Restoration of myocardial glucose uptake with facilitated myocardial glucose transporter 4 translocation contributes to alleviation of diabetic cardiomyopathy in rats after duodenaljejunal bypass

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ABSTRACT

Aims/Introduction: Duodenal-jejunal bypass (DJB) surgery has been reported to effectively relieve diabetic cardiomyopathy (DCM). However, the specific mechanisms remain largely unknown. The present study was designed to determine the alterations of myocardial glucose uptake (MGU) after DJB and their effects on DCM.

Materials and Methods: Duodenal-jejunal bypass and sham surgeries were carried out in diabetic rats induced by a high-fat diet and a low dose of streptozotocin, with chow-diet fed rats as controls. Bodyweight, food intake, glucose homeostasis and lipid profiles were measured at indicated time-points. Cardiac function was evaluated by transthoracic echocardiography and hemodynamic measurement. Cardiac remodeling was assessed by a series of morphometric analyses along with transmission electron microscopy. Positron-emission tomography with fluorine-18 labeled fluorodeoxyglucose was carried out to evaluate the MGU *in vivo*. Furthermore, myocardial glucose transporters (GLUT; GLUT1 and GLUT4), myocardial insulin signaling and GLUT-4 translocation-related proteins were investigated to elucidate the underlying mechanisms.

Results: The DJB group showed restored systolic and diastolic cardiac function, along with significant remission in cardiac hypertrophy, cardiac fibrosis, lipid deposit and ultrastructural disorder independent of weight loss compared with the sham group. Furthermore, the DJB group showed upregulated myocardial insulin signaling, hyperphosphorylation of AKT substrate of 160 kDa (AS160) and TBC1D1, along with preserved soluble N-ethylmaleimide-sensitive factor attachment protein receptor proteins, facilitating the GLUT-4 translocation to the myocardial cell surface and restoration of MGU.

Conclusions: The present findings provide evidence that restoration of MGU is implicated in the alleviation of DCM after DJB through facilitating GLUT-4 translocation, suggesting a potential choice for treatment of human DCM if properly implemented.

INTRODUCTION

Diabetic cardiomyopathy (DCM) substantially increases heart failure risk and leads to poor prognosis¹. However, there is

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currently no specific clinical intervention, and novel therapies are required.

Bariatric surgery has been proved to provide effective remission in type 2 diabetes and associated complications^{2,3}. It has attracted more attention because of its effectiveness in ameliorating diabetic cardiac dysfunction^{4,5}. However, most previous

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© 2018 The Authors. Journal of Diabetes Investigation published by Asian Association for the Study of Diabetes (AASD) and John Wiley & Sons Australia, Ltd This is an open access article under the terms of the Creative Commons Attribution-NonCommercial-NoDerivs License, which permits use and distribution in any medium, provided the original work is properly cited, the use is non-commercial and no modifications or adaptations are made. studies were carried out on the background of obesity, and remarkable weight loss after surgery would inevitably confound the results.

Duodenal-jejunal bypass (DJB) is a commonly used experimental model to investigate the antidiabetic effect of bariatric surgery independent of weight loss^{6,7}. However, there has been limited study regarding the therapeutic effects of DJB on DCM to date. Our previous research showed that DJB could ameliorate diabetic cardiac dysfunction in a rat model⁸. However, more studies are required to elucidate the underlying mechanism.

Insulin resistance is one of the main triggers of DCM⁹. Although multiple downstream signaling is involved, impaired insulin-mediated myocardial glucose uptake (MGU) is consistently observed^{10–12}. The compromised glucose utilization facilitates a substrate shift toward increased free fatty acid oxidation, which would in turn lead to cardiac damage¹³. Conversely, normalizing MGU and cardiac metabolism could reverse DCM¹⁴.

At the cellular level, MGU is dependent on the transmembrane glucose gradient and the content of cell surface glucose transporters (GLUT)¹⁵. GLUT-1 is mainly localized in the sarcolemma and regulates the basal MGU, whereas GLUT-4 takes effect relying on a sophisticated translocation from intracellular storage sites to the cell surface through a cascade activation of the insulin signaling pathway in the presence of stimulation^{16,17}.

To our knowledge, the alterations in MGU after DJB and its effects on DCM have not been reported. In the present study, we carried out DJB and sham surgeries on diabetic rats induced by a high-fat diet (HFD) and a low dose of streptozotocin, with chow-diet fed rats as controls. We aim to investigate the alterations in MGU after DJB and the associated mechanisms, as well as its morphological and functional effects on DCM.

