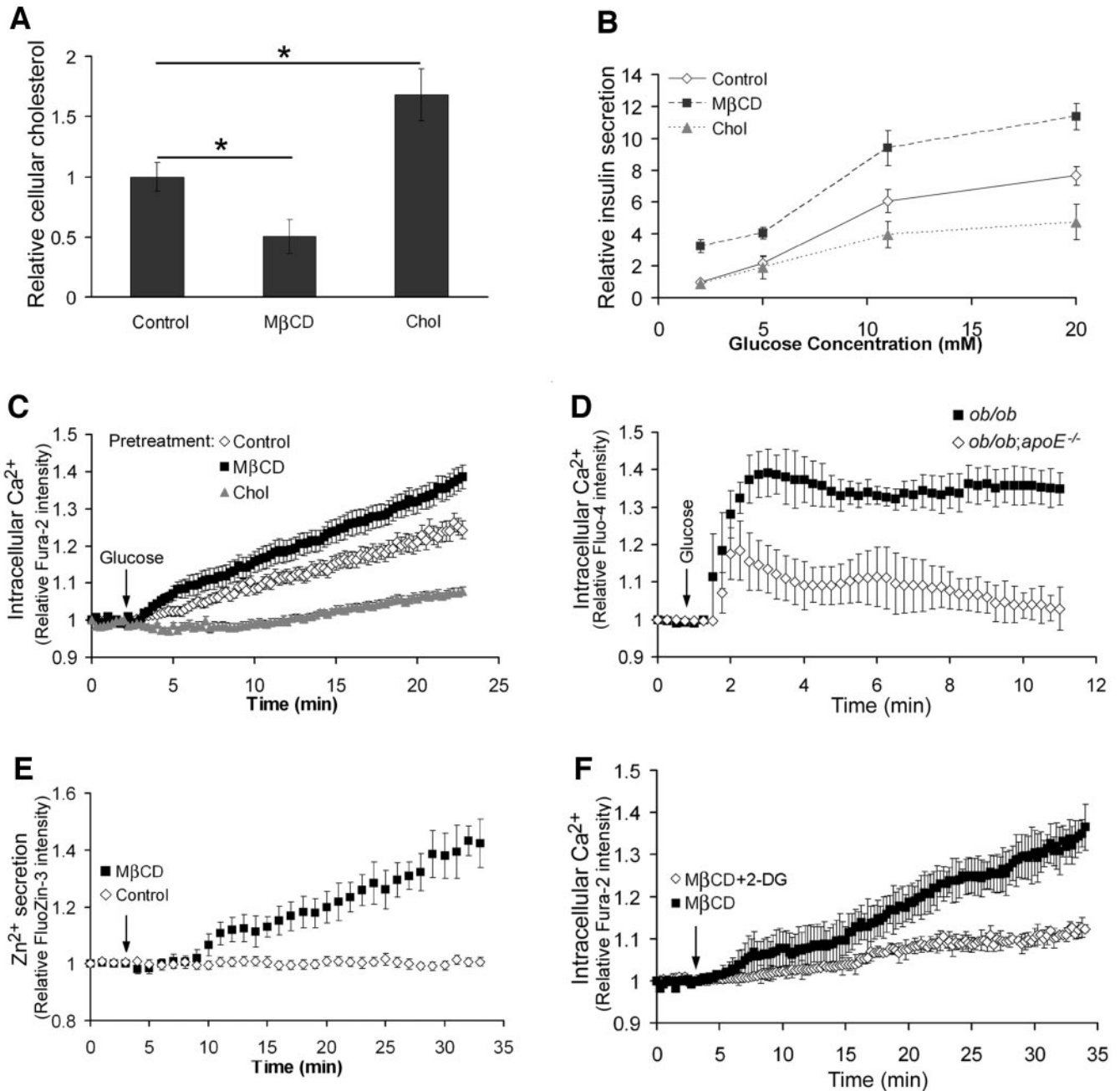
**FIG. 2.** Enhanced GSIS from wild-type pancreatic islets with reduced cholesterol levels. Two methods of cholesterol depletion were used on islets isolated from C57BL/6J: acute depletion by MβCD ( $n = 6$ ) and metabolic depletion by mevastatin ( $n = 7$ ).  $n = 12$  mice for control islets. Each sample consisted of 10 islets, and the results were taken from an average of triplicate samples. GSIS was measured using the method described in Fig. 1. **A:** Islet cholesterol levels were determined from lipids extracted from each sample and normalized to the protein content in the same sample,  $*P < 0.05$ . **B:** Total islet insulin levels were measured from Triton X-100-extracted samples and normalized to the protein content in the same sample. **C:** Insulin secretion in response to 3 and 20 mmol/l glucose from each sample, expressed as fractional release,  $*P < 0.05$ . **D:** GSIS (20 mmol/l glucose) plotted against islet cholesterol levels. Each data point represents an average of triplicate samples from one mouse.

islets from *ob/ob;apoE<sup>-/-</sup>* versus *ob/ob* mice (Fig. 3D). To determine whether cholesterol depletion alone is sufficient to stimulate insulin secretion, we used a fluorescent zinc indicator (FluoZin-3) to monitor insulin release from MβCD-treated βTC3 cells. FluoZin-3 has been successfully used to study insulin release in pancreatic β-cells because insulin and Zn<sup>2+</sup> are co-released by exocytosis (25). When 10 mmol/l MβCD is added to βTC3 cells kept in 2 mmol/l glucose, a rise in FluoZin-3 fluorescence indicates the initiation of insulin secretion after cholesterol removal, even in the absence of other insulin secretagogues (Fig. 3E).

