

PARP-1 and SIRT-1 are Interacted in Diabetic Nephropathy by Activating AMPK/PGC-1 α Signaling Pathway

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Introduction: Diabetic nephropathy (DN) is a metabolic disorder characterized by the accumulation of extracellular matrix (ECM). This study aims to investigate whether exists an interplay between poly (ADP-ribose) polymerase 1 (PARP-1) and sirtuin 1 (SIRT-1) in DN via AMP-activated protein kinase (AMPK)/peroxisome proliferator-activated receptor gamma coactivator 1- α (PGC-1 α) signaling pathway.

Methods: Eight-week-old male obese leptin-resistant (db/db) mice and nondiabetic control male C57BLKs/J (db/m) mice were used in this study. Body weight and blood glucose were evaluated after 6 h of fasting, which continues for 4 weeks. The kidney tissues were dissected for Western blot, immunofluorescence (IF) assay. Besides, PARP activity assay, MTT assay, NAD⁺ qualification, Western blot and IF were also performed to detect the level and relation of PARP-1 and SIRT-1 in mouse mesangial cells (MCs) with or without high glucose followed by inhibiting or elevating PARP-1 and SIRT-1, respectively.

Results: Western blotting shows PARP-1 and ECM marker fibronectin (FN) are upregulated while SIRT-1 is downregulated in db/db mice ($p < 0.05$) or in mouse MCs with high glucose ($p < 0.05$), which are significantly restored by PARP-1 inhibitor (PJ34) ($p < 0.05$) and SIRT-1 lentiviral transfected treatment ($p < 0.05$), or worsened by SIRT-1 inhibitor EX527 ($p < 0.05$). PJ34 treatment ($p < 0.05$) or SIRT-1 overexpression ($p < 0.05$) could increase PGC-1 α and p-AMPK levels, concomitant with down expression of FN, however, were reversed in the presence of EX527 ($p < 0.05$).

Discussion: Our results suggest an important relationship between PARP-1 and SIRT-1 through AMPK-PGC-1 α pathway, indicating a potential therapeutic method for DN.

Keywords: PARP-1, SIRT-1, diabetic nephropathy, AMPK/PGC-1 α signaling pathway

Introduction

Diabetic nephropathy (DN), also referred to as diabetic kidney disease (DKD), is the leading cause of end-stage renal disease (ESRD).^{1,2} Characterized by high prevalence and poor prognosis, DN has become a research priority for scientists around the world.³ However, the precise pathogenesis of DN has not yet been fully elucidated. Recently, evidences show that the accumulation of extracellular matrix (ECM) has been involved in the pathogenesis of glomerulosclerosis, which is a characteristic pathological feature of DN.⁴ When subjected to metabolic, immunologic or hemodynamic injury, the kidney mesangial cells (MCs) are activated and then synthesize excessive ECM proteins.⁵ Hence, exploring the mechanism that regulates the accumulation of ECM could contribute to restoring kidney function. Previous study has shown that

hyperglycemia is the major driving force of the progression from DN to ESRD.^{6,7} However, the link between higher levels of glucose and ECM accumulation is multifactorial and may include the overactivation of poly (ADP-ribose) polymerase 1 (PARP-1),^{7,8} and the reduction of sirtuin 1 (SIRT-1).^{9,10}

PARPs constitute a family of several nicotinamide adenine dinucleotide (NAD⁺)-dependent nuclear enzymes which function as DNA damage sensor and signaling molecule.¹¹ The activated PARP can recognize DNA breaks promptly along with the consumption of NAD⁺ and subsequently catalyze the synthesis of ADP-ribose polymers (PAR), which serves to initiate the DNA repair machinery and maintain genome integrity.¹² However, overactivation of PARP-1 in the context of pathological extensive oxidative stress could reverse the protective role of PARP-1, resulting in cell necrosis or apoptosis.¹³ Increasing evidences show that the activation of PARP, especially of PARP-1 is involved in the development of multiple diseases, such as stroke,¹⁴ myocardial infarction,¹⁵ heart failure,¹⁶ vascular dysfunction,¹⁷ and mesenteric damage,¹⁸ muscle dysfunction,^{19,20} or renal ischemia-reperfusion injury,²¹ etc. Additionally, both pharmacological and genetic inhibition or deletion of PARP-1 produces striking reno-protection across several renal disease models.^{22–24} Thus, PARP-1 could be an attractive target for mechanistic and therapeutic investigation in DN.²⁵

