

# Upregulation Of Renal GLUT2 And SGLT2 Is Involved In High-Fat Diet-Induced Gestational Diabetes In Mice

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**Introduction:** Gestational diabetes mellitus (GDM) is a metabolic disorder during mid- to late-pregnancy characterized by hyperglycemia, insulin resistance and fetal mal-development. Glucose transporter type 2 (GLUT2) and sodium-coupled glucose cotransporters 2 (SGLT2) in the proximal tubules play a critical role in the reabsorption of glucose and have been linked to the occurrence of type 2 diabetes mellitus (T2DM). Our study was designed to investigate the role of GLUT2 and SGLT2 in the pathogenesis of GDM, which is considered a forerunner of T2DM, and investigate the related molecular mechanism.

**Methods:** High-fat diet (HFD) was utilized to build a GDM mouse model that closely induces metabolic abnormalities similar to human GDM. Body weight, blood glucose and serum insulin were recorded in the experimental process. Glucose tolerance was determined by the use of an intraperitoneal glucose tolerance test (IPGTT). In addition, levels of GLUT2 and SGLT2 were evaluated to further explore the underlying mechanism of GDM.

**Results:** HFD feeding induced abnormal glucose metabolism as manifested by increased levels of blood glucose and insulin and prominent glucose intolerance. Additionally, fetal mice from mother feed on HFD showed higher mean body weight. Furthermore, HFD feeding led to an increase in the number of positive cells of GLUT2 and SGLT2 in the renal proximal tubule and the expressions of renal GLUT2 and SGLT2 mRNA and proteins in mice. However, no obvious change was observed in renal morphology.

**Conclusion:** Our study demonstrates a potential involvement of renal GLUT2 and SGLT2 in GDM pathology in an HFD-induced GDM mouse model, which further supports the role of renal GLUT2 and SGLT2 not only in T1DM and T2DM but also in GDM.

**Keywords:** gestational diabetes mellitus, insulin resistance, renal threshold for glucose, GLUT2, SGLT2

## Introduction

Gestational diabetes mellitus (GDM) is defined as abnormal glucose metabolism that is first discovered or diagnosed during pregnancy.<sup>1</sup> As most patients with GDM usually do not show obvious clinical manifestations, GDM screening has become a routine prenatal examination using a glucose tolerance test (OGTT) during the second trimester of pregnancy. It has been reported that GDM affects 5% to 10% of pregnant women depending on different screening methods, diagnostic criteria and the population screened over the past decades.<sup>2,3</sup> GDM is not only associated with higher risk of maternal postpartum type 2 diabetes mellitus (T2DM) but also related to an increased frequency of potential adverse outcomes and long-term

metabolic dysregulation in offspring.<sup>4-7</sup> However, related pathological mechanisms underlying GDM are still not understood.

Similar to T2DM, previous reports have suggested that the main pathophysiologic mechanisms behind GDM are primarily  $\beta$ -cell dysfunction and insulin resistance.<sup>8</sup> While recent studies report that women with GDM may suffer dysfunction of physiological decline in the renal threshold for glucose ( $RT_G$ ) during pregnancy.<sup>9</sup>  $RT_G$  is defined as the point of glucose concentration in the blood at which the kidney will excrete glucose (see glycosuria), which is normally stable within a certain range<sup>10</sup> and may change in some cases, such as T1DM and T2DM.<sup>11,12</sup> The kidney is critical to the balance of glucose homeostasis through filtration and reabsorption of glucose. Normally, the kidneys filter approximately 180g of glucose per day, all of which is reabsorbed into the circulation through glucose transporter proteins in cell membranes within the proximal tubules. Glucose carrier proteins mainly involve 2 major classes: the glucose transporters (GLUTs) and the sodium-coupled glucose cotransporters (SGLTs).<sup>13,14</sup> Of the various GLUT and SGLT proteins in the kidneys, GLUT2 and SGLT2 are considered to be more crucial in the glucose regulation. SGLT2 is responsible for reabsorbing approximately 90% of the glucose filtered at the glomerulus in the proximal tubular cells, and GLUT2 promotes reabsorption glucose back into circulation.<sup>15</sup> The role of the kidney in glucose control has been increasingly recognized, due to its involvement in the occurrence of T2DM and its complications. Most prior evidence has mainly concentrated on the role of GLUT2 and SGLT2 in T2DM.<sup>13,16</sup> Recently, it is suggested that upregulation of renal SGLT2 may be associated with GDM.<sup>9</sup>

Stable and mature animal model of GDM is the basis of exploring the pathophysiologic aspects of human GDM. Previous studies have reported several animal models available to study GDM, including induced, spontaneous and transgenic animal models.<sup>17,18</sup> Spontaneous and transgenic animal models commendably develop hyperglycemia, which is close to human GDM.<sup>19</sup> While the method is not widely used in research due to expensive animal cost, limited species and individual differences. Streptozotocin (STZ) induces a high or moderate hyperglycemia by the destruction of pancreatic  $\beta$ -cells depending on different dose, which is commonly used in animal study in the past.<sup>20</sup> However, it is difficult to control the dose of STZ during the molding process to induce an expected blood glucose level. In recent years, increasing evidence

suggests the availability of high-fat diet (HFD) in the establishment of GDM animal model.<sup>21</sup> The model is prevalent due to its convenience and similarity to GDM in human.

