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Involvement of peroxisome proliferator-activated receptors in cardiac and vascular remodeling in a novel minipig model of insulin resistance and atherosclerosis induced by consumption of a high-fat/cholesterol diet

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Abstract

Background: A long-term high-fat/cholesterol (HFC) diet leads to insulin resistance (IR), which is associated with inflammation, atherosclerosis (AS), cardiac sympathovagal imbalance, and cardiac dysfunction. Peroxisome proliferator-activated receptors (PPARs) and nuclear factor κB (NF-κB) are involved in the development of IR-AS. Thus, we elucidated the pathological molecular mechanism of IR-AS by feeding an HFC diet to Tibetan minipigs to induce IR and AS.

Methods: Male Tibetan minipigs were fed either a normal diet or an HFC diet for 24 weeks. Thereafter, the minipigs were tested for physiological and biochemical blood indices, blood pressure, cardiac function, glucose tolerance, heart rate variability (HRV), and PPAR-associated gene and protein expression levels.

Results: HFC-fed minipigs exhibited IR through increased body weight, fasting blood glucose levels, plasma cholesterol and its composition, and insulin and free fatty acid (FFA) levels; decreased insulin sensitivity; impaired glucose tolerance; and hypertension. Increased C-reactive protein (CRP) levels, cardiac dysfunction, depressed HRV, and the up-regulation of PPAR expression in the abdominal aorta concomitant with down-regulation in the heart tissue were observed in HFC-fed minipigs. Furthermore, the levels of NF-κBp65, IL-1β, TNF-α, MCP-1, VCAM-1, ICAM-1, MMP-9, and CRP proteins were also significantly increased.

Conclusions: These data suggest that HFC-fed Tibetan minipigs develop IR and AS and that PPARs are involved in cardiovascular remodeling and impaired function.

Keywords: Tibetan minipig, Insulin resistance, Atherosclerosis, Myocardial ischemia, PPARs, NF-кВ, High-fat/cholesterol diet, Heart rate variability

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Background

Hyperlipidemia is a common predicament in modern society owing to changes in lifestyle and food consumption in humans. Diet plays an important role in the management of lipoprotein and lipid concentrations in blood. Long-term, high-fat consumption leads to insulin resistance (IR), because saturated fatty acids interfere with the action of insulin [1].

IR is a precursor and the primary characteristic of type 2 diabetes. Based on an epidemiological study, the incidence of diabetes is rapidly increasing with the population aging, and approximately 9.7% of adults have diabetes in China [2]. Moreover, the mortality of heart disease is 4-fold higher in diabetes patients [3]. In fact, myocardial ischemia (MI), caused by atherosclerosis (AS) of coronary arteries, more frequently occurs without prior symptoms in these patients [4]. Therefore, the pre-diabetic pathological state of IR is concerning.

IR has been implicated in AS promotion and impaired endothelial-dependent diastolic function [5]. Data from patients with IR or elevated fasting blood glucose, increased heart rate, reduced heart rate variability (HRV) [6,7], and cardiac sympathovagal imbalance [8] and data from animal experimental and clinical studies all confirm that IR is associated with impairment of selective signaling pathways and that IR leads to peripheral vascular and myocardial structural and functional changes [9-11]. Therefore, it is necessary to establish an ideal animal model to have a better understanding of the pathological process involved in IR, hyperlipidemia, and AS, which will be helpful to further develop new therapeutic agents.

Pigs are a potentially useful animal model because, unlike mice and rabbits, their anatomy, physiology, feeding and sleeping habits, neo-intimal formation, and thrombosis are very similar to those of humans. Pigs can recapitulate the formation of diabetes- and inflammationinduced AS lesions [12,13], which are currently, mostly induced by toxicity to drugs (e.g., streptozotocin and alloxan), causing pancreatic β-cell ablation, and AS lesions are also induced by feeding with an atherogenic diet [14,15], rather than natural development subsequent to metabolic syndrome or diabetes. In a recent study, Göttingen minipig fed a high-fat diet for 12 weeks develops obesity, IR, and lipid disorders [16], and the obesity-prone Ossabaw minipig fed a high-fat/cholesterol (HFC) diet for 9 weeks develops IR, impaired glucose tolerance, dyslipidemia, hypertension, and early coronary intimal hyperplasia [17]. However, these models do not exhibit chronic ischemia formation. Many minipig varieties in China are used in experimental studies, such as Bama minipig, WZS minipig, Guizhou minipig, and Tibetan minipig. In our previous studies using the Chinese minipig model of cardiovascular diseases, we found that HFC-fed Tibetan minipigs could also develop IR, impaired glucose tolerance, dyslipidemia, hypertension, AS, and cardiac dysfunction [18]. These conditions are very similar to IR caused by related complications in clinical cases; thus, this disease model is a novel minipig model of IR-AS and warrants further study.

Peroxisome proliferator-activated receptors (PPARs) belong to a superfamily of the type II nuclear hormone receptors, are ligand-activated transcription factors, which have three subtypes (PPAR α , β/δ , and γ), and may participate in lipid metabolism regulation and AS development [19]. PPARs also inhibit the activity of inflammationrelated transcription factors (such as NF-kB), which inhibit inflammatory effects of a variety of inflammatory diseases. PPARs are gaining research interest with regard to metabolic and cardiovascular diseases, and PPAR expression in the target organ may play an important role in changing organ structure and function; moreover, PPAR subtypes function in cardiovascular diseases and may also influence lesion development. However, the specific active mechanism of PPARs in cardiovascular remodeling and dysfunction is unclear.

