

Antioxidants, NF κ B Activation, and Diabetogenesis (44445)

EMILY HO AND TAMMY M. BRAY¹

Department of Human Nutrition, The Ohio State University, Columbus, Ohio 43210-1295

Abstract. Although many risk factors can trigger the development of insulin-dependent diabetes (IDDM), it is likely that reactive oxygen species (ROS) play a central role in β -cell death and disease progression. This review will focus on the role of antioxidant defense systems in the susceptibility to IDDM and on ROS as cellular messengers that regulate the expression of genes leading to β -cell death. Accumulating evidence indicates that increased antioxidant defense systems reduce the susceptibility to IDDM in animal models or in human study. It is suggested that pancreas-specific ROS productions play a critical role in signaling the cellular autoimmune/inflammatory response by activating the transcription factor, NF κ B. Various diabetogenic factors may lead to an increase in ROS production, which activates the redox-sensitive NF κ B. This may be the initial event for the expression of cytokines and chemotactic agents involved in the autoimmune/inflammatory response. It is believed that this cascade results in a cyclic amplification of ROS and eventually leads to apoptosis and/or necrosis of β cells. The specificity of antioxidants to inhibit NF κ B activation and the hyperglycemic response emphasizes the importance of selectivity in antioxidant therapy. Research in this area will contribute significantly to our understanding of the cellular and mechanistic role of ROS in the etiology of IDDM and will lead to the development of better prevention strategies. [P.S.E.B.M. 1999, Vol 222]

Type I diabetes or insulin-dependent diabetes mellitus (IDDM) is a complex, multifactorial disease involving severe destruction of the insulin-producing β cells. Occurring predominantly in young children, IDDM has an acute onset and often progresses to numerous secondary health complications. Multiple risk factors such as genetics, environmental stresses, viral infection, and diet can predispose an individual to IDDM. Regardless of the triggering factors, the disease is always characterized by a progressive destruction of the insulin-producing β cells in the pancreas. Although the sequence of events at the cellular level is still unknown, a growing body of research suggests that reactive oxygen species (ROS) are involved in β cell destruction (1-3). The finding that ROS production is associated with the same risk factors that cause susceptibility to IDDM increases the likelihood that ROS play an important role in the pathogenesis of IDDM.

The link between multiple trigger factors for IDDM and the production of ROS is illustrated as a unified pathway in our proposed model (Fig. 1). Our model allows us to predict that certain risk factors will result in increased ROS production, which in turn will lead to the destruction of β cells. Involvement of free radicals in the etiology of IDDM and in the pathophysiology of Type 2 diabetes has been extensively reviewed by others (4-11). This review will focus on the role of antioxidant defense systems in the susceptibility to IDDM and on ROS as cellular messengers that regulate the expression of genes leading to β cell death.

ROS and the Development of IDDM

Growing evidence points to the involvement of ROS in the pathogenesis of IDDM. As illustrated in Figure 1, several known diabetogenic factors can be linked to ROS generation. For example, although IDDM has traditionally been described as an autoimmune disorder, the autoimmune response may be mediated by ROS. Insulinitis, a prediabetic stage in which islets are infiltrated with immune cells, is accompanied by the release of cytotoxic ROS, such as superoxide ($O_2^{\cdot-}$) and hydrogen peroxide (H_2O_2), and reactive nitrogen species (RNS), such as nitric oxide (NO^{\cdot}) (12). It is known that certain viruses are associated with the onset of IDDM (13, 14). Viral infections, either through general in-

¹ To whom requests for reprints should be addressed at 347 Campbell Hall, Department of Human Nutrition, The Ohio State University, 1787 Neil Avenue, Columbus, OH 43210-1295. E-mail: bray.21@osu.edu

