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ORIGINAL RESEARCH

The effects of exercise training and caloric restriction on the cardiac oxytocin natriuretic peptide system in the diabetic mouse

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Background: Regular exercise training (ET) and caloric restriction (CR) are the frontline strategies in the treatment of type 2 diabetes mellitus with the aim at reducing cardiometabolic risk. ET and CR improve body weight and glycemic control, and experimental studies indicate that these paradigms afford cardioprotection. In this study, the effects of combined ET and CR on the cardioprotective oxytocin (OT)–natriuretic peptide (NP) system were determined in the db/db mouse, a model of type 2 diabetes associated with insulin resistance, hyperglycemia, and obesity. **Methods:** Five-week-old male db/db mice were assigned to the following groups: sedentary, ET, and ET + CR. Nonobese heterozygote littermates served as controls. ET was performed on a treadmill at moderate intensity, and CR was induced by reducing food intake by 30% of that consumed by sedentary db/db mice for a period of 8 weeks.

Results: After 8 weeks, only ET + CR, but not ET, slightly improved body weight compared to sedentary db/db mice. Regardless of the treatment, db/db mice remained hyperglycemic. Hearts from db/db mice demonstrated reduced expression of genes linked to the cardiac OT–NP system. In fact, compared to control mice, mRNA expression of GATA binding protein 4 (GATA4), OT receptor, OT, brain NP, NP receptor type C, and endothelial nitric oxide synthase (eNOS) was decreased in hearts from sedentary db/db mice. Both ET alone and ET + CR increased the mRNA expression of GATA4 compared to sedentary db/db mice. Only ET combined with CR produced increased eNOS mRNA and protein expression.

Conclusion: Our data indicate that enhancement of eNOS by combined ET and CR may improve coronary endothelial vasodilator dysfunction in type 2 diabetes but did not prevent the downregulation of cardiac expression in the OT–NP system, possibly resulting from the sustained hyperglycemia and obesity in diabetic mice.

Keywords: running, diabetes, GATA4, oxytocin, natriuretic peptides, db/db

Introduction

The association between a sedentary lifestyle and the risk of developing cardiovascular and metabolic diseases is well established. In recent decades, the dramatic rise in sedentary lifestyle behavior has resulted in an unprecedented increase in the prevalence of obesity, type 2 diabetes, and vascular diseases.^{1–3} Type 2 diabetes is a chronic metabolic disorder and, with its high prevalence worldwide, is considered as one of the leading health threats in developed nations. Decreasing this unfavorable association between a sedentary lifestyle and cardiometabolic outcome can be accomplished by effective nonpharmacological approaches, such as limiting the caloric intake and

Diabetes, Metabolic Syndrome and Obesity: Targets and Therapy 2017:10 27–36 27 So 2017 Broderick et al. This work is published and licensed by Dove Medical Press Limited. The full terms of this license are available at https://www.dovepress.com/terms. you hereby accept the ferms. Non-commercial uses of the work are permitted "who may further permission from Dove Medical Press Limited, provided the work is properly attributed. For permission for commercial use of this work, please see paragraphs 4.2 and 5 of our Terms (https://www.dovepress.com/terms.php). increasing the level of physical activity.⁴ Regular exercise is warranted for the treatment of type 2 diabetes for its positive outcome on cardiovascular risk factors. Epidemiological studies unequivocally indicate that exercise training (ET) is an effective method in improving blood glucose regulation, insulin resistance, blood lipids, and blood pressure.5-7 The limited exercise capacity and impaired energy metabolism can also be improved, delaying the onset and progression of cardiovascular disease in patients with type 2 diabetes.^{8,9} In addition to exercise, a great body of evidence has shown that limiting food intake is an effective frontline strategy to reduce cardiometabolic risk and disorders.^{10,11} Reports have indicated that caloric restriction (CR) attenuates the development of cardiomyopathy and decline in heart function normally associated with aging.12 Moderate CR decreases visceral fat and blood pressure and improves the lipid profile, glucose uptake, and insulin sensitivity, and hearts from CR rodents show an improved ischemic tolerance.¹³⁻¹⁶ Furthermore, CR preserves pancreatic mass and function by suppressing apoptosis in the db/db mouse,¹⁷ attenuates the development of hepatic steatosis in the db/db mice,18 and decreases adiposity in the genetically hyperphagic OLEFT model of type 2 diabetes.¹⁹ Taken together, it is clear that the benefits of ET and CR share a common favorable outcome on cardiovascular and metabolic health.20