Therefore, we hypothesized that IR-AS with spontaneous chronic ischemia in HFC-fed Tibetan minipig maybe associated with inflammation, IR, autonomic dysfunction, and activation of PPARs and the NF-κB signaling pathway. In this work, we successfully recapitulated the IR-AS Tibetan minipig model induced by an HFC diet. We examined blood biochemical indices, blood pressure, cardiac function, glucose tolerance, HRV, and expression levels of genes and proteins associated with PPARs and the NF-κB signaling pathway. We aimed to elucidate whether HFC-fed Tibetan minipig developed IR-AS and to understand the cause of cardiac impairment and pathological molecular mechanisms. Consequently, our results provide an experimental basis for the relationship between IR and cardiovascular diseases and drug treatment.

Methods

Animal model

Ten male Tibetan minipigs at 3 months of age, weighing 12–14 kg, were purchased from the Experimental Animal Center of Southern Medical University, China. After adaptation to the environment for 8 weeks, Tibetan minipigs were randomly divided into 2 groups: control (Ctr) and HFC. Ctr minipigs (n = 4) were fed a diet of regular chow, and HFC minipigs (n = 6) were fed an atherogenic diet of HFC chow consisting of 1.5% cholesterol, 15% shortening oil, 10% egg yolk powder, and 73.5% regular chow (diet compositions are described in Table 1). All Tibetan minipigs in both groups received the same amount of food (2.5% of body weight) and were fed twice daily on a restricted schedule with regular chow or atherogenic diet

Table 1 Diet components

Component	Control diet		High-fat/high-cholesterol diet	
	g	kcal	g	kcal
Protein	17.4	22	15.9	19
Carbohydrate	55.2	70	41.1	50
Fat	6.3	8	25.2	31
Total		100		100
kcal/g	3.39		4.46	

for 24 weeks until they were euthanized. All animals were housed and fed in individual pens under a 12-h light: dark cycle. Water was provided *ad libitum*. All experiments were performed according to the guidelines proposed by the Laboratory Animal Research Center of Zhejiang Chinese Medical University (SYXK, Hangzhou, 2008-0116, China) and approved by the Institutional Animal Care and Use Committee of the Zhejiang Chinese Medical University.

Blood biochemical index measurements

At 24 weeks, blood samples were drawn from minipigs in a supine position by performing anterior vena cava puncture within 30 s after a fasting period of 12 h [20]. Blood samples were then separated to analyze total cholesterol (TC), high-density lipoprotein cholesterol (HDL-C), lowdensity lipoprotein cholesterol (LDL-C), triglycerides (TG), and fasting blood glucose (FBG), which were measured by an automatic biochemical analyzer (7020, HITACHI, Japan) using kits corresponding to each component (Shanghai Shenneng-DiaSys Diagnostic Technology Co., Ltd., China) for each test. The serum free fatty acid (FFA), insulin, and C-reactive protein (CRP) levels were measured using homologous specific ELISA kits (Shanghai Xinran Biotech Co, Ltd., China) according to the manufacturer's instructions. The homeostasis model assessment IR (HOMA-IR) method was used to evaluate insulin resistance. HOMA-IR was calculated using the following formula: FBG × insulin/22.5 [21].

Intravenous glucose tolerance test (IVGTT)

The IVGTT was performed in each minipig after 8 and 24 weeks of assigned diet consumption. Minipigs were acclimated to restraint in a specialized fixed sling for 3 days before IVGTT. They were anesthetized with isoflurane (maintained at 2–3% by mask with supplemental O₂), and then right jugular veins were catheterized percutaneously [22]. Minipigs were allowed to recover for 3 h before IVGTT to avoid any effects of isoflurane on glucose levels [23]. Blood samples were obtained at the baseline time point (0 min), then glucose (0.5 g/kg) was injected intravenously over 2 min, and further samples were obtained at 15, 30, 60, 90, and 120 min after injection for measurement of blood glucose and insulin

concentrations. From the IVGTT, the insulin sensitivity index (S2) for minipigs was calculated as previously described [24].

Blood pressure and cardiac function measurement

Minipigs were acclimated to a low-stress sling for blood pressure measurements. Non-invasive blood pressure was monitored using a non-invasive physiological signal telemetry system (EMKA Technologies S.A.S., France) by using a tail cuff method at the beginning of the experiment and 12 weeks into the experiment. Mean blood pressure (MBP), systolic blood pressure (SBP), and diastolic blood pressure (DBP) were analyzed by ECGAUTO software (EMKA Technologies). After 24 weeks, all minipigs were anesthetized with 10 mg/kg of intramuscular xylazine hydrochloride (Huamu Animal Health Products Co., Ltd., Jilin, China) and intubated, and anesthesia was maintained by inhalation of 0.5-2% isoflurane combined with oxygen and room air [25]. During anesthesis, minipigs were placed on heating pads maintained at 37°C, and the core temperature was measured via a rectal probe (Life Window 6000; Digicare Animal Health, Boynton Beach, FL, USA). All minipigs were equipped with two polyethylene blood vessel catheters. One catheter was inserted into the left ventricle via the right common carotid artery to record cardiac function parameters, and another catheter was inserted into the left femoral artery to record blood pressure parameters. After stabilizing for 20 min (i.e., after intramuscular injection of xylazine hydrochloride) at 80 min, when the effects of xylazine hydrochloride on blood pressure are negligible [26], the signals were recorded continuously by using a non-invasive physiological signal telemetry system (EMKA Technologies). MBP, SBP, DBP, left ventricular systolic pressure (LVSP), left ventricular end diastolic pressure (LVEDP), and the maximum rate of left ventricular pressure rise and fall $(\pm dp/dt_{max})$ were analyzed by ECGAUTO software (EMKA Technologies).