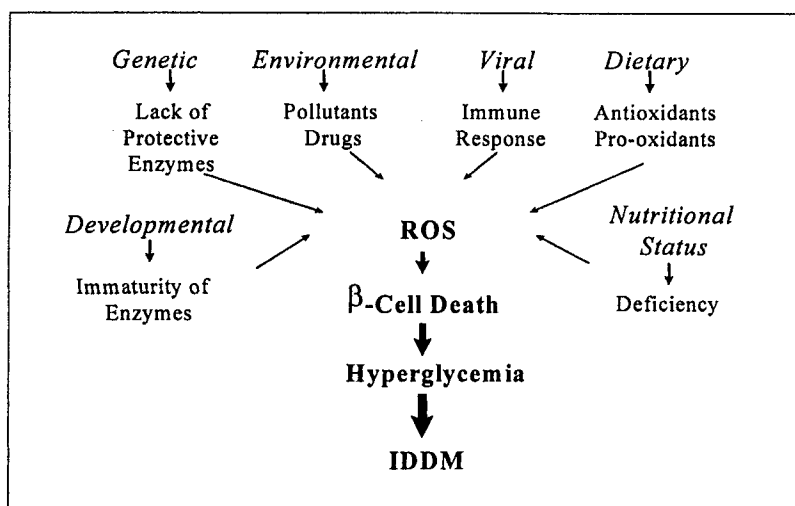


Figure 1. Linking various risk factors to ROS generation in the development of IDDM. ROS, reactive oxygen species; IDDM, insulin-dependent diabetes mellitus.

flammation or direct damage to the β cells, also initiate an immune response that culminates in the release of ROS. In addition to immunologic factors, accidental chemical induction of diabetes, particularly *via* N-nitroso compounds, is well documented in humans (15). Many investigators have demonstrated that poor nutritional status and diets high in nitrites and nitrates are correlated with an increased incidence of IDDM (5, 16, 17). Poor nutritional status produces an environment in which antioxidant defenses are often low resulting in a higher susceptibility to oxidative damage. In all these cases, the link between ROS and pathogenesis of IDDM is apparent; however, it is still unclear if ROS production directly causes β -cell death or is simply the consequence of the disease progression.

Antioxidant Status in IDDM Patients

Patients with IDDM appear to have significant defects in antioxidant protection compared to healthy, nondiabetic controls. Several studies have demonstrated a significant reduction in total antioxidant status in both plasma and serum samples from IDDM patients compared to age-matched controls (18). Diabetic children have also shown significant decreases in erythrocyte glutathione peroxidase (GPx), total glutathione, plasma α -tocopherol, and plasma β -carotene (19). This reduction in antioxidant activity is coupled with significant increases in lipid hydroperoxides, conjugated dienes, and protein carbonyls, which are markers for oxidative stress (20). This suggests that low antioxidant defenses predispose IDDM patients to enhanced oxidative stress. More recent studies have established that in the prediabetic condition, antioxidant status appears to be compromised (21). Islet cell antibodies (ICA) serve as a serological marker for risk for IDDM and have been used in population studies for the prediction of IDDM. Total plasma antioxidant status was assessed in both ICA-positive and ICA-negative, first-degree relatives of patients with IDDM. Antioxidant status was significantly lower in ICA-positive subjects compared

to ICA-negative relatives and healthy unrelated subjects. Hence, disturbances in antioxidant defenses appear even in the prediabetic state. These observations suggest that antioxidant status is another contributing risk factor in the development of IDDM.

In the past, the proposed genetic component of IDDM has focused on diabetes-susceptible HLA-haplotypes found in the major histocompatibility complex (MHC) (22, 23). Individuals who possess these susceptible alleles are more predisposed to developing the autoimmune response resulting in IDDM. However, not all diabetics possess a susceptible haplotype, and not all individuals with this haplotype go on to develop the disease (24). In fact, twin studies have only shown 30% concordance in developing diabetes (25). This suggests that both environmental and genetic factors play a role in the development of IDDM. If ROS contribute to β -cell death and dysfunction, individuals that possess relatively lower endogenous levels of key antioxidant enzymes may be more susceptible to developing IDDM.

