Adipocytes play an etiological role in the podocytopathy of high-fat diet-fed rats

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Abstract

Obesity is a risk factor that promotes progressive kidney disease. Studies have shown that an adipocytokine imbalance contributes to impaired renal function in humans and animals, but the underlying interplay between adipocytokines and renal injury remains to be elucidated. We aimed to investigate the mechanisms linking obesity to chronic kidney disease. We assessed renal function in high-fat (HF) diet-fed and normal diet-fed rats, and the effects of preadipocyte- and adipocyte-conditioned medium on cultured podocytes. HF diet-fed and normal diet-fed Sprague Dawley rats were used to analyze the changes in plasma BUN, creatinine, urine protein and renal histology. Additionally, podocytes were incubated with preadipocyte- or adipocyte-conditioned medium to investigate the effects on podocyte morphology and protein expression. In the HF diet group, 24 h urinary protein excretion (357.5 ± 64.2 mg/day vs 115.9 ± 12.4 mg/day, \(P<0.05\)) and the urine protein/creatinine ratio were significantly higher (1.76 ± 0.22 vs 1.09 ± 0.15, \(P<0.05\)), increased kidney weight (3.54 ± 0.04 g vs 3.38 ± 0.04 g, \(P<0.05\)) and the glomerular volume and podocyte effacement increased by electron microscopy. Increased renal expression of desmin and decreased renal expression of CD2AP and nephrin were also seen in the HF diet group \((P<0.05)\). Furthermore, we found that adipocyte-conditioned medium-treated podocytes showed increased desmin expression and decreased CD2AP and nephrin expression compared with that in preadipocyte-conditioned medium-treated controls \((P<0.05)\). These findings show that adipocyte-derived factor(s) can modulate renal function. Adipocyte-derived factors play an important role in obesity-related podocytopathy.
Introduction

Overweight and obesity now constitute a global public health crisis. They associate with metabolic syndrome, abdominal obesity, glucose intolerance, insulin resistance, lipid profile abnormalities and elevated blood pressure (Cerezo et al. 2013). The dominant risk factors for metabolic syndrome are abdominal obesity and insulin resistance. Epidemiological studies have revealed a significant association between insulin resistance and obesity, pathological hyperplasia and/or hypertrophy of adipose tissue (Galic et al. 2010). Adipose tissue is an important source of metabolically active secretory products, including free fatty acids, leptin, tumor necrosis factor-α (TNF-α), interleukin-6 (IL-6), plasminogen activator inhibitor-1, resistin and adiponectin, which can lead to the development of cardiovascular diseases (Galic et al. 2010, Van de Voorde et al. 2013).

An imbalance of adipokines in patients with chronic kidney disease (CKD) is a consequence of impaired renal function, abnormal hormone plasma levels and altered regulatory mechanisms caused by the uremic milieu (Mills et al. 2013). Leptin, a small peptide (16 kDa) belonging to the IL-6 family of cytokines, acts as a coactivator with transforming growth factor (TGF)-β to promote renal tubulointerstitial fibrosis (Kümpers et al. 2007). Adiponectin, another small peptide (30 kDa) secreted almost entirely by adipocytes, is the strongest predictor of stages 3–5 CKD according to the K/DOQI guidelines, showing a significant nonlinear inverse association (Nanayakkara et al. 2009). Resistin is an independent factor for CKD in nondiabetic obese patients (Stepien et al. 2013). Recent studies have established that visceral fat associates with cardiometabolic risk caused by dysregulation of adipokine secretion and the associated inflammatory state (Després et al. 2008, Phillips & Prins 2008).

The strong clinical association between metabolic syndrome and the risk of renal impairment is well recognized (Locatelli et al. 2006, Ninomiya & Kiyohara 2007, Heo et al. 2010). Several studies have shown that obesity carries an increased risk of CKD and development of end-stage renal disease (ESRD) (Galber et al. 2005, Obermayr et al. 2008, Satirapoj et al. 2013). CKD is defined as the presence of microalbuminuria (Locatelli et al. 2006) and a decrease in the glomerular filtration rate to <60 mL/min per 1.73 m², which is calculated using the abbreviated modification of diet in the renal disease equation (Levey et al. 1999). In addition, there is a high prevalence of insulin resistance in nondiabetic CKD patients, both in children and adults (Pham et al. 2011, Canpolat et al. 2012), and in chronic dialysis patients (Hung & Ikizler 2011), and its etiology is complex and multifactorial. It is unclear whether insulin resistance is a cause or a consequence of CKD, and more convincing mechanistic evidence is needed to clarify this issue.