SIRT-1, a member of the silent information regulator 2 family, is the founding member of NAD⁺-dependent deacetylases, which is best known for its acknowledged relation to longevity regarding caloric restriction (CR).^{26,27} Recently, several investigators have found that SIRT-1 is associated with many other pathophysiological processes, particularly in age-related diseases, such as neurodegenerative disorders and osteoarthritis.^{28,29} In addition, evidences have shown that SIRT-1 can modulate the expression of target proteins and genes such as p53,^{29,30} peroxisome proliferator-activated receptor gamma coactivator 1 α (PGC-1 α)³¹ and nuclear factor kappa B (NF- κ B),³² which may participate in cellular pathways involving in metabolic diseases such as diabetes,^{33–35} although the contradictory consults which suggest the down-expression of SIRT-1 promotes the development of DN are also reported.³⁶ Karbasforooshan et al³⁷ and Wang et al^{32,38} elucidated again recently that the activity and expression of SIRT-1 were significantly decreased in diabetes and SIRT-1 may relate to the process of diabetes or diabetes-related diseases, indicating that SIRT-1 may also serve as a possible therapeutic target for DN. However, the way in

which SIRT-1 contributes to kidney ECM accumulation and then leads to the development of DN remains to be determined.

The pathophysiological overactivation of PARP-1 consumes a large amount of NAD⁺, leading to NAD⁺ depletion sufficient to inhibit SIRT-1 and lead to cell death due to the same NAD⁺ they are competing for.³⁹ Previously, Waldman et al⁴⁰ illustrated that inhibiting PARP-1 by PARP-1 inhibitor can protect diabetic cardiomyocytes from extensive oxidative stress through the activation of SIRT-1-PGC-1 α pathway. In addition, Lu et al¹⁴ also found that PARP-1-mediated NAD⁺ depletion inhibited the activity of SIRT-1, thereby leading to PGC-1 α hyperacetylation that impacted mitochondrial function and subsequently caused neuron cell death. However, there are no current studies have confirmed whether exists a concrete cross talk between PARP-1 and SIRT-1 in DN. According to the researches, the AMPK/PGC-1 α signaling pathway is thought to be the key mechanism to regulate the mitochondrial function, lipid metabolism and cell growth, which plays a crucial role in the pathophysiological mechanism of DN.^{41,42} Accumulating studies on AMPK/PGC-1 α signaling pathway have shown that the treatment of many drugs significantly activated AMPK in type 2 diabetes (T2D) induced nephropathy, thereby activating PGC-1 α , which was otherwise inhibited by T2D, such as 4-O-methylhonokiol,⁴³ extracellular superoxide dismutase,⁴⁴ okra,⁴⁵ etc. However, evidences regarding the effects of PARP-1 and SIRT-1 through activation of AMPK/PGC-1 α in DN are poorly understood.

In the current study, we demonstrated that there was an interesting relationship between PARP-1 and SIRT-1 that participated in the pathogenesis of DN via SIRT-1-AMPK-PGC-1 α axis, which may open new avenues to help develop novel drugs for the prevention of DN.

Materials and Methods

Reagents

D-glucose was purchased from Sigma-Aldrich (St. Louis, MO, USA). Fetal bovine serum, DMEM/F12 medium and TRIzolTM reagent were purchased from Thermo Fisher scientific Invitrogen (Carlsbad, CA, USA). Rabbit anti-collagen-4 polyclonal antibody (ab6586) and rabbit anti-fibronectin polyclonal antibody (ab2413) were from Abcam (Cambridge, MA, USA). Rabbit anti-PARP-1 (46D11) polyclonal antibody (#9532), SIRT-1 antibody sampler Kit (#9787), Phospho-AMPK α (Thr172) (D4D6D) rabbit mAb (#50,081), AMPK α (D63G4) rabbit mAb (#5832),

Acetylated-Lysine antibody (#9441) and β -Actin (8H10D10) mouse mAb (#3700) were from Cell Signaling Technology (Beverly, MA, USA). Rabbit anti- PGC-1 α (AF5395) was from Affinity Biosciences (Cincinnati, OH, USA). PJ-34 hydrochloride hydrate/P4365 and EX-527/E7034 were from Sigma-Aldrich Chemical Co. (St. Louis, MO, USA). SIRT-1 lentiviral (sirtuin 1 Gene ID: 93,759, LV5 EF-1aF/GFP& Puro) was from gene pharma (Jiangsu, China). Cell lysis, mouse anti-rat β -actin monoclonal antibody and horseradish peroxidase-labeled secondary antibody against rabbit were purchased from Cell Signaling Technology (Beverly, MA, USA). HRP-labeled rabbit anti-mouse IgG were purchased from Beyotime (Shanghai, China). The Revert Aid™ First Strand cDNA reverse transcription kit was purchased from Fermentas (Canada). The Universal PARP Colorimetric Assay Kit (4684-096-K) was purchased from Trevigen Inc. (Gaithersburg, MD, USA).