HRV analysis

The physiological telemetry test was performed for each minipig every 8 weeks. All minipigs were acclimatized to the experimental jackets for 3 days before each test. Minipigs were shaved, and surface ECG electrodes were attached. Then, specially designed jackets were placed on each minipig to protect the ECG electrodes and lead wires, and minipigs were returned to their cages. After stabilization for 2 h, vital signs of each animal (Lead I and II ECG) were monitored for 24 h and recorded using a Non-Invasive Physiological Signal Telemetry System (EMKA Technologies). ECG signals were recorded at a sampling rate of 500 Hz by EMKA IOX data acquisition software. ECG-AUTO software (EMKA Technologies) was used for ECG, activity, and HRV analyses. Data from the 24 h sampling period were divided into

30-minsegments of RR data. R waves were detected, and the RR interval (RRI) tachogram was calculated as raw HRV in sequence order. Similarly, from the same recordings, HRV was analyzed for time and frequency domains. The time-domain parameters obtained were as follows: mean of normal RRIs, standard deviation of all normal RRIs (SDNN), and square root of the mean square successive differences between successive normal intervals (rMSSD). From this tachogram, data sets of 1024 points were re-sampled at 286 ms. A Hamming window was applied to each data set to minimize spectral leakage. A fast Fourier transform was used to obtain the power spectrum of the fluctuation. We identified frequency bands of very low frequency (VLF; 0.003-0.01 Hz), low frequency (LF; 0.01–0.07 Hz), and high frequency (HF; 0.07–1.0 Hz) according to a previous study [27]; the sympathetic and vagal modulation ratio balance was defined by the LF/HF ratio.

Body composition

The percentage of carcass fat was calculated according to the following equation: carcass fat weight/body weight \times 100%. Body mass index was measured as in humans [28] and was calculated by the following equation: (body weight in kg)/(square of the minipig length from the end of the snout to the base of the tail).

Myocardium ischemia size (MIS) determination

MIS was assessed by nitrotetrazolium blue chloride (NBT; Sinopharm Chemical Reagent Co., Ltd., China) staining, as described previously [29]. Briefly, at 24 weeks, the heart was excised, washed in chilled saline, weighed immediately, and then dissected to remove the atria and right ventricle. The left ventricle was frozen at – 20°C, cut in a semi-frozen state into 5 equally thick sections, and incubated in 0.1% NBT solution for 15 min at 37°C. The non-ischemia area stained blue, whereas the ischemia area was not stained. The area of ischemia was measured in each slice with Image J software (version 1.4, National Institutes of Health, USA) and was expressed as the percentage of the total left ventricular area.

Pathological examination

Upon anesthesis at 24 weeks, the chest of each minipig was opened to achieve euthanasia. Samples of the aorta, arteries, heart, pancreas, liver, and adipose tissues were removed from each animal. The thoracic and abdominal portions of the aortas were longitudinally incised and fixed with 10% buffered formalin for 24 h, followed by staining with Oil Red O stain for lipid deposition analysis. The Oil Red O-stained area relative to the entire surface was calculated using Image J software. The heart, pancreas, abdominal aorta, and left branch of the coronary artery were fixed overnight with 10% phosphate-

buffered formalin, dehydrated, embedded in paraffin, sectioned into 5- μ m-thick slices, and stained with hematoxylin and eosin (H&E). The intima-media thickness (IMT) was measured at the point of greatest thickness in sections from the standardized sample sites by using Image J software.

RNA extractions and quantitative real-time PCR (qPCR)

Total RNA was extracted from the abdominal aorta and myocardium tissue using TAKARA RNAiso Plus Reagent according to the instructions. For each sample, first-strand cDNA was synthesized using the Prime Script RT Reagent Kit (TaKaRa, Dalian, China). The sequences of the primers were as follows: porcine GAPDH sense, 5'-AGGTCG GAGTGAACGGATTTG-3', GAPDH anti-sense 5'-ACC ATGTAGTGGAGGTCAATGAAG-3'; PPAR-α sense, 5'-GCTGCTCAGCACCAAGGT-3', PPAR-α anti-sense, 5'-ACCTCCTTCATGCTCTCTCTAT-3'; $PPAR-\beta/\delta$ sense 5'-CTACAAGTGCCAGTGTGAGGAG-3', PPAR-β/δ anti-sense 5'-GGTTGGTGAAGAAGAGGTAGGC-3'; and PPAR-y sense 5'-GACGCAGATTTTGTAGACG-3', PPAR-y anti-sense 5'-CCCTACTGGTTTCTGGATG-3'. The expected sizes of synthesized cDNA were 125 bp for *GAPDH*, 115 bp for *PPAR-\alpha*, 95 bp for *PPAR-\beta/\delta*, and 81 bp for PPAR-y. qPCR was performed in an automated thermal cycler (Minioption™ Real-time PCR System; BIO-RAD) in a final volume of 25 μL, containing 2 μL of cDNA solution, 12.5 μL of SYBR° Premix Ex Taq (TaKaRa, Dalian, China), 1 µL of each primer (10 µmol/L), and 8.5 μL of dH₂O. The cycling reaction was performed according to the manufacturer's instructions via a

Table 2 Phenotypic characteristics of Tibetan minipigs fed a control (Ctr) or high-fat/cholesterol atherogenic (HFC) diet