METHODS

Animals

All experiments were approved by the Ethics Committee on Experiment Animal of Shandong University Qilu Hospital. Eight-week-old male Sprague-Dawley rats (the Laboratory Animal Center of Shandong University, Jinan, China) were randomly assigned to the control group or diabetic group. The control group (n = 10) was fed with standard rodent chow diet (15% of calories as fat, Laboratory Animal Center of Shandong University) throughout the study period. Type 2 diabetes was induced as previously described¹⁸. Briefly, rats in the diabetic group received HFD (40% calories as fat, Huafukang Biotech Company, Beijing, China) for 4 weeks to induce insulin resistance. After a 12-h fast, they were injected intraperitoneally with 2% of streptozotocin (Sigma, St. Louis, MO, USA; 35 mg/ kg dissolved in ice cold citrate buffer, pH 4.5), while the control group received citrate buffer alone. Three days later, rats with non-fasting glucose ≥16.7 mmol/L were considered diabetic (n = 20). After 16 weeks of diabetes, transthoracic echocardiography was carried out to evaluate the cardiac function before surgery. Then diabetic rats (n = 20) were randomly allocated to undergo corresponding surgery in the DJB group (n = 10) or the sham group (n = 10). The study design is summarized in Figure S1.

Surgical Procedures

DJB surgery was carried out as previously reported¹⁹. Sham surgery consisted of midline laparotomy with the same exposure, but not removal of stomach tissue. The operation time was prolonged similar to that of DJB to ensure equivalent anesthetic and surgical stress. Access to water was given from 2 h postoperatively. Rats were fed with 10% of Ensure (Abbott Laboratories, Abbott Park, IL, USA) from 24 h after surgery for 3 days, followed by standard rodent chow diet until the end of the study.

Oral Glucose Tolerance Test and Insulin Tolerance Test

The oral glucose tolerance test (OGTT) and insulin tolerance test (ITT) were carried out preoperatively, and at both 2 and 8 weeks after surgery, as previously described¹⁸. Briefly, after a 12-h fast, blood glucose of rats was monitored from the tail vein at baseline, 10, 30, 60 and 120 min after administration of 20% glucose (1 g/kg) by intragastric gavage for OGTT, or insulin lispro (0.5 IU/kg) by intraperitoneal injection for ITT. ITT was carried out 2 days after OGTT to ensure recovery.

Blood Sampling and Analysis

Blood samples were collected from the retro-orbital plexus of rats under light ether anesthesia after an overnight fast. Levels of serum triglyceride, total cholesterol, fasting plasma glucose (FPG), high-density lipoprotein cholesterol, low-density lipoprotein cholesterol and non-esterified fatty acid were analyzed by the Roche Cobas 8000 modular analyzer system (Roche Diagnostics, Indianapolis, IN, USA). Plasma fasting insulin and glucagon-like peptide-1 (GLP-1) were quantified using the Ultra Sensitive Rat Insulin ELISA Kit (Crystal Chem, Elk Grove, IL, USA) and multispecies GLP-1 ELISA kit (Millipore, Billerica, MA, USA), respectively. The homeostasis model assessment of basal insulin resistance was calculated as FPG (mmol/L) \times fasting insulin (mIU/L) / 22.5²⁰.

Echocardiographic Evaluation

Before and at 8 weeks after surgery, transthoracic echocardiography was carried out with a VEVO 2100 imaging system (VisualSonics, Toronto, ON, Canada). Rats were lightly anesthetized with inhaled isoflurane- O_2 . Two-dimensional and M-mode imaging was carried out to evaluate cardiac structure by determining left ventricular end diastolic diameter and left ventricular end systolic diameter. The left ventricular systolic function was assessed according to ejection fraction and fractional shortening. Mitral inflow was recorded by pulsed-wave Doppler at the apical position; peak velocities of early filling (E) were measured. Tissue Doppler imaging of mitral annulus was obtained in the apical four-chamber view at the highest possible frame rate and peak early diastolic velocities (e') were measured. The ratio of E/e' was calculated to evaluate the left ventricular diastolic function²¹.

Hemodynamic and Blood Pressure Measurements

At 8 weeks after surgery, under deep anesthesia, a fluid-filled catheter was inserted from the right carotid artery into the left ventricle (LV) of rats. The left ventricular end diastolic pressure was measured.