ROS Production in Animal Models of IDDM

In animal models of IDDM, the involvement of free radicals in the development of the disease has been clearly demonstrated. For example, alloxan, a derivative of uric acid, is a known diabetogenic agent and has been used for over 50 years to induce diabetes in laboratory animals (26). Alloxan-induced diabetes was the first described chemical model for the disease, yet still little is known about its precise cytotoxic mechanism or its selectivity for the pancreatic β cells. Research has suggested that alloxan-induced β -cell damage is mediated through the generation of ROS (27, 28). During the metabolism of alloxan, it is reduced to dialuric acid. ROS species are produced during alloxan metabolism *via* auto-oxidation of dialuric acid back to alloxan. The production of $O_2^{\cdot -}$ can lead to the production of other ROS such as H_2O_2 and hydroxyl radical ($\cdot OH$) *via* Fenton reactions (29).

Another commonly used diabetogenic agent is the N-nitroso compound, streptozotocin (STZ). Several lines of research suggest that NO• production contributes to the cytotoxicity of STZ to the pancreatic β cells. Biochemical evidence for nitric oxide formation in pancreatic islets has also been demonstrated (6, 30, 31). An increase in nitrite, a stable end product of NO•, can be detected in islets treated with STZ. In STZ-induced diabetes, the incidence of hyperglycemia is decreased when NO synthase (NOS) inhibitors are used (32). In addition, transgenic mice deficient in the inducible isoform of NOS (iNOS) have also demonstrated a reduced sensitivity to STZ-induced diabetes (33), suggesting that NO• is involved in the mechanism of the disease.

In diabetes-prone animals such as the nonobese diabetes (NOD) mouse and the Biobreeding (BB) rat, islet cell infiltration of immune and inflammatory cells occurs prior to β -cell death. Both of these strains are thought to mimic the autoimmune destruction of β cells seen in the human disease. Destruction of the insulin-producing β cells may result from direct exposure to free radicals produced by the immune cells (34). In addition, free radicals may be produced due to production of pro-inflammatory cytokines such as interleukin-1 β (IL-1 β), tumor necrosis factor α (TNF- α), and γ -interferon (IFN- γ), which can be cytotoxic to cells through mechanisms involving free radical production (12, 35).

Antioxidant Defense and Protection Against Diabetogenesis. ROS and RNS scavengers have been shown to prevent β -cell death induced by diabetogenic drugs such as alloxan and STZ and prevent β -cell death induced by pro-inflammatory cytokines. *In vitro* addition of scavenging agents such as SOD, catalase, hydroxyl radical scavengers, and metal chelators prior to alloxan (36–38), STZ (39–42), and cytokine exposure (43) has prevented β -cell death in isolated islets. Our lab has demonstrated *in vivo*, that overexpression of CuZnSOD in transgenic animals provides some protection against the development of alloxan- and STZ-induced diabetes (44, 45). Transgenic strains that either ubiquitously overexpress CuZnSOD (TgHS mice) or contain β cell-specific CuZnSOD overexpression (RIPSOD mice) have significantly lower fasting blood glucose levels following injection of alloxan or STZ compared with nontransgenic control mice. Supplementation with antioxidants such as N-acetylcysteine (NAC) and α -phenyl-t-butyl nitrone (PBN) also protect against the development of experimentally induced diabetes. Dietary supplementation of NAC, a glutathione precursor, effectively attenuated the hyperglycemic response and weight loss associated with alloxan-induced diabetes (46). Co-injection of PBN with multiple STZ injections similarly prevented β -cell death and attenuated hyperglycemia and weight loss in STZ-induced diabetes (47). Similarly, antioxidant supplementation is also beneficial in reducing spontaneous β -cell loss and hyperglycemia in NOD mice and BB rats. A variety of antioxidants including metal chelators, nicotinamide, SOD, and α -tocopherol have been shown to

give some protection against the development of IDDM in those diabetes-prone animals (48–54).

