The Effect of Tribbles-Related Protein 3 on ER Stress-Suppressed Insulin Gene Expression in INS-1 Cells

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Background: The highly developed endoplasmic reticulum (ER) structure in pancreatic beta cells is heavily involved in insulin biosynthesis. Thus, any perturbation in ER function inevitably impacts insulin biosynthesis. Recent studies showed that the expression of tribbles-related protein 3 (TRB3), a mammalian homolog of Drosophila tribbles, in various cell types is induced by ER stress. Here, we examined whether ER stress induces TRB3 expression in INS-1 cells and found that TRB3 mediates ER stress-induced suppression of insulin gene expression.

Methods: The effects of tunicamycin and thapsigargin on insulin and TRB3 expression in INS-1 cells were measured by Northern and Western blot analysis, respectively. The effects of adenovirus-mediated overexpression of TRB3 on insulin, PDX-1 and MafA gene expression in INS-1 cells were measured by Northern blot analysis. The effect of TRB3 on insulin promoter was measured by transient transfection study with constructs of human insulin promoter.

Results: The treatment of INS-1 cells with tunicamycin and thapsigargin decreased insulin mRNA expression, but increased TRB3 protein expression. Adenovirus-mediated overexpression of TRB3 decreased insulin gene expression in a dose-dependent manner. A transient transfection study showed that TRB3 inhibited insulin promoter activity, suggesting that TRB3 inhibited insulin gene expression at transcriptional level. Adenovirus-mediated overexpression of TRB3 also decreased PDX-1 mRNA expression, but did not influence MafA mRNA expression.

Conclusions: This study showed that ER stress induced TRB3 expression, but decreased both insulin and PDX-1 gene expression in INS-1 cells. Our data suggest that TRB3 plays an important role in ER stress-induced beta cell dysfunction.

Keywords: Diabetes; Endoplasmic reticulum stress; Insulin; TRB3

INTRODUCTION

The endoplasmic reticulum (ER), a membrane component that is located near the nucleus, is an organelle where amino acids synthesized by mRNA become mature proteins after going through folding, assembly, glycosylation, disulfide bonding, and post-translational modifications [1]. ER stress is induced by a large influx of immature proteins that the ER cannot manage or the depletion of calcium by physiological or pathological environment [1-3]. The ER is well developed in endocrine cells such as pancreatic beta cells in which proteins are synthesized and secreted. Active insulin is made in the ER from proinsulin, which is converted to proinsulin by post-translational modification and disulfide bond formation, and then removal of C-peptide [4,5]. Because the ER of pancreatic beta cells plays an important role in synthesizing and secreting active insulin, any ER stress by pathological conditions can impair the biosynthesis of insulin [6].

When ER stress occurs, cellular defense mechanisms related to ER stress response are activated for the survival of the cells.
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The ER stress response is made up of: 1) activation of protein kinase RNA (PKR)-like ER kinase (PERK) and reduction of mRNA to protein translation by phosphorylation of eIF2α (eukaryotic translation initiation factor 2α subunit) [7], 2) activation of inositol-requiring 1 (IRE1)/X-box binding protein 1 (XBP-1) and transcription factor (ATF6) through the increased expression of ER chaperones and subsequent increase of ER folding capacity [8,9], 3) ER stress-associated protein degradation, which degrades unfolded or improperly folded proteins [10], and 4) apoptosis by the activation of CCAAT/enhancer-binding homologous protein (CHOP) [2,11].

In addition, many intracellular signal transduction systems are generated by ER stress; recent studies indicate that the expression of tribbles-related protein 3 (TRB3) is increased by ER stress-activated ATF4 and CHOP [12]. TRB3 contributes to insulin resistance by physically interrupting phosphorylation of Akt/PKB, which is important for insulin transduction [13]. During fasting, activation of cAMP response element binding protein (CREB) in liver induces the expression of peroxisome proliferator-activated (PPAR)-gamma coactivator-1α (PGC-1α) and caused gluconeogenesis. In this situation, activated CREB/PGC-1α increases TRB3 expression and causes insulin resistance by interrupting insulin action, which increases during food intake [14]. Additionally, TRB3 enters the nucleus by binding other transcription factors, which affects cellular growth, differentiation, and metabolism [13,15-17]. TRB3 in the nucleus is involved in the differentiation of adipocytes by activating CCAAT/enhancer-binding protein β (C/EBP) [18,19], and is also involved in lipid metabolism of adipocytes by activating E3 ligase constitutive photomorphogenic 1 (COP1) [20]. However, so far, there is little information on the role of TRB3 in pancreatic beta cells.

