Effects of treadmill exercise on PI3K/AKT/GSK-3β pathway and tau protein in high-fat diet-fed rats

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INTRODUCTION

Obesity, which is defined as a state of having excessive fat tissue that is caused by overnutrition, a sedentary lifestyle, and other environmental factors, is an important risk factor for various metabolic diseases1-3. Several studies have shown that obesity is closely related to metabolic diseases such as type 2 diabetes, hypertension, and cardiac disease. Recently, obesity has additionally been linked with various neurological diseases including sleep apnea, depression, and anxiety4-6 as well as Alzheimer’s disease (AD) and other cognitive dysfunctions7.

The functions of insulin in peripheral tissues are well understood, while the role of insulin in the central nervous system, especially within the brain, is being re-evaluated. In the brain, insulin is important in the production, development, and activation of neurons; thus, it affects memory and learning in humans7. Recent findings have suggested that brain insulin dysfunction might be a cause of AD7,8. Under normal circumstances, brain insulin affects the development and activation of neurons and improves cognitive function8. However, overeating-induced obesity results in impaired brain insulin signaling, which may lead to cognitive dysfunction and ultimately neurodegenerative disease. Obesity has been found to reduce insulin/insulin-like growth factor-1 activation and increase the levels of tumor necrosis factor-α in the brain, which then increases neural inflammation9-11. In other words, oxidative stress12 and inflammatory responses are enhanced owing to food-induced obesity, and cerebrospinal insulin resistance is triggered through various pathological phenomena originating from these peripheral tissues. Results from previous studies showing that inflammatory responses in brain tissues became stronger after the extended feeding of a high-fat diet (HFD) support the close association of obesity with brain health parameters including memory and cognitive function13.

The molecular mechanisms underlying the relationship between brain insulin dysfunction and neurodegenerative disease are not yet clear. However, brain insulin dysfunction is expected to lead to increased amyloid-β production via the mitogen-activated protein kinase pathway as well as tau hyperphosphorylation via the insulin receptor (IR) and phosphoinositide 3-kinase (PI3K)/protein kinase B (Akt)/glycogen synthase kinase-3β (GSK-3β) pathways. Notably, GSK-3β is associated with the production of hyperphosphorylated tau, which is found in paired helical filaments. Hyperphosphorylated tau is located downstream of the Akt signaling pathway, and tau protein exhibits decreased activity following phosphorylation by Akt14,15; thus, an association is expected between tau hyperphosphorylation and insulin. Hence, if brain insulin resistance is a cause of AD, then detailed studies of brain insulin signaling might yield information...
regarding the alleviation of brain insulin resistance, which could lead to improvements in the prevention and treatment of AD.

Recent studies have shown increased amyloid-β deposition, tau hyperphosphorylation, and memory impairments after obesity was induced by an HFD in transgenic mouse models of AD\textsuperscript{[46-48]. Importantly, these findings suggest that an HFD accelerates AD. However, because these studies used transgenic animal models that regularly display symptoms of dementia, they were not able to evaluate the effects of obesity-induced insulin dysfunction.

Therefore, to clearly evaluate the effects of obesity on cerebral health in the present study, we used a long-term HFD (20 weeks) to induce obesity in rats; then, we investigated the effects of obesity and subsequent treadmill exercise (TE) on brain insulin signaling and tau hyperphosphorylation.

**METHODS**

**Experimental animals**

This study protocol was approved by the Animal Ethics Review Board of H-University (KNSU-IACUC-2015-08). The experimental animals were 8-week-old (w.o.) male Sprague–Dawley rats (Koatech Co., Ltd., Gyeonggi-do, Korea) that were reared to 24 w.o. in the animal rearing room at H-University (temperature, 22 ± 2°C; humidity, 50 ± 5%; light-dark cycle, 12 h). When the rats (n = 24) were 24 w.o., they were supplied with water and a high-fat feed (D12482; carbohydrates: 20%, fat: 60%, protein: 20%; Central Lab. Animal, Inc., Seoul, Korea) \textit{ad libitum} for 20 weeks to induce obesity. The rats were assigned to one of the following three groups: Normal Diet-Control (ND-Con, n = 12), HFD-Control (HFD-Con, n = 12), or HFD-TE (n = 12).