Pancreatic Islets Are Highly Susceptible to Oxidative Stress

Several key enzymes in ROS defense are unusually low in pancreatic islets compared with other tissues, suggesting that the islet cells are uniquely susceptible to oxidative stress-induced damage. Gene expression and activity of several key antioxidant enzymes such as CuZnSOD, MnSOD, GPx, and catalase are all markedly decreased compared with other tissues such as liver (55). Studies found that the gene expression levels of the cytoplasmic CuZnSOD and the mitochondrial MnSOD in the pancreatic islet cells were 30%–40% of levels found in liver. GPx expression was 15%, and catalase gene expression was not detectable in pancreatic islets (56). Corresponding protein and activity levels were also found to be markedly lower in the islet cells. The low levels of antioxidant enzyme gene expression may account for the exquisite sensitivity of the β cells to ROS and free radical-induced damage leading to β -cell death and IDDM. For example, the uptake of alloxan by both liver and pancreatic islets has been observed (57). However, cytotoxicity of alloxan is apparent only to pancreatic islets. Research suggests that most chemical-induced β -cell damage is mediated through the generation of ROS (2, 58, 59). If increased ROS production is a causative factor in the destruction of the β cells, why is generalized oxidative stress solely detrimental to the pancreatic islet cells and not other tissues? Some of this specificity may be attributed to the low antioxidant capacity of the pancreatic islet cells.

ROS as Signaling Molecules

What is the mechanism that causes ROS to destroy these cells? The missing piece in the pathway lies in the connection between ROS production and β -cell death. Whereas higher concentrations of ROS may result in massive, indiscriminate oxidative damage, it is possible that low concentrations of ROS are sufficient to activate specific genes and cause the inappropriate activation of autoimmune or inflammatory responses leading to β -cell death. We will present some evidence that production of ROS can activate the transcription factor NF κ B. ROS-mediated activation of NF κ B may be the key modulator in the pathway that begins with the triggering of ROS production by multiple factors and leads to the ultimate destruction of β cells and the development of IDDM.

The majority of the research has concentrated on how ROS production causes direct cellular damage by oxidizing nucleic acids, protein, and membrane lipids to induce the disease. However, increasing evidence shows that ROS also play roles as physiologically important cellular messengers. ROS appear to play a critical role in regulating the expression of genes that encode for proteins involved in inflammation, immune response, and cell death (60, 61). Excess ROS production may be involved in disease progression

through inappropriate activation of genes involved in cellular defenses, excess NO• production, and apoptosis. These observations have launched a revolution in the understanding of the role of ROS in many human inflammatory and autoimmune diseases. Although IDDM is largely considered to be an autoimmune disease, this link between ROS and cellular response is a relatively unexplored area of research, leaving a large gap in the understanding of the mechanisms that signal β -cell death.

NF κ B, a Transcription Factor Sensitive to Oxidative Stress

ROS are produced during normal cellular metabolism. However, under certain stresses, an increase in the production of oxygen free radicals may overwhelm our antioxidant defenses, resulting in oxidative stress. Several researchers have now discovered that there are several "redox sensitive" biological molecules important in cell signaling that are sensitive to low concentrations of ROS. One potential target of ROS activation is the nuclear transcription factor, NF κ B.

NF κ B was first discovered by Baltimore and Sen as B-cell specific nuclear protein that bound to a site in the immunoglobulin κ light chain gene enhancer (62). Since then, NF κ B has gained widespread attention in many fields of research. It is now known that NF κ B is present in many cell types and controls the expression of numerous gene products. NF κ B appears to play a central role in regulating immune and inflammatory responses (63). For example, many inflammatory response factors such as pro-inflammatory cytokines, chemokines, adhesion molecules, colony-stimulating factors and inflammatory enzymes (Table I) are products of genes regulated by NF κ B. Thus, dysregulation or aberrant activation of NF κ B could initiate inappropriate autoimmune and inflammatory responses.