The present study establishes a mouse model of GDM using HFD and explores the role of renal GLUT2 and SGLT2 in the development of GDM.

## Materials And Methods

### Animals And Study Design

All experimental procedures involving animals were in accordance with the NIH guidelines for the Care and Use of Laboratory Animals, and the study protocols were approved by Animal Care and Use Committee of Shanghai Jiao Tong University.

Female clean healthy C57BL/6 mice aged 6 weeks and weighing 16–18 g were employed for the present study. The animals were housed in a controlled environment ( $21 \pm 2^\circ\text{C}$ ,  $50 \pm 10\%$  humidity, 12 hrs light/dark cycle) with food and water available ad libitum. After adaptive feeding for 1 week, female mice were randomly divided into two experimental groups ( $n = 30$  per group): control or HFD group. Mice consumed control rodent diet (10% kcal fat; Research Diets, New Brunswick, NJ) or HFD (45% kcal fat; Research Diets, New Brunswick, NJ) (Table 1). After 6-week dietary intervention, mice in control group were divided into two subgroups ( $n = 15$ ): control virgin

**Table 1** Formula And Nutrient Of Normal And High-Fat Diets

Formula (g)	Normal Diet	High-Fat Diet
Casein	200	200
L-Cystine	3	3
Corn Starch	315	72.8
Maltodextrin 10	35	100
Sucrose	350	172.8
Cellulose	50	50
Soybean Oil	25	25
Lard	20	177.5
Mineral Mix S10026	10	10
DiCalcium Phosphate	13	13
Calcium Carbonate	5.5	5.5
Potassium Citrate	16.5	16.5
Vitamin Mix V10001	10	10
Choline Bitartrate	2	2
Nutrient (g%)		
Protein	19.2	24
Carbohydrate	67.3	41
Fat	4.3	24

group (CV) and control pregnant group (CP), and mice in HFD group were divided into HFD virgin group (HV) and HFD pregnant group (HP) ( $n = 15$ ). Female mice in CP and HP groups were mated with males of the same genotype in a ratio of 1:2. The next morning, female rats were observed for the presence or absence of vaginal suppositories, which were taken with a cotton swab and observed further under the microscope. If sperms were found in three different fields, the female rate was marked as positive for pregnancy, and the date was marked as gestation day (GD) 0. The mating procedure lasted for 1 week which comprised approximately one estrous cycle. Non-pregnant female mice in this period were considered infertile and excluded from the study. Fortunately, all 30 female rates were found pregnant. Then, the mice in HV and HP groups continued feeding HFD until GD 18.

## Measurement Of Body Weight, Blood Glucose, And Serum Insulin

Body weight, blood glucose, and serum insulin were monitored at different time points, including before dietary intervention, after 6 weeks of HFD, and on GD 0, 10, and 18. Blood glucose and insulin levels were determined from tail venipuncture blood samples. Blood glucose concentration was measured immediately using a blood glucose meter and strips (OneTouch Ultra; Johnson & Johnson, Langhorne, PA). The blood samples were then centrifuged at low speed (4°C, 5000 rpm, 15 mins) within 1 hr; the supernatant was harvested and stored at -80°C for measuring serum insulin level by enzyme-linked immunosorbent assay (ELISA; Mercodia AB, Uppsala, Sweden) according to the manufacturer's instructions. In addition, urine volume and fluid intake were observed daily.

## Glucose Tolerance Test

Glucose tolerance was determined by an intraperitoneal glucose tolerance test (IPGTT). Mice were fasted for 12 hrs with free access to water and intraperitoneally injected with glucose (1.5 g/kg body weight). Blood samples were collected from the tail vein at 0, 15, 30, 60 and 120 mins after glucose administration. Blood glucose and insulin levels were measured instantly using methods as mentioned above. Meanwhile, area under the curve (AUC) of blood glucose and insulin was calculated.

## Hematoxylin And Eosin (HE) Staining

On GD 18, all of the mice were anesthetized by intraperitoneal injection of overdose Nembutal and sacrificed. We

chose GD 18 as the time point because the general gestation period of mice is 17–19 days.<sup>17,22</sup> Then cesarean section was performed, and the number and weights of fetuses in CP and HP groups were recorded. In each group, 7 out of 15 mice were sacrificed to obtain the kidney for HE staining or immunohistochemistry examination, while the rest were subjected to mRNA or protein analysis. The protocol was described as previously.<sup>23</sup> Morphologic changes of glomeruli and renal tubules were observed under light microscopy (BX50, Olympus Corporation, Tokyo, Japan).