In terms of blood pressure measurement, the heart rate, systolic blood pressure, diastolic blood pressure and mean arterial pressure were measured by a tail-cuff automatic sphygmomanometer (Softron, Tokyo, Japan) as described previously²².

Positron-Emission Tomography Scan and Image Processing

Three rounds of positron-emission tomography (PET) scan were carried out at 8 weeks after surgery to evaluate the MGU in vivo with an Inveon dedicated PET system (Siemens, Bonn, Germany). Briefly, the first round was carried out regularly after fluorine-18 labeled fluorodeoxyglucose (18F-FDG) injection from the tail vein. To ensure the maximal stimulation of MGU, the second round was carried out after an intravenous glucose load. To fully simulate the postprandial hyperglycemia state, the third round was carried out after an intragastric glucose load instead. To determine the MGU, multiple regions-of-interest were drawn manually on at least five representative midventricular transverse planes by visualizing the heart margin as they appeared as bumps under the skin²³. The mean standard uptake value was calculated to identify the radioactivity of the heart. The detailed method and image processing are listed in Appendix S1.

Organ Harvesting

At the end of the study (8 weeks after the surgery), human insulin (Velosulin, 10 U/kg bodyweight) was injected into the caudal vena cava of rats. Three minutes later, rats were euthanized and the hearts were rapidly removed and washed in cold phosphate-buffered saline. Then, 1-mm cubes were quickly cut from the LV and fixed in 2.5% glutaraldehyde for electron microscopy analysis. The remaining part of the LV was sliced into three pieces, one was fixed in 4% paraformaldehyde, and the other two were put into liquid nitrogen and stored at - 80°C for further analysis.

Morphometric and Histological Analysis

LV tissues fixed in 4% paraformaldehyde were embedded in paraffin, and 4- μ m thick sections were produced for hematoxylin–eosin staining and Masson's trichrome staining. The myocyte cross-sectional area was calculated under ×400 magnification from hematoxylin–eosin-stained sections. Collagen volume fraction and perivascular collagen area/luminal area ratio were quantified from Masson-stained sections. LV frozen sections (5 μ m) were stained with Oil Red O and then counterstained with hematoxylin to detect neutral lipid. All quantitative morphometry was carried out with Image-Pro Plus 6.0 (Media Cybernatics, Bethesda, MD, USA).

Transmission Electron Microscopy

LV samples (1-mm cubes) were fixed in 2.5% glutaraldehyde overnight at 4°C and post-fixed in 1% osmium tetroxide, then dehydrated through a graded ethanol series and embedded in epoxy resin. Ultrathin sections double-stained with uranyl acetate and lead citrate were examined by a transmission electron microscope (FEI/Philips CM-100; Thermo Fisher Scientific, Waltham, MA, USA).

Immunohistochemistry and Western Blotting

Image-Pro Plus 6.0 software (Media Cybernatics) was used to calculate the mean integrated optical density per stained area (μ m²) for immunohistochemistry. In terms of Western blotting, the total and membrane target protein were normalized relative to β -actin and Na+/K+ ATPase1, respectively. The detailed methods are listed in Table S1 and Appendix S2.

Statistical Analysis

Quantitative data are presented as mean \pm standard deviation, and were analyzed using SPSS software (ver. 18.0; SPSS Inc., Chicago, IL, USA). The area under the curve (AUC) for OGTT (AUC_{OGTT}) and ITT (AUC_{ITT}) was calculated by trapezoidal integration. Bodyweight, food intake, FPG, fasting insulin and homeostasis model assessment of basal insulin resistance over time were analyzed using two-way (repeated-measures) analysis of variance (ANOVA). The results are reported differences as ^A*P* by group, ^B*P* over time and ^C*P* due to the interaction of the two factors. The Bonferroni post-hoc test was carried out where appropriate. The other datasets were compared using one-way ANOVA followed by Bonferroni or Dunnett's T3 correction. P < 0.05 was taken to show statistical significance.

RESULTS

General Characteristics of Diabetic Rats Before Surgery

After 16 weeks of diabetes, the diabetic group showed higher water intake and urine volume than the control group, as expected. The two groups showed similar levels of bodyweight, systolic blood pressure, diastolic blood pressure and mean arterial pressure (Table S2). However, the diabetic group showed significantly higher left ventricular end diastolic diameter, left ventricular end systolic diameter and E/e', along with lower ejection fraction and fractional shortening (Figure 1), indicating both systolic and diastolic cardiac dysfunction before surgery.