To date, few studies have investigated the effects of combined ET and CR on cardiometabolic protection in type 2 diabetes. Recent work has demonstrated that the obesity and insulin resistance induced by a high-fat diet (diet-induced obesity [DIO]) in mice were reversed with exercise and CR.²¹ Using the same model of obesity, Cui et al²² reported that CR and treadmill running improved common cardiovascular risk factors, decreased plasma leptin, increased plasma adiponectin, and activated autophagyrelated pathways in the cardiac muscle. Decreasing the caloric intake and increasing the energy expenditure by voluntary exercise in the hypoleptinemic ob/ob mouse reduced fasting hyperglycemia and hyperglucagonemia but failed to reduce plasma insulin levels.^{23,24} In the db/db mouse, a model characterized by insulin resistance, hyperglycemia, and hyperleptinemia as a result of a mutation that inactivates the leptin receptor, combining exercise and dietary energy restriction improved fasting blood glucose.²⁵ The effects of this combined intervention approach on metabolic control using the db/db mouse are unclear. The fact is that, overall, there is limited literature examining the effects of this combined nonpharmacological approach in the db/db mouse, particularly for cardioprotective genes.

Oxytocin (OT) is a cardiovascular hormone with robust cardioprotective effects. In heart, OT exerts direct cardioprotection through the OT receptor (OTR), or indirectly via the stimulation of natriuretic peptides (NPs; atrial NP [ANP], brain NP [BNP], and C-type) and nitric oxide (NO) synthesis.26,27 This functional OT-NP-NO system is downregulated in hearts from db/db mice.^{28,29} The contribution of this system to the development of diabetic cardiomyopathy is indicated by the fact that chronic administration of OT and BNP prevented cardiac dysfunction and heart remodeling naturally developing in the db/db mice.^{30,31} We have demonstrated that ET improves expression of the OT-NP system in ovariectomized rats³² but fails to reverse most components of this cardioprotective system in the diabetic mouse.^{28,33} It is unknown to what extent the development of cardiomyopathy in exercised db/db mice is associated with binge eating that is common in the diabetic db/db mouse.³⁴ Some light on this problem may be obtained by limiting the excess of food intake during ET. In fact, while cardiac levels of the OT, ANP, and BNP remained low, ET only partially reversed the defect in NO synthesis,^{28,33} while other studies indicate an improved lipid profile and vascular function.35,36 Considering the supportive role of reduced dietary intake in metabolic control in diabetes, the present study examined the effects of combined ET and CR on the OT-NP system in the heart from the db/db mouse.

Methods

Mouse model of diabetes

This study was approved by the Midwestern University Research and Animal Care Committee. All animals used in this study were cared in accordance with the recommendations in the Guide for the Care and Use of Laboratory Animals, National Institute of Health, publication number 85-23, 1986. Five-week-old male db/db mice (C57BL/6J background strain; B6.V-Lep^{db} strain) were obtained from Jackson Laboratory (Bar Harbor, ME, USA). The strain displays both the metabolic alterations and cardiac dysfunction seen in diabetes and obesity.³⁷ The onset of diabetes in this strain is gradual and characterized by hyperphagia with a subsequent development of hyperglycemia as a result of two mutant copies of the leptin receptor gene. The lean littermate, which is characterized by only one mutation of the leptin gene, was used as control.