Parameters	Ctr group	HFC group	<i>P</i> -value
Starting body weight (kg)	16.73 ± 0.72	16.30 ± 0.54	NS
End body weight (kg)	33.43 ± 1.29	49.40 ± 1.07^{b}	< 0.0001
End body mass index (kg/m²)	41.98 ± 1.35	57.05 ± 1.93 ^b	0.0002
Total cholesterol (mg/dL)	96.08 ± 7.96	406.60 ± 42.26^{b}	0.0006
HDL-C (mg/dL)	33.77 ± 3.42	120.60 ± 5.34^{b}	<0.0001
LDL-C (mg/dL)	42.76 ± 5.55	186.30 ± 21.93^{b}	0.0009
Triglycerides (mg/dL)	23.61 ± 2.31	25.38 ± 3.69	NS
Free fatty acids (nmol/L)	0.51 ± 0.01	0.61 ± 0.01^{b}	0.0002
Fasting blood glucose (mmol/L)	4.64 ± 0.10	5.47 ± 0.20^{b}	0.0076
Fasting insulin (mU/L)	18.05 ± 0.74	21.61 ± 0.49^{b}	0.0083
HOMA-IR	3.71 ± 0.09	5.24 ± 0.17^{b}	< 0.0001

Ctr: control group animals (n = 4); HFC: high-fat/cholesterol atherogenic diet animals (n = 6); HDL-C: high-density lipoprotein cholesterol; LDL-C: low-density lipoprotein cholesterol; HOMA-IR: homeostasis model assessment of insulin resistance. Values are shown as mean \pm SEM. ^a, P < 0.05 and ^b, P < 0.01 versus control group. NS: not significant.

standard two-step PCR. *GAPDH* was used as a house-keeping gene to normalize target gene expression, and the expression levels were determined by the comparative CT method.

Abdominal aorta and myocardial protein extraction and western blotting analysis

Abdominal aorta and myocardial proteins were extracted according to the method provided by the manufacturer of the extraction kit (Key Gen Bio TECH, Nanjing, China). The concentrations of total protein in the supernatant were determined by the BCA method, and then 2% sodium dodecyl sulfate (SDS) and 5% 2-mercaptoethanol were added to denature proteins at 100°C for 10 min. Fifty micrograms of protein was separated by SDS-

polyacrylamide gel electrophoresis and transferred to polyvinylidenedifluoride membranes. The membranes were blocked with 5% non-fat milk in 20 mmol/L Trisbuffered saline (pH 7.4) containing 500 mmol/L NaCl and 0.05% Tween 20 (TBST) and then incubated with primary antibody (1:150 dilution, PPAR- α , SC-1982; PPAR- γ , SC-1981; and GAPDH, SC-166545: Santa Cruz Biotechnology, USA; PPAR- β/δ , ab23673: Abcam, USA) overnight at 4°C. Thereafter, the membranes were washed three times with TBST and incubated with a 1:10,000 dilution of horseradish peroxidase-conjugated secondary antibody for 1 h at room temperature. Membranes were again washed three times in TBST. Finally, signals were detected using the Odyssey Infrared Fluorescent Scanner (LI-COR, USA) and quantified by Image J software.

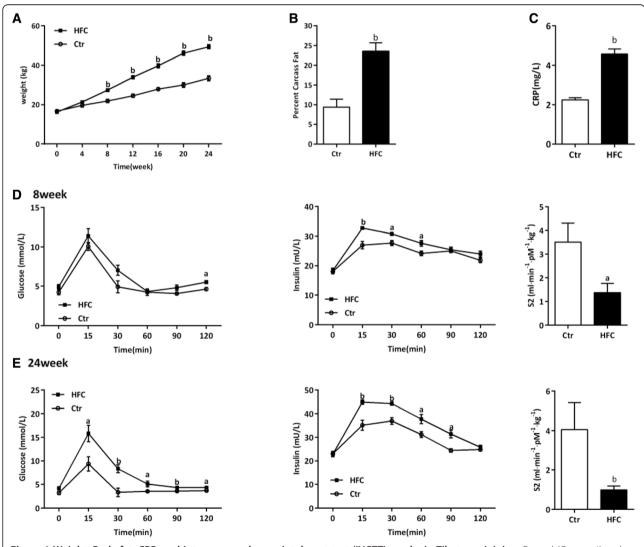


Figure 1 Weight, Body fat, CRP, and intravenous glucose-intolerant test (IVGTT) results in Tibetan minipigs. Control (Ctr, n = 4) and high-fat/cholesterol atherogenic diet minipigs (HFC, n = 6). (A) Weight, (B) percentage of carcass fat, (C) C reaction protein (CRP), (D) simultaneous measurement of glucose and insulin responses during IVGTT and insulin sensitivity index (S2) at 8 week, (E) simultaneous measurement of glucose and insulin responses during IVGTT and S2 at 24 week. Data are presented as mean \pm SEM. ^a, P < 0.05 and ^b, P < 0.01 versus Ctr.

Immunohistochemistry

Fresh abdominal aortas were sliced into 7-um-thick frozen sections, blocked with 3% hydrogen peroxide solution for 10 min to eliminate the activity of endogenous peroxidase, and then incubated overnight at 4°C in the following polyclonal antibody solutions: anti-NF-κB p65, antiinterleukin 1 beta (IL-1β), anti-tumor necrosis factor alpha (TNF-α), anti-monocyte chemoattractant protein-1 (MCP-1), anti-vascular cell adhesion molecule 1 (VCAM-1), anti-intercellular adhesion molecule 1 (ICAM-1), antimatrix metalloproteinase 9 (MMP-9), and anti-CRP (Zhongshan GoldenBridge, Beijing, China). Subsequently, samples were washed with phosphate- buffered saline three times, and the sections were incubated with secondary antibodies for 1 h. After incubation, the sections were stained using 3, 3'-diaminobenzidine tetrahydrochloride (DAB) as a color reagent and counterstained with hematoxylin. The stained slides were viewed with a light microscope (Nikon Eclipse 80i), and digital images of six randomly selected, high-power (×400 magnification) fields were captured on NIS- ElementS.F.2.30 software (Nikon United Kingdom Ltd., Surrey, UK). Protein expression was quantified with Image Pro Plus 5.0 software. This software discriminates between positive- and negative-stained pixels. A visual check was also performed to ensure accurate discrimination of immunolabeled regions. Positive expression is represented as the percentage of the total area.