Bodyweight and Food Intake After Surgery

Due to perioperative food restriction and surgical stress, both bodyweight and food intake in the sham and DJB groups decreased sharply during the first week after surgery, and increased gradually thereafter (Figure 2a,b). No significant difference was detected between these two groups in terms of bodyweight and food intake after surgery (Figure 2a,b).



Figure 1 | Evaluation of cardiac function in the control and diabetic groups before the surgery. (a) Representative M-mode (left), pulsed-wave Doppler mode (middle) and tissue Doppler mode (right) echocardiogram along with measurements of left ventricular end diastolic diameter (LVEDD), left ventricular end systolic diameter (LVESD), peak velocities of early filling (E) and peak early diastolic velocities (e'). (b–g) Sequential evaluations of (b) heart rate, (c) LVEDD, (d) LVESD, (e) ejection fraction (EF), (f) fractional shortening (FS) and (g) E/e'. Data are expressed as mean \pm standard deviation. n = 10 in the control group and n = 20 in the diabetic group. *P < 0.05, **P < 0.01 diabetic versus control.

DJB Surgery Significantly Improved Glucose Homeostasis and Serum Lipid Profiles

The DJB group showed lower levels of FPG than the sham group from 2 weeks after surgery (Figure 2c). Though the fasting insulin level of the two groups was similar, the DJB group showed improved homeostasis model assessment of basal insulin resistance from 2 weeks after surgery (Figure 2d,e). In addition, lower levels of AUC_{OGTT} and AUC_{ITT} were detected in the DJB group at 2 and 8 weeks after surgery than the sham group (Figure 2f,g). These findings suggested that DJB could reverse the impaired glucose tolerance and insulin sensitivity in diabetic rats. Furthermore, the DJB group had higher levels of plasma GLP-1 than the sham group at both 2 and 8 weeks after surgery (Figure 2h).

Compared with the sham group, the DJB group showed lower serum triglyceride and non-esterified fatty acid, along with higher serum high-density lipoprotein cholesterol at 2 weeks after surgery (Table 1). At 8 weeks postoperatively, the DJB group showed significantly lower serum triglyceride, total cholesterol, low-density lipoprotein cholesterol and non-esterified fatty acid, as well as higher serum high-density lipoprotein cholesterol, which were at the same levels as in the control group (Table 1). Thus, DJB surgery could effectively improve the serum lipid profiles of diabetic rats.



Figure 2 | Bodyweight, food intake and glucose homeostasis of rats. (a) Bodyweight, (b) food intake, (c) fasting plasma glucose (FPG), (d) fasting insulin (FINS) and (e) homeostasis model assessment of basal insulin resistance (HOMA-IR) before and after surgery. (f) The area under the curve (AUC) for the oral glucose tolerance test and (AUC_{OGTT}) and (g) insulin tolerance test (AUC_{ITT}) before and at 2 and 8 weeks after surgery. (h) Level of plasma glucogon-like peptide-1 (GLP-1) before and at 2 and 8 weeks after surgery. Data are expressed as mean ± standard deviation for n = 10 per groups. ^AP, difference by groups; ^BP, difference over time; ^CP, difference due to interaction of groups and time in two-way ANOVA. *P < 0.05, **P < 0.01 sham versus control; ^{##}P < 0.01 duodenal-jejunal bypass versus sham; [&]P < 0.05, ^{&&}P < 0.01 duodenal-jejunal bypass versus control.

DJB Surgery Reversed Diabetic-Induced Cardiac Remodeling

As shown in Figure 3a, the myocardial cells in the sham group lined up in disorder with irregular nucleus and uneven cytoplasm distribution, and the myocardial fibers were fragmented and disrupted. The DJB group showed well-arranged myocardial cells, which was similar as those in the control group. The DJB group showed a 29% decrease in myocyte size compared with the sham group (Figure 3f), indicating a significant amelioration of the cardiomyocyte hypertrophy. In addition, the Oil Red O staining results showed less myocardial lipid deposition in the DJB (Figure 3c). Furthermore, both interstitial and perivascular cardiac fibrosis were alleviated in the DJB group, with a lower collagen volume fraction and perivascular collagen area/luminal area ratio than the sham group (Figure 3b,g,h). This was further confirmed by the significantly lower collagen I and III content in the DJB group (Figure 3d,i). In terms of myocardial ultrastructure, the sham group showed randomly distributed swollen mitochondria and lipid droplets between poorly organized myofibrils, whereas the DJB group showed layers of mitochondria intervened between myofibrils consisting of regular and continuous sarcomeres (Figure 3e). These results showed that DJB could reverse the diabetes-induced cardiac remodeling remarkably.