Activation of NF κ B, Inflammatory Response, and β -Cell Death

NF κ B is usually stored in the cytosol in its inactive form bound to the inhibitory unit I κ B α , which prevents DNA binding and nuclear uptake of the factor. Degradation of I κ B α is critical for NF κ B activation. Extracellular stimuli such as ROS signal the degradation and release of the inhibitory unit I κ B α through a complex but rapid cascade of events resulting in a rapid translocation of active NF κ B to the nucleus (Fig. 2) (64–66). NF κ B-inducing agents will initiate the phosphorylation of I κ B α on its N-terminal serine residues (Ser32 and Ser36) (67, 68). Currently, the kinase cascade that directly phosphorylates I κ B α has not been clearly identified. Recently, NF κ B-inducing kinase (NIK) and I κ B α kinase (IKK) have been identified (69–72), but the precise pathway from inducing agent to phosphorylation of I κ B α still remains unknown.

This phosphorylation of I κ B α induces polyubiquitination of I κ B α at multiple sites, tagging the subunit for degradation by the 26S proteasome complex. The free

Table I. Genes Regulated by NF κ B

Class	Target genes
Cytokines	Tumor necrosis factor- α (TNF- α) TNF- β Interferon- β (IFN- β) Interleukin-1 β (IL- β) IL-2 IL-6 IL-12 IL-8
Chemokines	Gro α , β , γ RANTES Macrophage chemotactic protein-1 (MCP-1)
Adhesion molecules	ICAM-1 E-selectin V-CAM
Colony stimulating factors	Granulocyte-macrophage colony stimulating factor (GM-CSF) Granulocyte colony stimulating factor (G-CSF) Macrophage colony stimulating factor (M-CSF)
Immunoreceptors	Ig κ light chain Major histocompatibility complex (MHC) class I T-cell receptor β chain
Inflammatory enzymes	Inducible nitric oxide synthase (iNOS) Cyclooxygenase-2 (COX-2) 12-Lipoxygenase

Note. Adapted table, see references (101, 103).

NF κ B unit is now able to translocate into the nucleus and bind to consensus DNA binding sites in target genes. In the nucleus, NF κ B binds to DNA and modulates the expression of several genes, including the genes controlling inflammatory and autoimmune process (66, 73, 74). Unlike many other systems, NF κ B is already present in the cytosol; thus, no new protein synthesis is required for its activation. This unique activation system allows NF κ B to regulate immune and inflammatory processes in a rapid and very efficient manner.

NF κ B activation is responsible for both the initiation and amplification of immune and inflammatory responses in the cell. An increase in NF κ B activation is followed by an increase in the release of cytokines and other chemotactic factors involved in inflammation (75). NF κ B also up-regulates the expression of adhesion molecules critical to leukocyte migration into target tissues (76). NF κ B binding sites are also present in the HLA class I genes and other immunoreceptors involved in cell-mediated immune responses. At the same time, more ROS and other reactive species such as NO• are generated as a result of the infiltration of immune cells such as macrophages and leukocytes that produce ROS and NO• as a killing or defense mecha-

