Inflammation in Diabetic Encephalopathy is Prevented by C-Peptide

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Abstract

Encephalopathy is an increasingly recognized complication of type 1 diabetes. The underlying mechanisms are not well understood, although insulin deficiency has been implicated. The spontaneously diabetic BB/Wor-rat develops neurobehavioral deficits and neuronal cell death in hippocampus and frontal cortex, which can be prevented by insulinomimetic C-peptide. Here we examined whether contributing factors such as activation of innate immune mediators are responsive to C-peptide replacement. Seven-month diabetic BB/Wor-rats and those treated with full C-peptide replacement were compared to age-matched control rats. Hippocampi of diabetic rats showed upregulation of RAGE and NF-κB, the former being localized to proliferating astrocytes. These changes were associated with increased expression of TNF-α, IL-1β, IL-2 and IL-6 in hippocampi of diabetic rats. Full C-peptide replacement, which did not induce hyperglycemia, resulted in significant prevention of upregulation of RAGE expression, activation of NF-κB and activation of pro-inflammatory factors. In conclusion, impaired insulin activity is associated with upregulation of RAGE and pro-inflammatory factors, and these are likely to contribute to previously described oxidative and apoptotic neuronal cell death. Replacement of insulinomimetic C-peptide significantly prevents this cascade of events.

Keywords: type 1 diabetes · BB/Wor-rat · encephalopathy · inflammation · C-peptide · hippocampus · NF-κB · RAGE

Introduction

Diabetic encephalopathy is an increasingly recognized complication in type 1 diabetes [1]. It appears to be correlated with age of onset, duration of diabetes, and quality of glycemic control and leads to impaired learning abilities, memory and intelligence in affected patients [2, 3]. Experimental rodent models display impaired spatial learning and memory associated with impaired hippocampal long-term potentiation. These complications can be prevented by insulin treatment [4]. In long-term type 1 diabetic BB/Wor-rats, we previously demonstrated cognitive defects associated with apoptotic neuronal cell death in hippocampus and frontal cortex, changes that were prevented by pro-insulin C-peptide replacement [5]. Although the mechanisms underlying diabetic encephalopathy are not fully understood, the consequences of hyperglycemia and impaired insulin signaling are likely to play important pathophysiological roles [6-8]. Central insulin action has a modulating effect on the expression of other neurotrophic factors such as IGF-1 and NGF and their receptors; it inhibits oxidative stress-induced DNA damage by 8-OHdG and has significant beneficial effects on several caspases including active caspase 3 [5].
On the other hand, increased exposure of the brain to glucose and subsequent autodissociative glycosylation, oxidative-nitrosative stress and polyol-pathway products like carboxymethyl lysine (CML), pentosidine and fructose [9-13] are associated with increased accumulation of advanced glycation endproducts (AGE) and upregulation of its multi-ligand receptor RAGE, both in humans [7] and in experimental models of diabetes [8-10, 14, 15].

In this study, we examined the expression of RAGE and NF-κB and the associated activation of pro-inflammatory factors in hippocampi of chronically diabetic BB/Wor rats and the preventive effect of pro-insulin C-peptide replacement.

Research design and methods

Animals

Fourteen pre-diabetic male BB/Wor rats and seven age-matched non-diabetes prone BB/Wor rats were obtained from Biomedical Research Models (Worcester, MA). Diabetes-prone rats developed diabetes at 73 ± 4 days of age and were given small daily substitution doses (0.5-3.2 IU) of protamine zinc insulin (Blue Ridge Pharma, Greensboro, NC) to maintain a steady hyper glycemic level of approximately 25 mmol/l and to prevent ketoacidosis (Table 1). Blood glucose levels were monitored weekly. The animals had free access to rat chow and drinking water. In seven diabetic animals, rat C-peptide (75 mmol × kg⁻¹ × day⁻¹, Sigma Genosys, The Woodlands, TX) was delivered by subcutaneously implanted osmopumps (Alzet, Palo Alto, CA) from the onset of diabetes throughout the observation period. At the time of sacrifice, after 7 months of diabetes, blood samples were collected from all animal groups for assessment of blood glucose, glycated hemoglobin levels, insulin and C-peptide levels.