Because cholesterol directly affects GSIS, we examined the glucose-stimulated signal transduction pathway to see where cholesterol action occurs. To determine whether cholesterol is coupled to proximal glucose metabolism, we monitored the effect of MβCD treatment on intracellular Ca<sup>2+</sup> in the presence or absence of a nonmetabolizable glucose analog, 2-deoxyglucose (2-DG). As shown in Fig. 3F, adding MβCD to βTC3 cells in 2 mmol/l glucose raises intracellular Ca<sup>2+</sup>. However, the increase in intracellular Ca<sup>2+</sup> caused by cholesterol depletion is blunted when glucose is replaced by 2-DG, suggesting that glucose metabolism is required for initiating insulin secretory response in cholesterol-depleted cells. We also used

NADPH autofluorescence to assay cellular redox state (26). Pretreating βTC3 cells with MβCD leads to a slightly higher NADPH production on 20 mmol/l glucose stimulation compared with the control cells (Fig. 4A). Overloading with cholesterol before glucose stimulation decreases NADPH production (Fig. 4A). Compared with *ob/ob* mice, NADPH production in islets from *ob/ob;apoE<sup>-/-</sup>* mice is significantly reduced (Fig. 4B). As with Zn<sup>2+</sup> secretion (Fig. 3E) and intracellular Ca<sup>2+</sup> (Fig. 3F), cholesterol depletion with MβCD alone initiates NADPH production (Fig. 4C). NADPH fluorescence does not increase in response to MβCD when 2-DG is present or when soluble cholesterol is added to the cells (Fig. 4D). These results suggest that the effect of cholesterol depletion using MβCD on intracellular Ca<sup>2+</sup> increase, NADPH production, and insulin secretion (indicated by Zn<sup>2+</sup> release) involves proximal glucose metabolism.

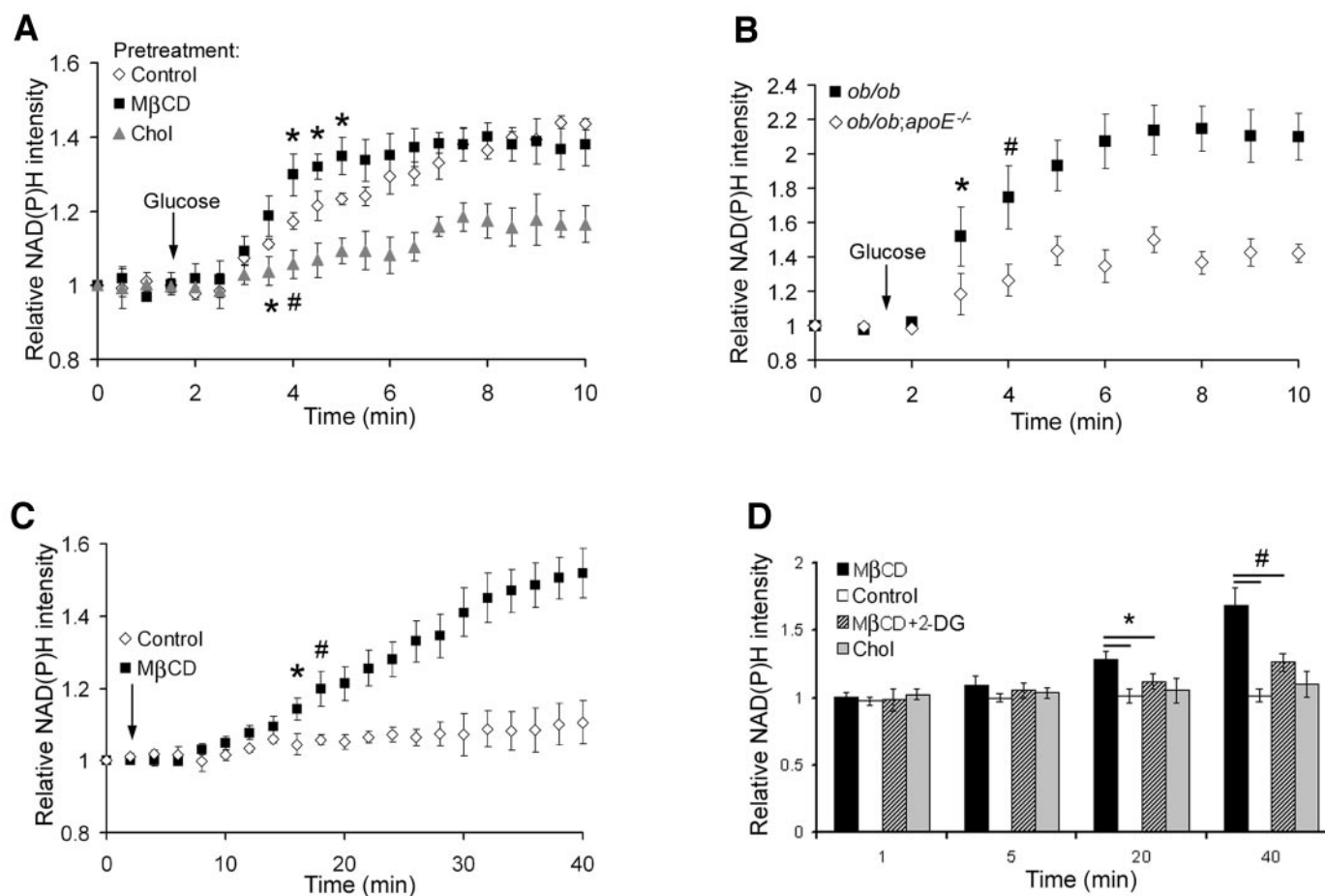
**Cholesterol is involved in nNOS regulation of GK.** GK critically regulates GSIS by exerting tight control over glucose metabolism. Its unique glucose-sensing property underlies the ability of pancreatic β-cells to respond to plasma glucose fluctuations. Because of this, even small changes in GK activity have an impact on the rate of GSIS (27). This is supported by many studies of inactivating mutations in the gene encoding GK in humans with