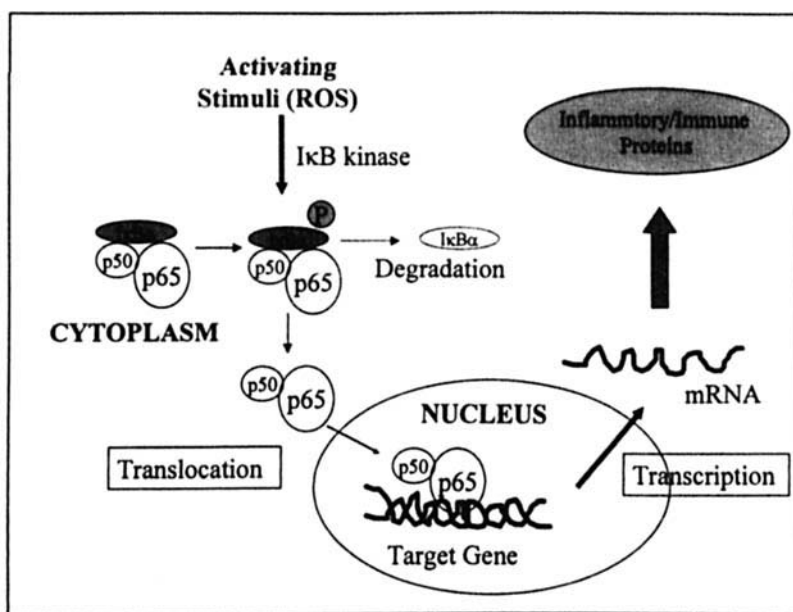


Figure 2. Schematic illustration of ROS-mediated NFκB activation. In response to extracellular inducers, such as ROS, IκB kinase (IKK) is activated and phosphorylates the IκBα subunit associated with the NFκB p50/p65 heterodimer. The phosphorylated IκBα then becomes ubiquitinated. This results in subsequent degradation of the IκBα subunit by proteasomes. Degradation of IκBα releases the p50/p65 complex, allowing the complex to translocate into the nucleus where it binds to the κB-binding sites of gene promoters and induces their transcription. Many of the target genes regulated by NFκB include several immune and inflammatory factors.

nism (63). NFκB has also been shown to stimulate the production of NO• by activating the inducible form of NO synthase (iNOS) directly (75). Many of these inflammatory cytokines in turn also activate NFκB, thus amplifying ROS production and creating a vicious cycle. In the development of IDDM, it is possible that ROS-mediated NFκB activation in the pancreas is the central signal that initiates and propa-

gates the inflammation and autoimmune processes responsible for β-cell death (Fig. 3). If this is true, agents that can inhibit this process should protect against the development of the disease.

Although IDDM is largely considered to be an autoimmune disease, this link between ROS and cellular response is a relatively unexplored area of research, leaving a large

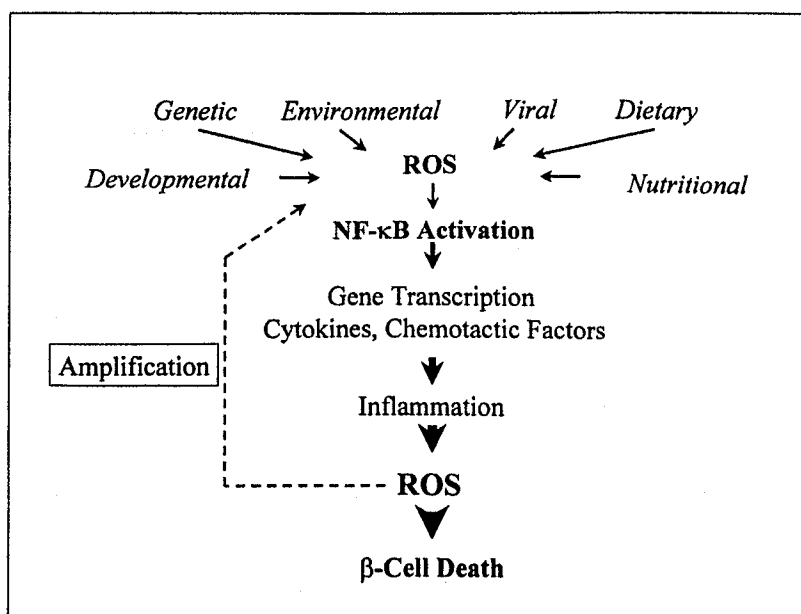


Figure 3. Initiation and amplification of the immune/inflammatory response by ROS-induced NFκB activation in β-cell death and IDDM. ROS production induced by diabetogenic factors cause the activation of NFκB. This in turn induces the transcription of autoimmune and inflammatory factors. Initiation of immune or inflammatory responses results in the production of more ROS and further activation of NFκB. This pathway acts as a positive loop, amplifying ROS production and the immune response, ultimately destroying the pancreatic β cells.

gap in the understanding of the mechanisms that signal β -cell death. More recently, NF κ B activation has been detected in rat insulinoma cell lines following exposure to cytokines such as IL-1 (77, 78). These studies confirm the possible link between NF κ B activation, inflammation, and iNOS production in a cell culture model system of diabetes.