## Immunohistochemistry Analysis

Bouin-fixed kidney from each group was processed, paraffin embedded, and sectioned at 5  $\mu$ m for GLUT2 and SGLT2 immunohistochemistry. Sections were dewaxed in xylene and hydrated using a graded series of ethanol. Antigenic retrieval was performed by immersing mounted tissue sections in 0.01 mM sodium citrate buffer (pH 6.0) and heating in an autoclave (121°C) for 5 mins. Deparaffinized sections were blocked for 1 hr in normal goat serum, followed by incubation in rabbit anti-GLUT2 and -SGLT2 polyclonal antibodies (1:100; Santa Cruz Biotechnology, Santa Cruz, CA, USA) overnight in a humidified chamber at 4°C. Subsequently, the tissue slides were incubated with peroxidase-conjugated secondary antibody (Vector Laboratories, Burlingame, CA, USA) at room temperature for 30 mins and visualised by incubation with a 3,3'-diaminobenzidine tetrahydrochloride (DAB; Vector Laboratories, Burlingame, CA, USA) at room temperature for 5 mins. The sections were then counterstained with hematoxylin (Merck KGaA, Darmstadt, Germany) and observed by light microscopy (BX50, Olympus Corporation, Tokyo, Japan).

## Western Blot Analysis

Kidney tissues were harvested and were homogenized on ice in the presence of protease and phosphatase inhibitors. Homogenates were centrifuged at 12,000  $\times$  g at 4°C for 15 mins. Protein concentration in supernatants was quantified by the BCA method using bovine serum albumin (BSA) as the standard. Proteins were analyzed by 10% SDS-PAGE and transferred to PVDF membranes that were incubated in 5% non-fat milk at room temperature for 1 hr, then incubated with appropriate primary and secondary antibodies. Membranes were washed and proteins were detected by enhanced chemiluminescence (ECL) using a LAS-4000 lumino-image analyzer (Fuji Film, Tokyo, Japan). Bands

were digitally scanned and analyzed using ImageJ software (NIH Image, National Institutes of Health, Bethesda, MD, USA).

## Real-Time Quantitative Polymerase Chain Reaction

Total RNA was extracted from the kidney using TRIzol® Reagent (Invitrogen, Carlsbad, CA, USA). cDNA was synthesized from 1 µg of RNA using the PrimeScript™ RT Reagent Kit (Takara Biotechnology Co., Ltd., Dalian, China) according to the manufacturer's instructions. Amplification was performed on a 7500 Real-Time PCR System (Applied Biosystems, Foster City, CA, USA) using SYBR® Premix Ex Taq™ (Takara Biotechnology Co.) according to the manufacturer's instructions. Primers used for quantitative PCR were shown as follows:

GLUT2 (forward: GGAGCCTCCAGTAAGAAGTCTG 5'-3', reverse: TGGCAGGTAGAATTAGTCTCAGG 5'-3'); SGLT2 (forward: TTTTGGTGGTTGTGCTGG 5'-3', reverse: CTCGGAAGATGTGGAAGG 5'-3'); *Glyceraldehyde 3-phosphate dehydrogenase* (GAPDH) (forward: CCCTCTG GAAAGCTGTGG 5'-3', reverse: AGTGGATGCAGGGA TGATG 5'-3'). Relative changes in gene expression were determined using the  $2^{-\Delta\Delta Ct}$  method, with the housekeeping gene GAPDH as an internal control.

## Statistical Analysis

All data were calculated as means  $\pm$  SD and checked using the Kolmogorov–Smirnov (KS) test before further analysis. Statistical significance between two datasets was assessed using the Student's *t*-test. Multiple groups were compared using one-way ANOVA followed by Tukey multiple comparison testing. A *P* value of  $<0.05$  was considered statistically significant. All statistical tests were performed using GraphPad Prism Version 6.0 (GraphPad Prism Software, Inc. CA, USA).

## Results

### Changes Of Body Weight, Blood Glucose, And Serum Insulin In Mice

Body weight was determined at different time points for all groups. As indicated in Figure 1A, the body weight showed an increasing trend after 6 weeks of HFD, and a rapid elevation of body weight was found in the mice of HP group compared to the moderate increase in CP group ( $P < 0.05$ ). The weight at GD 18 and total weight gain of mice in HV and HP groups were significantly higher than that of CV and CP

groups ( $P < 0.05$ , Figure 1B). Next, we examined the blood glucose and serum insulin in these mice. Blood glucose levels exhibited a gradual upregulation at the end of 6-week HFD feeding and during pregnancy in the mice of HV and HP groups but not CV or CP group ( $P < 0.05$ , Figure 1C). Similar to the trend of blood glucose change, serum insulin levels were enhanced at the end of HFD feeding in HV and HP groups and continued to elevate throughout pregnancy ( $P < 0.05$ , Figure 1E). Moreover, blood glucose and serum insulin at GD 18 were significantly advanced in HV mice in contrast to the CV mice ( $P < 0.05$ , Figure 1D and F). Interestingly, blood glucose levels were deregulated at the end of pregnancy in CP mice but further increased in HP mice ( $P < 0.05$ , Figure 1D), whereas insulin levels were increased in both CP and HP mice and showed a more pronounced enhancement in the mice of HP group ( $P < 0.05$ , Figure 1F). In addition, the urine volume and fluid intake were increased significantly during mid- to late-pregnancy in HP mice compared to CP mice. However, there was no significant change in the mice of CV or HV group.