Statistical analysis

Data are expressed as means \pm SEM. SPSS 17.0 software (SPSS, Chicago, IL, USA) was used to perform statistical analysis. Data were compared using unpaired student's t tests. A two-tailed P value <0.05 was considered statistically significant.

Results

General characteristics of Ctr and HFC-fed Tibetan minipigs

Tibetan minipigs were fed an HFC diet for 24 weeks to induce IR-AS. HFC-fed minipigs had significantly increased body weight, body mass, TC, HDL-C, LDL-C, FFA, FBG, insulin, and HOMA-IR index compared with Ctr minipigs (P < 0.01, Table 2), and the TG level also showed a trend of increasing (P > 0.05, Table 2). Carcass fat and CRP level of HFC-fed minipigs were significantly higher than those of Ctr-fed minipigs (P < 0.01, Figure 1B and C). The IVGTT showed that HFC-fed minipigs had

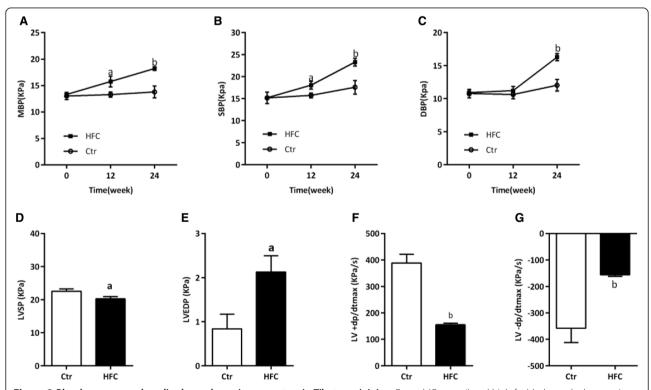


Figure 2 Blood pressure and cardiac hemodynamic parameters in Tibetan minipigs. Control (Ctr, n = 4) and high-fat/cholesterol atherogenic diet minipigs (HFC, n = 6). **(A)** Mean blood pressure (MBP), **(B)** systolic blood pressure (SBP), **(C)** diastolic blood pressure (DBP), **(D)** left ventricular systolic pressure (LVSP), **(E)** left ventricular end diastolic pressure (LVEDP), **(F)** LV + dp/dt_{max}, and **(G)** LV -dp/dt_{max}. Data are presented as mean \pm SEM. ^a, P < 0.05 and ^b, P < 0.01 versus Ctr.

significantly higher blood glucose and insulin levels during the 120 min after 0.5 mg/kg intravenous injection of glucose at 8 or 24 weeks, and the S2 was also decreased significantly (P < 0.05, P < 0.01, Figure 1D and E). In addition, HFC-fed minipigs showed significantly elevated MBP and SBP at 12 or 24 weeks (P < 0.05, P < 0.01, Figure 2A and B) and elevated DBP at 24 weeks (P < 0.01, Figure 2C). Cardiac hemodynamic results showed that HFC-fed minipigs had significantly lower LVSP and LV \pm dp/dt_{max} but significantly higher LVEDP than Ctr minipigs (P < 0.05, P < 0.01, Figure 2D–G).

HRV analysis

HFC-fed minipigs exhibited significantly reduced SDNN, total power (TP), and HF power at 8 weeks after experiment initiation compared with the Ctr group (P < 0.05, P < 0.01, Figure 3B,D, and G). rMSSD and VLF and LF

power levels were also significantly reduced at 16 weeks (P < 0.05, P < 0.01, Figure 3C,E, and F); however, the heart rate and LF/HF ratio were significantly high at 8 weeks (P < 0.05, P < 0.01, Figure 3A and H). In addition, the frequency domain analysis also showed that the LF and HF power levels were reduced in the HFC group compared with the Ctr group (Figure 3I).

Histopathological assessment

Oil Red O staining analysis demonstrated that aortic atherosclerotic lesions significantly increased in the HFC group compared with the Ctr group (P < 0.01, Figure 4A and B), and we also observed that the atherosclerotic lesions developed first at the aortic arch and abdominal aorta and progressed to involve the entire aorta in the HFC group. H&E staining showed that abdominal aorta and coronary artery atherosclerotic

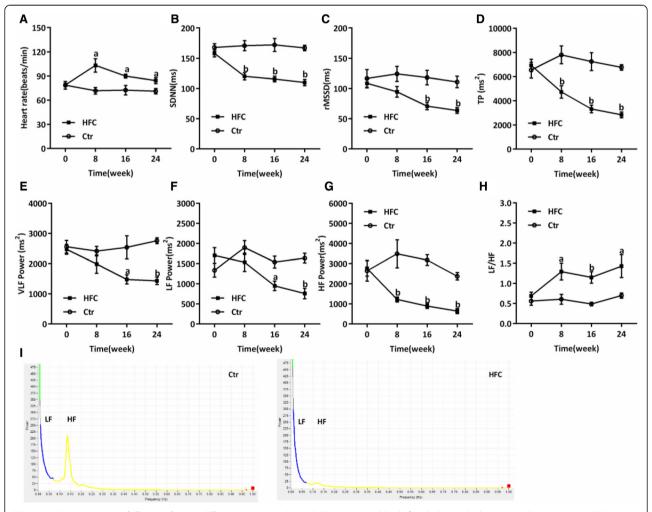


Figure 3 Heart rate variability analysis in Tibetan minipig. Control (Ctr, n = 4) and high-fat/cholesterol atherogenic diet minipigs (HFC, n = 6). **(A)** Heart rate, **(B)** standard deviation of all normal RR intervals (SDNN), **(C)** square root of the mean square successive differences between successive normal intervals (rMSSD), **(D)** total power (TP), **(E)** very low frequency (VLF) power, **(F)** low frequency (LF) power, **(G)** high frequency (HF) power, **(H)** LF/HF ratio, **(I)** graph of frequency domain analysis. Data are presented as mean \pm SEM. ^a, P < 0.05 and ^b, P < 0.01 versus Ctr.