Behavioral testing

The Morris water maze paradigm was used for behavioral testing and performed as originally described [16]. The testing of animals started two weeks prior to sacrifice. The animals were placed in a circular pool of water measuring 2.04 m in diameter and 0.4 m in height. The water temperature was kept at 28°C. A platform was submerged 3 cm below the water, 10 cm from the pool’s edge. The pool was divided into four quadrants (Q1-Q4). The animals were trained to reach the platform from the four quadrants for two consecutive days each week for two weeks prior to final testing. On each day they were given three acquisition trials. The final test was performed three days after the last training swim. The latencies to reach the platform from the four quadrants were recorded in seconds. Rats that failed to reach the platform in 100 s were lifted out and given a latency score of 100 s. Increased latencies reflect impaired learning and spatial memory.

Tissue collection

Animals were anesthetized and the whole brain was removed and bi-sected. One hemisphere was snap-frozen in liquid nitrogen and stored at -80°C for protein isolation. The other hemisphere was fixed in buffered PBS (pH 7.4) 4% paraformaldehyde, paraffin embedded and 6 µm thick sections were prepared for immunohistochemistry.

Western blotting

Hippocampi were lysed in detergent lysis buffer (50 mmol/l Tris-HCl, pH 7.4, 150 mmol/l NaCl, 1 mmol/l EDTA, 1% Triton X-100, 1 mmol/l phenylmethylsulfonyl fluoride, 1 µg/ml leupeptin and 1 µg/ml aprotinin). Protein concentrations were measured using bicinchoninic acid protein assay reagent (Pierce, Rockford, IL). Ten to 40 µg was separated by 7.5-12% SDS-PAGE and transferred to polyvinylidine fluoride membranes (Bio-Rad, Hercules, CA). Membranes were blocked with Tween 20-Tris buffered saline containing 5% non-fat dry milk before incubation with primary antibodies. They were mouse anti-RAGE, mouse anti-NF-κB (Santa Cruz Biotechnology, Santa Cruz, CA), mouse anti-TNF-α and goat anti-IL-

<table>
<thead>
<tr>
<th>Parameter</th>
<th>C-peptide-replaced BB/Wor-rats (n = 7)</th>
<th>BB/Wor rats (n = 7)</th>
<th>Control (n = 7)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Body weight (g)</td>
<td>385.7 ± 12.0*</td>
<td>378.0 ± 6.3*</td>
<td>496.2 ± 12.8</td>
</tr>
<tr>
<td>Glucose (mmol/l)</td>
<td>24.3 ± 2.3*</td>
<td>25.2 ± 1.5*</td>
<td>5.0 ± 0.2</td>
</tr>
<tr>
<td>Glycated Hb (%)</td>
<td>11.4 ± 1.2*</td>
<td>11.3 ± 1.0*</td>
<td>3.3 ± 0.2</td>
</tr>
<tr>
<td>Serum insulin (pmol/l)</td>
<td>58.0 ± 7.4*</td>
<td>62.2 ± 5.7*</td>
<td>451.2 ± 18.0</td>
</tr>
<tr>
<td>Serum C-peptide (pmol/l)</td>
<td>731.3 ± 106.0*</td>
<td>&lt; 25.0*</td>
<td>982.0 ± 109.0</td>
</tr>
</tbody>
</table>

Legend: * p < 0.001 vs. control rats. ** p < 0.001 vs. type 1 BB/Wor rats.
β, anti-IL-2 (Sigma Aldrich, St. Louis, MO) and goat anti-IL6 (Pierce, Rockford, IL).

Antigen detection was performed using chemiluminescence (Amersham Pharmacia Biotech, Piscataway, NJ) with the appropriate horseradish peroxidase conjugated secondary antibodies. Membranes were exposed to Biomax film (Kodak, Rochester, NY) and densities were determined by Bio-Rad Fluoro-S multimeter. Protein expression was corrected for actin expression.

Immunofluorescent immunostaining

Tissue sections were blocked with 10 mg/ml BSA and incubated with a mixture of rabbit anti-GFAP antibody (1:200, Sigma, MO) and mouse anti-RAGE antibodies (1:50, Santa Cruz) at room temperature for 1 hour at 4°C overnight. Sections were then washed in PBS and incubated with a mixture of FITC conjugated anti-rabbit secondary antibody (1:50, Sigma) and rhodamine conjugated anti-mouse secondary antibody (1:50, Chemicon, Teurecula, CA) at room temperature for 1 hour. The sections were mounted with Gel Mount (Biomedia, Foster City, CA) and examined using a Leica microscope (DMLB) equipped with SPOT RT image system and V 3.0 software (Diagnostic Instruments Inc, Sterling Heights, MI).