Regardless of the human or experimental model of glomerular damage, podocyte loss associates with progressive glomerular diseases, and glomerular volume acts as a key determinant of the total glomerular filtration surface (Fukuda et al. 2012, Weil et al. 2012). Podocytes serve as the final defense against the loss of serum albumin, and podocyte injury is closely related to proteinuria (Mathieson 2011). A reduction in podocyte number and density has been suggested to play an important role in the progression of podocyte dysfunction and the development of proteinuria (Tharaux & Huber 2012). Desmin is a sensitive marker for acute podocyte damage in the glomerulus (Herrmann et al. 2012) and glomerular podocyte damage during the early stages of diabetic nephropathy (Kakimoto et al. 2014). In kidney podocytes, the slit diaphragm proteins nephrin and CD2-associated protein (CD2AP) contribute to the filtration barrier, which prevents the loss of urinary proteins (Asanuma et al. 2007, Xiao et al. 2013). Knight and coworkers found that the expression of glomerular nephrin expression decreases and the urinary albumin level increases in a high-fat (HF) diet model of obesity. Furthermore, hypertension and renal inflammation, which was detected as an increase in renal macrophage-specific CD68-positive staining, were present in the HF diet-fed animals (Knight et al. 2010). Recently, obesity was shown to lead to decreased adiponectin levels, resulting in podocyte injury (Sharma 2009). However, the relationship between obesity and podocyte function has not been fully evaluated.

Body weight gain has been shown to be an independent risk factor for CKD (Nguyen & Hsu 2007, Hsu et al. 2009). In addition, studies have shown that the development of obesity-related CKD is mediated by cardiovascular risk factors (Foster et al. 2008, Munkhaugen et al. 2009). Although there have been several large epidemiological studies of obesity-related CKD, the association between obesity and CKD and the underlying interplay between adipokotkines and renal injury remain to be elucidated. The purpose of this study was to clarify the mechanism linking obesity to CKD.
Materials and methods

Study design

Male Sprague Dawley rats, weighing 200–250 g, were housed four to a cage in an air-conditioned room (22±2°C) on a 12 h light cycle (06:00–18:00 h). Except when scheduled for an oral glucose tolerance test (OGTT) or killing, the animals had free access to food and water. The rats in the HF diet group (n=8) were fed a HF diet (60% fat), while those in the control (C; n=8) were fed standard chow. The experiment lasted for 3 months, and changes in the levels of plasma glucose, insulin, insulin sensitivity, triglycerides, cholesterol, blood urea nitrogen (BUN), creatinine, urine creatinine and albumin, as well as the albumin/creatinine ratio, were measured. Insulin sensitivity was measured using the OGTT. After the 3-month experimental period, the body fat distribution of each rat was evaluated by computed tomography (CT). The animals were then killed, the fat pad weight was measured and adipocytes were isolated for the analysis of adipocyte size and size distribution. We removed both kidneys from each rat and measured their weight. We then removed the kidney cortex for histological analysis and Western blotting. We calculated the mean glomerular volume by the point counting method at a magnification of 400× on hematoxylin and eosin-stained sections as described previously (Adamczak et al. 2003). In addition, the levels of nephrin, CD2AP and desmin in the kidneys were measured by Western blotting. We calculated the mean glomerular volume in each field (magnification, 200×). Three random fields in each section were chosen, and five sections from each rat were used for analysis. Representative images were captured using a digital spot camera. The glomerular volume was calculated as the total area of the glomerular tuft (A) × (1.5) × 1.382/1.01 (1.382 = shape coefficient, 1.01 = size distribution coefficient) (Adamczak et al. 2003).

Cell cultures

Conditionally immortalized mouse podocytes, kindly provided by Prof. Jochen Reiser, MD, PhD (Rush University Medical Center, Chicago, IL, USA), were cultured as described previously (Mundel et al. 1997). These immortalized mouse podocytes can be cultured under different conditions to generate two different phenotypes; under permissive conditions (33°C), they proliferate and maintain an epithelial phenotype with a cobblestone-like morphology (immature podocytes), whereas under nonpermissive conditions (37°C), they stop proliferation and undergo conversion to arborized cells (differentiated or mature podocytes). This differentiation results in the sprouting of long processes from the cell bodies and in the formation of spindle-like projections from these primary processes.