**Treadmill exercise**

The HFD-TE group underwent acclimation for 30 min/day for 4 days on a rodent treadmill (DJI2-242; Daejong Instrument Industry Co., Ltd., Seoul, Korea) that was fixed at an incline of 0%. After acclimation, the rats underwent training for 5 days per week for 8 weeks. The incline was fixed at 0%, and the load was gradually increased (first 5 min: 8 m/min; next 5 min: 11 m/min; last 20 min: 14 m/min). The TE protocol used in this study was identical to the moderate-intensity exercise program presented by Kang et al.\textsuperscript{[13].

**Passive avoidance task**

A passive avoidance task was performed before and after TE to evaluate cognitive function. The passive avoidance task was conducted in a light chamber (18 × 18 × 25 cm), which was lit by a bright white light, and a dark chamber (18 × 18 × 25 cm), which was completely dark. The floor of the dark chamber was fitted with a steel shock floor, and a 4-cm-diameter circular hole in the wall between the two chambers was covered with a guillotine door. Each rat was placed by itself in a different cage for 1 min before the experiment. The rat was then placed in the light chamber and allowed to acclimate for 10 s before the guillotine door was opened to allow the rat to explore freely. Once the rat had all four feet inside the dark chamber, the door was swiftly closed. The time taken for the rat to enter the dark chamber (latency time) was measured, and an electric shock (0.5 mA) was given for 2 s. After 5 s, the rat was removed and placed back in the rearing cage. After 72 h, the experiment was repeated using the same methods, and the latency time (up to 300 s) was recorded.

**Body weight and oral glucose tolerance test**

The bodyweights (BW) of the rats were measured twice per week throughout the study, using laboratory scales. The oral glucose tolerance test was performed 12 h after the completion of the 8 weeks of TE. When the rats were in a fasted state, a 30% glucose solution (1 g/kg) was administered orally. Thereafter, blood was collected from the tail vein five times (at 0, 30, 60, 90, and 120 min), and blood glucose was measured using a glucose meter. The blood glucose data were used to calculate the total area under the glucose response curve (AUC\textsubscript{0–120}, mg/dL/min), which was analyzed using the NCSS 2007 program (NCSS, Kaysville, UT, USA).

**Tissue sample collection**

Tissue samples were collected 24 h after the completion of the 8 weeks of TE. Anesthesia was induced in seven rats by intraperitoneal injections of a ketamine/xylazine mixture (2:1, 10 mg/kg) and brain tissue, blood, and abdominal tissue were sampled to measure protein expression. Brain tissue samples that were collected from the cerebral cortex were immediately frozen in liquid nitrogen and stored at −80°C until analysis. Blood was collected from the heart; then, serum was isolated by centrifugation for the glucose and insulin analyses. The abdomen was opened and abdominal visceral fat (AVF) samples were collected, washed in cold physiological saline, and dried with gauze before weighing. For immunohistochemical analysis, five rats were anesthetized and the thoracic cavity was opened. Phosphate-buffered saline (PBS; 50 mM) was perfused via the left ventricle for 10 min, which was followed by perfusion fixation with 4% paraformaldehyde in 100 mM PBS. After this perfusion fixation, each rat’s brain was extracted and submerged in 4% paraformaldehyde fixative for 4 h at 4°C. The fixed brain tissue was then submerged in a 30% sucrose solution for 2 days. The anatomical location (from −4.88 to −1.76 mm from the bregma) including the cortex of the fixed cerebral cortex and hippocampus was extracted using Rodent Brain Matrix (RBМ-4000C, 1 mm coronal section; ASI Instruments, Inc., Warren, MI, USA) and continuous coronal slices 40 μm in thickness were obtained using a freezing microtome (Leica, Nussloch, Germany).