The objectives of this study were to elucidate whether ER stress mediates the expression of TRB3 in INS-1 rat insulinoma cell line (INS-1 cell) and to determine the effect of TRB3 on the expression of insulin gene using adenovirus containing TRB3.

METHODS

Materials
Tunicamycin and thapsigargin used in the study were purchased from Sigma (St. Louis, MO, USA). The TRB3 antibodies used in the Western blot analysis were purchased from Calbiochem (La Jolla, CA, USA). Anti-rabbit antibodies and the radio-labeled probe ([α-32P]dCTP) was purchased from Amersham Bioscience (Little Chalfont, UK). Mouse TRB3 cDNA and recombinant TRB3 adenovirus was provided by Professor Seong-Hoi Koo of the Sungkyunkwan University College of Medicine.

Cell incubation
Insulin dependent rat host-cells, INS-1 cells, with subculture numbers between 20 and 30, were maintained in RPMI 1640 medium (Giboco-BRL, Grand Island, NY, USA) with 10% fetal bovine serum, 1 mM pyruvate, 10 mM HEPES, 50 mM 2-mercaptoethanol, 100 units/mL penicillin, 100 μg/mL streptomycin and incubated in 5% CO2/95% air at 37°C. The medium was changed every 3-4 days and was successively cultured using trypsin-EDTA.

Preparation of recombinant adenovirus
LacZ expressing adenovirus was used as a control for recombinant TRB3 adenovirus. The cDNA encoding a LacZ was inserted into the pAdTrack-CMV shuttle vector. The vector construct was then electroporated into BJ5138 cells, and a recombinant vector was generated using the AdEasy adenoviral vector system. The recombinant viruses were amplified in HEK-293 cells and purified by CsCl (Sigma) gradient centrifugation. Viral preparations were collected and desalted, and titers were determined using Adeno-X rapid titer (BD Bioscience, San Jose, CA, USA), according to the manufacturer’s instructions.

Northern blot analysis
INS-1 cells were treated with tunicamycin and thapsigargin, and infected with adenoviral vectors expressing TRB3. Cells were harvested at the indicated times, total RNA was isolated using Trizol reagent (Invitrogen, Carlsbad, CA, USA) according to the manufacturer’s instructions, and 20 μg of total RNA from each sample was used. The probes for insulin, PDX-1 and MafA were labeled with [α-32P]dCTP using a random-primer DNA-labeling system (Amersham Biosciences, Little Chalfont, UK).

Western blot analysis
INS-1 cells were treated with tunicamycin and thapsigargin for various times and then harvested in lysis buffer (50 mmol/L Tris-HCl [pH 8.0], 150 mmol/L NaCl, 1 mmol/L EDTA, 1% Nonidet P-40, 0.25% Na-deoxycholate) containing proteinase inhibitors. The proteins were resolved by SDS-PAGE and then transferred electrophoretically to a polyvinyl difluoride mem-
brane (Millipore, Bedford, MA, USA). The membrane was blocked by incubation in blocking buffer, incubated with anti-TRB3 antibody and was then developed using an ECL Western blot detection kit (Amersham Biosciences). The membrane was rebotted with anti-actin antibody to verify equal loading of protein in each lane. Densitometry was used to quantitate the results, using the digitalized scientific software program UNSCAN-IT (Skik Scientific Co., Orem, UT, USA).