In this study, differentiated podocytes were cultured at 37°C for 8–10 days without IFN-γ in Dulbecco’s modified Eagle’s (DME) medium containing 5.5 mmol/L glucose, 100 units/mL penicillin, 100 μg/mL streptomycin (all from Gibco BRL) and 10% fetal bovine serum (Biological Industries, Kibbutz Be’er Ha’Emek, Israel) in 5% CO₂ and 95% humidified air. The podocytes were plated at 0.2 × 10⁶ cells/well on 24-well plates (Corning) and serum-starved overnight. They were incubated for 24 h with conditioned medium from 3T3-L1 preadipocytes or 3T3-L1 adipocytes, prepared as described below and washed twice with phosphate-buffered saline; RNA was isolated for real-time polymerase chain reaction (RT-PCR) studies.

3T3-L1 preadipocytes (American Type Culture Collection) were seeded on 6-well plates (Falcon, Becton Dickinson, NJ, USA) and grown in complete medium (DME high glucose (25 mmol/L) medium containing 100 U/mL penicillin, 100 μg/mL streptomycin, and 10% fetal bovine serum) in 10% CO₂, with the medium replaced...
every 3 days. After the cells reached confluency by 3 days, they were induced to differentiate for 3 days in complete medium containing 0.5 mM isobutylmethylxanthine, 0.5 µM dexamethasone and 1.7 µM insulin (all from Sigma). The medium was then replaced with complete medium, which was changed every 3 days until the cells were fully differentiated. By day 10, >95% of the preadipocytes differentiated into adipocytes as determined by staining for lipid accumulation using Oil Red O. For the collection of conditioned medium, 3T3-L1 preadipocytes or well-differentiated 3T3-L1 adipocytes were incubated for 24 h in serum-free DMEM medium containing 5.5 mmol/L glucose and 1% bovine serum albumin. Cultured podocytes were incubated with preadipocyte-cultured medium (PCM), adipocyte-cultured medium (ACM) or adipocyte-cultured medium with an antibody against adiponectin (PA1-054, Thermo Fisher Scientific), leptin (5367-100, BioVision), resistin (sc-22315, Santa Cruz Biotechnology), TNF-α (NB600-587, Novus Biologicals, Littleton, CO, USA) or the IgG isotype for 24 h. Podocyte function was then evaluated in vitro. Podocyte injury was evaluated by the expression of nephrin, CD2AP and desmin in the slit diaphragm, which was measured using RT-PCR.

**Computer tomography**

For each rat, CT was performed on one slice from each of the three regions, namely, the sacroiliac joint (inguinal level), the upper portion of the iliac crest (L4-L5, upper iliac crest level) and 1.5 cm above the second slice (L1-L2, kidney level) as described previously with minor modifications (Hillebrand et al. 2010) using the ultra-high resolution setting of a SOMATOM Sensation 16 scanner (Siemens AG). A density range between −150 and −40 Hounsfield units was used to define an area of fat, and a range between −40 and 250 Hounsfield units was used to define an area of muscle (water density = 0). All CT scans were performed in the prone position under anesthesia. The total adipose and muscle tissue in these three regions was calculated using a computer. The intra-abdominal cavity, including the retroperitoneal space, was outlined manually with a cursor and the area of adipose tissue was expressed as a percentage of the total area calculated. Subtraction of the intra-abdominal area from the total abdominal area gave the subcutaneous area. The average visceral fat area/subcutaneous fat area ratio in the slices from the three regions was used for statistics.

**Oral glucose tolerance test**

Rats were subjected to an OGTT after overnight fasting. A 0 min blood sample was taken from each rat, and the rats were immediately given a glucose solution (concentration, 0.2 g/0.1 mL; 0.1 mL/100 g body weight) by gavage, and four additional blood samples were collected at 30, 60, 90 and 120 min. The concentration of plasma insulin was determined using a commercial rat insulin enzyme-linked immunosorbent assay (ELISA) kit (Mercodia AB, Uppsala, Sweden), and the plasma glucose level was measured using a glucose analyzer (Model 23A, Yellow Springs Instrument, Yellow Springs, OH, USA).