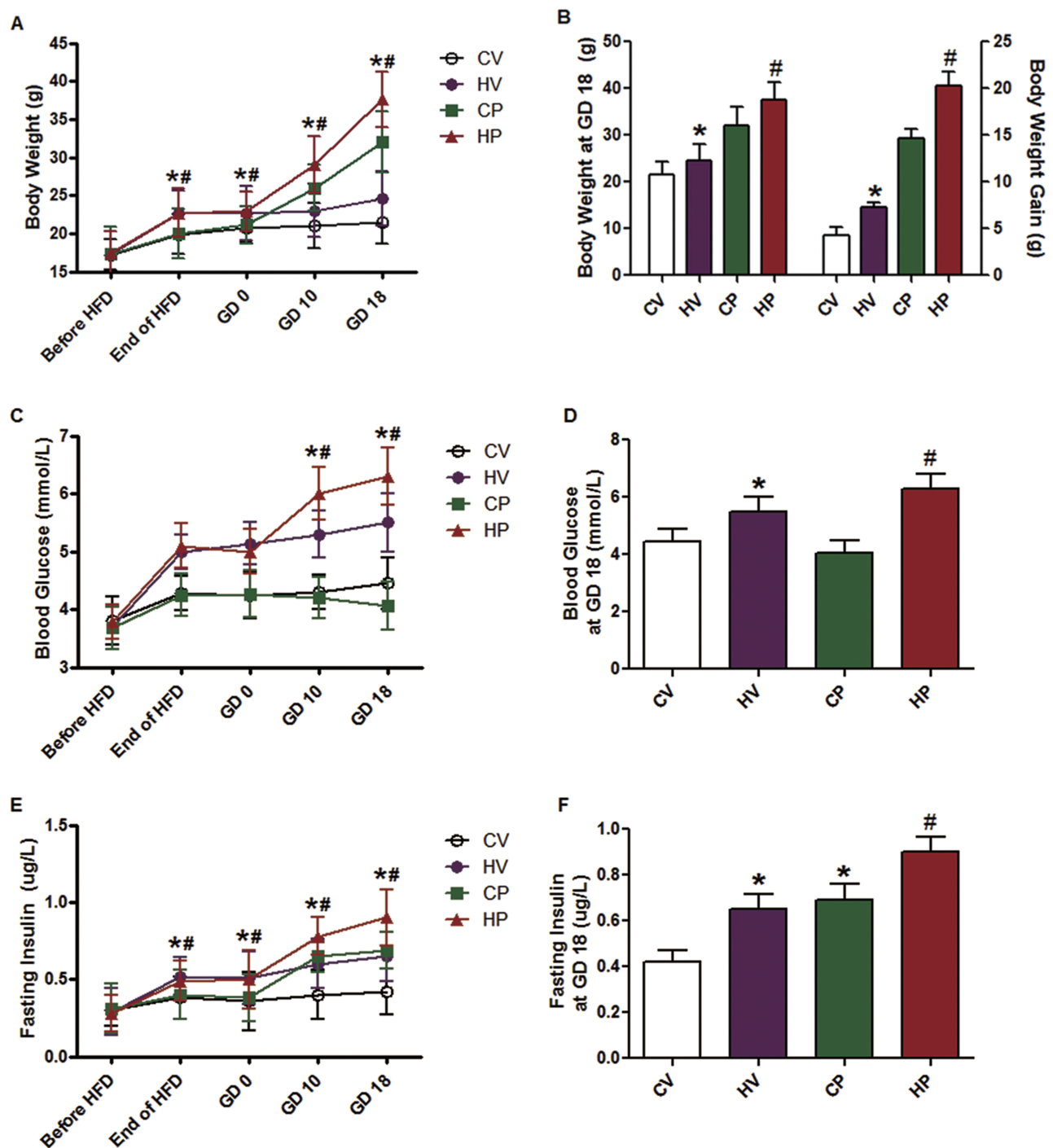
### Glucose Tolerance Test In Mice

By performing a glucose tolerance test at the end of HFD feeding, we found that blood glucose and insulin levels of mice in HV and HP groups indicated a slight increase without statistically significant differences, suggesting HFD for 6 weeks did not cause dramatical changes in either glucose or insulin levels (Figure 2A and B). Then, we conducted the same test at GD 18. As shown in Figure 2C, compared with CP group, the mice in HP group showed impaired glucose tolerance, as manifested by obviously increased glucose levels after glucose injection ( $P < 0.05$ , 15 and 30 mins). Consistently, insulin levels of HP group mice were also dramatically higher than that of CP throughout the test ( $P < 0.05$ , Figure 2D). These results suggested successful establishment of a mouse model of GDM. However, no significant upregulation was observed in the glucose or insulin levels in HV mice compared with CV mice.