Experimental groups and ET protocol

Diabetic db/db mice were assigned to the following three groups: sedentary, diabetic runners (ET), and diabetic runners with CR (ET + CR). Exercise was performed on a treadmill

(Exer 3/6 treadmill; Columbus Instruments, Columbus, OH, USA) at moderate intensity of 5 days/week for 8 weeks, as previously reported.²⁸ The training regimen consisted of the following 3-week graded increase in exercise duration and intensity: week 1, 10 minutes at 10 m/minute; week 2, 20 minutes at 10 m/minute; week 3, 30 minutes at 12 m/minute. From weeks 4 to 8, the intensity was increased 30 minutes at 15 m/minute, corresponding to an estimated submaximal VO₂ of ~50 mL/kg/minute.³⁸ In mice assigned to the ET + CR group, the amount of food provided on a daily basis was reduced by 30% of the amount consumed by the sedentary group. Both ET and CR were maintained for a period of 8 weeks. Mice in groups not subjected to restricted food intake were provided with food ad libitum. All mice were maintained in a room with alternating 12-hour light/dark cycle and kept at 22°C.

Blood glucose status of mice

In the late morning at the end of each week, overnight fasted mice were weighed and then placed on a warm heating pad for a period of 30 minutes. Mice were then placed in a restraining chamber with the tail exposed for the collection of blood by puncturing the tip of the tail using a 23 G needle. Blood glucose was measured collected using a commercially available kit (Wako Chemicals USA, Richmond, VA, USA). At the end of the protocol, and 48 hours after the last exercise session, overnight fasted mice were euthanized by CO_2 gassing between 8 and 11 am. Hearts were rapidly removed and frozen with clamps that were precooled with light nitrogen for gene analysis.

Real-time PCR

Total RNA was extracted from freeze-clamped hearts with Trizol reagent (Invitrogen Life Technologies, Burlington, ON, USA) according to the manufacturer's protocol. To remove

Table I PCR	primer	sequences
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genomic DNA, RNA samples were incubated with 2 U of deoxyribonuclease I (DNase I; Invitrogen Life Technologies)/ µg of RNA for 30 minutes at 37°C. Polymerase chain reaction (PCR) was carried out in the iCycler IQ[™] Real-Time PCR Detection System (Bio-Rad Laboratories, Hercules, CA, USA), using the SYBR® Green chemistry. The samples were analyzed in duplicate or triplicate. For amplification, 2 µL of diluted cDNA was added to a 20 µL of reaction mixture containing 1×iQ SYBR® Green Supermix (Bio-Rad Laboratories) and 200 nM forward and reverse primers. The thermal cycling program was 95°C for 2 minutes, followed by 40 cycles of 95°C for 30 seconds, 60°C for 30 seconds, and 72°C for 30 seconds. The primers were purchased from Invitrogen Life Technologies. Primer sets served to generate amplicons (Table 1). Optical data were recorded during the annealing step of each cycle. After PCR, the reaction products were melted for 1 minute at 95°C, and the temperature was lowered to 55°C and then gradually increased to 95°C in 1.0°C increments, 10 seconds per increment. Optical data were collected over the duration of the temperature increments, with a dramatic drop in fluorescence occurring. This was done to ensure that only one PCR product was amplified per reaction.

The relative expression of the RT-PCR products was determined by the $\Delta\Delta$ Ct method. This method calculates relative expression using the following equation: fold induction =2^{-[$\Delta\Delta$ Ct]}, where Ct is the threshold cycle, ie, the cycle number at which the sample's relative fluorescence rises above the background fluorescence, and $\Delta\Delta$ Ct = [Ct gene of interest (unknown sample) – Ct glyceraldehyde-3-phosphate dehydrogenase [GAPDH] (unknown sample)] – [Ct gene of interest (calibrator sample) – Ct GAPDH (calibrator sample)]. One of the control samples was chosen as the calibrator sample and tested in each PCR. Each sample was run in