**Biochemical measurements**

The levels of plasma total cholesterol, high-density lipoprotein cholesterol, low-density lipoprotein cholesterol, triglycerides and free fatty acids were measured using commercial EIA kits from Diagnostic Systems GmbH (Holzheim, Germany). The levels of plasma BUN and creatinine were measured using kits from ARKRAY, Inc. (Minami-ku, Kyoto, Japan). Plasma data of leptin, adiponectin and TNF-α levels were measured using ELISA kits purchased from R&D Systems.

**Enzyme-linked immunosorbent assay**

Samples were collected in 0.1 M HCl buffer and mixed by pipetting. After centrifugation (1000 g, 10 min), the supernatant was decanted and collected for measurement of the intracellular cGMP concentration in accordance with the product specification.

**Transmission electron microscopy**

A fragment of the kidney cortex was cut into 1 mm³ pieces and fixed in 2.5% glutaraldehyde in Millonig solution, and then embedded in PolyBed 812 (Polysciences, Inc, Warrington, PA, USA) for transmission electron microscopy (TEM) analysis (Sharma et al. 2008).

**Western blotting**

Whole cell lysates were prepared by sonication in ice-cold lysis buffer (1% Triton X-100, 50 mM KCl, 25 mM HEPES, pH 7.8, supplemented with 10 µg/mL leupeptin, 20 µg/mL aprotonin, 125 µM dithiothreitol, 1 mM phenylmethylsulfonyl fluoride and 1 mM sodium...
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Fluorescent microscopy

To investigate the adipokine effects of PCM or ACM, differentiated podocytes cultured on 12-mm cover slips in a 24-well plate were treated with PCM or ACM with the IgG isotype as the control or with an antibody against adiponectin, leptin, resistin or TNF-α. The immunofluorescent staining of desmin (green), nephrin (green), CD2AP (green), F-actin (red) and DAPI (blue) was detected using a Leica DM6000B fluorescent microscope (Leica Microsystems). Each slide was evaluated in eight randomly selected fields from four slides per sample (magnification, 100×) by two different investigators who were blinded to the sample identities.

Quantitative analysis of desmin, nephrin and CD2AP transcripts by RT-PCR

Total RNA was extracted from cultured mouse podocytes using TRI Reagent (Sigma-Aldrich). The extraction was carried out as described previously (Seow et al. 2011). TaqMan Gene Expression Assays (FAM dye-labeled MGB probe; Applied Biosystems) containing specific primers (assay ID for desmin: Mm00802455_m1; nephrin: Mm00497828_m1; CD2AP: Mm01277622_g1; GAPDH: Mm99999915_g1), the TaqMan MGB probe (FAM dye-labeled), TaqMan Fast Universal PCR Master Mix and 100 ng of cDNA were used to detect and quantify the mRNA levels of desmin, nephrin and CD2AP in cultured mouse podocytes. GAPDH served as the internal control. The Ct value of GAPDH was subtracted from that of desmin, nephrin and CD2AP (ΔCt). The ΔCt value of the ACM group was compared with that of the PCM group.

Table 1  Body weight, fat pad weight, kidney weight, metabolism-related biochemical parameters and lipid profiles in controls (C) and high-fat diet-fed rats (HF) (n=8).