Several lines of evidence suggest that NF κ B is also a redox-sensitive transcription factor. Many different agents can activate NF κ B, including phorbol esters, inflammatory cytokines, UV light, γ rays, viral and bacterial proteins, and lipopolysaccharide (75, 79). All of these agents produce oxidative stress. Thus, despite the diverse stimuli, ROS appear to serve as the common intracellular agents involved in the activation of NF κ B (80). Secondly, direct exposure to oxidants such as H₂O₂ activates NF κ B (81). Antioxidants have also been known to have the ability to inhibit NF κ B activation, both *in vitro* and *in vivo*. *In vitro*, NF κ B activation can be inhibited by addition of various antioxidants including lipoic acid (61), NAC (82), vitamin E derivatives (83), pyrrolidine dithiocarbamate (PTDC) (84), selenoproteins (85), and vitamin C (86). Antioxidant supplementation in animals has also been shown to inhibit NF κ B activation *in vivo*. Administration of NAC can inhibit NF κ B activation by systemic endotoxin treatment in the lung of rats (87). NAC has also been shown to inhibit NF κ B activation by the diabetogenic drug alloxan in the pancreas of CD1 mice (46). The production of ROS appears to be the key factor in initiating NF κ B activation. Thus, strategies to enhance antioxidant status may be beneficial in inhibiting inappropriate activation of NF κ B and preventing any downstream pathological events.

We have demonstrated in our laboratory, that in alloxan-induced diabetes, there appears to be a specific activation of NF κ B in the pancreas, not in the liver, within 30 min of alloxan injection. In addition, supplementation with NAC can inhibit NF κ B activation in pancreas by alloxan and concurrently provide some protection against the development of the disease. NAC-supplemented animals have significantly reduced hyperglycemia and weight loss compared to unsupplemented controls. NAC supplementation also inhibited the expression of the inducible nitric oxide synthase (iNOS), a downstream inflammatory enzyme under regulation by NF κ B. These results confirm that the role of ROS in the pathogenesis of IDDM may be far more complex than originally thought. The benefit of antioxidant therapy may not be simply to protect the β cells from oxidative damage as a result of ROS produced in the immune response. Instead antioxidant therapy may be used to stop the initiation and propagation of immune and inflammatory responses directly through inhibition of NF κ B activation. If this is true, the efficacy of antioxidant therapy will depend on the ability of antioxidants to modulate cellular response pathways that have been activated inappropriately by excess ROS.

The Future of Antioxidant Therapy in IDDM

In vitro studies using defined chemical or enzymatic sources of NO \bullet or ROS have demonstrated that islet cells are very susceptible to free radical-induced islet cell death (88–90). Furthermore, addition of antioxidants and NO \bullet inhibitors protect against this damage (38, 91). Because this inhibition of β -cell death occurs *in vitro* with antioxidants and NO \bullet inhibitors, we expect to see a protective effect *in vivo*. However, the translation of this phenomenon to *in vivo* studies is less clear. *In vivo* antioxidant intervention studies in both human and animal models have shown conflicting results (92). For example, in some studies, supplementation with NO synthase inhibitors such as N-nitro-L-arginine-methylester (NAME) has reduced the hyperglycemic response in STZ-induced diabetes (6, 93). However, other studies have shown no effect with inhibitors specific to the iNOS (94, 95). It is clear that excess production of free radical species (both ROS and RNS) is detrimental to the pancreatic islet cells. However, attempts to use antioxidant therapy in a clinical setting have shown questionable results. Currently, large scale, multinational, randomized, placebo-controlled intervention studies are underway to test the efficacy of nicotinamide to prevent IDDM. Generation of free radicals, DNA strand breaks, activation of polyADP-ribose polymerase (PARP), and depletion of NAD appear to be common events in β -cell death (96). Nicotinamide at high doses has shown to be a free radical scavenger, an inhibitor, and protector against depletion of intracellular NAD. Based on these functions, and promising *in vitro* and animal studies, nicotinamide appeared to be an excellent candidate for clinical studies in ICA-positive first-degree relatives and children at high risk for IDDM. So far, results indicate only a modest decrease or delay in diabetes development with nicotinamide (97, 98). In some cohorts, no benefit in disease progression was seen at all (99). It is possible that nicotinamide supplementation targets the repair of oxidative damage that occurs late in the diabetogenic process. A more effective therapy may need to inhibit oxidative signaling processes much earlier in the diabetogenic pathway to be effective.