Gene	Sense primer (5'-3') Antisense primer (5'-3')		Accession number	
ANP	CCTGTGTACAGTGCGGTGTC	GTGTC CCTAGAAGCACTGCCGTCTC		
BNP	CTGAAGGTGCTGTCCCAGAT	GTTCTTTTGTGAGGCCTTGG	NM_008726	
GATA4	CACTATGGGCACAGCAGCTCC	TTGGAGCTGGCCTGCGATGGTC	NM_008092	
IRAP	CAAAGACCGAGCCAACCTGATC	GCTAAAGAGGAACAACCAGCC	NM_172827	
NPR-A	AGTACGCCAACAACCTGGAG	AAGAGCTGTAAAGCCCACGA	NM_008727	
NPR-B	TCATGACAGCCCATGGTAAA	GGTGACAATGCAGATGTTGG	NM_173788	
NPR-C	TGACACCATTCGGAGAATCA	TTTCACGGTCCTCAGTAGGG	NM_010933	
eNOS	AACCAGCGTCCTGCAAAC	AACCAGCGTCCTGCAAAC	NM_008713	
ОТ	CCTACAGCGGATCTCAGACTGA	TCAGAGCCAGTAAGCCAAGCA	NM_011025	
OTR	CGACTCAGGACGAAGGTGGAGGA	AAGATGACCTTCATCATTGTTC	NM_001081147	
Actin	ACCAACTGGGACGATATGGAGAAGA	TACGACCAGAGGCATACAGGGACAA	NM_007393	
GAPDH	TTCACCACCATGGAGAAGGC	GGCATGGACTGTGGTCATGA	NM_008084	

Abbreviations: ANP, atrial natriuretic peptide; BNP, brain natriuretic peptide; eNOS, endothelial nitric oxide synthase; GAPDH, glyceraldehyde-3-phosphate dehydrogenase; GATA4, GATA binding protein 4; IRAP, insulin-regulated aminopeptidase; NPR-A, natriuretic peptide receptor type A; NPR-B, natriuretic peptide receptor type B; NPR-C, natriuretic peptide receptor type C; PCR, polymerase chain reaction; OT, oxytocin; OTR, oxytocin receptor.

duplicate, and the mean Ct was taken in the $\Delta\Delta$ Ct equation. GAPDH was chosen for normalization because this gene showed a consistent expression relative to other housekeeping genes among the treatment groups in our experiments.

Western blot analysis

The analysis of protein levels was performed as previously reported.²⁸ Heart samples (~100 mg) were prepared by homogenization in modified radioimmunoprecipitation assay (RIPA) buffer (1× phosphate-buffered saline, 1% Igepal CA-630, 0.5% sodium deoxycholate, 0.1% SDS, 10 mg/mL phenylmethylsulfonyl fluoride, aprotinin, 100 mM sodium orthovanadate, and 4% protease inhibitor). After 2 hours in constant agitation at 4°C, the samples were centrifuged at $10,000 \times g$ for 20 minutes at 4°C. The supernatants were collected, and the protein concentration was determined by modified Bradford assay. Thirty micrograms of total protein was applied to each well of 10% sodium dodecyl sulfate-polyacrylamide gel and electrophoresed for 2 hours at 130 V along with a set of molecular weight markers (RPN800; Amersham Biosciences, Piscataway, NJ, USA). The resolved protein bands were then transferred onto polyvinylidene difluoride membranes (Hybond-C; Amersham Biosciences) at 20 V for 60 minutes at room temperature using a transfer buffer (25 mmol/L Tris base, 192 mmol/L glycine, and 20% methanol). The blots were blocked overnight at 4°C with blocking buffer (5% nonfat milk in 10 mmol/L Tris, pH 7.5, 100 mmol/L NaCl, and 0.1% Tween 20) (Amersham Pharmacia). The membranes were then probed with specific primary antibodies for GATA binding protein 4 (GATA4) (1:500, sc-25310) and endothelial nitric oxide synthase (eNOS) (1:1000, sc-654) overnight at 4°C. As an internal control, blots were reprobed with an anti-B-GAPDH antibody (1:20,000; G9545-200UL; Sigma-Aldrich, Oakville, ON, CA). Blots were then washed using tris-buffered saline washing buffer (10 mmol/L Tris, pH 7.5, 100 mmol/L NaCl, and 0.1% Tween 20) and incubated with horseradish peroxidase-conjugated immunoglobulin G (IgG) during 1 hour at room temperature. The blots were finally detected by chemiluminescence detection system (RPN2132; Amersham Biosciences) and visualized by exposure to Kodak X-Omat film. Densitometric measurement of the bands was performed using Photoshop 7 software.