**FIG. 3.** Cellular cholesterol levels affect GSIS from cultured pancreatic  $\beta$ -cells. Cholesterol depletion and overloading in cultured  $\beta$ -cells were done using 10 mmol/l M $\beta$ CD and 10 mmol/l soluble cholesterol (Chol), respectively, for 1 h at 37°C before taking measurements. **A** and **B**: Cellular cholesterol levels and GSIS, respectively, in 832/13 INS-1 cells after cholesterol treatment. GSIS was measured using the method described in Fig. 1 from cells plated in 12-well plates. **A**:  $n = 8$ ,  $*P < 0.05$ . **B**:  $n = 5$ ,  $P < 0.01$  between M $\beta$ CD-treated and control cells for all glucose concentrations;  $P < 0.05$  between cholesterol-treated and control cells for 11 and 20 mmol/l glucose. **C**: Glucose-stimulated intracellular Ca<sup>2+</sup> response of  $\beta$ TC3 cells with different levels of cellular cholesterol. Cells kept in glucose-free buffer underwent cholesterol treatment before loading with 2  $\mu$ mol/l Fura-2. A 340- to 380-nm intensity ratio of Fura-2 was obtained from the entire image after background correction. The arrow points to when 20 mmol/l glucose was added.  $n = 6$ ,  $P < 0.05$  between M $\beta$ CD-treated and control cells and  $P < 0.01$  between cholesterol-treated and control cells for all time points 3 min postglucose addition. **D**: Glucose-stimulated intracellular Ca<sup>2+</sup> response of islets from *ob/ob* and *ob/ob;apoE<sup>-/-</sup>* mice. Cultured islets were labeled with 4  $\mu$ mol/l Fluo-4 in 2 mmol/l glucose at room temperature for 1 h. The arrow indicates when 20 mmol/l glucose was added.  $n = 5$ ,  $P < 0.01$  for all time points 2 min after glucose addition. **E**: Time course detection of Zn<sup>2+</sup> secretion in M $\beta$ CD-treated  $\beta$ TC3 cells. Images were taken in the presence of FluoZin-3 added to the extracellular medium of cells kept in 2 mmol/l glucose. Fluorescence quantification was done on three small regions in the extracellular medium, as determined by the differential interference contrast image. The arrow indicates the addition of buffer (control) or M $\beta$ CD (10 mmol/l final concentration).  $n = 7$ ,  $P < 0.01$  for all time points 11 min after the start of the experiment. **F**: Changes in intracellular Ca<sup>2+</sup> in response to M $\beta$ CD with and without a nonmetabolizable glucose analog (2-DG). Fura-2-loaded  $\beta$ TC3 cells were kept in either 2 mmol/l glucose or 5 mmol/l 2-DG throughout. The arrow indicates the addition of M $\beta$ CD.  $n = 6$ ,  $P < 0.01$  for all time points 15 min after the start of the experiment.

maturity-onset diabetes of the young (28). It has been shown on glucose stimulation that GK dissociates from the insulin granules and is released to the cytoplasm, where it is activated (29), as depicted in Fig. 5A. In this study,

pretreating INS-1 cells with M $\beta$ CD enhances GK activity under both basal (3 mmol/l) and high glucose (20 mmol/l) concentrations (Fig. 5B). In contrast, cholesterol overloading decreases GK activity on glucose stimulation (Fig. 5B).