An important research goal is to understand how the balance between ROS and antioxidants determines susceptibility to the disease. We hypothesize that this critical balance is defined by the ability of ROS to activate cellular responses, such as inflammation and other immune processes. Specifically, an imbalance between ROS and antioxidants (i.e., an excess of ROS or inadequacy of antioxidants) induces the activation of the transcription factor, NF κ B. This activation results in an increase in inflammatory and immune responses and leads to an amplification of ROS and NO \bullet production which, in turn, ultimately leads to the destruction of the β cells, hyperglycemia, and the development of IDDM.

Currently, inhibition of NF κ B activation to diminish the expression of pro-inflammatory and immune response

genes has been discussed as a therapeutic approach in several other immune or inflammatory-related diseases including inflammatory bowel disease, inflammatory response syndrome, septic shock, asthma, and rheumatoid arthritis (100, 101). Despite the obvious link of NF κ B to regulation of immune responses and the link between immune dysregulation and the development of IDDM, the link between NF κ B activation and diabetogenesis has still not been fully established. Strategies to limit inappropriate activation of NF κ B may prove to be a very effective approach in preventing the disease. We have focused primarily on the role of antioxidants in preventing NF κ B activation. Several other compounds also have the ability to inhibit NF κ B activation. These include proteasome inhibitors, corticosteroids, and agents that can block NIK and IKK. However, use of these compounds may have limited therapeutic effect. Proteasome degradation is critical for normal protein turnover and regulates normal cellular function and cell cycling. Thus, the use of proteasome inhibitors clinically may have serious side effects *in vivo*. Long-term glucocorticoid therapy also has limited benefit due to side effects associated with its influence on endocrine function and metabolism. Currently, agents that can block NIK and IKK selectively, and thus block I κ B α degradation, have not been identified. Phosphorylation cascades are a widespread regulatory modification throughout cellular metabolism. So far, less specific kinase inhibitors have shown conflicting results.

The key to successful antioxidant therapy in IDDM will rely both on effective targeting to the islet cells and on dose. Studies investigating the effect of NAC in animal models of acute respiratory disease (ARDS) have found that NAC is beneficial in ARDS possibly through inhibition of NF κ B activation. However, the effect of NAC only proved to be beneficial at lower supplementation levels, whereas at high doses lung injury was exacerbated (102). Thus, dose response relationships with antioxidant therapy will also need to be considered.

Summary

Although there is substantial evidence that free radicals are involved in the etiology of diabetes, the precise cellular mechanisms leading to β -cell death and the development of IDDM remains unknown. The understanding of the pathways leading to IDDM is essential before effective treatment and prevention strategies can be developed. Since IDDM is primarily a childhood disease, developing effective preventative strategies will produce considerable savings in terms of long-term health care costs, and immeasurable savings in the human anguish associated with this disease. Currently, there is little understanding of the cellular events leading to the development of diabetes. Activation of NF κ B by acute oxidative stress may be the critical signal initiating the cascade of events leading to β -cell death and IDDM. Thus, understanding these ROS-induced signal pathways in the immune and inflammatory response be-

comes essential in finding preventive treatments. Antioxidant therapy that can target this activation of NF κ B may prove to be an effective therapeutic tool in finding treatments to prevent IDDM.

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