Statistics

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Data are presented as mean \pm standard error of the mean (SEM). Group mean difference was determined using analysis of variance (ANOVA analysis of variance), followed by a Tukey–Kramer comparison for post hoc analysis between treatment groups. A value of *P*<0.05 was considered significant.

Results Effect of ET and CR on body weight and plasma glucose

Earlier studies have demonstrated that treadmill ET induces a modest improvement in body weight and blood glucose control in the db/db mouse.^{28,35,36} Restricting the caloric intake also has beneficial effects on body weight and metabolic control.^{17–19} Based on these observations, we therefore examined the effects of combined treadmill training and CR on the diabetic state in the leptin-resistant mouse. As expected, body weight was higher in the db/db mice than in the lean control mice (Table 2). ET alone had no effect on body weight in the db/db mice, but when combined with CR, a reduction was observed after 8 weeks. Plasma glucose levels were also elevated in the db/db mice, confirming the diabetic state. However, plasma glucose levels remained elevated after ET and were not different compared with sedentary mice. ET in combination with CR also had no beneficial effect on plasma levels of glucose.

Effect of ET and CR on cardiac GATA4 expression

GATA4 is a transcription factor expressed in heart and known to regulate the synthesis of structural and contractile genes, as well as genes of the cardioprotective OT-NP system.^{39,40} In diabetic myocardium, reduced expression of the OT-NP system is associated with apoptosis, accumulation of collagen and reduced utilization of glucose, and peripheral effects, including impaired vasodilation and diuresis.26,28,29,37 As shown in Figure 1, mRNA and protein expression of GATA4 was reduced in hearts from sedentary db/db mice compared with control mice, confirming our earlier studies.^{28,33} Compared with sedentary mice, ET of db/db mice resulted in increased mRNA expression of GATA4. ET in combination with CR further increased mRNA expression by 25%, such that the expression of GATA4 was reversed. However, GATA4 protein expression was not correspondingly increased by ET alone or in combination with CR.

Parameter	Control	Sedentary	Runners	Runners + CR
Initial body weight (g)	23.9±0.5	34.6±0.6ª	34.1±1.3ª	34.9±0.5ª
Final body weight (g)	26.4±0.5⁵	34.7±2.1ª	33.5±1.4ª	31.6±1.1ª,b
Initial glucose	6.45±0.33	18.82±1.88ª	17.84±1.59ª	14.63±1.89ª
Final glucose	5.09±0.18°	19.56±1.32ª	21.68±1.66ª	22.65±2.91 ^{a,c}

Notes: Values are reported as mean \pm SEM for 7–10 mice for each group. Glucose values are expressed in millimole per liter. ^aCompared to control mice. ^bCompared to initial body weight. ^cCompared to initial blood glucose. **Abbreviation:** CR, caloric restriction.



Figure I The effects of diabetes, ET, and CR on cardiac GATA4 mRNA expression (A) and protein expression (B).

Notes: Values are reported as mean \pm SEM for six mice per group. *Compared to lean control mice. **P*<0.05, compared to sedentary db/db mice.

Abbreviations: CR, caloric restriction; ET, exercise training; GATA4, GATA binding protein 4.

Effect of ET and CR on cardiac OTR, OT, and NP expression

OT is a cardiovascular hormone linked to the synthesis of ANP and NO, and the expression of OT is reduced under diabetic conditions.⁴¹ We next examined whether ET or combined ET and CR improved the synthesis of OT and associated NPs. Figure 2 shows that the expression of OTR was significantly reduced in hearts from sedentary db/db mice compared with control mice. Slight increases in OTR expression following ET and ET + CR were observed. Indeed, compared with sedentary db/db mice, the expression of OTR was increased by ~25% with ET and by ~50% with ET + CR.