### Reproductive Outcome Of Pregnant Mice

The number of fetal mice was counted for each female from CP and HP groups. However, there was no significant differences in average litter size between CP and HP groups (Figure 3A). Moreover, we determined the body weight of fetal mice and found that mean body weight of fetal mice born by HP mothers was significantly higher than those by CP mothers ( $P < 0.05$ , Figure 3B).



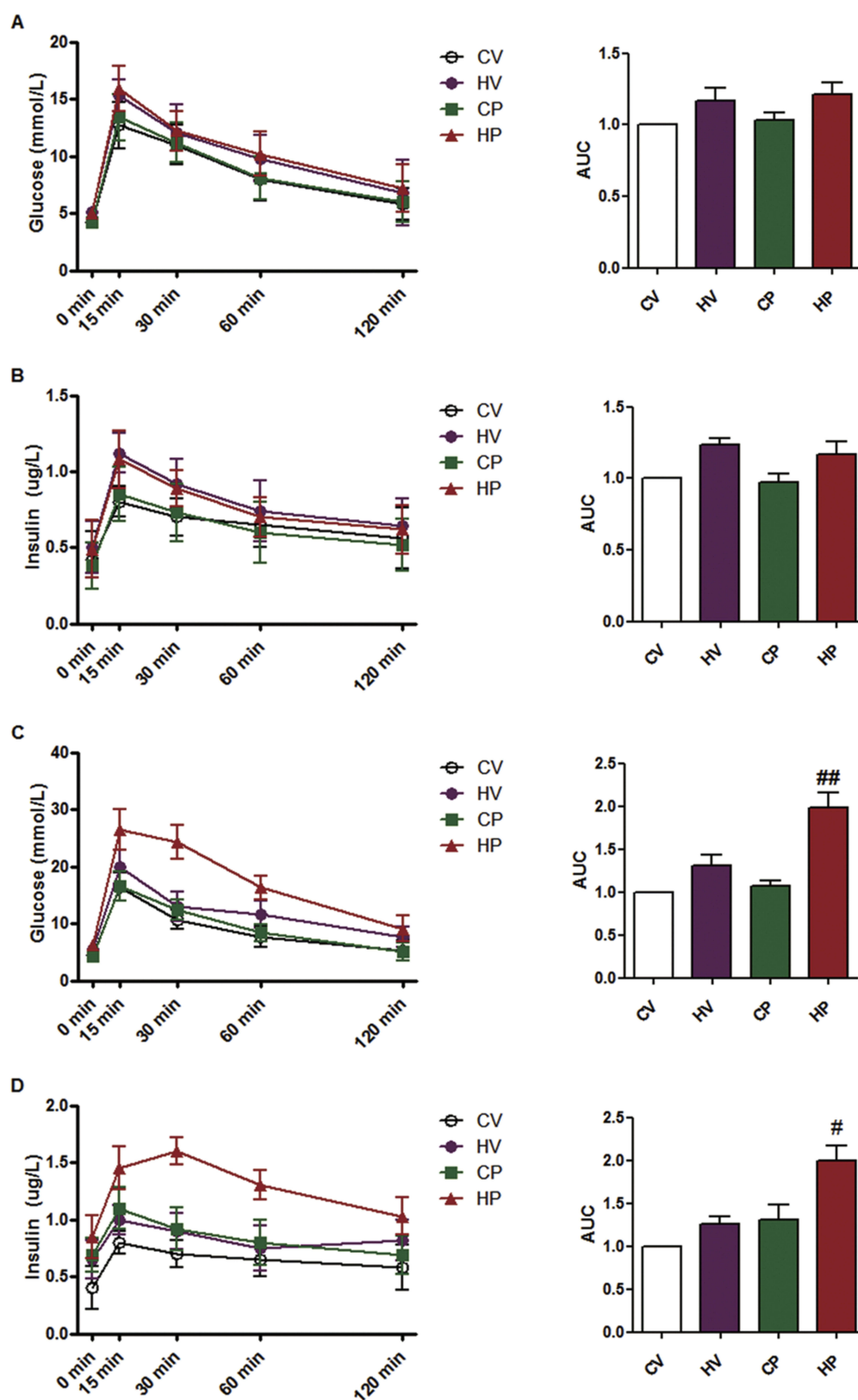


**Figure 1** Changes of body weight (A, B), blood glucose (C, D), and serum insulin (E, F) in mice of different groups before dietary intervention, after 6 weeks of HFD, and on GD 0, 10, and 18. Data are expressed as the mean  $\pm$  SD values ( $n = 15$ ). \* $P < 0.05$  vs CV group, # $P < 0.05$  vs CP group.