Statistical methods

All data are presented as means ± SD. The standard statistical package SPSS, V 13.0 was used to perform the analyses. Parametric statistics and analysis of variance (ANOVA) were used and Scheffes post hoc test was applied for group comparisons.

Results

Clinical data

Seven-month diabetic rats showed significant weight loss, hyperglycemia, elevated glycated hemoglobin, insulin deficiency and non-measurable C-peptide levels (all p < 0.001; Table 1). C-peptide treatment restored C-peptide levels to normal but did not alter body weight, glucose, glycated hemoglobin, or insulin levels in diabetic rats (Table 1).

Behavioral studies

The latencies of diabetic BB/Wor-rats in the Morris water maze were significantly prolonged in the four quadrants (Figure 1). The latencies of C-peptide replaced diabetic rats were not significantly different from those of non-diabetic control rats in quadrants Q1, Q2 and Q4. Although the latency of C-peptide replaced diabetic rats in quadrant Q3 was significantly improved (p < 0.005), it remained prolonged (p < 0.01) compared to that of control rats (Figure 1).

Figure 1. Results of behavioral tests. The figure shows latencies in seconds for controls, non-treated BB/Wor-rats and C-peptide replaced diabetic BB/Wor-rats in the four quadrants of the Morris water maze. * p < 0.01, ** p < 0.001 vs. control rats. # p < 0.05, §§ p < 0.005, *** p < 0.001 vs. untreated BB/Wor-rats.

Statistical methods

All data are presented as means ± SD. The standard statistical package SPSS, V 13.0 was used to perform

Figure 2. Immunocytochemistry of hippocampal dentate gyrus in control, diabetic and diabetic C-peptide-replaced animals. Increased staining for RAGE (red) and GFAP (green) indicates activation of astrocytes and increased RAGE expression. Merged images confirm the localization of RAGE mainly to astrocytes and their processes. Diabetic rats that received full replacement of rat C-peptide (Table 1) showed prevention of RAGE expression and only minimal GFAP positivity indicative of decreased astrocyte proliferation. Magnification 250X.
Immunocytochemistry

An increased number of astrocytes in hippocampal pyramidal cell layer, dentate fascia (Figure 2) and temporal white matter as identified by glial fibrillary acidic protein (GFAP) positivity was observed in diabetic animals. Immunopositive RAGE co-localized with GFAP in astrocytes (Figure 2) was increased in hippocampal pyramidal cells as well as in white matter oligodendrocytes.

C-peptide treatment prevented proliferation of GFAP-positive astrocytes and the increased immunopositive expression of RAGE in hippocampal regions including the dentate fascia (Figure 2).

Western blotting

The expression of NF-κB increased significantly in diabetic hippocampi (p < 0.01). This pattern was associated with increased expression of RAGE, TNF-α (both p < 0.01), IL-1β (p < 0.01) IL-2 and IL-6 (both p < 0.001) (Figure 3). Full C-peptide replacement in diabetic rats prevented significantly the upregulation of NF-κB, RAGE, TNF-α and IL-1β, IL-2 and IL-6 (Figure 3).

Discussion

In this study, we found that pro-inflammatory factors were upregulated in hippocampi of diabetic BB/Wor-rats and that replacement of insulinomimetic C-peptide prevented the upregulation of RAGE and NF-κB and the downstream activation of pro-inflammatory factors such as TNF-α, IL-1β, IL-2 and IL-6. These findings were associated with the prevention of proliferating astrocytes and significant prevention of deficits in learning and spatial memory as indicated by the Morris water maze paradigm.