DJB Surgery Ameliorated Diabetic-Induced Cardiac Dysfunction

The three groups showed similar heart rate, systolic blood pressure, diastolic blood pressure and mean arterial pressure at both

Determination	2 weeks after surgery			8 weeks after surgery		
	Control ($n = 10$)	Sham (<i>n</i> = 10)	DJB (<i>n</i> = 10)	Control ($n = 10$)	Sham (<i>n</i> = 10)	DJB (n = 10)
HR (b.p.m.)	371.46 ± 44.72	393.06 ± 37.29	382.53 ± 45.31	378.23 ± 40.75	391.74 ± 52.33	380.28 ± 45.62
SBP (mmHg)	121.53 ± 6.46	117.70 ± 8.36	119.14 ± 11.51	122.37 ± 8.16	119.55 ± 9.61	123.67 ± 10.32
DBP (mmHg)	99.62 ± 6.37	101.21 ± 6.38	98.64 ± 8.12	95.47 ± 7.41	95.32 ± 7.93	97.61 ± 8.19
MAP (mmHg)	105.14 ± 5.74	106.87 ± 7.51	104.94 ± 6.52	103.29 ± 7.48	104.11 ± 8.16	106.16 ± 6.87
TG (mmol/L)	1.05 ± 0.31	2.69 ± 0.72**	1.92 ± 0.65 ^{#,&&}	1.11 ± 0.34	2.95 ± 0.84**	1.43 ± 0.65 ^{##}
TC (mmol/L)	1.80 ± 0.35	2.32 ± 0.49*	1.96 ± 0.32	1.87 ± 0.44	3.60 ± 1.16**	$2.19 \pm 0.73^{\#}$
HDL-c (mmol/L)	1.15 ± 0.27	0.62 ± 0.20**	0.96 ± 0.39 [#]	1.21 ± 0.21	0.52 ± 0.25**	$1.04 \pm 0.31^{\#}$
LDL-c (mmol/L)	0.42 ± 0.05	0.54 ± 0.16	0.47 ± 0.15	0.39 ± 0.08	0.64 ± 0.13**	0.46 ± 0.09 ^{##}
NEFA (µmol/dL)	22.54 ± 3.39	48.42 ± 9.81**	30.83 ± 6.75 ^{##,&&}	24.27 ± 4.31	62.43 ± 10.28**	27.51 ± 6.20 ^{##}

Table 1 Heart rate, blood pressure and serum lipid profiles of rats at	2 and 8	3 weeks after si	urgery
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Data are expressed as mean \pm standard deviation, n = 10 per group. *P < 0.05, **P < 0.01 sham versus control; "P < 0.05, ""P < 0.05, ""P < 0.01 duodenaljejunal bypass (DJB) versus sham; ^{&&} P < 0.01 DJB versus control. DBP, diastolic blood pressure; HDL-c, high-density lipoprotein cholesterol; HR, heart rate; LDL-c, low-density lipoprotein cholesterol; MAP, mean arterial pressure; NEFA, non-esterified fatty acid; SBP, systolic blood pressure; TC, total cholesterol; TG, triglyceride.

2 and 8 weeks after surgery (Table 1). Compared with the sham group, the DJB group showed significant reductions in left ventricular end diastolic diameter, left ventricular end systolic diameter and E/e', along with higher levels of ejection fraction and fractional shortening (Figure 4a–e). In addition, the hemodynamic measurement showed lower left ventricular end diastolic pressure in the DJB group than in the sham group (Figure 4f). There was no significant difference between the DJB group and control group, except for the E/e' ratio (Figure 4e). These results showed that DJB could improve both systolic and diastolic LV function of diabetic rats to levels close to or similar to those observed in the control group.

DJB Surgery Restored MGU of Diabetic Rats

The sham group showed remarkably impaired MGU in all three rounds of PET scans (Figure 4g,h). However, the DJB group showed a significantly higher standard uptake value than the sham group in all three rounds, although it did not reach the level of the control group (Figure 4i). These results showed that DJB could alleviate diabetes-induced MGU impairment in both basal and insulin-stimulated states.