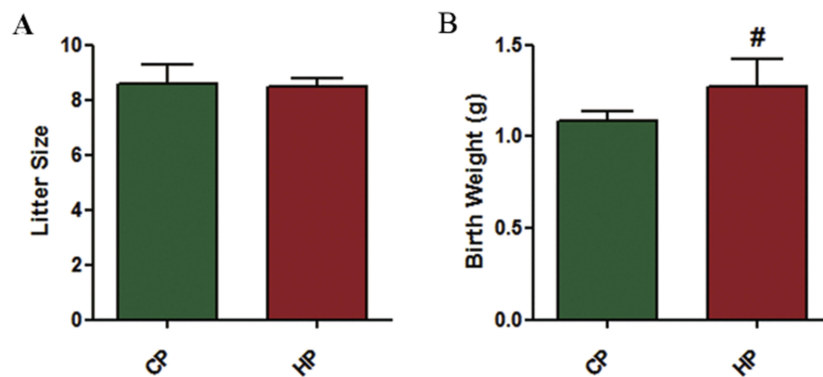
## Changes Of Renal Morphology In Mice

Representative graphs of renal morphology of the four groups are shown in Figure 4. It revealed that glomeruli of the mice in the CV group were normal with clear Bowman capsule and normal thickness of the glomerular basement membrane. The mesangial matrix, mesangial

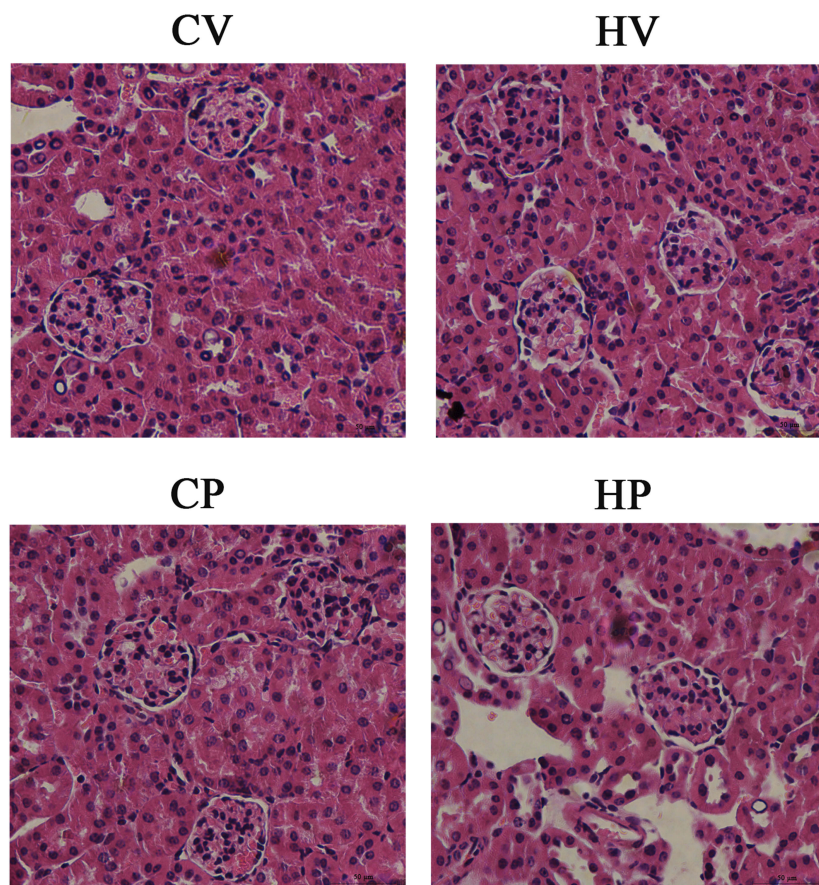
cells and endothelial cells showed no significant proliferation. Renal tubular epithelial cells displayed mild swelling, and renal interstitial vascular showed mild congestion with no obvious inflammatory cell infiltration. Renal morphology in the other three groups did not demonstrate significant changes compared with CV group.



**Figure 2** Glucose tolerance test. Blood glucose (A) and serum insulin levels (B) during intraperitoneal glucose tolerance test in the four groups after 6 weeks of HFD. Blood glucose (C) and serum insulin levels (D) during intraperitoneal glucose tolerance test in the four groups on GD 18. Area under the curve (AUC) of blood glucose and serum insulin was calculated. Data are expressed as the mean  $\pm$  SD values ( $n = 15$ ). <sup>#</sup> $P < 0.05$ , <sup>##</sup> $P < 0.01$  vs CP group.



**Figure 3** Reproductive outcome of pregnant mice. Litter size (A) and body weight at birth (B) of fetal mice by each female mouse from CP and HP experimental groups were recorded. Data are expressed as the mean  $\pm$  SD values ( $n = 15$ ). <sup>#</sup> $P < 0.05$  vs CP group.



**Figure 4** Changes of renal morphology. The kidney was stained with HE for morphologic analysis under light microscopy. Morphologic changes of glomeruli and renal tubules were observed under a high magnification field ( $\times 400$ ).