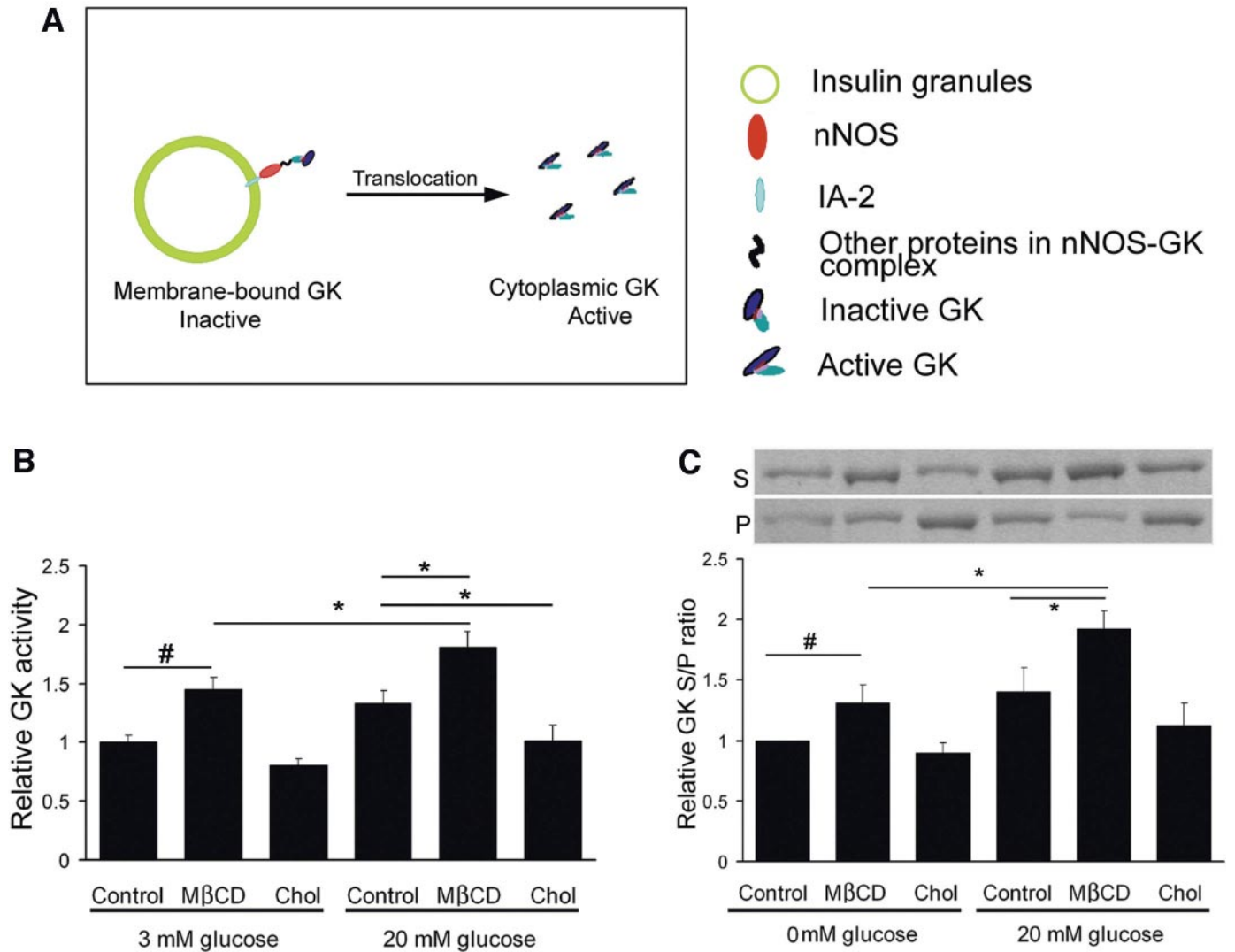
**FIG. 4.** Cholesterol-dependent regulation of GSIS involves glucose metabolism. **A:** Time-course imaging of NADPH response of  $\beta$ TC3 cells stimulated with 20 mmol/l glucose. Glucose-starved cells underwent cholesterol treatment before NADPH imaging ( $40\times$  Plan Neofluar 1.3 NA objective lens, 710-nm mode-locked Ti:Saph laser and a nondescanned detector with a custom 380- to 550-nm filter). The arrow indicates the addition of 20 mmol/l glucose.  $n = 5$ ,  $*P < 0.05$  vs. control.  $\#P < 0.01$  vs. control for this and all subsequent time points. **B:** NADPH response of islets from *ob/ob* and *ob/ob;apoE<sup>-/-</sup>* mice. Cultured islets were incubated in 2 mmol/l glucose for 1 h before imaging. The arrow indicates the addition of 20 mmol/l glucose.  $n = 5$ ,  $*P < 0.05$  and  $\#P < 0.01$  for this and all subsequent time points. **C:** NADPH response to M $\beta$ CD treatment in  $\beta$ TC3 cells kept in 2 mmol/l glucose. The arrow indicates the addition of buffer (control) or M $\beta$ CD (10 mmol/l final concentration).  $n = 7$ ,  $*P < 0.05$  and  $\#P < 0.01$  for this and all subsequent time points. **D:** NADPH response in  $\beta$ TC3 cells kept in 2 mmol/l glucose or 5 mmol/l 2-DG and subjected to different cholesterol treatments. The X-axis shows the time elapsed since the addition of indicated reagents.  $n = 5$ ,  $*P < 0.05$  and  $\#P < 0.01$ .