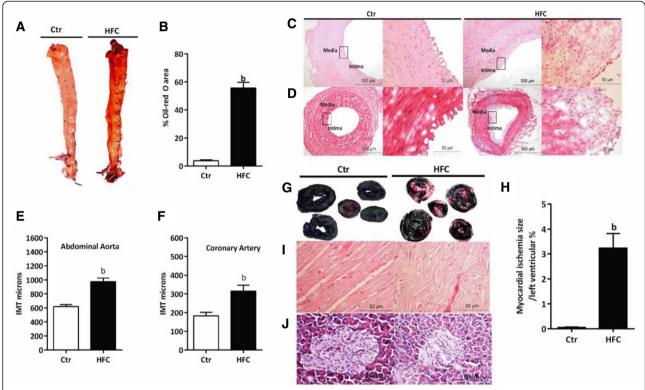


Figure 4 Histopathological assessment in Tibetan minipigs. Control (Ctr, n = 4) and high-fat/cholesterol atherogenic diet minipigs (HFC, n = 6). **(A)** Aortic lipid analysis by Oil Red O staining, **(B)** percent of Oil Red O area, **(C)** H&E-stained abdominal aorta (left is H&E 40x, right is H&E 400x), **(D)** HE-stained coronary artery (left is H&E 40x, right is H&E 400x), **(E)** IMT of abdominal aorta, **(F)** IMT of coronary artery, **(G)** myocardial tissue with NBT staining, **(H)** myocardial ischemia size/left ventricular, **(I)** H&E-stained myocardial tissue (H&E 400x), **(J)** H&E-stained pancreatic tissue (H&E 400x). Data are presented as mean \pm SEM. ^b, P < 0.01 versus Ctr.

lesions were mainly located in the intima. The lesions were composed of foam cell infiltration, thickening of the intima, and protrusion into the lumen, and the elastic membrane of the plaque bottom was destroyed with infiltration of inflammatory cells (Figure 4C and D). The IMTs of the abdominal aorta and coronary artery were significantly increased in the HFC group compared to the Ctr group (P < 0.01, Figure 4E and F). These data strongly suggest that HFC-fed minipigs can form obvious atherosclerotic lesions. In addition, we were surprised to find that HFC-fed minipigs experienced spontaneous myocardial infarction and islet cell changes. NBT staining demonstrated that MIS was significantly increased in the HFC group compared with the Ctr group (P < 0.01, Figure 4G and H). H&E staining further showed that HFC-fed minipigs had disorderly arranged myocardial fibers with a wave-like appearance, narrowed muscle gaps, and hypertrophic cardiomyocytes, while no changes were observed in the Ctr group (Figure 4I). Moreover, partial islet cell ablation and mild cloudy swelling were observed in the HFC group, but no changes were observed in the Ctr group (Figure 4J).

Up-regulation of PPAR gene and protein expression in the abdominal aorta of HFC-fed Tibetan minipigs

PPAR gene and protein expression levels in the abdominal aorta were assessed to confirm whether activated PPARs participate in an important pathological mechanism in the Tibetan minipig model of IR-AS. Compared with the Ctr group, HFC-fed minipigs had significantly higher levels of $PPAR-\alpha$, $PPAR-\beta/\delta$, and $PPAR-\gamma$ mRNAs and corresponding proteins in abdominal aortic tissues (P < 0.05, P < 0.01, Figure 5A–G).

Down-regulation of PPAR gene and protein expression in the heart of HFC-fed Tibetan minipigs

qPCR and western blotting showed that mRNA and protein levels, respectively, of PPAR- α and PPAR- β/δ were significantly reduced in the heart tissue of HFC-fed minipigs (P < 0.05, Figure 6A–B, D–F).

NF-κB pathway activation and up-regulation of inflammatory cytokines in the abdominal aorta of HFC-fed Tibetan minipigs

NF-κB protein levels increased in the HFC group compared with the Ctr group (Figure 7A and B). Moreover,

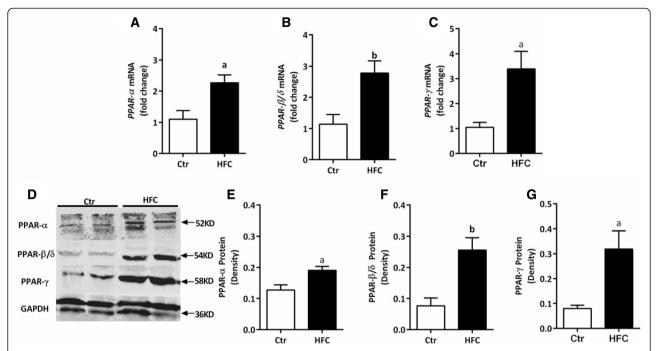


Figure 5 Gene expression and protein levels in the abdominal aorta tissue in Tibetan minipigs. Gene expression was measured by qPCR, and protein levels were measured by western blot assay in control (Ctr) and high-fat/cholesterol atherogenic diet minipigs (HFC). **(A)** *PPAR-α* mRNA, **(B)** *PPAR-β/δ* mRNA, **(C)** *PPAR-γ* mRNA, **(D)** western blot of PPAR proteins, and quantification of **(E)** PPAR-α, **(F)** PPAR-β/δ, and **(G)** PPAR-γ proteins. Data are presented as mean \pm SEM. a , P < 0.05 and b , P < 0.01 versus Ctr group.