**Glucose and insulin concentration analyses**

Glucose and insulin concentrations were analyzed using a glucose hexokinase kit (Bayer, Pittsburgh, PA, USA) and a rat insulin kit (Merckodia AB, Uppsala, Sweden), respectively, in accordance with the manufacturers’ instructions. The homeostatic model assessment (HOMA)-insulin resistance (IR) index was calculated using the glucose and insulin concentrations, according to the following equation: HOMA-IR = (fasting serum glucose (mmol/L) × fasting serum insulin (μU/L) / 22.5).
## Western blotting

The cerebral cortex was isolated from the previously harvested rat brain tissue, homogenized in lysis buffer, and then centrifuged at 13,000 x g for 15 min at 4°C. The supernatant was collected and the concentration of total protein was measured using the Bradford method. Protein samples (30 µg) were subjected to electrophoresis in 7% and 10% sodium dodecyl sulfate-polyacrylamide gels before being transferred to polyvinylidene difluoride membranes (GE Healthcare Bio-Sciences, Pittsburgh, PA, USA). Membranes were then blocked for 1 h at room temperature in a Tris-buffered saline with Tween® 20 (TBS-T) solution containing 3% bovine serum albumin and incubated with the relevant primary antibody (Table 1) for at least 12 h at 4°C. The next day, the membrane was washed three times for 10 min each time in TBS-T before incubation with the relevant secondary antibody for 1 h at room temperature; this was followed by three washes for 10 min each in TBS-T. Finally, the western blotting luminol reagent (sc-2048; Santa Cruz Biotechnology, Inc., Dallas, TX, USA) was added to the membrane. After incubation for 1 min to facilitate chemiluminescence, the membrane was scanned using an image analysis system (ChemiDoc™ XRS; Bio-Rad Laboratories, Inc., Hercules, CA, USA); Quantity One 1-D analysis software (Bio-Rad Laboratories, Inc.) was used to calculate the concentration of protein.

### Immunohistochemical analyses

Five animals were used in the immunohistochemical analyses, and eight slices were selected from each group. Sections were analyzed using the free-floating method. After washing the sections three times for 10 min each time in 0.01 M PBS, each slice was incubated for 8 min at 80°C in a beaker containing 0.01 M sodium citrate, followed by blocking for 40 min in 10% normal donkey serum. After blocking, each slice was incubated with the primary antibody (p-Tau, Ser202/Thr205) for 12 h overnight at 4°C, followed by three washes for 5 min each time in 0.01 M PBS. The sections were then incubated with a horseradish peroxidase-conjugated goat anti-mouse antibody for 1 h at room temperature. The samples were washed an additional three times for 5 min each in 0.01 M PBS, then incubated for 30 min in VECTASTAIN®Elite® ABC kit reagent (Vector Laboratories, Inc., Burlingame, CA, USA) at room temperature and washed a further three times for 5 min each time in 0.01 M PBS. Finally, using a DAB peroxidase substrate kit (SK-4100; Vector Laboratories), 3,3'-diaminobenzidine tetrahydrochloride was diluted to 0.02% in 0.05 M Tris buffer (pH 7.6). Next, 0.02% hydrogen peroxide was added and the samples were stained for 5 min. The stained tissue slices were mounted and analyzed using a light microscope (DM-2500, Leica Microsystems, Wetzlar, Germany).

### Data processing

The data in this study are presented as the mean ± standard deviation and were calculated using SPSS for Windows (version 18.0; IBM Corporation, Armonk, NY, USA). Differences between the groups were analyzed using one-way analysis of variance (ANOVA), and significant differences were verified using Bonferroni post-hoc testing. The level of statistical significance was set at α=.05.