We next examined GK translocation by immunoblotting of the pellet (membrane) and soluble (cytoplasmic) fractions of digitonin-permeabilized cells under different conditions. Translocation of GK from the membrane bound to the cytoplasmic fraction is observed on cholesterol depletion (Fig. 5C). Further, glucose stimulation results in an additional increase in GK activity and translocation in M $\beta$ CD-treated cells (Fig. 5B and C), which may suggest that cholesterol depletion and glucose stimulation utilize separate mechanisms for GK regulation.

Previous evidence indicates that glucose-stimulated GK translocation involves S-nitrosylation of by nNOS (29). To determine whether nNOS activity impacts GK in cholesterol-depleted cells, we examined nNOS activity using a fluorescent NO indicator, DAF-FM (4-amino-5-methylamino-2',7'-difluorofluorescein). Figure 6A shows that cholesterol depletion before insulin stimulation prevents NO production observed in control cells. Insulin was used here as the stimulant because it caused a more rapid change in nNOS activity than glucose (29). The inhibition in NO production is consistent with the observation that disruption of lipid microdomains by M $\beta$ CD results in reduced NO release from insulin-secreting cells (15). Add-

ing M $\beta$ CD to  $\beta$ TC3 cells in 2 mmol/l glucose leads to a decrease in DAF-FM fluorescence, indicating that cellular NO content is reduced on cholesterol depletion (Fig. 6B). These results suggest that nNOS activity is inhibited by cholesterol depletion.

Since cholesterol modulation using M $\beta$ CD occurs within 30 min (30), its effect on nNOS and GK activity is unlikely to work through changes in gene transcription or protein stability. To determine how cholesterol depletion might decrease nNOS activity and release GK from insulin granules, we looked at the role of cholesterol-rich membrane microdomains in determining nNOS properties. nNOS is localized to insulin granules through its interaction with the transmembrane protein insulinoma-associated protein 2 in pancreatic  $\beta$ -cells (31). Characterization of nNOS has shown that the native protein is a homodimer (32), and dimerization is absolutely required for nNOS activity (33). Thus, we hypothesized a model where cholesterol depletion disrupts nNOS dimers, which in turn weakens the association between nNOS and GK, thereby releasing GK to the cytoplasm, where it becomes activated. Excess cholesterol, on the other hand, stabilizes nNOS dimers and



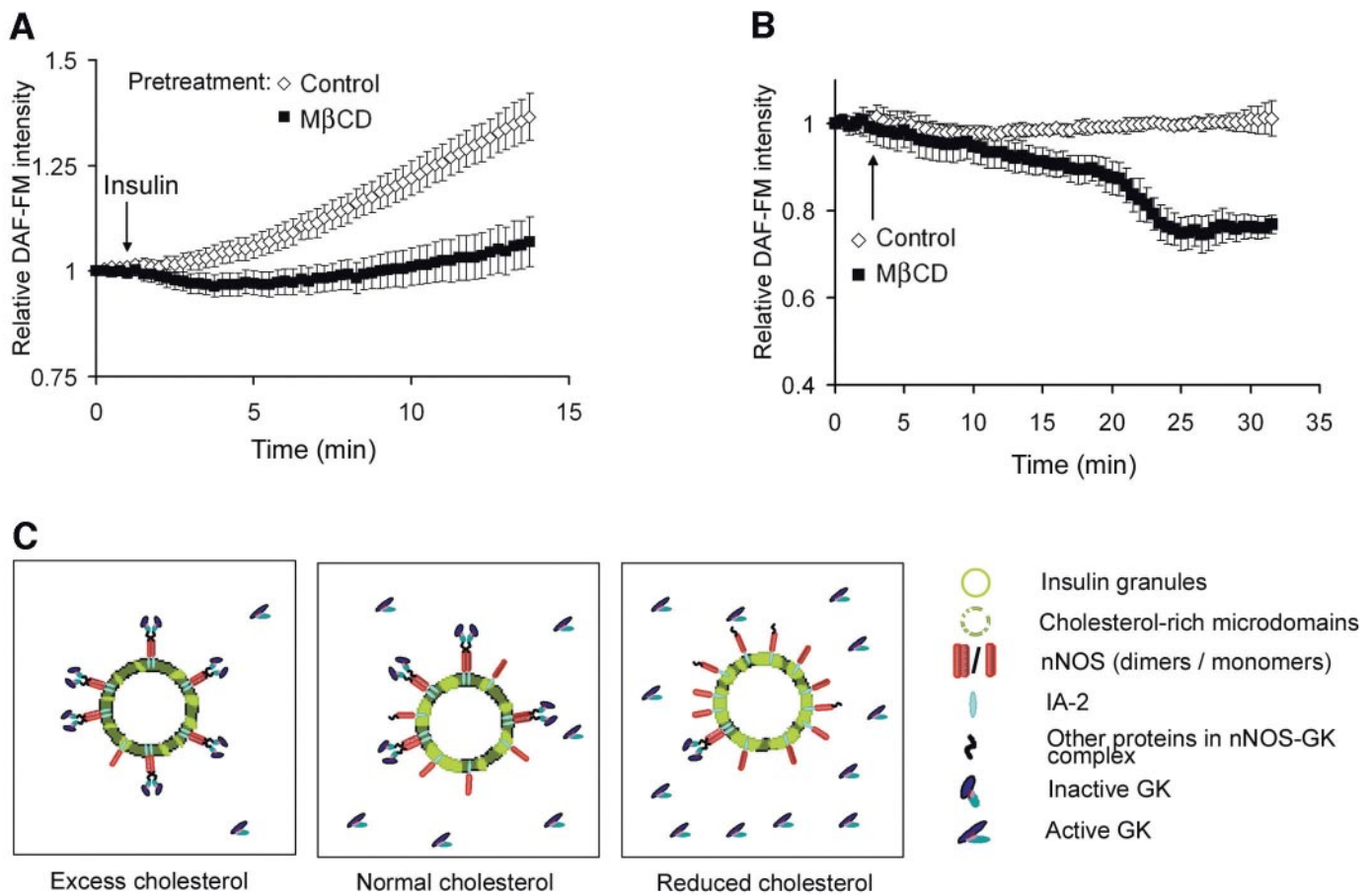
**FIG. 5.** Cholesterol modulation changes GK activity and localization. **A:** Schematic drawing of how GK activity is related to its cellular localization (see ref. 29). Glucose stimulation results in translocation of GK from insulin granules to the cytoplasm where GK is activated. **B:** GK enzymatic activity in 832/13 INS-1 cells under different conditions and normalized to that measured in control cells under basal glucose.  $n = 8$ ,  $*P < 0.05$  and  $\#P < 0.01$ . **C:** Translocation of GK from the membrane-bound to the cytoplasmic fraction in 832/13 INS-1 cells under different conditions. Distribution of GK was quantified by taking a ratio of the amount of GK present in the digitonin-soluble (S, cytoplasmic) fraction to that in the pellet (P, membrane-bound) fraction, and normalized to that in glucose-starved control cells.  $n = 6$ ,  $*P < 0.05$  and  $\#P < 0.01$ .