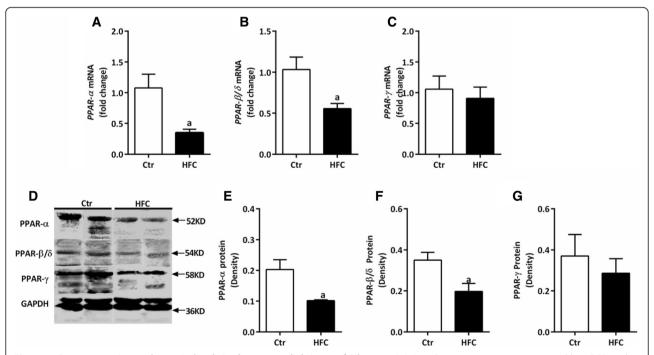


Figure 6 Gene expression and protein levels in the myocardial tissue of Tibetan minipigs. Gene expression was measured by qPCR, and protein levels were measured by western blot assay in control (Ctr) and high-fat/cholesterol atherogenic diet minipigs (HFC). (A) *PPAR-α* mRNA, (B) *PPAR-β/δ* mRNA, (C) *PPAR-γ* mRNA, (D) western blot of PPAR proteins, and quantification of (E) PPAR-α, (F) PPAR-β/δ, and (G) PPAR-γ proteins. Data are presented as mean \pm SEM. a , P < 0.05 versus Ctr.

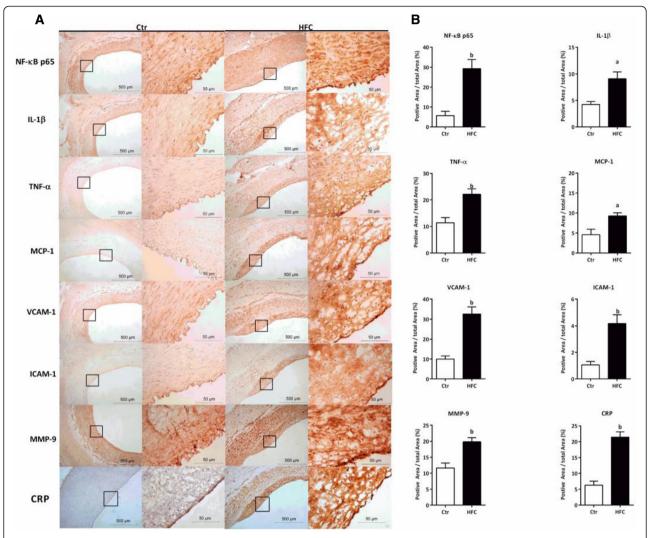


Figure 7 Immunohistochemical staining in the abdominal aorta of control (Ctr) and high-fat/cholesterol atherogenic diet minipigs (HFC). (A) Immunohistochemical staining for inflammatory cytokines in the abdominal aorta, NF-κB p65: nuclear factor κB p65; IL-1β: interleukin 1 beta; TNF-α: tumor necrosis factor alpha; MCP-1: monocyte chemoattractant protein-1; VCAM-1: vascular cell adhesion molecule 1; ICAM-1: intercellular adhesion molecule 1; MMP-9: matrix metalloproteinase 9; CRP: C-reactive protein; left, 40×; and right, 400×. (B) NF-κB p65, IL-1β, TNF-α, MCP-1, VCAM-1, ICAM-1, MMP-9, and CRP protein levels were quantified in HFC and Ctr samples. Data are presented as mean \pm SEM. a , P < 0.05 and b , P < 0.01 versus the Ctr group.

immunohistochemical analysis showed that positive staining for IL-1 β , TNF- α , MCP-1, VCAM-1, ICAM-1, MMP-9, and CRP proteins significantly increased in the HFC group compared with the Ctr group (P < 0.05, P < 0.01, Figure 7B), and staining was primarily located in the endothelium of intimal lesions and within the fibrous lesions (Figure 7A).

Discussion

In the current study, we aimed to establish a Tibetan minipig model of IR-AS by feeding an HFC diet, which resulted in significant weight gain, dyslipidemia, glucose intolerance, hypertension, inflammation, and activated the PPARs and NF- κ B signaling pathways. These

abnormalities are known to develop in IR and AS; thus, the Tibetan minipig model may be helpful to further understanding of human IR and cardiovascular disease development.

Other investigators have induced IR in pig models by using high-energy diets enriched with cholesterol and fat, but IR reportedly develops at different times, e.g., 2 weeks [30] and 16 weeks [31,32], which may be related to composition of diet and pig strains. A recent report stated that the IVGTT-derived S2 is more sensitive to evaluate insulin sensitivity, which has been shown to correlate with clamp insulin sensitivity in pigs [30]. Interestingly, we also found that 8 weeks of feeding an HFC diet were sufficient to induce glucose intolerance and IR in Tibetan minipigs.