### Changes Of Immunohistochemical Staining And Protein Expression Of Renal GLUT2 And SGLT2 In Mice

We speculate the reason for increased blood glucose in HV and HP mice could be attributed to elevated expression of renal GLUT2 and SGLT2. Therefore, we sacrificed the

mice at GD 18 and harvested their kidneys to detect the expression changes of GLUT2 and SGLT2 by immunohistochemistry, RT-qPCR and Western blot analysis. The positive staining of GLUT2 and SGLT2 is mainly located in the renal proximal tubule. As expected, positive cells of GLUT2 and SGLT2 were increased in the renal proximal

tubule of HV and HP mice compared with CV and CP mice (Figure 5A). Consistent with the trend of immunohistochemical staining, the results also demonstrated significantly higher expressions of SGLT2 and SGLT2 mRNA and protein in mice of HV and HP groups than that of CV and CP groups ( $P < 0.05$ , Figure 5B and C).

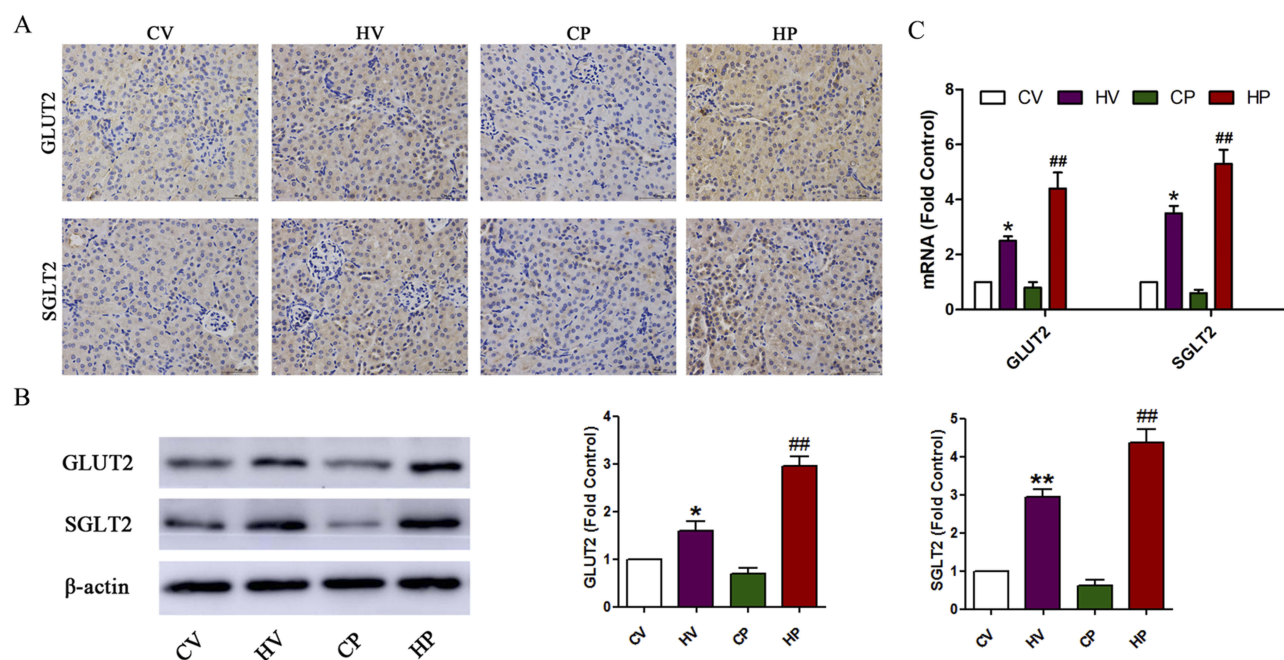
## Discussion

In the present study, we determined the role of renal GLUT2 and SGLT2 in the pathophysiologic mechanisms of GDM through employing an HFD-induced GDM mouse model. The outcome of this study showed that HFD feeding before and during pregnancy in mice may lead to physiological and pathological performances close to human GDM. HFD intervention increased body weight and induced abnormal glucose metabolism as manifested by increased blood glucose and serum insulin and prominent glucose intolerance in mice, leading to various acute or long-term adverse consequences in offspring. Moreover, HFD upregulated levels of renal GLUT2 and SGLT2, which demonstrates the potential role of GLUT2 and SGLT2 in the pathogenesis of GDM in mice.

GDM is defined as any degree of glucose intolerance with onset or first recognition during pregnancy.<sup>1</sup> In the second and third trimester, due to the physiological insulin

resistance, maternal insulin secretion is elevated to maintain blood glucose levels. Impairment of pancreatic  $\beta$ -cell function or compensatory increases of insulin secretion or both leads to GDM.<sup>4</sup> Given the prevalence of GDM, a growing body of research has carried out to investigate the physiological and pathological mechanisms of GDM both in animal and human.<sup>9,24–28</sup> However, due to the specificity of pregnancy, it is difficult to carry out some invasive research, limiting further understanding of GDM. Therefore, an ideal GDM animal model will undoubtedly provide a good research foundation and platform for further study of etiology, pathogenesis and effective prevention and treatment of GDM.