Establishment of Animal Model

Six 8-week-old male db/db (Lepr db/db) mice (36.5–38.9g) and six nondiabetic control male db/m mice C57BLKs/J (Jackson Laboratories, Bar Harbor, ME) (18.6–20.6g) were purchased from Nanjing Model Animal Center, Jiangsu, China. All animals were housed at proper room temperature (26°C) and humidity (70%) under a controlled light/dark cycle and had free access to water. All animals should be adaptively fed.⁴⁶ The db/m mice and the db/db mice were fed with standard diet (12% fat, 28% protein and 60% carbohydrates). Blood glucose was randomly measured with Glucometer Elite (Bayer) after 6 h of fasting. Then, DN can be diagnosed when the urine albumin creatinine ratio (ACR) of the db/db mice was over 50 μ g/mg and random blood glucose of the db/db mice was over 16.7 mM.⁴⁶ After collecting the data of blood glucose and 24-hour urine for 4 weeks, animals were anesthetized and killed, then the renal tissues were dissected for the following experiments, respectively. All animal experiments were guided in accordance with the guidelines of the Care and Use of Laboratory Animals by the National Institutes of Health, and all efforts were made to minimize animal suffering.

Cell Culture and in vitro Experimental Design

Mouse MCs line SV40 MES13 that were purchased from the cell bank of Chinese Academy of Sciences were kindly

provided by Associate Prof. Hu Wenxue, which was approved by the Department of Science and Technology of the First Affiliated Hospital of Nanchang University.

The cell was cultured in Dulbecco's Modified Eagle Medium (DMEM, Gibco, Grand Island, CA, USA) supplemented with 10% fetal bovine serum (FBS) (Gibco, Grand Island, CA, USA), 100 U/mL penicillin and 100 mg/mL streptomycin under standard culture conditions (37°C in 5% CO₂). For all following experiments, cells were cultured in serum-free conditions for 24 hours when they reached 80% confluence. Growth-arrested MCs were incubated with normal glucose (Control, 5 mM) or high glucose (HG, 25 mM) medium for 48 hours, respectively (see [Supplementary data 1 and Figure 1](#)). And then the HG-treated-cells were pretreated with the PARP-1 inhibitor PJ34 (3 μ L) and/or the SIRT-1 inhibitor EX527 (2 μ L) for 2 h (See [Supplementary data 2 and Figure 2](#)). Moreover, the HG group should be transfected with SIRT-1 lentiviral (See [Supplementary data 3 and Figure 3](#)) with or without PJ34 (3 μ L). All doses and times of these treatments were determined in our pre-experimental results and previous study.⁴⁷ As described above, MCs were then divided into the following groups for subsequent experiments: G5, G25, G25+PJ34, G25+EX527, G25+PJ34+EX527, G25+SIRT-1 lentivirus transfected (G25+Lv-SIRT-1), G25+SIRT-1 lentivirus transfected+PJ34 (G25+Lv-SIRT-1+PJ34).

Lentiviral Treatment of MC Cultures

To achieve SIRT-1 overexpression in MCs, we transfected HG-treated-MCs with SIRT-1 overexpression-lentiviral (purchased from gene pharma) according to the manufacturer's instructions.⁴⁸ Control cells were transfected with negative controls. MCs were infected at 1 d in vitro with a multiplicity of infection (MOI) of 1 in the presence of 1 μ g/mL polybrene for 6 h. Then, the transfection medium was replaced with fresh DMEM. Subsequent assays were conducted 48 h after transfection. The overexpression of SIRT-1 was confirmed by fluorescence-inverted microscope.

