Heat shock protein induction in rat pancreatic islets by recombinant human interleukin 1β

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Summary. Interleukin 1 β , potentiated by tumour necrosis factor α , is cytotoxic to pancreatic Beta cells in vitro. We have hypothesized that interleukin 1ß induces oxygen free radicals in Beta cells. Since cytotoxicity induced by free radicals and by heat may activate the same cellular repair mechanism (the heat shock response), the aim of this study was to investigate the pattern of protein synthesis in isolated islets after exposure to interleukin 1 β (150 pg/ml, 24 h), tumour necrosis factor α (50 ng/ml, 24 h), heat shock (43 °C, 30 min) and H₂O₂ (0.1 mmol/l, 20 min). By polyacrylamide gel electrophoresis, autoradiography, Western-blot analysis and partial peptide mapping of ³⁵S-methionine labelled islets, interleukin 1β was found to induce a 73 kilodalton protein belonging to the heat shock protein family heat shock protein 70, a heat shock protein 90, and haem oxygenase. A minor induction of heat shock protein 73 and haem oxygenase was seen after H_2O_2 . Interleukin 1ß did not induce heat shock proteins in rat thyroid cells, rat mesangial cells or in human monocytes. Tumour necrosis factor α did not induce selective protein synthesis. Pre-exposure of islets to heat, tumour necrosis factor α , or H₂O₂ did not prevent the impairment of glucose-stimulated insulin release seen after 24 h of interleukin 1 β exposure. The data are compatible with free radical induction by interleukin 1 β . However, the heat shock response is not specific for oxidative injury, and previous studies have shown discrepant effects as to a protective effect of free radical scavengers against interleukin 1 β -mediated beta-cytotoxicity. Thus, a role for free radicals in this context is not definitely proven.

Key words: Interleukin 1β , islets of Langerhans, heat shock proteins, haem oxygenase, free radicals, Type 1 (insulin-dependent) diabetes mellitus.

The cytokine interleukin 1 β (IL-1), mainly produced by the monocyte/macrophage cell lineage, is cytotoxic to Beta cells in isolated pancreatic islets [1, 2]. The long-term effects of IL-1 involve decreased islet insulin biosynthesis and release, decreased glucose oxidation, oxygen uptake, protein synthesis and islet content of insulin and DNA [2]. IL-1-exposed islets present signs of Beta-cell necrosis such as condensation of chromatin, vacuolization of the endoplasmic reticulum and cytolysis [3, 4]. Tumour necrosis factor α (TNF) potentiates some of the IL-1 effects, but has no effect on Beta cells in itself [5, 6]. Since the first cells to appear in the insulitis process are monocytic cells [7], IL-1 may be responsible for the initial Beta-cell damage eventually resulting in Type 1 (insulin-dependent) diabetes mellitus [8].

Why IL-1 is Beta-cell cytotoxic is not clear. We hypothesized that Beta-cell death was caused by oxidative injury due to IL-1-mediated induction of oxygen free radicals [8]. Eukaryotic and prokaryotic cells possess a stress-response mechanism [9], which is activated under several

stress conditions, e.g. exposure of cells to heat, oxygen free radicals, heavy metals, and ultraviolet radiation [9]. A hallmark of this stress response is a decrease of cellular protein synthesis, coinciding with the induction of a specific set of proteins called stress proteins or heat shock proteins (HSPs) [9]. For this cellular response pattern the term "heat shock response" is generally used. HSPs are believed to play a role in repair and protection against cellular stress. HSPs (induced by non-lethal heat shock) protect a TNFsensitive tumour cell line from TNF cytotoxicity [10], and, in turn, oxidative stress can under certain circumstances induce thermotolerance [11]. Moreover, TNF raises a defense against oxidative stress by way of the induction of manganous superoxide dismutase (MnSOD) in several human epithelial cell types [12], and we recently found that IL-1 induces new protein synthesis in ratislets [13]. Against this background we studied the pattern of protein synthesis in islet cells after exposure to IL-1 and/or TNF. The effect was compared to that of heat shock and oxidative stress (H₂O₂). In addition, we investigated whether pre-exposure