### RESULTS

#### The effects of TE on the BW, AUC, HOMA-IR, and AVF/BW of the HFD-fed rats

The changes in BW that occurred during the HFD and TE periods are shown in Table 2. First, the initial BWs during the HFD period (at 24 w.o.) were not significantly different among the ND-Con, HFD-Con, and HFD-TE groups \([F(2,35)=1.147, \ p=.330]\). However, the final BWs during the HFD period (at 43 w.o.) exhibited significant differences among the groups \([F(2,35)=19.771, \ p=.001]\). The BWs in the first week of TE (at 44 w.o.) also exhibited significant differences among the three groups \([F(2,35)=21.227, \ p=.001]\); likewise, significant differences in BWs were observed among the groups after eight weeks of TE (at 51 w.o.) \([F(2,35)=20.956, \ p=.001]\). AUC results that were obtained using an oral glucose tolerance test revealed significant differences among the ND-Con, HFD-Con, and HFD-TE groups \([F(2,35)=8.728, \ p=.001]\). HOMA-IR results also showed significant differences among the three groups \([F(2,35)=24.271, \ p=.001]\). Finally, the AVF/BW ratio also exhibited significant differences among the three groups \([F(2,35)=80.823, \ p=.001]\).

#### The effects of TE on the cognitive function (working memory) of the HFD-fed rats

The results of the passive avoidance test that was used to evaluate cognitive function are shown in Figure 1. First, passive avoidance testing before TE revealed significant differences among the ND-Con, HFD-Con, and HFD-TE groups \([F(2,35)=173.363, \ p=.001]\). Moreover, passive avoidance testing after TE also showed significant differences among the three groups \([F(2,35)=363.775, \ p=.001]\).
Table 2. Comparisons of body weight, AUC, HOMA-IR and AVF/body weight

| Metabolic parameter | ND-Con | HFD-Con | HFD-TE |
|---------------------|--------|---------|--------|
| **Initial BW (g)**  | 454.9±18.87 | 464.0±23.52 | 469.4±27.85 |
| **Final BW (g)**    | 699.2±63.90* | 725.4±68.29* | 698.8±73.49* |
| AUC (g)             | 39436.25 | 54021.25 | 45137.50 |
| HOMA-IR             | 1.57±0.37 | 5.62±1.11* | 3.99±1.11* |
| AVF/body weight (%) | 5.22±0.77 | 8.88±0.49 | 7.17±0.22* |

Values are presented as means ± SD, ND: Normal diet, HFD: High fat diet, TE: treadmill exercise, HOMA-IR: homeostasis model assessment-insulin resistance, AVF: abdominal visceral fat. *Denotes statistical difference from the ND-Con group. **Denotes statistical difference from the HFD-Con group (p < 0.05).

The effects of TE on the expression of insulin signaling-related proteins in the cerebral cortex of the HFD-fed rats

We performed ANOVAs to compare the phosphorylation of insulin-signaling-related proteins (IR, PI3K, Akt, and GSK3β) (Figure 3). We observed statistically significant differences among the ND-Con, HFD-Con, and HFD-TE groups in terms of IR phosphorylation [F(2,20)=316.150, p=0.001], PI3K phosphorylation [F(2,20)=193.496, p=0.001], Akt phosphorylation [F(2,20)=509.365, p=0.001], and GSK-3β phosphorylation [F(2,20)=59.876, p=0.001].

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DISCUSSION

In the present study, we induced obesity by feeding rats a long-term (20 weeks) HFD; we then investigated the effects of TE (eight weeks) on various pathological phenomena. First, we found that TE reduced AVF and improved IR. In addition, TE restored the expression of brain insulin signaling-related proteins and alleviated the hyperphosphorylation of tau, which suggested that TE was able to prevent cognitive dysfunction. Here, we discuss these results in further detail.

The long-term administration of an HFD caused increases in BW and AVF in the experimental animals, and these changes were accompanied by a worsening of the insulin resistance marker HOMA-IR and blood glucose metabolism (represented by the AUC calculated from the oral glucose tolerance test data), as compared with the ND-Con group. Ultimately, the long-term HFD negatively impacted energy metabolism, as compared with the ND. However, following the HFD, eight weeks of TE resulted in decreases in BW and AVF, as well as improvements in the HOMA-IR and AUC. These results were consistent with those of previous studies. Therefore, our results support the contention that obesity promotes metabolic disease, whereas aerobic exercise can reduce BW via fat loss and improve insulin resistance by improving blood glucose metabolism. Our results demonstrated that diet-induced pathological energy metabolism can be prevented and ameliorated by TE.