Western Blot Analysis

Protein samples were extracted from cells or kidney tissues as described previously.²⁴ Briefly, treated cells or kidney tissues were lysed in RIPA lysis buffer containing protease inhibitor and the protein concentrations were finally determined by BCA kit. Then, the protein samples were fractionated by SDS-PAGE (10–15% polyacrylamide gels) and then transferred to PVDF membranes, which were blocked and then incubated with the appropriate primary antibodies

and secondary antibody. Primary antibodies were used to detect PARP-1 (1:1000; Cell Signaling Technology, Beverly, MA), SIRT-1 (1:1000; Cell Signaling Technology, Beverly, MA), p-AMPK (Thr172) (1:1000; Cell Signaling Technology, Beverly, MA), FN (1:500; Abcam, Cambridge, MA), PGC-1 α (1:500; Affinity Biosciences, Cincinnati, OH).³⁵ β -Actin (1:1000; Cell Signaling Technology, Beverly, MA) was used as an internal control. The protein expression levels of the target genes were quantified by relative densitometry performed by ImageJ (National Institutes of Health) and normalized to the protein expression levels of β -actin. The values were shown as changes relative to those of the control sample.⁴⁹

Immunofluorescence (IF) Assay

MCs on 35 mm glass dishes were treated as described previously.⁵⁰ The plates were washed with cold PBS and fixed with 4% paraformaldehyde for 20 min at room temperature. After permeabilizing with 0.3% of triton, the unspecific sites of the cells were blocked by 1% BSA + 0.3% triton. Then, the cells were incubated with primary antibody experimenters used before,³⁵ rabbit anti-SIRT-1 diluted 1:10 (Santa Cruz Biotechnology) and mouse anti-PARP-1 diluted 1:10 (Enzo Life Sciences, Farmingdale NY) overnight at 4°C and incubated with fluorescence conjugated secondary antibodies (Beyotime, China) counterstained with DAPI next day. Images were analyzed under a fluorescence-inverted microscope.

PARP Activity Assay

PARP activity was tested by detecting the quantity of biotinylated-NAD⁺ converting into nicotinamide and biotinylated-ADP ribose polymers via the universal colorimetric PARP assay kit (Trevigen Inc., Gaithersburg, MD). The activity of PARP-1 was detected in accordance with the manufacturer's instructions. Cell lysates from MCs which contains about 50 μ g of protein were added into a 96-well plate coated with histones and biotinylated poly-ADP-ribose, after 1h-incubation, treated the cells with streptavidin-horse-radish peroxidase, and read at 450 nm in a spectrophotometer.^{51,52}

MTT Assay

Presently, the MTT (3-(4,5-dimethylthiazol-2-yl)-5-(3-carboxymethoxyphenyl)-2-(4-sulphophenyl)-2H-tetrazolium) assay is widely used to access the viability and the cytotoxicity of cells.⁵³ To detect MCs viability, MTT assay was performed according to the manufacturer's instructions.

MCs were seeded in 96-well culture plates at a density of 1×10^4 cells per well. After an-overnight adsorption, the cells were incubated with medium either containing 5 mM glucose or 25 mM glucose, with PJ34 and/or EX527, respectively. And 25 mM glucose medium containing SIRT-1 overexpression lentiviral with or without PJ34 was also considered as experimental groups. Followed by incubation with 100 μ L MTT solution (10 mg/mL in PBS) for 4h, the MCs were further lysed, and the purple formazan crystals were solubilized with 100 μ L DMSO (Sigma-Aldrich). The wells containing only medium without MCs were set as a blank. Finally, we used an enzyme-linked immunosorbent assay reader (Thermo LabSystems, OY, Helsinki, Finland) to analyze the absorbance at 570 nm.⁵⁰

NAD⁺ Qualification

NAD⁺ in db/db kidney tissues and HG-induced MCs was estimated colorimetrically using an NAD⁺/NADH assay kit (Abcam, Cambridge, MA) as described previously.⁵⁰ The standard was prepared according to the manufacturer's protocol.

Statistical Analysis

All data are presented as mean \pm SD deviation from at least three independent experiments. Differences between two groups were evaluated using Student's *t*-test, and multiple comparisons were performed using one-way analysis of variance (ANOVA) followed by a Student–Newman–Keuls (SNK) test. A *p* value <0.05 was considered statistically significant.