Table 1. Effects of cytokines, heat shock and H_2O_2 on islet function and islet contents of insulin and DNA

	IRI release to medium $ng/islet \times 24 h$	IRI secretion ng/islet × 2 h	IRI content ng/islet	DNA content ng/islet	IRI/DNA ng/ng
Culture condition		······································			······································
rIL-1β150 pg/ml, 24 h	54.3 ± 4.62^{a}	9.54±1.17°	82.1 ± 4.52^{a}	102 ± 5.52	$82.1 + 4.94^{\circ}$
TNF 50 ng/ml, 24 h	127 ± 7.82	88.4 ± 11.5	83.1 ± 3.29	102 ± 4.65	83.3 ± 6.49
H ₂ O ₂ 0.1 mmol/l, 20 min	_	108 ± 17.2	109 ± 18.3	97.5 ± 12.0	127 ± 35.7
heat shock 43°C, 30 min	-	79.7 ± 12.5	97.2 ± 8.95	100 ± 8.97	96.8 ± 8.70

Islets were cultured in RPMI 1640 under conditions as noted above. After culture, duplicate groups of 50 islets were incubated in Krebs-Ringer bicarbonate buffer (KRB) [2] containing 0.2% bovine serum albumin and 1.67 mmol/l of glucose. After 1 h of incubation at 37 °C, the buffer was removed and replaced by KRB containing 16.7 mmol/l glucose. Islets were finally incubated at 37 °C for 2 h. Data originate from different series of experiments, individually

Table 2. Islet cell synthesis of 32 kilodalton (kD) and 72/73 kD proteins during a 4 h labelling period (³⁵S-methionine) after exposure to cytokines and heat shock

Culture condition	32 kD protein	72/73 kD protein	п
	(% of total protein synthesis)		
Control, 24 h	0.5 ± 0.2	0.8 ± 0.5	23
IL-1β150 pg/ml, 24 h	7.1 ± 0.6^{b}	7.4 ± 0.6^{b}	23
Heat shock, 43 °C, 30 min	1.2 ± 0.4	14.1 ± 2.4^{a}	6
Heat shock before IL-1β	$8.7 \pm 1.1^{\circ}$	2.5 ± 1.9	6
TNFa 50 ng/ml, 24 h	0.6 ± 0.3	0.4 ± 0.4	8
$TNF\alpha + IL-1\beta$	$7.7 \pm 1.2^{\circ}$	7.3 ± 0.3^{a}	5
TNFα before IL-1β	7.3 ± 0.1^{a}	7.3 ± 0.6^{a}	7
H ₂ O ₂ 0.1 mmol/l, 20 min	1.8 ± 0.9	2.0 ± 1.0	9
H_2O_2 before IL-1 β	$6.8 \pm 0.5^{\circ}$	6.4 ± 0.5^{a}	5

Quantification of protein synthesis was performed by densitometry of autoradiograms of labelled islet cell proteins. Data (mean \pm SEM) express the percentage of total islet cell protein synthesis. *n* is number of experiments. ^a and ^b denote p < 0.05 and p < 0.01, respectively, compared with control culture.

 $IL-1\beta$ = interleukin-1 β , TNF = tumour necrosis factor

of islets to heat, oxidative stress (H_2O_2) or TNF would modulate the effects of IL-1 on Beta cells.