The PI3K/Akt/GSK-3β pathway, which is a brain insulin signaling pathway downstream of IR and a tau phosphorylation pathway in AD, is thought to be involved in insulin signaling, tau hyperphosphorylation, and, by extension, the onset of AD. Beginning with IR, cell survival and metabolism are controlled by the PI3K/Akt signaling pathway. GSK-3β plays an important role in the control of tau phosphorylation, and a decrease in the activity of GSK-3β following its phosphorylation by Akt kinase results in reduced tau phosphorylation. Notably, GSK-3β targets the Ser199, Thr231, Ser396, and Ser413 phosphorylation sites in tau; moreover, it is the primary tau kinase that is involved in tau overexpression. The brain insulin signaling-related proteins IR, PI3K, Akt, and GSK-3β all exhibited decreased phosphorylation in the HD-Con group, as compared with the ND-Con group. However, TE reversed these changes, thus leading to an increased expression of these brain insulin-related pathways.
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Fig. 2. Effects of treadmill exercise on the expression of phosphorylated tau protein in HFD-fed rats. (A) Photomicrographs showing phospho-tau (Ser202/Thr205) immunoreactivity in the cerebral cortex of HFD-fed rats in each group. (B) Quantitative analysis of phospho-tau (Ser202, Thr205) protein in the cerebral cortex. The data shown for immunohistochemistry represent the means from five rat brains. (C) Representative western blots of the Ser404, Ser396, Thr231, and Ser199/202 proteins. (D) Densitometric analysis of the western blot bands, normalized to total tau. The data shown in the western blot represent the means from seven rat brains. α-Tubulin was probed as an internal control. Bonferroni’s post-hoc test was performed after ANOVA. Values are presented as mean ± SD. (Aa–c) Scale bar = 200 μm. (Ad–f) Scale bar = 100 μm.

Fig. 3. Effects of treadmill exercise on the expression of brain insulin signaling-related proteins in the cerebral cortex of HFD-fed rats. (A) The total abundances of the proteins IR, PI3K, AKT, and GSK-3β in cerebral cortex extracts were measured using western blotting and (B–E) quantitative analysis. The phosphorylation levels of IR, PI3K, AKT, and GSK-3β were normalized to total IR, PI3K, AKT, and GSK-3β. α-Tubulin was used as a control for total loading. Statistical analysis was calculated by one-way ANOVA followed by Bonferroni’s post-hoc test.

proteins. Moreover, the long-term HFD resulted in increased tau phosphorylation [(Ser202/Thr205, Ser202, Ser306, Thr231, and Ser199/202], whereas TE reduced tau phosphorylation. These results were consistent with those of previous studies that reported a positive effect of TE on tau phosphorylation. Takashima reported that tau hyperphosphorylation via GSK-3β was associated with insulin resistance in the central nervous system; this suggested, as implied by our study, that impaired brain insulin signaling could contribute to tau hyperphosphorylation. Hence, TE improves brain insulin signaling function, which results in a balanced phosphorylation and dephosphorylation of tau, thereby maintaining the neuronal skeleton and stabilizing the capillaries involved in the transport of neurotransmitters.

In the passive avoidance task used to evaluate memory impairments, the retention latency time until the rat entered the dark chamber was shorter in the HF-Con group than in the ND-Con group, which indicated memory impairments. Previous studies have shown that obesity increases peripheral insulin resistance and causes central insulin resistance by decreasing the levels of brain insulin; importantly, peripheral insulin resistance is associated with cognitive impairment. In rodents, a HFD results in cognitive and memory impairments. Thus, obesity, which can affect cognitive functions either directly or indirectly, is a major cause of cognitive impairment.

In the present study, we investigated brain insulin signaling and tau expression in experimental animals that exhibited obesity induced by a long-term HFD, rather than by genetic mutations or drug treatments. The long-term HFD resulted in pathological phenomena including brain insulin dysfunction, tau hyperphosphorylation, and memory impairments. Thus, HFD-induced obe-
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