**Luciferase activity measurement**

INS-1 cells were plated at a density of $3 \times 10^5$ cell per well in a 12-well plate and subcultured for 2 days in INS-1 medium. The promoter constructs (300 ng/well) and other DNAs were transiently transfected by using Lipofectamine™ 2000 transfection reagent (Invitrogen). β-galactosidase plasmids were co-transfected as an internal control. Cells were transfected for 4 hours, washed to remove plasmids. Cells were harvested approximately 24 hours after transfection for luciferase and β-galactosidase assays. 20 μL of cell lysate containing 15 μg of protein was analyzed by using the Luciferase assay system according to the manufacturer’s instructions (Promega, Madison, WI, USA). Luciferase activity was detected using a SIRUS Luminometer (Berthold, Pforzheim, Germany). The luciferase activity was normalized by using the β-galactosidase activity.

**Induction of ER stress**

Tunicamycin and thapsigargin were used to induce ER stress in INS cells. Insulin expression was measured after 24 hours of treatment with 2 μg/mL tunicamycin, and after 5 hours of treatment with 1 μM thapsigargin. Additionally, TRB3 protein expression was measured at an indicated time point after treatment of tunicamycin and thapsigargin.

**Statistical analysis**

The results are given as the average ± standard deviation, and Duncan’s test was used to analyze variables. Cases with a $P$ value lower than 0.05 were statistically significant for determination, and all trials were independently run over three times.

**RESULTS**

**The effect of ER stress on insulin gene expression in INS-1 cells**

To determine the effect of ER stress on the insulin gene expression, INS-1 cells were treated with the ER stress-inducing sub-

**Fig. 1.** The effects of tunicamycin and thapsigargin on insulin mRNA expression in INS-1 cells. Northern blot analysis of insulin mRNA expression in INS-1 cells treated with tunicamycin (A) and thapsigargin (B). INS-1 cells were treated with tunicamycin (2 μg/mL) for 24 hours or thapsigargin (1 μM) for 5 hours. 18S rRNA levels were analyzed as an internal control. Data in bar graph are the mean ± SEM of three independent measurements. $^aP < 0.01$ and $^bP < 0.001$ compared to control.
pressing adenoviruses were prepared. After infecting INS-1 cells with TRB3-expressing adenovirus at varying concentrations, the proportional increase in TRB3 expression was confirmed (Fig. 3). Control INS-1 cells showed high expression of insulin mRNA. However, when TRB3 overexpression was induced with adenovirus, there was a dose-dependent decrease in insulin mRNA expression (Fig. 3). The decrease in insulin gene expression caused by TRB3 is not an effect of adenovirus because insulin mRNA expression did not decrease in response to the control LacZ-expressing adenovirus (Fig. 3).

The effects of TRB3 on insulin promoter activity

In order to determine the effect of TRB3 on insulin gene transcriptional activity, a human insulin promoter-expressing vector and TRB3 expressing vector were cotransfected in INS-1 cells. With this approach, we measured the activity of luciferase under the control of the insulin transcriptional promoter. As shown in Fig. 4, insulin promoter luciferase activity decreased in a dose-dependent manner with the TRB3 expressing vector. Thus, TRB3 appears to suppress gene expression that is regulated by the insulin gene transcriptional promoter region.

The effects of TRB3 on the expression of MafA and PDX-1 in INS-1 cells

Finally, we examined the effects of TRB3 on the expression of the pancreatic beta cell specific transcription factors, PDX-1 and MafA, which stimulate the expression of insulin mRNA. At baseline, INS-1 cells expressed high levels of PDX-1 and MafA mRNA, but when infected with adenovirus TRB3, the cells demonstrated significantly decreased PDX-1 mRNA expression but had no effect on the expression of MafA mRNA (Fig. 3).