keeps GK bound to insulin granules, where it is less active (Fig. 6C).

To test this hypothesis, we immunoprecipitated nNOS from cell lysates and probed by immunoblot for GK. As shown in Fig. 7A, cholesterol depletion by MβCD treatment results in dissociation of GK from nNOS under both low and high glucose. In contrast, cholesterol overloading increases association of GK with nNOS on glucose stimulation. There is an additional weakening of the interaction between GK and nNOS when MβCD-treated cells are stimulated with high glucose (Fig. 7A), again possibly suggesting that nNOS regulation of GK by cholesterol depletion is different from that by glucose stimulation. Our hypothesis is further supported by the fact that GSIS is greatly enhanced in nNOS knockout mice (Fig. 7B), consistent with the idea that a weaker association of GK with insulin granules (due to the lack of granule-associated nNOS) leads to greater GK activity in β-cells. Using low-temperature SDS-PAGE (34), we show that there is less nNOS dimerization in cholesterol-depleted cells com-

pared with untreated cells and that the opposite is observed with cholesterol overloading (Fig. 7C). We also measured the nNOS dimer and monomer distribution in the membrane-bound and cytoplasmic fractions using digitonin permeabilization. In both control and cholesterol-depleted cells, there are more nNOS dimers in the pellet fraction, suggesting that the dimer structure is enriched in the insulin granule membranes (Fig. 7D). Furthermore, the nNOS oligomeric state dramatically shifts from predominantly dimers to mostly monomers in the membrane-bound fraction on cholesterol depletion. These results show that cholesterol depletion disrupts nNOS dimers associated with insulin granules and leads to decreased nNOS activity (Fig. 6A and B). This is consistent with the previous finding that disruption of the dimeric state of nNOS by its inhibitors affects the secretagogue-induced insulin response (35). Although the exact mechanism by which nNOS dimerization is affected by membrane structure remains unknown, it is conceivable that clustering of insulinoma-associated protein 2 in cholesterol-rich do-





**FIG. 6.** Cholesterol depletion inhibits nNOS activity. **A:** Reduced NO production in insulin-stimulated  $\beta$ TC3 cells pretreated with M $\beta$ CD, detected using a fluorescent indicator (DAF-FM). The arrow indicates the addition of 100 nmol/l insulin.  $n = 5$ . **B:** Decreased cellular NO content in response to M $\beta$ CD.  $\beta$ TC3 cells were kept in 2 mmol/l glucose and loaded with DAF-FM. The arrow indicates the addition of buffer (control) or M $\beta$ CD (10 mmol/l final concentration).  $n = 5$ . **C:** Proposed model of nNOS regulation of GK by its dimeric state at different cholesterol levels. Cholesterol depletion disrupts nNOS dimers and weakens the association between nNOS and GK, leading to the release of GK to the cytoplasm, where it is activated. Excess cholesterol stabilizes nNOS dimers and keeps GK bound to insulin granules, where it is less active.

mains effectively increases the local concentration of nNOS, thus promoting dimer formation.