<table>
<thead>
<tr>
<th>Parameters</th>
<th>C</th>
<th>HF</th>
<th>P-value</th>
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<tbody>
<tr>
<td>Body weight (g)</td>
<td>459 ± 9</td>
<td>532 ± 22</td>
<td>&lt;0.05</td>
</tr>
<tr>
<td>Fat pad weight (g)</td>
<td></td>
<td></td>
<td></td>
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<tr>
<td>Epididymal</td>
<td>8.5 ± 0.9</td>
<td>22.2 ± 2.5</td>
<td>&lt;0.05</td>
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<tr>
<td>Retroperitoneal</td>
<td>10.7 ± 0.5</td>
<td>31.5 ± 3.6</td>
<td>&lt;0.05</td>
</tr>
<tr>
<td>Inguinal</td>
<td>12.2 ± 0.9</td>
<td>23.4 ± 1.3</td>
<td>&lt;0.05</td>
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<tr>
<td>Kidney weight (g)</td>
<td>3.38 ± 0.04</td>
<td>3.54 ± 0.04</td>
<td>&lt;0.05</td>
</tr>
<tr>
<td>Glucose (mg/dL)</td>
<td>115 ± 2.4</td>
<td>138.8 ± 4.5</td>
<td>&lt;0.05</td>
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<tr>
<td>Insulin (μU/mL)</td>
<td>24.9 ± 3.8</td>
<td>45.5 ± 6.2</td>
<td>&lt;0.05</td>
</tr>
<tr>
<td>Triglyceride (mmol/L)</td>
<td>0.8 ± 0.1</td>
<td>1.1 ± 0.1</td>
<td>&lt;0.05</td>
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<tr>
<td>Cholesterol (mg/dL)</td>
<td>145.8 ± 8.9</td>
<td>244.7 ± 9.1</td>
<td>&lt;0.05</td>
</tr>
<tr>
<td>BUN (mg/dL)</td>
<td>12.4 ± 0.5</td>
<td>12.3 ± 0.8</td>
<td>NS</td>
</tr>
<tr>
<td>Creatinine (mg/dL)</td>
<td>0.23 ± 0.02</td>
<td>0.32 ± 0.01</td>
<td>&lt;0.05</td>
</tr>
<tr>
<td>Urine protein (mg/day)</td>
<td>115.9 ± 12.4</td>
<td>357.5 ± 64.2</td>
<td>&lt;0.05</td>
</tr>
<tr>
<td>Urine creatinine (mg/day)</td>
<td>120.4 ± 19.8</td>
<td>202.2 ± 18.9</td>
<td>&lt;0.05</td>
</tr>
<tr>
<td>Urine protein:creatinine</td>
<td>1.09 ± 0.15</td>
<td>1.76 ± 0.22</td>
<td>&lt;0.05</td>
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</table>

Values are expressed as mean ± s.d. Statistical analysis was performed using Mann–Whitney U test with repeated measurements. A value of P<0.05 was considered significant.

BUN, blood urea nitrogen; NS, not significant.
The reactions were performed at 95°C for 10 min, followed by 40 cycles at 95°C for 15 s and 60°C for 1 min.

Statistical analysis
The cell studies were performed four times. In the animal experiments, each group consisted of eight rats. The results were expressed as the mean ± standard deviation (s.d.). Statistical significance was assessed by one-way analysis of variance or the Mann–Whitney U test. A P-value of <0.05 was considered statistically significant.

Results

Insulin resistance in HF diet-fed rats
As shown in Table 1, the body weight increased significantly after 3 months in rats fed the HF diet compared with those fed the standard chow (C). In addition, the fasting levels of plasma glucose, insulin, triglycerides and cholesterol were significantly higher in the HF diet group than those in the control. To further evaluate the change in insulin sensitivity, an OGTT was performed, and the results for the two groups at the end of the last experimental week are shown in Fig. 1. There was no difference in the baseline plasma glucose level between the two groups; however, the plasma glucose level increased significantly in rats of the HF diet group compared with that in rats of the control at 30, 60, 90 and 120 min after oral glucose challenge (Fig. 1A; P<0.05). As shown in Fig. 1B, the HF diet-fed rats had a significantly higher baseline plasma insulin level than the standard chow-fed rats. In response to oral glucose challenge, both groups responded with an increase in the plasma insulin level, except that the increase in the HF diet-fed rats was significantly higher than that in the standard chow-fed rats (P<0.05).

Visceral fat predominance in HF diet-fed rats
As shown in Table 1, the weights of the epididymal, retroperitoneal and inguinal adipose deposits were significantly higher in rats of the HF diet group than those in rats of the control. The fat content of each area in the two groups was measured as the percentage of the total area occupied by fat as assessed by CT. As shown in Fig. 2 and Table 2, the fat content in the HF diet group was significantly higher than that in the control. Table 3A shows hematoxylin and

<table>
<thead>
<tr>
<th>Locations</th>
<th>C</th>
<th>HF</th>
<th>P-value</th>
</tr>
</thead>
<tbody>
<tr>
<td>Kidney level</td>
<td>31.4±1.9</td>
<td>52.3±4.3</td>
<td>&lt;0.05</td>
</tr>
<tr>
<td>Upper iliac crest level</td>
<td>19.7±1.5</td>
<td>29.3±1.3</td>
<td>&lt;0.05</td>
</tr>
<tr>
<td>Inguinal level</td>
<td>5.6±0.4</td>
<td>7.6±0.4</td>
<td>&lt;0.05</td>
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</table>