In recent years, much evidence has been accumulated to indicate that dietary factors play a critical role in the pathogenesis of GDM.<sup>29–31</sup> As part of a classical western lifestyle, HFD has been indicated to be closely connected with glucose disturbance and insulin resistance, which subsequently leads to T2DM.<sup>32</sup> And HFD-induced GDM animal model is more consistent with the clinical features of diabetes, which replicates the pathogenesis of diabetes by changing diet structure to increase insulin secretion, reduce insulin sensitivity, and ultimately result in insulin resistance.<sup>4</sup> Accordingly, in the current study, we employed HFD intervention in an effort to develop an ideal GDM model in mice. Moreover, NV and



**Figure 5** Changes of renal GLUT2 and SGLT2 levels. **(A)** Immunohistochemical staining of renal GLUT2 and SGLT2. Coronal kidney sections were stained with GLUT2 and SGLT2 antibodies and observed under a high magnification field ( $\times 400$ ). **(B)** Representative immunoblot and quantification of GLUT2 and SGLT2 in the kidney of mice from the four groups. **(C)** mRNA expression levels of GLUT2 and SGLT2 in the kidney of mice from the four groups. Data are expressed as the mean  $\pm$  SD values ( $n = 15$ ). \* $P < 0.05$ , \*\* $P < 0.01$  vs CV group, ### $P < 0.01$  vs CP group.



HV groups were established to exclude the influences of placental secretion of various hormones and enzymes in pregnancy and further demonstrate HFD-induced insulin resistance in pregnancy.

Obesity and overweight take the top spot in the various risk factors for the development of diabetes, and weight gain during pregnancy increases the incidence of T2DM in pregnant women.<sup>33</sup> Consistently, our results indeed confirmed that HP mice exhibited dramatically elevated weight gain compared to CP mice. Along with the increase of body weight, blood glucose and insulin showed a corresponding elevation, especially in mid- and late-pregnancy, during which GDM usually occurs. Meanwhile, we found that HFD feeding for 6 weeks did not cause significant insulin resistance, although it induced an increase in body weight and serum insulin. However, insulin resistance was evident during late pregnancy as indicated by prominent glucose intolerance. Increased insulin resistance, the so-called physiological insulin resistance, is common during pregnancy to maintain the glucose for the fetus, resulting in elevated insulin levels.<sup>4</sup> Elevated levels of various pregnancy-related hormones may account for this phenomenon.<sup>8,34</sup> However, failure to upregulate insulin production to overcome the further aggravated degree of insulin resistance leads to hyperglycemia or finally GDM during pregnancy.<sup>35</sup> Our results demonstrated that HFD feeding contributed to pathologic performances in mice close to the human condition of GDM.

Insulin resistance induced by hormonal and adipokine secretion from the placenta contributes to the development of GDM.<sup>8</sup> Exploration of potential pathophysiological mechanisms and new therapeutic targets is of great importance to the prevention and treatment GDM. Recent study suggested that impairment in the decrease of the  $RT_G$  may have a place in the development of GDM.<sup>9</sup>  $RT_G$  is dependent on the renal glucose reabsorptive capacity, which is associated with GLUTs and SGLTs in the proximal tubules. Emerging evidence suggests that expression of GLUT2 or SGLT2 or both was increased in hyperglycemic rodents and humans,<sup>13,14,16</sup> which may lead to increased reabsorption of renal glucose in patients with T1DM and T2DM.<sup>11,36</sup> On the contrary,  $RT_G$  has been indicated to be lower in pregnant women than non-pregnant women,<sup>37,38</sup> which may be explained by high renal blood flow and decreased renal glucose transporter expression.<sup>37,38</sup> Similarly, recent study demonstrated that  $RT_G$  in normal glucose tolerance pregnancy was significantly lower than that in the GDM women, as well as the healthy controls.<sup>9</sup> And the relatively high  $RT_G$  in GDM may attribute to upregulation of renal glucose transporter expression including GLUT2 and

SGLT2.<sup>9</sup> Consistent with the findings, in an HFD-induced GDM model, our results suggested that expressions of GLUT2 and SGLT2 were enhanced significantly in HP mice as indicated by increased number of positive cells of GLUT2 and SGLT2 in the renal proximal tubule and levels of renal GLUT2 and SGLT2 mRNA and protein. The enhancement of renal glucose transporter expression may contribute to the elevated blood glucose level, which further aggravates insulin resistance and eventually leads to GDM.

## Conclusions

In conclusion, the present study indicated a potential involvement of renal GLUT2 and SGLT2 in the pathology of GDM in an HFD-induced GDM mouse model. As far as we know, our study is the first study which explores a probable connection between renal glucose transporter and GDM. The current findings display new evidence supporting the role of renal GLUT2 and SGLT2 in diabetes. A better understanding of the renal involvement in GDM may provide new evidence for the pathogenesis of GDM and open new horizons for pharmaceuticals developments.

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## Disclosure

The authors declare no conflicts of interest, financial or otherwise in this work.

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