## DISCUSSION

Our results indicate that excess cellular cholesterol plays a direct role in pancreatic islet dysfunction and may well be a key factor underlying the progression of type 2 diabetes. Using different animal models, we show that elevated serum cholesterol leads to increased cholesterol in pancreatic islets. More importantly, islet cholesterol levels directly and significantly impact the extent of GSIS, independent of FFA levels. This has great implications because it indicates the existence of a novel mechanism linking hyperlipidemia and the pathogenesis of type 2 diabetes that is independent of FFAs. It also suggests that the regulation of cellular cholesterol may be a potential target for therapeutic intervention aimed at preserving or improving GSIS function in pancreatic  $\beta$ -cells.

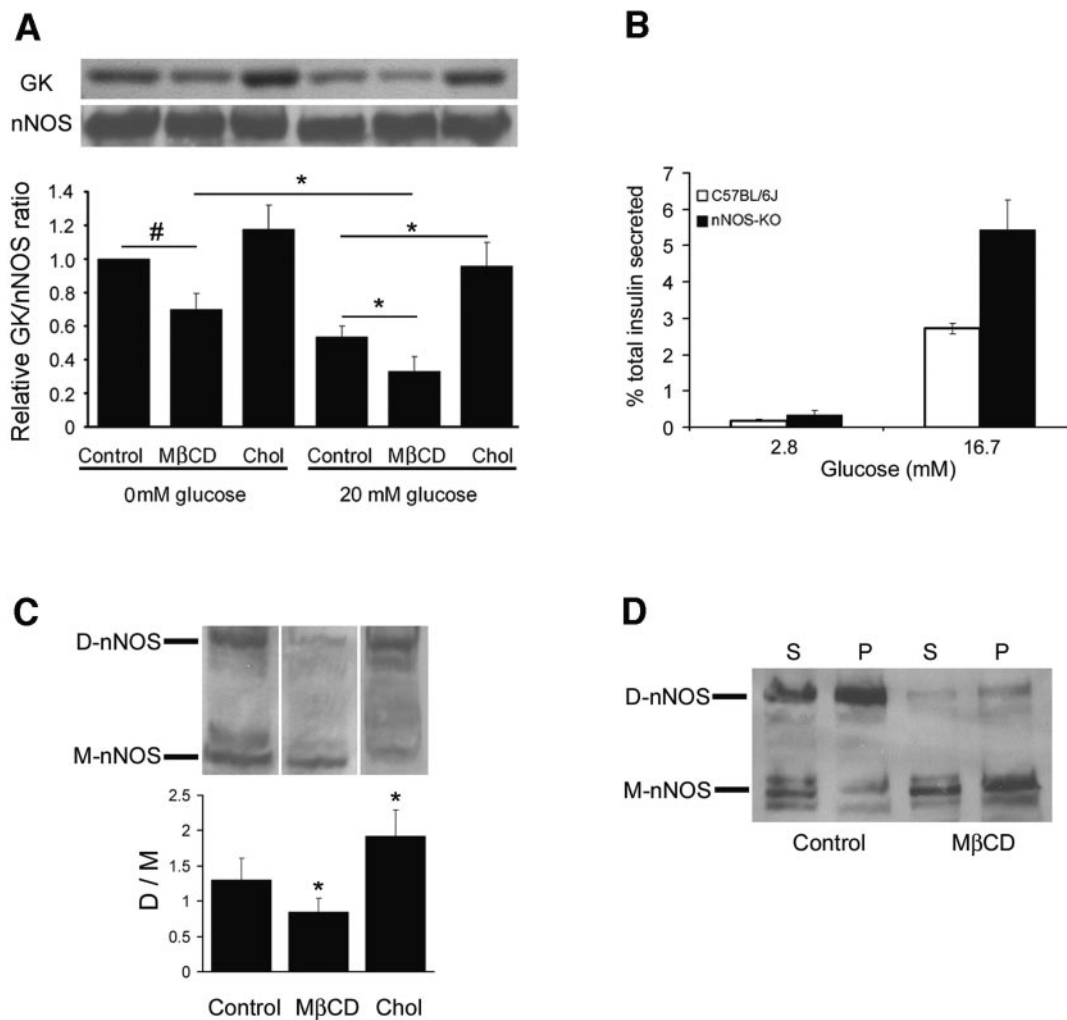
It has been shown that VLDL is markedly increased in *apoE*<sup>-/-</sup> mice (36), and VLDL exposure leads to a decrease in insulin mRNA levels in  $\beta$ TC3 cells (37). However, we did not find a significant difference between control and *apoE*<sup>-/-</sup> mice in either plasma insulin levels (Table 1) or total islet insulin content (Fig. 1B). The discrepancy might be due to the difference in the cells used (cultured insulinoma cells vs. islets), cholesterol-to-triglyceride ratio of the VLDL particles (triglyceride-rich human VLDL vs.

triglyceride-poor *apoE*<sup>-/-</sup> VLDL), or the length of exposure time (short term vs. chronic). ApoE deficiency is shown to induce greater insulin sensitivity (38). This may explain the relatively normal glucose tolerance in *apoE*<sup>-/-</sup> mice despite the reduced GSIS observed in our study.