DISCUSSION

Several studies have demonstrated that in pancreatic beta cells with well developed ERs, beta cell dysfunction caused by ER stress can contribute to diabetes [21-25]. Initially, in obesity or
type 2 diabetes, in order for the body to overcome insulin resistance in peripheral tissues and maintain normal blood glucose levels, hyperinsulinemia occurs. Thus, in long term continued cases of insulin resistance, demands for insulin synthesis and secretion continuously increase preproinsulin protein influx into the ER of pancreatic beta cells to maintain homoeostatic glucose metabolism and cause ER stress [2]. Therefore, the ER plays an important role in beta cell function. It can be assumed that beta cell dysfunction caused by ER stress can cause diabetes. Our previous study indicated that exposing INS-1 cells to high glucose conditions for a long time can reduce insulin gene expression due to glucose toxicity [26]. Additionally, chronic hyperglycemia causes ER stress in INS-1 cells and the pancreas islets of OLETF rats [6]. Our previous study indicated that the mechanism of ER stress-induced insulin gene suppression is related to the activation of ATF6 [6]. This study also showed that treating INS-1 cells with tunicamycin and thapsigargin results in a decrease in insulin gene expression, which agrees with results from our previous work. In addition, in this study, we confirmed that TRB3 reduces insulin gene expression in response to ER stress. TRB3 is expressed during fasting in the liver through the activation of the PPAR-α/PGC-1α pathway and causes insulin resistance by suppressing Akt/PKB activity [13,27]. Bi et al. [28] reported that TRB3 can cause insulin resistance in adipose tissue. In metabolic syndrome-induced rats, it has been reported that there is an increase in TRB3 and significant decrease in Akt phosphorylation in adipose tissue. Collectively, these data showed that TRB3 affects insulin resistance not only in liver tissue, but also in adipose tissue. It has been reported that ER stress induces the expression of TRB3 in a variety of cells. Corcoran et al. [29] reported that ER stress induces expression of TRB3 in the MCF-7 human breast cancer cell line, DU 145 human prostate cancer cell, and H1299 human lung cancer cell, etc.
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Additionally, it has been reported that tunicamycin increases the expression of TRB3 in human embryonic kidney cells and HepG2 liver cancer cells [12]. In this experiment, we confirmed that tunicamycin and thapsigargin induced ER stress and increased the expression of TRB3 in INS-1 rat insulinoma cells.

Recently, Qian et al. [30] reported that the increased expression of TRB3 as a result of glucose toxicity and ER stress induces apoptosis and dysfunction of the pancreatic beta cells. In this study, we investigated the effect of ER stress-induced TRB3 expression on insulin gene expression. When TRB3 was overexpressed in INS-1 cells by infecting the cells with TRB3 expressing adenovirus, the expression of the insulin gene was significantly decreased. Additionally, it was confirmed that when human insulin gene transcription promoter and TRB3 are cotransfected in INS-1 cells, insulin gene transcription promoter activation is significantly decreased. These results suggest that TRB3 suppresses the expression of the insulin gene in at the transcriptional level. However, in order to confirm that ER stress-induced TRB3 expression is directly related to insulin gene suppression, it is necessary to elucidate whether suppression of TRB3 expression prevents ER stress-induced suppression of insulin gene expression.

There are various transcription factors that regulate insulin gene expression at the insulin transcription promoter site. Among them, PDX-1 and MafA are important transcription factors that are involved in expression of insulin gene and are related to pancreatic beta cell function [31-36]. PDX-1 is an essential transcription factor for maintaining pancreatic development and beta cell differentiation and function [37]. MafA controls insulin transcription and also plays an important role in beta cell function [35]. Thus, inhibition of PDX-1 and MafA expression is closely related to beta cell dysfunction and diabetes [38]. In this experiment, we studied the effect that TRB3 on the expression of MafA and PDX-1. We confirmed that when TRB3 is overexpressed with adenovirus in INS-1 cells, MafA gene expression is not affected, while PDX-1 mRNA expression decreases. In order to verify whether the decrease in PDX-1 expression is directly related to the TRB3-dependent insulin gene suppression, additional studies are necessary to determine if the overexpression of PDX-1 offsets the effect of TRB3, which reduces insulin expression.

In summary, factors causing ER stress in the INS-1 cells increase TRB3 expression. When TRB3 is overexpressed, insulin gene expression is suppressed. The TRB3-induced decrease in insulin gene expression is associated with the TRB3-induced suppression of PDX-1.

ACKNOWLEDGMENT

This work was supported by Korean Diabetes Association Grant (2008) and the Korea Research Foundation Grant funded by the Korean Government (MOEHRD, Basic Research Promotion Fund) (KRF-2008-331-E00121).

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