AGE and nonenzymatically glycated or oxidated proteins, lipids or nucleic acids that are formed under conditions of oxidative stress and hyperglycemia [17] are traditionally regarded as the key activators of the

![Figure 3. Immunoblot analyses of control, diabetic and C-peptide replaced diabetic hippocampi. NF-κB (A) was significantly upregulated in diabetic hippocampus, and this was significantly prevented by C-peptide. RAGE (B) was, as expected, significantly upregulated in diabetic rats and was mainly localized to proliferating astrocytes (cf. Figure 1). Significant prevention of RAGE upregulation by C-peptide in the presence of unaltered hyperglycemia was observed (cf. Table 1). TNF-α (C) expression was increased in diabetic rats and significantly but not fully prevented by C-peptide replacement. Additional pro-inflammatory factors IL-1β (D), IL-2 (E) and IL-6 (F) were upregulated in diabetic hippocampi, activations which were significantly prevented by C-peptide replenishment. * p < 0.05, ** p < 0.01, *** p < 0.001 vs. non-diabetic control rats. # p < 0.05, ## p < 0.01 vs. diabetic BB/Wor-rats.](image-url)
multiligand RAGE receptor [18, 19]. However, in this study, C-peptide substitution had no effect on hyperglycemia, suggesting that other mechanisms are responsible for the upregulation of RAGE. We have previously demonstrated that C-peptide replacement has a beneficial effect on ROS-induced DNA damage in hippocampus in this model [5]. We suppose that this effect may provide an alternative source of RAGE upregulation. The most likely mechanism is that impaired insulin signaling, which is corrected by C-peptide [5, 20-22], directly or indirectly influences RAGE expression.

It has been shown that RAGE activation stimulates NF-κB [19, 23], although the exact signaling mechanism is not known [13]. An important consequence of NF-κB activation, which also occurs as a consequence of impaired insulin signaling [24-27], is the upregulation of RAGE itself [19, 28], hence providing a self-perpetuating loop. The mechanism underlying such an effect is that the gene encoding RAGE contains functional binding elements for NF-κB [29].

RAGE was localized to activated astrocytes, which along with microglia play key roles in the production of inflammatory mediators. RAGE is a multiligand receptor and has also been identified as a receptor for β-amyloid and prion fibrils [19]. Indeed, β-amyloid does accumulate in brains of BB/Wor-rats [30], and this potentially contributes to increased RAGE signaling in this model. The present data suggest that impaired insulin signaling per se may play a prominent role in RAGE signaling activities. This interpretation is supported by data obtained from diabetic mice treated with intranasal insulin, which does not effect systemic hyperglycemia [8]. Upregulation of RAGE has been described in the CNS in this mouse model [15]. In our study, we found prevention of CNS insulin signaling intermediaries, cognitive decline and structural changes. In the light of previous data [5, 14], these findings suggest a role of impaired insulin signaling in RAGE regulation and in diabetic encephalopathy.

Existing evidence supports the role of RAGE in innate immune responses [13, 15, 28]. Apart from promoting NF-κB activation, ligand interaction of RAGE activates MAPK family members like JNK, p38 and ERK1/2 [13, 31], thereby fueling the apoptotic effects caused by aberration of the PI-3 kinase pathway [5]. It has been shown that interaction of RAGE with S100/calgranulin triggers macrophage activation through NF-κB resulting in the production of pro-inflammatory cytokines TNF-α and IL-1β [32].

The present findings showing an upregulation of several cytokines such as IL-2 and IL-6 is consistent with this cascade of signaling events [13, 33]. In addition to the already compromised hippocampal insulin receptor in the BB/Wor-rat [5, 34], activation of TNF-α inhibits Akt phosphorylation downstream of P13K in the insulin signaling pathway [35]. This inhibition is likely to impact on insulin’s anti-apoptotic effects. It is therefore obvious that upregulation of RAGE signaling and activation of NF-κB result in a series of sometimes self-perpetuating events contributing to activation of inflammatory events and compromising anti-apoptotic regulators eventually resulting in apoptotic cell death in diabetic CNS [5, 14, 36].

Conclusions

In summary, the present data demonstrate that full substitution of insulinomimetic C-peptide in the type 1 diabetic BB/Wor-rat corrects the upregulation of RAGE and NF-κB activation with downstream beneficial effects on pro-inflammatory factors such as TNF-α and interleukins. These findings are associated with a preventative effect on the proliferation of RAGE-positive astrocytes in hippocampus and prevention of deficits in spatial memory and learning.

We conclude that activation of innate immune responses contributes to the development of diabetic encephalopathy and that it can be effectively prevented by replacement of insulinomimetic C-peptide.

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Conflict of interest statement: The authors declare that they have no competing conflict of interests with respect to financial or other issues.

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