Results

PARP-1 Activation and SIRT-1 Inhibition in db/db Mice Induced by Hyperglycemia

To study the interplay between SIRT-1 and PARP-1 in vivo, we induced hyperglycemia by using a mouse model of insulin resistance, which has been shown to generate ROS and DNA damage in kidney tissues.²³ As shown in **Figure 1A** and **B**, the activity of PARP-1 increased markedly in the renal tissues from db/db mice (DN group) compared to control db/m mouse (NC group) (*p* < 0.05), which followed by increasing expression of ECM protein fibronectin (FN) (*p* < 0.001) and decreasing expression of SIRT-1 (*p* < 0.05). Considering the results of many researchers in explaining the association between overexpression of PARP-1 and down-expression of SIRT-1 in diabetic cells, it is reasonable to consider the decrease expression of SIRT-1 in the kidney of db/db mice

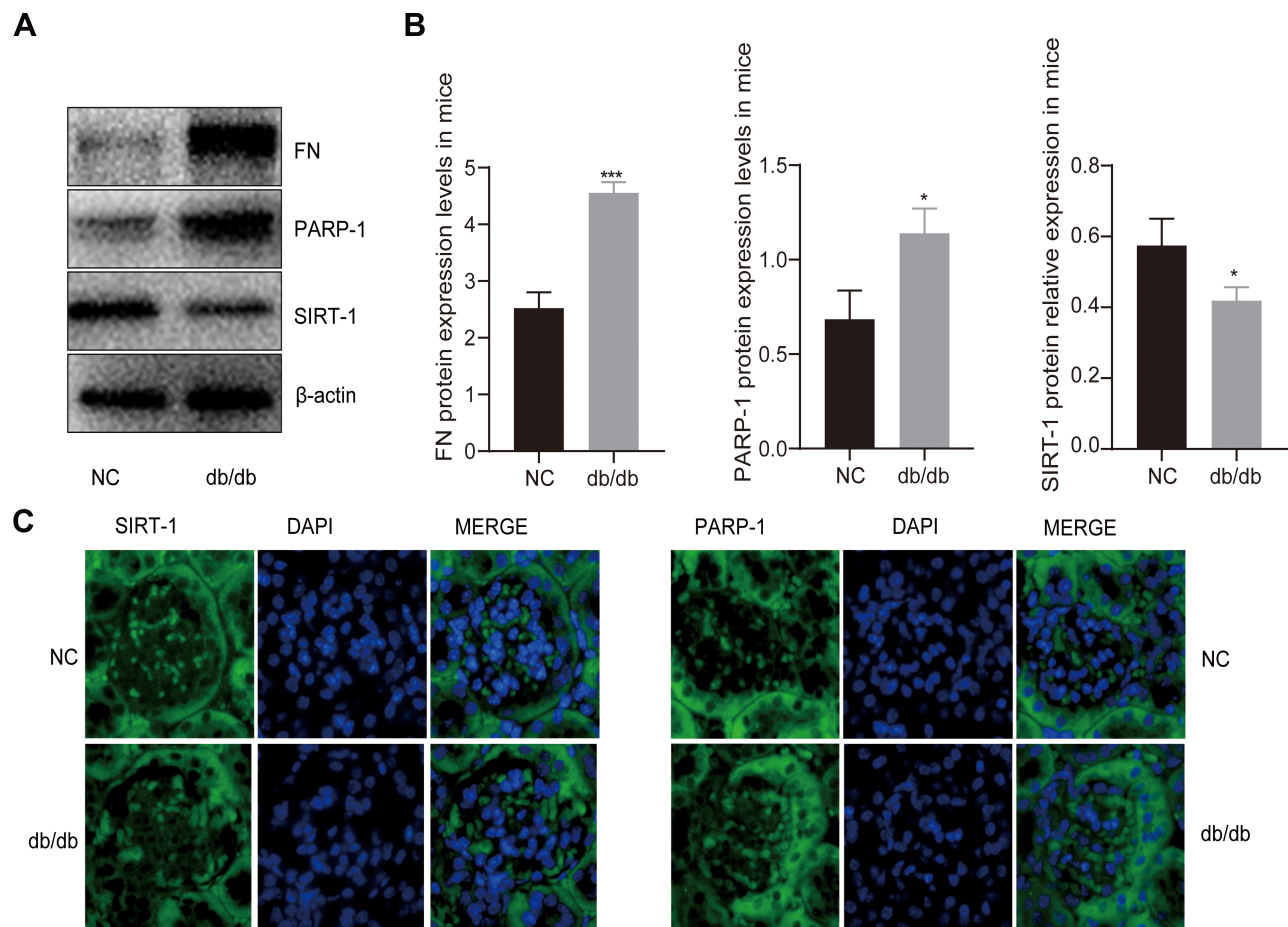


Figure 1 The expression of PARP-1 and SIRT-1 in db/db mice and db/m mice. **(A and B)** Western blotting results showed hyperglycemia decreased the levels of SIRT-1 but increased the activation of PARP-1 and the expression of FN in db/db mice compared to db/m group. **(C)** Immunofluorescence images showed that the expression of PARP-1 was increased in db/db group mice glomerulus compared with db/m group mice glomerulus, which was contrary to the expression of SIRT-1. Blue, nuclear staining (DAPI); green, target protein staining. Data are presented as mean \pm SD. We conduct 3 independent experiments. *Compared with NC, $P < 0.05$; ***compared with NC, $P < 0.001$.