Materials and methods

Reagents

Recombinant human interleukin 1 β (rIL-1 β) was provided by Novo Nordisk Ltd., (Bagsværd, Denmark). The specific activity was 400 U/ng [14]; Recombinant human tumour necrosis factor α (TNF) was purchased from Genzyme, (Boston, Mass., USA). The specific activity (cytotoxicity of L929 cells in the presence of Actinomycin-D) was 2×10^7 U/mg. H₂O₂ and sodium-dodecyl-hydrogensulphate (SDS) were purchased from Merck, (Darmstadt, FRG). 2-Mercapto-ethanol and Nonidet P40 were obtained from Fluka AG, (Buchs, Switzerland) and BDH Chemicals, (Poole, UK), respectively. ³⁵S-Methionine and Amplify were purchased from Amersham International (Amersham, UK). RPMI 1640 culture medium was purchased from Gibco (Paisley, Scotland).

Islet isolation and culture

Pancreatic islets from 5–7 day old outbred Wistar rats were isolated after collagenase exposure to the pancreas as previously described [15]. After pre-culture for 5 days in RPMI containing 10% newborn

performed in a paired design and are therefore presented as percentages (mean \pm SEM) of corresponding control islets cultured in RPMI 1640 at 37 °C.

n = 6 for all experiments; ^a denotes p < 0.05 vs 100% (Wilcoxon's matched pairs test).

IRI = immunoreactive insulin, TNF = tumour necrosis factor, rIL-1 β = recombinant human interleukin 1 β

calf serum islets were cultured in RPMI 1640 containing 11 mmol/l glucose plus 0.5% normal human serum. The following protocols for islet culture were followed: rIL-1 β 150 pg/ml, 24 h; TNF 50 ng/ml, 24 h; Heat shock (43 °C), 30 min; H₂O₂ 0.1 mmol/l, 20 min. To study the influence of islet pre-exposure to TNF and H₂O₂ on the rIL-1 β effect, islets were carefully washed twice after TNF or H₂O₂ exposure before being transferred to rIL-1 β -containing medium. In experiments studying the rIL-1 β effects after heat shock, islets were transferred to rIL-1 β containing mediately after heat shock. The reason for exposing islets to rIL-1 β immediately after the heat shock procedure was that the initial effect of rIL-1 β is stimulatory rather than cytotoxic to islet cells [8]. Thus, islets were considered to be sufficiently able to synthesize HSPs after heat shock during rIL-1 β exposure.

After culture, duplicate groups of 50 islets were washed once and incubated as described [2] in Krebs-Ringer bicarbonate buffer (KRB) containing 0.2% bovine serum albumin and 1.67 mmol/l of glucose. After 1 h of incubation at 37°C, the buffer was removed and replaced by KRB containing 16.7 mmol/l glucose. Islets were finally incubated at 37°C for 2 h. Insulin in KRB containing 16.7 mmol/l glucose was measured by RIA [16]. Subsequently, islets were washed once and suspended in distilled water. After ultrasonication, insulin and DNA [17] contents were measured in the aqueous islet cell homogenates to determine islet insulin and DNA contents.

Preparation of rat thyroid cells

The thyroid gland was excised from normal adult Wistar rats. After cutting the gland into pieces (diameter 1–2 mm), the tissue was exposed to collagenase (1 mg/ml) and dispase (1.2 U/ml) and shaken rapidly for 1 h at 37 °C. The subsequent washing procedure (four times) and cell culture were performed in media identical to those used for islets. Cells were tested for identity and function by measurement of the medium accumulation of thyroxin (data not shown). To maintain thyroid cell activity the following agents (all purchased from Sigma Chemical Co., St. Louis, Mo., USA) were added to the culture medium: hydrocortisone (10^{-1} mol/l), transferrin (5 µg/ml), somatostatin (10 ng/ml), glycyl-histidyl-lysin-acetate (20 ng/ml), TSH (1 mU/ml) and insulin (10 µg/ml). The cells were pre-cultured for six days before exposure to 150 pg/ml of rIL-1 β for one or three days.