Consistent with the diabetic state, the expression of OT was also significantly decreased.³³ ET and ET in combination with CR were not effective in reversing the effect of diabetes. While no effect of diabetes on ANP expression was associated with the reduced synthesis of OT, a significant reduction in the expression of BNP in hearts from sedentary db/db mice compared with lean control mice was observed. ET and ET with CR had no effect on the expression of both ANP and BNP.

Effect of ET and CR on cardiac NP receptor (NPR) gene expression

Gene expression of NPR type A (NPR-A), NPR type B (NPR-B), and NPR type C (NPR-C) is illustrated in Figure 3.



Figure 2 The effects of diabetes, ET, and CR on cardiac mRNA expression of the oxytocin receptor (A), oxytocin (B), ANP (C), and BNP (D). Notes: Values are reported as mean ± SEM for six mice per group. *Compared to lean control mice. Abbreviations: ANP, atrial natriuretic peptide; BNP, brain natriuretic peptide; CR, caloric restriction; ET, exercise training; GAPDH, glyceraldehyde-3-phosphate dehydrogenase; OT, oxytocin; OTR, OT receptor.



Figure 3 The effects of diabetes, ET, and CR on cardiac mRNA expression of NPR-A (**A**), NPR-B (**B**), and NPR-C (**C**). Notes: Values are reported as mean ± SEM for six mice per group. *Compared to lean sedentary control mice. Abbreviations: ANP, atrial natriuretic peptide; CR, caloric restriction; ET, exercise training; GAPDH, glyceraldehyde-3-phosphate dehydrogenase; NPR-A, natriuretic peptide receptor type A; NPR-B, natriuretic peptide receptor type B; NPR-C, natriuretic peptide receptor type C.

Expression of NPR-A and NPR-B was not altered by diabetes, ET, or ET and CR. However, gene expression of NPR-C was decreased in hearts from sedentary db/db mice. Similarly, a decrease in the expression of this receptor was also observed following ET and ET in combination with CR, compared with control mice.

Effect of ET and CR on cardiac eNOS expression

In addition to ANP and BNP, the expression of eNOS is also linked to the OT-NP system.²⁶ The expression of eNOS is reduced in hearts from db/db mice, and ET partially restores the levels of eNOS.28 Because the effects of ET in combination with CR on eNOS expression are unclear, this was examined. As shown in Figure 4, mRNA expression of eNOS was reduced by ~40% in hearts from db/db mice compared with control mice. This reduction was associated with a significant decrease in protein expression. While ET alone had no effect on either mRNA expression of eNOS or protein expression of eNOS, ET in combination with CR was clearly beneficial and increased both mRNA and protein expression of eNOS. In fact, eNOS protein expression was normalized by combining ET with CR. Taken together, our results indicate that ET with reduced caloric intake improves the expression of certain genes of the OT-NP system, namely the transcription factor GATA4, the OTR, and eNOS.



Figure 4 The effects of diabetes, ET, and CR on cardiac eNOS mRNA expression (A) and protein expression (B).

Notes: Values are reported as mean \pm SEM for six mice per group. *Compared to lean control mice. #P<0.05, compared to sedentary db/db mice. \pm P<0.5, compared to exercise-trained mice.

Abbreviations: CR, caloric restriction; eNOS, endothelial nitric oxide synthase; ET, exercise training.



Figure 5 The effects of diabetes, ET, and CR on cardiac IRAP mRNA expression. Notes: Values are reported as mean \pm SEM for six mice per group. *Compared to lean control mice.

Abbreviations: CR, caloric restriction; ET, exercise training; IRAP, insulin-regulated aminopeptidase.