The units are the percentage of the total cross-sectional area. Values are expressed as mean±s.d. Statistical analysis was performed using Mann–Whitney U test with repeated measurements. A value of P<0.05 was considered significant.
eosin-stained slides of the epididymal white adipose tissue. The mean adipocyte size was significantly greater in the HF diet-fed rats than that in the control (Fig. 3B), and the adipocyte size distribution curve shifted to the right in the HF diet-fed rats compared with that in the control (Fig. 3C). We also examined the levels of plasma adipocytokines in both groups and showed a decreased plasma adiponectin level and increased levels of plasma leptin, resistin and TNF-α in HF diet-fed rats compared with those in the control (Fig. 3D; *P < 0.05). From these data, we further investigated the interplay between these factors and glomerular podocytes.

Renal function impairment and podocyte injury in HF diet-fed rats

Table 1 shows that the kidney weight increased significantly in HF diet-fed rats compared with that in the control, as well as the levels of plasma creatinine (but not BUN), 24 h urinary protein excretion, creatinine and the protein/creatinine ratio. The immunohistochemical staining for desmin in the glomerulus and tubules showed much more brown pigmentation in the HF diet group than the control (Fig. 4A), and the optical density of the desmin staining in the glomerulus was significantly higher in the HF diet group (Fig. 4B, right panel). The histopathological staining for desmin showed an increase in glomerular damage in the HF diet group, but not in the control (Fig. 4A), and the calculated glomerular volume was also significantly higher in the HF diet group than that in the control (Fig. 4B, left panel). We also measured the levels of desmin, CD2AP and nephrin in the kidney by Western blotting. As shown in Fig. 4C, increased desmin expression, but decreased CD2AP and nephrin expression, was seen in the HF diet group compared with that in the control. When the renal cortical tissue was examined by TEM, obvious podocyte effacement was observed in the HF diet group, but not in the control (Fig. 4D). Because desmin is an injury marker, increased desmin expression associated with glomerular damage (Herrmann et al. 2012) and the downregulation of CD2AP and nephrin indicated podocyte injury in the kidney (Asanuma et al. 2007, Xiao et al. 2013).

Increased desmin expression and decreased CD2AP and nephrin expression in ACM-treated podocytes

Based on the Western blotting results, we found that the high-fat diet damaged podocytes. Therefore, we cultured mature podocytes and examined the effects of PCM and ACM with phase photography and immunofluorescent staining (Fig. 5A). We found ACM disrupted the cytoskeleton in podocytes. We also incubated podocytes for 24 h with PCM or ACM with/without the IgG isotype and an antibody against adiponectin, leptin, resistin or
TNF-α. We examined the effects of adipocyte-derived factors on podocyte function and measured the mRNA levels of desmin, CD2AP and nephrin by quantitative RT-PCR. As shown in Fig. 5B, increased desmin mRNA expression and decreased CD2AP and nephrin mRNA expression were seen in the ACM-treated group compared with that in the PCM-treated group (P<0.05). We also found that antibodies against leptin, resistin and TNF-α in the ACM-treated group were the three major correction factors, and the antibody against adiponectin in the ACM-treated group was a worse factor (P<0.05) compared with those in the ACM-treated group alone. Moreover, we also observed desmin, CD2AP and nephrin immunostaining of PCM-cultured and ACM-cultured podocytes with/without the IgG isotype as the control or with an antibody against adiponectin, leptin or resistin, or TNF-α. As shown in Fig. 5C, D and E, antibodies against resistin, TNF-α and leptin significantly decreased desmin expression and increased CD2AP and nephrin expression in ACM-treated podocytes compared with that in the ACM-treated group alone (P<0.05).