SDS polyacrylamide gel electrophoresis (SDS-PAGE), autoradiography, Western-blot analysis, and partial peptide mapping

Before islet protein separation, groups of 125 islets were washed twice in methionine-free RPMI 1640 and kept at 37 °C for 15 min in methionine-free RPMI 1640. Thereafter, islets were transferred to and kept at 37 °C for 4 h in 0.5 ml of RPMI 1640 with the addition of



Fig. 1 a, b. SDS-PAGE (10% gels) and autoradiography of lysed, ³⁵Smethionine-labelled, isolated and pre-cultured rat pancreatic islets [2]. Before labelling for 4 h, islets were kept under control culture conditions (a and b, *lane 1*) for 24 h or were exposed to recombinant human interleukin 1 β (rIL-1 β) tumour necrosis factor α (TNF α), heat shock or H₂O₂ according to the following protocol: rIL-1 β

0.001 mCi ³⁵S-methionine per islet. SDS-PAGE was performed according to Laemmli [18]: islets were washed in TRIS-buffer, and immediately afterwards exposed to boiling 2-Mercaptoethanol (1%) and SDS (0.3%). Nonidet P40 (1.5%) dissolved in glycin-buffer (pH 8.8) was finally added. After addition of SDS sample buffer, aliquots of islet proteins were separated on 0.5 mm 10% SDS-PAGE gels. After gel exposure for 20 min to Amplify, gels were dried, and autoradiography and densitometry were performed. Thyroid cells were treated identically to islets.

Western-blot analysis was performed with a mouse monoclonal anti-human HSP70 [19] and revealed by Auroprobe BLplus Immunogold reagent and intense silver enhancement system (Janssen Pharmaceutica, Beerse, Belgium) or by peroxidase conjugate mouse IgG (Sigma).

Partial peptide mapping was performed as previously described [20]; in short, an excised gel slice, containing the protein in question, was exposed to N-chlorosuccinimide, which selectively cleaves tryptophanyl peptide bonds in proteins. The protein product was subsequently re-analysed by SDS-PAGE and autoradiography.

Statistical analysis

Data are means \pm SEM. For each experimental condition, islets used for functional studies and protein synthesis analysis originated from the same batch of islets. Data originating from two series of experiments presented in Table 3 show some differences as to the islet

150 pg/ml, 24 h; TNF 50 ng/ml, 24 h; Heat shock: 43 °C, 30 min. H₂O₂: 0.1 mmol/l, 20 min. **a** and **b**, *lane 2*: rIL-1 β ; **a**, *lane 3*: heat shock; **a**, *lane 4*: heat shock before rIL-1 β ; **a**, *lane 5*: TNF α ; **a**, *lane 6*: rIL-1 β + TNF α ; **b**, *lane 3*: TNF α before 24 h control culture; **b**, *lane 4*: TNF α before rIL-1 β ; **b**, *lane 5*: H₂O₂ before 24 h control culture; **b**, *lane 6*: H₂O₂ before rIL-1 β ; **b**, *lane 7*: H₂O₂

content of insulin and DNA, presumably due to variability among islet preparations; therefore, all conclusions in this study originate from experiments performed in a paired design. Wilcoxon's matched pairs test was used for statistical analysis with p < 0.05 as the level of significance.

Results

Exposure for 24 h to 150 pg/ml rIL-1 β caused approximately 50% reduction in insulin release to the medium during rIL-1 β exposure and approximately 90% reduction of insulin release during a subsequent glucose challenge (exposure to KRB with 16.7 mmol/l of glucose) (Table 1). Islet insulin content was slightly, but significantly, decreased; islet DNA content was not influenced by rIL-1 β (Table 1). TNF, H₂O₂, and heat shock did not significantly influence any of the above parameters (Table 1).