significant.^{14,40} In addition, the fluorescence of PARP-1 was stronger in the glomeruli of DN mice; however, the fluorescence of SIRT-1 was stronger in the NC mice, which demonstrated that the expression of PARP-1 was significantly higher in the glomeruli of the untreated DN mice compared to that of NC mice, on the contrary, the level of SIRT-1 in DN group was significantly lower compared to the NC group (Figure 1C). These results strongly suggested that hyperglycemia may lead to the accumulation of FN and increase of PARP-1 along with the decline of SIRT-1, which indicates a potential link between PARP-1 and SIRT-1 in diabetic mice.

The Effect of PARP-1 and SIRT-1 in High Glucose-Induced Activating MCs

To further investigate the role of SIRT-1 and PARP-1 in high glucose-stimulated MCs in vitro, we applied PJ34,

EX527 or Lv-SIRT-1 to co-incubate MCs with high glucose according to the above research. We postulated that PARP-1 inhibition will lead to the activation of SIRT-1 since both PARP-1 and SIRT1 are NAD⁺ dependent (See [Supplementary data 4 and Figure 4](#)).³⁹ As is shown in [Figure 2A and B](#), high glucose (25 mM) over-activated PARP-1 ($p < 0.05$) and declined SIRT-1 protein expression ($p < 0.001$), thus leading to increase FN protein expression ($p < 0.05$), compared to that of normal glucose (5 mM). PARP-1 inhibitor PJ34 alleviated the protein level of PARP-1 ($p < 0.05$), promoted the expression SIRT-1 ($p < 0.05$) and FN ($p < 0.01$) while SIRT-1 inhibitor EX527 worsened the above change ($p < 0.05$) ([Figure 2A and B](#)). Moreover, when PJ34 was incubated with EX527, the changes in PARP-1, SIRT-1 and FN protein levels were between the PJ34 and EX527 groups ([Figure 2A and B](#)). In addition, Lv-SIRT-1 totally reversed the protein level of PARP-1,