Effect of ET and CR on cardiac insulinregulated aminopeptidase (IRAP) expression

Cardiac IRAP expression, involved in GLUT4 trafficking,⁴² in hearts from diabetic mice is illustrated in Figure 5. Expression of IRAP was decreased in hearts from sedentary db/db mice, and treatment had no effect on the expression of this intracellular vesicular transporter protein.

Discussion

The nonpharmacological management of type 2 diabetes and related metabolic disorders focuses on a healthy lifestyle, which includes increasing the level of physical activity and reducing the caloric intake. This maintains better glycemic control, and a weight loss from CR improves cardiometabolic health in patients with obesity and type 2 diabetes.¹¹ To date, most experimental studies have focused on the role of exercise or different exercise paradigms on cardioprotection in the db/db mouse model of diabetes without concurrent restricted caloric intake. The db/db mouse was selected because of its close representation to the phenotype observed in human diabetes and obesity. A feature of this model is hyperleptinemia from a mutation in the leptin receptor, rarely the culprit for the metabolic disturbances seen in human diabetes but nonetheless detected in diabetes.43 By including a dietary restriction component, we aimed to determine whether the defect in the OT-NP-NO system in the db/db heart could be prevented with ET.

Our results indicate that the benefits afforded by exercise and the reduced caloric intake in the db/db mouse were modest. A reduction in body weight was observed, but diabetic mice remained hyperglycemic after 8 weeks of ET combined

with CR. In addition, with the exception of protein and gene syntheses for eNOS, ET and CR failed to significantly improve gene expression of the OTR, OT, and BNP. Gene expression of NPR-A and NPR-B was not altered by the diabetic state, ET, or ET with CR, while the expression of NPR-C was reduced under all conditions. Considering the vast body of reports indicating that combining both exercise and limited food intake is beneficial on cardiometabolic health for the treatment of diabetes, our results showing only partial reversal of the defect in the OT system in diabetic mice were clearly unexpected. The reason for this response is unclear, although evidence suggests that impaired expression of this system may be related to observation that diabetic mice remained hyperglycemic, despite the exercise combined with CR.⁴⁴ Indeed, evidence indicates that hyperglycemia markedly inhibits the synthesis of OT, ANP, and BNP and structural proteins by suppressing the expression of cardiac GATA4, an essential transcription factor.44 The importance of GATA4 is highlighted by the observation that low GATA4 content is further associated with an increased ischemic injury and the risk of heart failure, both of which are evident in hearts from db/db mice.45 Although our results confirm earlier studies in the db/db mouse,²⁸ synthesis of the OT system is also reduced in the ob/ob mice and in hearts from mice with streptozotocin-induced type 1 diabetes or fed with high-fat diet.^{29,44,46} Hyperglycemia is present in all these rodent models, but it should be noted that hyperglycemia per se cannot solely account for low GATA4 expression in these models. The role of defective leptin receptor signaling, mutation of the ob gene, insulin resistance or deficiency, and calorie-rich food consumption are potential causes for the decreased expression of the OT, the OTR, and BNP. However, we have reported that hearts from db/db mice show increased gene and protein expression of GATA4 following 8 weeks of treadmill running without concomitant changes in the expression of OT and NP synthesis.³³ The observations that ET alone,²⁸ or in combination with CR, is associated with an increased protein expression of eNOS are of interest. Since eNOS is a component of the OT system as well as a downstream gene product of GATA4,²⁶ it is possible that other transcription factors are involved in the synthesis of OT, NPs, and eNOS. In fact, GATA6 not only exerts overlapping functional redundancy with GATA4 in the regulation of these peptides but also is a direct transcriptional regulator of eNOS.^{40,47} Endothelial GATA6 deficiency leads to decreased synthesis of eNOS, increased vascular injury, and worsened remodeling in the hypoxia- and monocrotaline-induced models of pulmonary hypertension.47