Effects of DJB Surgery on Myocardial Glucose and Fatty Acid Transporter

The total protein levels of GLUT-1 and GLUT-4 were similar among the three groups (Figure 5a,b,d). No significant difference was detected in plasma membrane GLUT-1 among the three groups (Figure 5b,e). However, the plasma membrane expression of GLUT-4 was significantly upregulated in the DJB group compared with the sham group (Figure 5b,e). The inconsistent intracellular distributions of GLUT-4 were further confirmed by immunohistochemical staining, which showed that most myocardial GLUT-4 in the control and DJB groups was gathered near the cell membrane (Figure 5a). The plasma membrane cluster of differentiation 36 (CD36), the major myocardial fatty acid transporter, was significantly attenuated after DJB compared with the sham group, although the total protein of which remained similar (Figure 5c–e). These results showed that the DJB group had more GLUT-4 and less CD36 in the plasma membrane compared with the sham group, but there was no significant difference in GLUT-1 (Figure 5f).

Effects of DJB Surgery on Myocardial Insulin Signaling and GLUT-4 Translocation

The DJB group showed a higher phospho/total ratio of insulin receptor substrate 1, phosphatidylinositol 3-kinase (PI3K) and protein kinase B (AKT/PKB) than the sham group (Figure 6a–d). In terms of proteins involved in GLUT-4 translocation, the DJB group showed upregulated phosphorylation of AKT substrate of 160 kDa (AS160) and TBC1D1 compared with the sham group (Figure 6e–h). No significant difference was detected in the expression of vesicle-associated membrane protein 2, synaptosome-associated protein of 23 kDa or syntaxin 4 among the three groups (Figure 6e,i). These results showed that DJB could facilitate myocardial insulin signaling and GLUT-4 translocation.

DISCUSSION

Diabetic cardiomyopathy is defined as LV dysfunction in patients with diabetes in the absence of coronary artery disease and arterial hypertension²⁴. It is characterized by morphological, functional and metabolic changes in the heart²⁵. The present study was carried out to further explore the therapeutic effects of DJB on DCM and the associated mechanisms.

In the present study, HFD combined with low-dose streptozotocin was used to induce diabetes in rats. This method has been accepted by many researchers, as it simulates the natural history and metabolic characteristics of patients with type 2 diabetes^{26–28}. Furthermore, this model is commonly used in research of DCM, and it was reported that 12–16 weeks of



Figure 3 | Effect of duodenal-jejunal bypass (DJB) on diabetes-induced cardiac remodeling. (a) Representative (a1) longitudinal and (a2) transverse sections of left ventricle stained with hematoxylin–eosin. Scale bar, 50 μ m. (b1) Interstitial (scale bar, 100 μ m) and (b2) perivascular (scale bar, 50 μ m) fibrosis, as shown by Masson's trichrome stain. Collagen fibers are dark blue and myocardium is red. (c) Representative Oil Red O-stained myocardial sections, the red droplets indicate neutral lipid staining. Scale bar, 20 μ m. (d) Representative immunohistochemical images of (d1) collagen I and (d2) III. Brown staining is considered positive. Scale bar, 2 μ m (magnification, ×10,000) (f) The myocyte size of the three groups. (g) The collagen volume fraction (CVF) of the three groups. (h) The perivascular collagen area/luminal area ratio (PVCA/LA) ratio of the three groups. (i) Semi-quantification of collagen I and III staining indicated by integrated optical density/area relative density, the value of the control group was used as the standard (100%). Data are expressed as mean ± standard deviation for n = 10 per groups. **P < 0.01 sham versus control; ##P < 0.01 DJB versus sham; ^{&&} P < 0.01 DJB versus control.

diabetes was sufficient to induce DCM in this model^{29–31}. In the present study, after 16 weeks of diabetes, the diabetic rats showed both systolic and diastolic LV dysfunction before surgery, which makes it a suitable experimental model for investigating the effects of DJB on DCM.

The present study provided direct evidence that DJB could markedly alleviate DCM morphologically and functionally

independent of weight loss. We further showed that DJB could ameliorate the MGU defects in diabetic rats, which was associated with facilitated myocardial insulin signaling and GLUT-4 translocation (Figure S2).