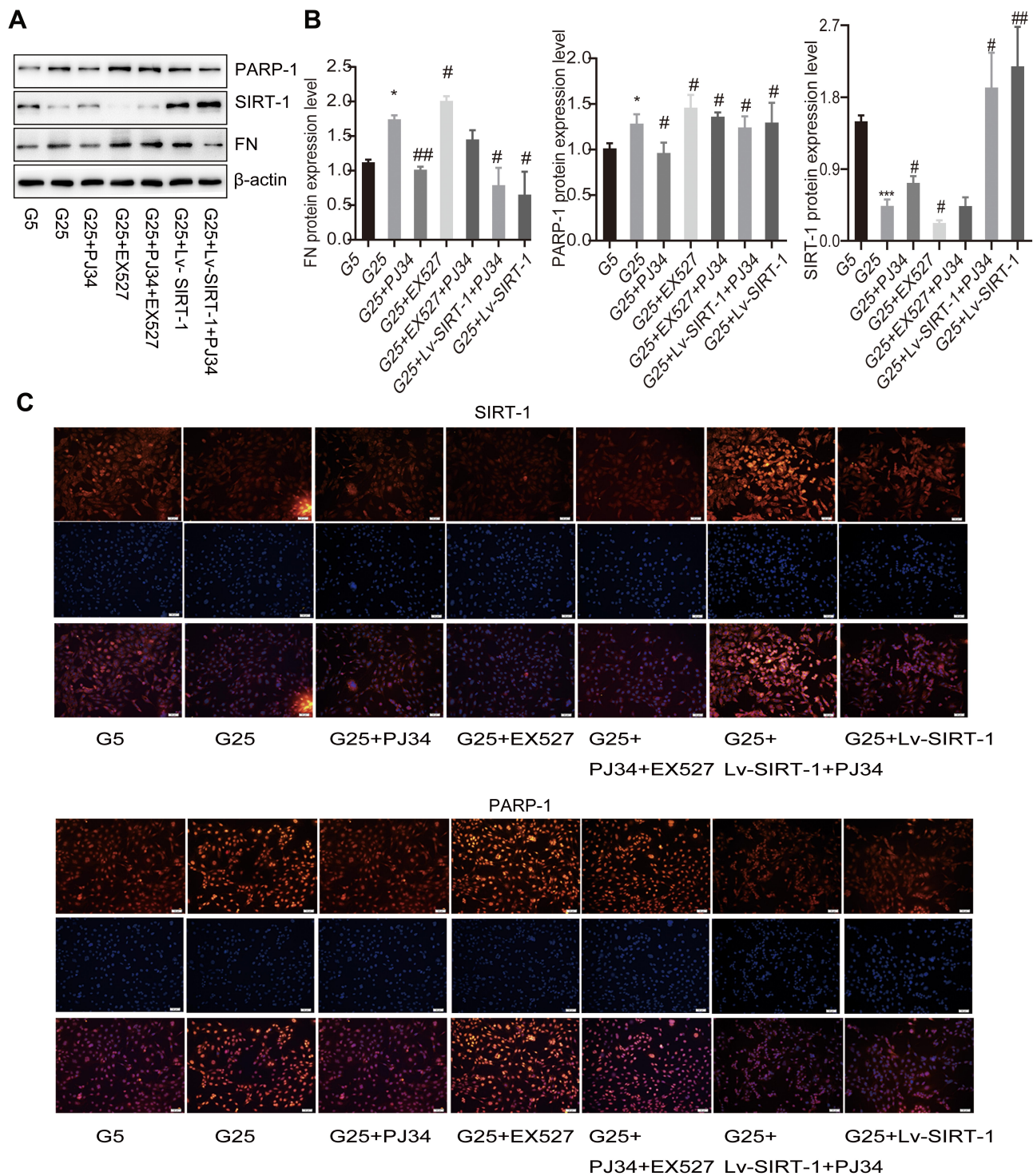


Figure 2 The effect of PARP-1 and SIRT-1 in MCs. (A and B) Proteins expression in MCs in high glucose condition or low glucose condition incubated with different reagents by Western blotting. β -Actin was used as control of protein loading. Bars represent mean \pm SD of three independent experiments. *Compared with G5, $P < 0.05$; ***compared with G5, $P < 0.001$. #Compared with G25, $P < 0.05$; ##compared with G25, $P < 0.01$. (C) Immunofluorescence images showed the change of protein SIRT-1 and PARP-1 expression in different MCs groups. Blue, nuclear staining (DAPI); green, target protein staining. Images were analyzed by confocal microscopy under a 50- μ m ruler.

SIRT-1 and FN, while incubation PJ34 with Lv-SIRT-1 did even better ($p < 0.05$) (Figure 2A and B). The above results are consistent with the results in

immunofluorescence (Figure 2C), which indicates that PARP-1 and SIRT-1 may exert a completely different role in DN.

The Change of Cell Viability is Related to the Activity of PARP-1

In the meanwhile, we examined the viability of hyperglycemia induced-MCs by MTT assay in different groups (Figure 3B). Hyperglycemia can strongly elevate the viability of MCs compared to the low glucose ($P < 0.01$) and this promotion can be increased or declined by EX527 ($p < 0.001$) and PJ34 ($p < 0.001$), respectively. Overexpression SIRT-1 also cause a decline in cell viability ($p < 0.001$). In addition, when extra PJ34 and EX527 are added to high glucose to stimulate MCs, the changes in cell viability were between the PJ34 and EX527 groups (Figure 3B). The likely reason is that PARP-1 exposure to high glucose levels induces a promotion of cell viability in the MCs, while SIRT-1 possesses a reverse role. To further determine the effect of PARP-1 on hyperglycemia-induced MC activation, the activity of PARP-1 was observed (Figure 3A). The trend and outcomes were much the same between the PARP-1 activity and cell viability in each group, which further demonstrated PARP-1 and SIRT-1 had an opposite effect in diabetic kidney tissues.