While treadmill running has positive gains on metabolic risk factors and vascular function in the db/db mouse,^{35,36} this form of exercise is known to induce hyperglycemia by disrupting the hypothalamic-pituitary-adrenal axis, stimulate adrenal gland hypertrophy, and increase the secretion of catecholamines.39,48 We have demonstrated that db/db mice remain hyperglycemic after forced treadmill exercise, a response caused by excessive corticosterone and catecholamine production.³⁴ Excessive norepinephrine secretion during acute exercise is reported in hypertensive and nonhypertensive type 2 diabetic patients, and poor metabolic control is associated with postexercise hyperglycemia and hyperinsulinemia in patients with type 2 diabetes.^{49,50} Forced treadmill running increases the production of endogenous glucocorticoid and the expression of glucocorticoid receptor and gluconeogenic enzymes in liver from db/db mice, leading to increased hepatic glucose production.⁴⁸ In addition, 11β-hydroxysteroid dehydrogenase type 1, which converts inactive cortisone to physiologically active corticosterone, is abundantly expressed in liver from db/db mice after acute exercise, contributing to an increased glucocorticoid plasma pool and insulin resistance and further explaining the hyperglycemia noted in the db/db mouse after training.^{34,48,51} Despite these changes with treadmill exercise, alternating moderate-intensity exercise with high-intensity exercise in each exercise session affords cardioprotection in the db/db mouse but fails to improve the hyperglycemia and hyperinsulinemia.52 On the other hand, the metabolic disturbances linked to treadmill running are largely mitigated by voluntary wheel running exercise, supporting the use of voluntary exercise as a more physiologically gentle approach to reach desirable outcomes.34,51 Taken together, our results show that forced treadmill exercise may not always produce desirable effects with concurrent CR. The benefits of combining exercise with CR on metabolic control are better manifested with voluntary ET, as recently reported. Improved insulin sensitivity, increased neural function and synaptic plasticity, and reversal of mitochondrial dysfunction and oxidative stress have been reported in the db/db mouse and under high-fat diet conditions.^{21,24,25} In these studies, one could also argue that the beneficial effects are attributed to a more restricted diet with decreases in caloric intake by 35-40%,^{21,25} and by 50%,²¹ resulting in a normalized mitochondrial function. Hence, increasing the exercise intensity by forced means does not always correlate with better metabolic control in the db/db mouse compared to low-intensity voluntary wheel running and using voluntary running as an exercise paradigm, while subjecting mice to a more restricted caloric intake

provides greater protection against the sequelae associated with diabetes.

Inactivity and excessive food consumption contribute to the development of obesity and diabetes. Diabetes induces cardiomyopathy, and evidence indicates that this is explained by defective synthesis of the cardiac OT-NP system.²⁸ Here, we demonstrate that exercise in the form of forced moderateintensity ET with moderate CR does not significantly reverse the defect in the cardioprotective OT-NP-NO system in the db/ db mouse. However, we found that exercise and CR improved eNOS, suggesting that some protection is afforded by this strategy. The consequences of this disrupted system in the diabetic heart have been reported, including apoptosis, collagen accumulation in fibroblasts, lipid accumulation, decreased glucose uptake, and cardiomyocyte hypertrophy,26,29,53-55 and low plasma OT, ANP, and BNP are linked to increased sympathetic tone, impaired diuresis, increased volume, and reduced glucose utilization in the obese patient.56,57 Based on the significance of the cardiac OT system on cardiac function and metabolism, and that cardiovascular diseases account for ~70% of diabetesrelated deaths, further studies are needed to develop strategies for the treatment of diabetes and obesity, as well as in patients with leptin deficiency or resistance.43

Conclusion

We show a mild effect of ET in combination with CR on cardioprotection in the db/db mouse. Therefore, the development of exercise techniques and novel nutritional therapies to stimulate pathways that are different from those induced by diabetes and leptin resistance is essential for cardiometabolic health.

Acknowledgments

The study was funded by a grant from the Diabetes Action Research and Education Foundation (TLB), the Midwestern University Office of Research and Sponsored Programs (TLB), and the Canadian Institute for Health Research (JG and MJ).

Disclosure

The authors report no conflicts of interest in the work.

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