As a central trigger of DCM, the impaired MGU in diabetic hearts has been reported^{10,12}. However, these studies suffered from a drawback that the measurements were carried out



Figure 4 | Effect of duodenal-jejunal bypass (DJB) on cardiac function and myocardial glucose uptake. (a–e) Sequential evaluations of (a) left ventricular end diastolic diameter (LVEDD), (b) left ventricular end systolic diameter (LVESD), (c) ejection fraction (EF), (d) fractional shortening (FS) and (e) the ratio peak velocities of early filling/peak early diastolic velocities of (E/e') at 8 weeks after surgery. (f) Result of hemodynamic measurement indicated by left ventricular end diastolic pressure (LVEDP). (g) Representative transverse axis images of the heart showing ¹⁸F-FDG distribution in the first (g1), second (g2), and third (g3) rounds of PET scan, the yellow cross indicates the position of heart. (h) Representative three-dimensional reconstruction of the third round of fluorine-18 labeled fluorodeoxyglucose scan, the green square indicates the position of the heart. The first round was carried out regularly after fluorine-18 labeled fluorodeoxyglucose injection, whereas the second round was carried out after an intravenous glucose load, and the third round was carried out after an intragastric glucose load. (i) The statistical results of standard uptake value (SUV). Data are expressed as mean \pm standard deviation for n = 10 per groups. **P < 0.01 sham versus control; "P < 0.05, "#P < 0.01 DJB versus control.

in vitro using isolated working hearts. Recently, PET has been used to evaluate myocardial energy substrate uptake, with the advantage of fully reflecting the physiological state *in vivo*^{29,32}.

We carried out three rounds of ¹⁸F-FDG PET in this study. The first round was carried out in the basal state, whereas the second one was carried out under conditions of hyperglycemia



Figure 5 | Impact of duodenal-jejunal bypass (DJB) on myocardial glucose and fatty acid transporter. (a) Representative immunohistochemical staining of glucose transporter 4 (GLUT-4) in (a1) longitudinal (scale bar, 100 μ m) and (a2) transverse (scale bar, 50 μ m) myocardial sections. Brown staining is considered positive. (b) Representative western blot of myocardial total and plasma membrane GLUT-1 and GLUT-4. (c) Representative western blot of myocardial total and plasma membrane cD36. (d–f) Comparisons of (d) total expression, (e) plasma membrane expression, and (f) membrane/total ratio of GLUT-1, GLUT-4 and cluster of differentiation 36 (CD36), the value of control group was used as standard (100%). Data are expressed as mean \pm standard deviation for n = 10 per groups. **P < 0.01 sham versus control; ${}^{#}P < 0.05$, ${}^{##}P < 0.01$ DJB versus sham.

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Figure 6 | Impact of duodenal-jejunal bypass (DJB) on myocardial expression of proteins involved in insulin signaling and glucose transporter 4 (GLUT-4) translocation. (a) Representative western blot of proteins in myocardial insulin signaling pathway. (b–d) Comparisons of (b) phospho/total ratio of insulin receptor substrate 1 (IRS1), (c) phosphatidylinositol 3-kinase (PI3K) and (d) protein kinase B (AKT). (e, f) Representative western blot of proteins involved in GLUT-4 translocation. (g) Comparisons of phospho/total ratio of AKT substrate of 160 kDa (AS160). (h) Comparisons of phospho/total ratio of TBC1D1. (i) Comparisons of vesicle associated membrane protein 2 (VAMP2), synaptosome-associated protein of 23 kDa (SNAP23) and syntaxin 4 expression, the value of the control group was used as the standard (100%). Data are expressed as mean \pm standard deviation for n = 10 per groups. ^{**}P < 0.01 sham versus control; [#]P < 0.05, ^{##}P < 0.01 DJB versus sham; [&]P < 0.05 DJB versus control.

and hyperinsulinemia to ensure maximal stimulation of myocardial glucose transport. The third round was carried out to better simulate postprandial hyperglycemia. The results confirmed that DJB significantly restored the MGU in diabetic rats.

GLUT-1 and GLUT-4 are responsible for basal and insulinstimulated MGU, respectively¹⁶. However, the alterations of GLUTs in DCM remained inconsistent^{10,33}. In the present study, similar levels of total GLUT-1 and GLUT-4 were observed among groups. However, we found attenuated levels of plasma membrane GLUT-4 in the sham group, which was consistent with previous studies^{12,32}. Furthermore, we had a novel finding that DJB could restore the myocardial expression of membrane GLUT-4. As the total GLUT-4 levels were similar among groups, the GLUT-4 defect in the sham group and rehabilitation after DJB should occur at the translocation process. PI3K/AKT is the central hub of signal transduction in the myocardial insulin signaling pathway³⁴, it receives the upstream signal of IRS family and plays a central role in promoting GLUT-4 translocation³⁵. The sham group in the present study had serious myocardial insulin resistance, which is consistent with previous studies^{5,31,36}. Furthermore, the DJB group showed restored insulin sensitivity with increased insulin receptor substrate 1 phosphorylation and intensified PI3K/AKT activation, showing DJB could alleviate myocardial insulin resistance at the molecular level.