Because of the profound effects of cholesterol on lipid organization and cellular functions, cells have formulated comprehensive mechanisms to maintain membrane cholesterol levels within a narrow range (39). Among all the signal transduction pathways in which cholesterol may be involved in  $\beta$ -cells, we have focused on the role of cholesterol-rich membrane domains in GSIS and one molecular mechanism in particular involving nNOS regulation of GK. Our data show that cholesterol depletion results in a change of nNOS dimer-to-monomer ratio, which in turn causes translocation of GK from insulin granules to the cytoplasm, where it is activated. In contrast, excess cholesterol prevents GK activation by keeping GK bound to insulin granules. Further studies are underway to provide additional details for other GSIS pathways that may also be affected by cholesterol modulation.

Membrane microdomains have raised great interest because they are proposed to be involved in many cellular functions, including signal transduction, protein trafficking, and cell polarization, as well as pathogenesis of many diseases (40). Cholesterol is an essential component of many putative membrane domains, and removal of chole-





**FIG. 7.** Cellular cholesterol levels affect the interaction between GK and nNOS through altering nNOS dimeric state. **A:** Association of GK with nNOS in 832/13 INS-1 cells under different conditions. Pellets immunoprecipitated from cell lysates using anti-nNOS pre-conjugated to Sepharose beads were analyzed by immunoblot using anti-nNOS and anti-GK antibodies. GK association with nNOS was quantified by taking the ratio of GK to nNOS signal and expressed as fraction of that in glucose-starved control cells.  $n = 6$ .  $*P < 0.05$  and  $\# P > 0.01$ . **B:** GSIS from islets isolated from nNOS knockout (nNOS-KO) mice. GSIS was measured using the method described in Fig. 1.  $n = 7$ . **C:** Detection of nNOS dimers (D-nNOS) and monomers (M-nNOS) in control, cholesterol-depleted (MβCD), and overloaded (Chol) 832/13 INS-1 cells using low-temperature SDS-PAGE.  $n = 5$ .  $*P < 0.05$  vs. control. **D:** Detection of nNOS dimers and monomers in control and cholesterol-depleted 832/13 INS-1 cells permeabilized by digitonin. Pellet (P, membrane-bound) and soluble (S, cytoplasmic) fractions were analyzed by low-temperature SDS-PAGE.

terol disrupts the function of proteins residing in these domains. It has been shown that MβCD treatment results in enhanced GSIS from HIT-T15 hamster insulinoma cells, due to redistribution of voltage-gated  $K^+$  channel and soluble *N*-ethylmaleimide-sensitive factor attachment protein receptor proteins of the putative membrane microdomains (13). However, our results indicate that glucose metabolism is directly affected by cholesterol depletion in addition to any changes in plasma membrane protein distribution that may occur. Our data also suggest that the function of two proteins critically involved in GSIS, nNOS, and GK depends on the cholesterol-enriched lipid environment of insulin granules. nNOS is found to be associated with granule membranes in many cell types. In neurons, MβCD redistributes nNOS from the Triton X-100 insoluble to the soluble fraction, and cholesterol repletion reverses nNOS localization (41). Several lines of evidence have shown that some intracellular membranes not only contain high amounts of cholesterol but also form detergent insoluble domains (42). For example, cholesterol makes up ~70% of the lipids in bovine pituitary secretory granule

membranes (43). Cholesterol offers the rigidity necessary for membrane curvature during secretory granule biogenesis. Insulin is packaged into dense core granules, which bud from the *trans*-Golgi network containing cholesterol and sphingolipid-rich microdomains (44).

Although the relationship between elevated cholesterol and diabetes has not been extensively examined, there is some evidence supporting a potential link between the two. First, a locus on chromosome 9 is associated with both the control of cholesterol levels and diabetes (45). Second, blood lipoprotein levels are an indication of the plasma cholesterol content and have been confirmed as important predictors for the onset of type 2 diabetes (46). In a recent study (46) aimed at examining the link between the use of a statin, which lowers serum cholesterol concentration, and subsequent risk of developing diabetes, it was shown that preventative administration of pravastatin reduced the occurrence of type 2 diabetes by 30%. However, results from this type of study are thus far inconclusive. Even though obesity has been identified as an important risk factor for type 2 diabetes, only a small

percentage of obese patients will eventually develop the disease. Research suggests that predisposition to the development of type 2 diabetes may be due to genetic differences, although the specific genes involved have not yet been identified. Type 2 diabetes has been associated with high synthesis and low absorption of cholesterol, indicating a link between glucose and cholesterol metabolism (47–50). Indeed, our study opens the possibility that genetic variations relating to cellular cholesterol management may play a role in this genetic predisposition for type 2 diabetes.

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