Discussion

Excess visceral fat is an independent cardiovascular risk factor (Matsushita et al. 2014, Okamoto et al. 2014). In this study, the HF diet induced obesity, insulin resistance, hyperinsulinemia and dyslipidemia in rats. A significant increase in visceral fat was seen in the HF diet group (Fig. 2 and Table 2). In addition, male Sprague Dawley rats fed a HF diet for 12 weeks in our study developed morphological and functional kidney disorders as same as past articles (Pinhal et al. 2013, Jenkin et al. 2016). Previous studies showed that obesity carries an increased risk of CKD and ESRD (Gelber et al. 2005, Obermayr et al. 2008, Satirapoj et al. 2013), and a recent report showed that perirenal visceral adiposity strongly associates with renal inflammation and diabetic nephropathy (Ndisang et al. 2014). Our data demonstrated that the levels of plasma creatinine, 24h urinary protein and creatinine excretion and the protein/creatinine ratio were significantly elevated in the HF diet group: glomerular changes that are consistent with kidney injury. Moreover, the visceral
Fat level also associates with coronary artery calcification in nondialysis-dependent CKD patients (Cordeiro et al. 2013). Our results suggested that increased visceral adiposity might enhance renal damage and cardiovascular risk in obese individuals. The limitation of this study was the small number of rats, which was not indicative of a normal distribution, and we plan on using more animals in future studies.

Past research has manifested morphological and histological changes of obesity related to glomerulopathy, such as glomerular hypertrophy with/without focal segmental glomerulosclerosis (Kambham et al. 2001, ...
In addition, there is other evidence of obesity-related glomerulopathy demonstrating vacuolar degeneration in renal tubules, increased glomerular ECM accumulation (Pan et al. 2014) and increased expression of TGFβ1, AT1 receptor, transcription factor ZEB2, fibronectin and desmin in glomerular and tubular cells (Pinhal et al. 2013). In this study, we focused on changes in the glomerulus because of its important physiological function as a barrier in humans. Our data indicated increased proteinuria, glomerular hypertrophy, increased desmin expression in the glomeruli and tubules and decreased CD2AP and nephrin expression of the glomeruli in HF diet-fed rats. Resveratrol and CB2 receptor agonism were also studied to alleviate renal injury in diet-induced obesity rats (Jenkin et al. 2016, Pan et al. 2014). Unlike previous studies, we hypothesized that adipocyte-derived factors act on podocytes in vitro and tried to search for the individual effect of each adipocyte-derived factor on podocytes.

Ndisang and coworkers (2014) showed increased proinflammatory/oxidative mediator levels, macrophage infiltration and profibrotic/extracellular matrix protein levels in perirenal adipose tissue, and decreased renal function in Zucker diabetic fatty rats after using hemin (a heme oxygenase inducer) to reduce perirenal visceral adiposity, renal inflammation and diabetic nephropathy. Knight and coworkers (2010) showed that the expression of glomerular nephrin expression decreases and the urinary albumin level increases in a HF diet model of obesity and hypertension. In this study, we observed renal cortical tissue injury, which was observed as an increase in desmin expression and a decrease in CD2AP and nephrin expression, in the HF diet-fed Sprague Dawley rats. Results of TEM revealed obvious podocyte effacement in the HF diet group. Our data show that adipocyte-derived factors were very harmful to podocytes. Adipocytes, therefore, play a pathological role in obesity-associated podocytopathy.

In the cultured podocyte experiments, we demonstrated that the treatment of mature podocytes with adipocyte-conditioned medium resulted in increased desmin expression and decreased CD2AP and nephrin expression compared with podocytes treated with PCM, especially enriched with leptin, resistin and TNF-α (Fig. 5B). Furthermore, decreased protein expression of desmin and increased protein expression of CD2AP and nephrin were observed by immunostaining after resistin, leptin and TNF-α antibody treatment of the ACM group (Fig. 5C, D and E), consistent with RT-PCR results. These findings showed that adipocyte-derived factor(s) can affect structural protein expression in podocytes, which impairs the function of the renal defense barrier. We also provided evidence that the adipocyte factors may play a pathological role in obesity-associated podocytopathy. Our data show that adipocyte-derived factors were very harmful to podocytes. Adipocytes cause podocytopathy

Declaration of interest
The authors declare that there is no conflict of interest that could be perceived as prejudicing the impartiality of the research reported.

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Author contribution statement
The study was conceived and designed by J Y C and C C J. The experiments and data collection were performed by J Y C, Y W C, C C L, Y T L, C H T, L K C, T C H, H M H, Y T W, T T K, L Y W and S W H. Data analysis and interpretation were performed by J Y C, D Y J, L K C, L Y W and C C J. All authors were involved in writing the paper and approved the submitted and published versions.

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