After islet exposure to rIL-1 β , the synthesis of three proteins of 32, 73, and 90 kilodalton (kD) were induced or enhanced (Fig. 1, Table 2). By Western-blot analysis the 73 kD protein was identified as a protein belonging to the HSP70 family (Fig. 2). By exposing the 32 kD protein to N-

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Fig.2 a, b. Rat islets (a, *lane 1*; b, *lane 1 and 2*) were exposed to recombinant human interleukin 1β (a, *lane 1*; b, *lane 2*) or maintained under control conditions (b, *lane 1*) as described, then lysed and proteins resolved on 10% one dimensional polyacrylamide gels. Western blotting was performed with a mouse monoclonal anti-human heat shock protein (HSP)70[29] and revealed by Auroprobe BLplus Immunogold reagent and intense silver enhancement system (a) or by peroxidase conjugate mouse IgG (b). *Lane 2*: a rat mesangial cells, heat shock; *lane 3*: a and b U937 cells (a human premonocytic line), heat shock; *lane 4*: a U937 cells, control. Note that the monoclonal antihuman HSP70 did not react at all with control U937 (a, *lane 4*) cells, whereas it did with control islet cells (b, *lane 1*). The figures presented are representative of four separate experiments

chloro-succinimide, three distinct cleavage products of approximately 26,24, and 12 kD were obtained (Fig. 3), a pattern characteristic of haem oxygenase [20, 21]. The molecular weight of the rIL-1 β induced 90 kD protein was identical to that of a 90 kD protein induced after heat shock before labelling. Additionally, heat shock induced HSPs of 32, 65, 72, 73, and 110 kD (Fig. 1). After exposure of islets to

0.1 mmol/l of H_2O_2 for 20 min before labelling, the synthesis of the 32 and 73 kD was slightly enhanced (Fig. 1, Table 2). TNF did not induce selective protein synthesis (Fig. 1, Table 2). Pre-exposure of islets to heat suppressed the rIL-1 β -mediated induction of the 73 and 90 kD proteins, whereas the induction of the 32 kD protein was unchanged (Fig. 1, Table 2). Pre-exposure of islets to TNF or H_2O_2 did not influence the rIL-1 β -mediated induction of protein synthesis. Pre-exposure of islets to heat shock, TNF or H_2O_2 neither significantly prevented nor potentiated the rIL-1 β effect on parameters of Beta-cell function and islet content of insulin and DNA (Table 3).

rIL-1 β did not induce selective protein synthesis in thy-rocytes (Fig. 4).

Discussion

The present study demonstrated that rIL-1ß activates a heat shock response in rat islets, under conditions where rIL-1 β is strongly inhibitory to glucose-stimulated insulin release. The three rIL-1\beta-induced stress proteins had molecular weights of 90, 73 and 32 kD. Our preliminary finding of rIL-1 β also inducing a 80 kD protein [13] was not confirmed in this study, presumably due to a former misinterpretation of the molecular weight of an rIL-1ß induced 90 kD protein. The 73 kD protein was identified as a protein belonging to the HSP70 family, the 32 kD as haem oxygenase, and the 90 kD protein had the same molecular weight as that of one of the proteins induced by heat shock. Haem oxygenase is induced by oxidative stress, but also by ultraviolet irradiation, and sodium arsenite in human cells [20, 21]. Further, haem oxygenase is a heat shock protein in the rat [22]. The heat shock response induced by rIL-1 β (and H₂O₂) had distinct characteristics compared to that induced by heat: The total amount of stress protein synthesis was smaller after rIL-1ß (and H_2O_2), the HSPs 65, 72, and 110 were not detected after exposure to rIL-1 β , whereas rIL-1 β caused a more

Table 3. Influence of pre-exposure of rat islets to heat shock, tumour necrosis factor (TNF) and H_2O_2 on recombinant human interleukin-1 β (rIL-1 β) effects on rat islets