The detailed mechanism regarding how exactly DJB interferes with myocardial insulin signaling pathway remains unclear. The present results showed increased levels of plasma GLP-1 after DJB, which was considered to be due to the accelerated nutrient delivery to the distal ileum where GLP-1 is secreted³⁷. A previous study reported that GLP-1 had direct effects on MGU and GLUT-4 translocation during reperfusion³⁸. More recent studies further confirmed that GLP-1 carried out its cardioprotective role in alleviating myocardial insulin resistance by activating the PI3K/AKT pathway^{39–41}. Taken together, GLP-1 probably plays an important role in the alleviation of myocardial insulin sensitivity after DJB. Future studies are warranted to confirm this hypothesis.

There is still a gap between upstream insulin signaling and downstream GLUT-4 translocation⁴². Two novel substrates of AKT, AS160 and TBC1D1, have been the major candidates potentially bridging this gap⁴³. They are essentially functioning as a brake on GLUT-4 vesicle exocytosis. The phosphorylation of AS160 and TBC1D1 by AKT in response to insulin results in the conversion of some downstream Rab proteins to the active guanosine-5'-triphosphate-bound form, and thus triggers the GLUT-4 translocation to the plasma membrane^{43,44}. Studies have confirmed the role of AS160 and TBC1D1 in skeletal muscle^{45,46}. However, there has been limited research into their role in DCM to date. In the present study, the phosphorylation and activation of AS160 were restored after DJB, which could play a pivotal role in the facilitation of GLUT-4 translocation.

The docking and fusion of GLUT-4 vesicles with the cell membrane represent the final step of GLUT-4 translocation. This process is mainly driven by soluble N-ethylmaleimide-sensitive factor attachment protein receptor (SNARE) complex assembly between the GLUT-4 vesicle-localized v-SNARE vesicle-associated membrane protein 2 and the plasma membrane-localized t-SNARE syntaxin 4, along with synaptosome-associated protein of 23 kDa^{17,47}. In the present study, the three groups showed similar levels of vesicle-associated membrane protein 2, Syntaxin4, and synaptosome-associated protein of 23 kDa, showing that both impaired MGU in diabetic rats and restored MGU after DJB are independent of the quantity of SNARE proteins. Further studies are required to validate the physiological relevance of the current findings to humans.

As glucose and fatty acid metabolism could be tightly coupled and inversely regulated in the myocardium⁴⁸, it is likely that DJB results not only in the improvement in MGU, but also in a decrease in lipotoxicity. CD36 acts as a major facilitator of myocardial fatty acids uptake in humans and rodents⁴⁹. Persistent relocation of CD36 to the sarcolemma is observed in the diabetic myocardium⁵⁰. The present results are consistent with previous report and further show a net internalization of CD36 after DJB compared with Sham surgery, which could account, at least in part, for the alleviation of myocardial lipid accumulation. Further studies are warranted to explore the effects of DJB surgery on myocardial fatty acid metabolism.

One potential limitation of the present study was the lack of detailed myocardial metabolic alterations. Although recovery in MGU and the balance of cardiac metabolism have been proved to effectively reverse DCM¹⁴, there are still gaps between enhanced myocardial GLUT-4 on the membrane and the final amelioration of cardiac function. Future studies focusing on myocardial metabolic profiling after DJB and associated effects on DCM would be of vital significance.

In conclusion, the present study showed that DJB surgery could restore the MGU in diabetic rats by facilitating myocardial GLUT-4 translocation, thus contributing to the morphological and functional alleviation of DCM.

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DISCLOSURE

The authors declare no conflict of interest.

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SUPPORTING INFORMATION

Additional supporting information may be found online in the Supporting Information section at the end of the article.

- Appendix S1 | Detailed method of positron-emission tomography scan and image processing.
- Appendix S2 | Detailed method of immunohistochemistry and western blotting.
- Figure S1 | Summary of the study protocol.
- Figure S2 | A potential model for the mechanism of upregulated myocardial glucose uptake after duodenal-jejunal bypass.
- Table S1 | Detailed information of primary antibodies used.
- Table S2 | Animal characteristics for the control and diabetic groups before surgery.