Additionally, HFC-fed Tibetan minipigs had significantly increased plasma FFA concentration and blood pressure and depressed HRV with increased sympathetic activities. Wang et al. [33] also reported that a high-fat/ high-carbohydrate diet fed to pigs could induce artery IR, vascular dysfunction, and hypertension. Accumulating evidence links elevated plasma FFA with IR, hypertension [34,35], and activation of the sympathetic nervous system and aggravates MI [36]. FFAs are regarded as potential biochemical markers of post-infarct myocardial remodeling [37]. Further, IR is also associated with left ventricular remodeling [38] and often is accompanied by hemodynamic stress, such as systemic hypertension, which could cause left ventricular pressure overload and result in heart failure [11]. Therefore, a long term HFC diet increases plasma FFA and aggravates IR, which leads to severe cardiac autonomic nervous dysfunction. Specifically, the interaction between the sympathetic nervous system and inflammation plays an important role in the development of vascular stiffening and AS [39]. Moreover, increased FFA was associated with augmented myocardial oxygen consumption [40], limited myocardial blood supply, and cardiac dysfunction, all of which further promote chronic MI in HFC-fed minipigs.

The most interesting, and possibly still largely underestimated, role of FFAs in cell signaling is their ability to bind to nuclear receptors such as PPARs. PPAR- α , PPAR- β/δ , and PPAR- γ are the ligand-activated transcription factors that function as the master regulators of insulin sensitivity, glucose homeostasis, fatty acid and lipoprotein metabolism, inflammation, and AS [41-47]. Considering this function, any dysregulation of these metabolic pathways may result in obesity, diabetes, and cardiovascular diseases. Previous animal experiments have identified that *PPAR-* α or *PPAR-* β/δ deficiency could reduce myocardial fatty acid oxidation, increase glucose utilization and myocardial hypertrophy, and increase ischemic heart sensitivity

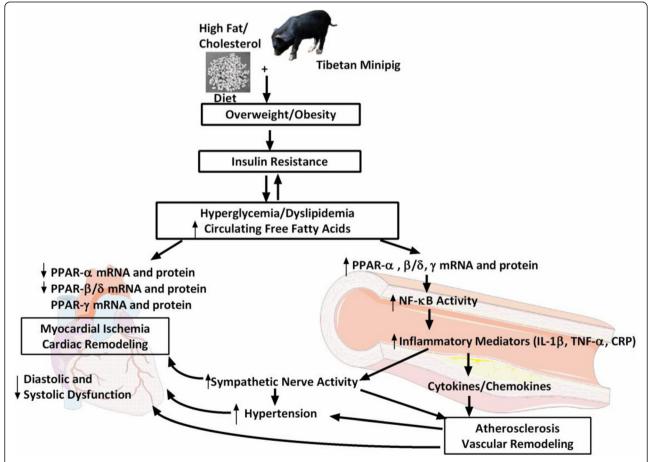


Figure 8 Schematic representation of insulin resistance and atherosclerosis mechanisms in Tibetan minipig. Conditions were induced by feeding with a high-fat/cholesterol diet in minipigs with chronic myocardial ischemia. PPARs, peroxisome proliferator-activated receptors; NF-κB, nuclear factor κΒ; IL-1β, interleukin 1 beta; and TNF-α: tumor necrosis factor alpha; CRP, C-reactive protein. Arrows pointing up or down indicate statistically significant increases or decreases (P < 0.05). Lack of arrows indicates no alteration. The heart and vessel figures were prepared using a template on the Servier Medical Art website (http:// www.servier.fr/servier-medical-art).

[42,48]. Similar to $PPAR-\alpha$ and $PPAR-\beta/\delta$ deficiency, $PPAR-\gamma$ -defective mice also exhibit severe myocardial hypertrophy [43,44]. Interestingly, $PPAR-\alpha$ and $PPAR-\beta/\delta$ expression in heart tissue of HFC-fed minipigs is significantly down-regulated (Figure 6). Thus, we speculate that $PPAR-\alpha$ and $PPAR-\beta/\delta$ may have dominant roles in regulating myocardial fatty acid metabolism, instead of $PPAR-\gamma$; further, $PPAR-\gamma$ may use an indirect mechanism to achieve cardioprotective effects, probably by mobilizing fat metabolism and directing glucose and fatty acid transport to the heart.

Moreover, PPAR genes are also expressed in vascular smooth muscle cells (VSMCs), endothelial cells, and monocytes/macrophages [45,46]. PPARs are highly expressed during monocyte to macrophage differentiation [47]. Some evidence implicates PPAR- α and PPAR- γ in atherogenesis and vascular remodeling of hypertension [47,49,50]. PPAR- β/δ is also expressed in VSMCs and up-regulated after vascular injury [51]. Meanwhile, PPARs can also repress gene expression by antagonizing the activities of NF-кB and activator proteins [46]. Our results also demonstrated that PPAR gene expression and protein levels were both up-regulated in AS lesions of the abdominal aorta, which is consistent with the results of previous clinical studies by Shchelkunova et al. [52]. Additionally, NF-κB protein and levels of its downstream inflammatory factors, such as IL-1β, TNF-α, ICAM-1, VCAM-1, MCP-1, MMP-9, and CRP were also increased in HFCfed minipigs (Figure 8). Thus, HFC-fed minipigs show obvious vascular remodeling and impaired function in the aorta. Therefore, it is clear that the PPAR and NFκB signaling pathways are involved in the development of IR-AS in HFC-fed minipigs.

Conclusions

Our findings describe a novel minipig model where by feeding Tibetan minipigs an HFC diet produced IR-AS, including dyslipidemia, IR, glucose intolerance, hypertension, and inflammatory and vascular AS lesions. PPARs are involved in cardiovascular remodeling and impaired function. We propose that this Tibetan minipig model may be helpful to further understanding of human IR and cardiovascular disease development.

Competing interests

The authors declare that they have no competing interests.

Authors' contributions

MC designed and executed the experiments. YP executed the experiments, interpreted data, and wrote the manuscript. ZC, YP, and LC performed the animal experiment and indexes measurements. KZ performed blood biochemical tests. FC performed pathology experiments and analysis. XX and QM performed data survey and analysis. All authors read and approved the final manuscript.

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