	IRI release to medium $ng/islet \times 24 h$	IRI secretion ng/islet ×2 h	IRI content ng/islet	DNA content ng/islet	IRI/DNA ng/ng
Culture condition					
RPMI 1640	20.3 ± 1.93	2.84 ± 0.29	13.3 ± 1.73	9.66 ± 0.80	1.43 ± 0.21
rIL-1β 150 pg/ml	$10.5\pm0.84^{\rm c}$	$0.30\pm0.09^{\rm c}$	10.7 ± 1.23°	9.55 ± 0.66	$1.16 \pm 0.14^{\circ}$
Heat shock (43 °C, 30 min) before rIL-1β 150 pg/ml	$11.4 \pm 0.99^{\circ}$	$0.45\pm0.13^{\rm c}$	11.0 ± 1.92^{a}	10.9 ± 1.30	1.07 ± 0.14^{a}
RPMI 1640	19.1 ± 2.56	3.67 ± 0.47	23.3 ± 0.87	14.0 ± 1.67	1.83 ± 0.19
rIL-1β150 pg/ml	$11.0 \pm 1.35^{\circ}$	$0.28\pm0.05^{\circ}$	$18.1 \pm 1.32^{\rm b}$	12.0 ± 1.36	1.57 ± 0.17^{a}
TNF 50 ng/ml for 24 h before rIL-1β 150 pg/ml	$6.61 \pm 0.93^{\circ}$	$0.61\pm0.18^{\rm c}$	20.8 ± 1.67	14.2 ± 1.21	1.53 ± 0.14
H ₂ O ₂ 0.1 mmol/l 20 min before rIL-1β 150 pg/ml	$9.38\pm1.39^{\text{b}}$	$0.24\pm0.06^{\text{a}}$	16.4 ± 0.98^{a}	9.72 ± 1.41	1.78 ± 0.22

Data are given for two sets of experiments. Islets were cultured for 24 h in RPMI 1640 with or without the addition of rIL-1 β preceded by heat shock or exposure to H₂O₂ and TNF as indicated above. Islets were subsequently treated as described under Table 1. Data are

means \pm SEM, n = 6-8. ^{a, b, c} denote p < 0.05, 0.02, 0.01, respectively, for a chance difference vs control islets cultured without rIL-1 β (Wilcoxon's matched pairs test). IRI = immunoreactive insulin



Fig. 3. Partial peptide mapping of recombinant human interleukin1 β (rIL-1 β) induced 32 kilodalton (kD) protein. *Lane A* represents proteins with known molecular weights (marker), *lane B* (representative of two separate experiments) represents rIL-1 β induced, N-chlorosuccinimide exposed rIL-1 β induced 32 kD protein

Fig. 4. SDS-PAGE (10% gels) and autoradiography of lysed, ³⁵S-methionine-labelled rat thyrocytes (*lane* 1–3) and islets (*lane* 4–6) cultured for 24 h with and without 150 pg/ml of rIL-1β. *Lane* 1: control culture; *lane* 2: rIL-1β; *lane* 3: heat shock; *lane* 4: control culture; *lane* 5: rIL-1β; *lane* 6: heat shock

pronounced induction of haem oxygenase. These different patterns of the heat shock response are in accordance with previous observations of oxidative injury inducing a heat shock response different from that induced by heat [23]. Although not specific for oxidative injury, the islet cell heat shock response induced by rIL-1 β may reflect the induction of free radicals. HSP induction by rIL-1 β was not seen in rat thyrocytes, rat mesangial cells (not shown) and normal human monocytes (not shown), all cells which are IL-1-responsive, but not susceptible to IL-1-mediated cytotoxicity. This is in agreement with previous studies showing that neither IL-1 nor TNF induced HSP synthesis in fibroblasts, Hep G2 and U937 cells [24], suggesting the activation of a unique response to IL-1 in islet cells.

At the chosen experimental conditions, pre-exposure of islets to neither heat, H₂O₂ nor TNF caused protection against or potentiation of the rIL-1ß effect on glucosestimulated insulin release (Table 2), although heat-induced HSPs were still present (demonstrable by Westernblot) in islets 24 h after the heat shock procedure (not shown). However, when is lets were pre-exposed to heat before the 24 h period of rIL-1 β exposure, less induction of the 73 and 90 kD protein was seen after rIL-1 β exposure. This suggests that the presence of heat-induced high molecular weight HSPs may generate a negative feed-back on rIL-1B-mediated HSP synthesis. It could also be considered that the immediate addition of rIL-1 β after heat shock would prevent the HSP synthesis after heat shock. However, HSP induction by heat during the first 4 h after heat shock was not influenced by the presence of rIL-1 β (not shown). Since heat shock pre-exposure suppressed the rIL-1 β -induced high molecular weight HSP synthesis, but not the inhibitory effect on insulin secretion, the enhanced synthesis of these HSPs might not be causally related to the toxic effect of rIL-1 β . Further, that HSPs did not protect against rIL-1 β may suggest that there is no causal relationship between the stress proteins and the cytotoxic rIL-1 β effect, thus maybe speaking against a role for oxygen free radicals.

The oxygen free radical species $O_2 - and H_2O_2$ are produced during oxidative metabolism in the mitochondria during electron transport [25]. We recently demonstrated that rIL-1 β increases the cytosolic concentration of free sodium in islet cells [26], possibly by the activation of the plasma membrane Na⁺/H⁺ exchange, which, in human neutrophils (however not yet demonstrated in islet cells), leads to intracellular alkalinization and increased O₂⁻ generation [27]. In analogy, rIL-1 β increases oxidative metabolism in short-term rIL-1 β -exposed islet cells [28], and the likely point of rIL-1\beta-mediated cytotoxicity (longterm rIL-1ß exposure) is the mitochondria [4]. Protection from oxygen free radical toxicity requires antioxidants such as superoxide dismutase (SOD), catalase, and glutathione peroxidase (GPO) [29, 30]. Compared to other cell types, islet cells contain small amounts of the oxygen free radical scavengers MnSOD and GPO [31], of which MnSOD is predominantly located to the mitochondria [29, 30]. Accordingly, islets were found to be most sensitive among different tissues examined for sensitivity to peroxide toxicity [32]. Thus, if rIL-1 β induces oxygen free

radicals in Beta cells, the Beta-cell cytotoxicity of rIL-1 β may reflect insufficient capability of these cells to produce protective proteins that are normally constitutively expressed or inducible. Consequently, Beta cells exposed to rIL-1 β may mount an alternative response (HSP and haem oxygenase synthesis) instead of the normal activation of MnSOD and glutathione peroxidase. In this case, Beta-cell survival will be highly dependent on an efficient heat shock response.

Other approaches have been taken as to the investigation of the hypothesis of oxygen free radicals being responsible for cytokine-mediated Beta-cell injury. Studies by others [4, 33–35] on the effect of scavengers of free radicals have given conflicting results, may be due to differences in the type and concentration of cytokines used, islet cell exposure time, and the islet cell preparation used.

Since proteins of the HSP70 family are considered important for cellular defense and repair, HSPs may be important determinants for Beta-cell destruction in Type 1 diabetes. In this context it is of interest that genes coding for the major inducible HSP70 in man and rat are located among the genes of the major histocompatibility complex (MHC) [36, 37]. A strong association between certain MHC genes and the susceptibility to develop Type 1 diabetes exists [38]. Thus, an HSP70 gene polymorphism [39] might be related to differences in Beta-cell capability of defense against IL-1, and might, through linkage disequilibrium between HSP70 genes and MHC-related Type 1 diabetes susceptibility genes like DQw8 [40] and the TNF β 10.5 kilobase allele [39], quantitatively contribute to the susceptibility to Type 1 diabetes.

In conclusion, the induction of HSPs in islet cells by rIL-1 β is compatible with an oxidative injury to islet cells. However, since the heat shock response is not specific for oxidative injury and since free radical scavengers have not been unequivocally shown to protect against rIL-1 β , the role for oxygen free radicals in beta-cytotoxicity by rIL-1 β is still not definitely proven.

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