PARP-1-Induced SIRT-1 Down-Regulation is Mediated by the SIRT-1-AMPK-PGC-1 α Signaling Pathway

Both PARP-1 and SIRT-1 affect chromatin remodeling and thus modulation of gene expression affecting inflammation

and diabetic related diseases.⁵⁴ In addition, the interaction of PARP-1 and SIRT-1 plays a role in the regulation of oxidative stress and diabetic related diseases.^{35,55} To clarify the molecular mechanism of how PARP-1 regulates SIRT-1 expression, we screened a variety of target proteins of which SIRT-1 can participate in the deacetylation, such as PGC-1 α and p-AMPK (Thr¹⁷²) (Figure 4A and B). We determined that an increase in SIRT-1 expression was associated with elevation of protein PGC-1 α ($p < 0.05$) and p-AMPK ($p < 0.05$) expression in the MCs, accompanied by a drop while inhibiting the level of protein SIRT-1 ($p < 0.05$) and reversed by an inhibition of PARP-1 (Figure 4A and B). In conclusion, these results suggest a direct relationship between SIRT-1 and PGC-1 α protecting renal tissues against oxidative stress and accumulation ECM which participate in the pathogenesis of DN disease. And inhibiting PARP-1 leading to an up-expression of SIRT-1 can be considered as an effective therapeutic target for DN (summarized in Figure 4C).

Discussion

The accumulation of ECM plays a critical role in the pathophysiology of glomerulosclerosis in DN, which can be triggered by various pathological conditions such as oxidative stress and autophagy.^{4,56} A plethora of evidences have shown a crucial role of SIRT-1 in retarding the ECM accumulation during MCs injury.^{57,58} In addition, Chen et al⁵⁹

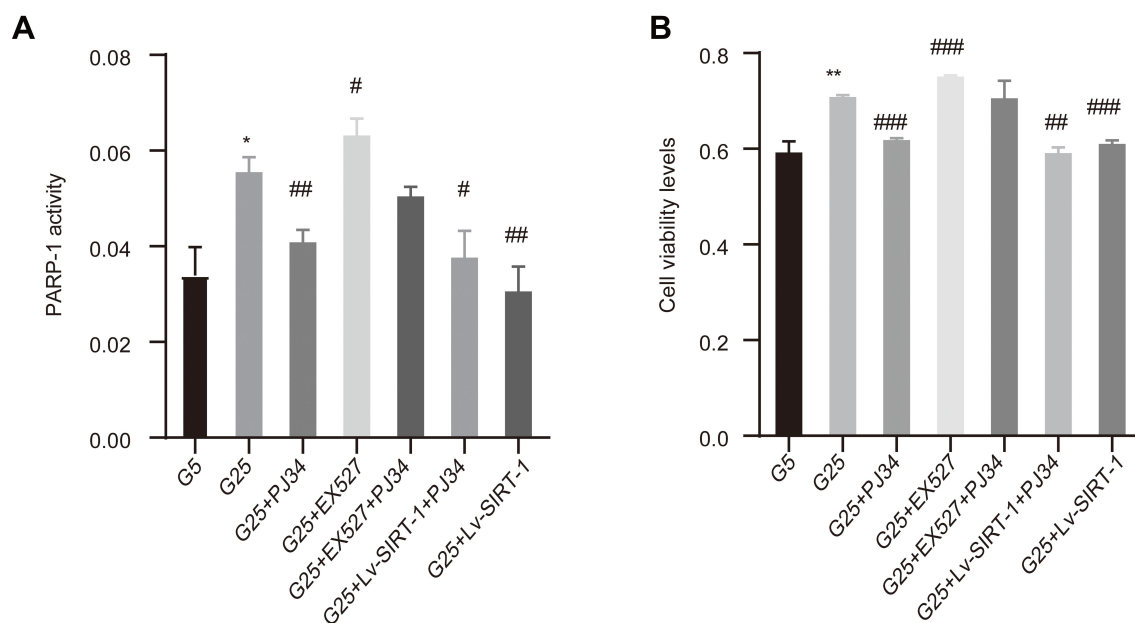


Figure 3 The result of PARP-1 activity and cell viability. (A) PARP-1 activity was measured by PARP-1 activity kit. (B) Cell viability measured by MTT assay upon 48 h treatment of different reagents. Data of three experiments are presented as mean \pm SD. *Compared with G5, $P < 0.05$; **compared with G5, $P < 0.01$. #Compared with G25, $P < 0.05$; ###compared with G25, $P < 0.01$; ####compared with G25, $P < 0.001$.

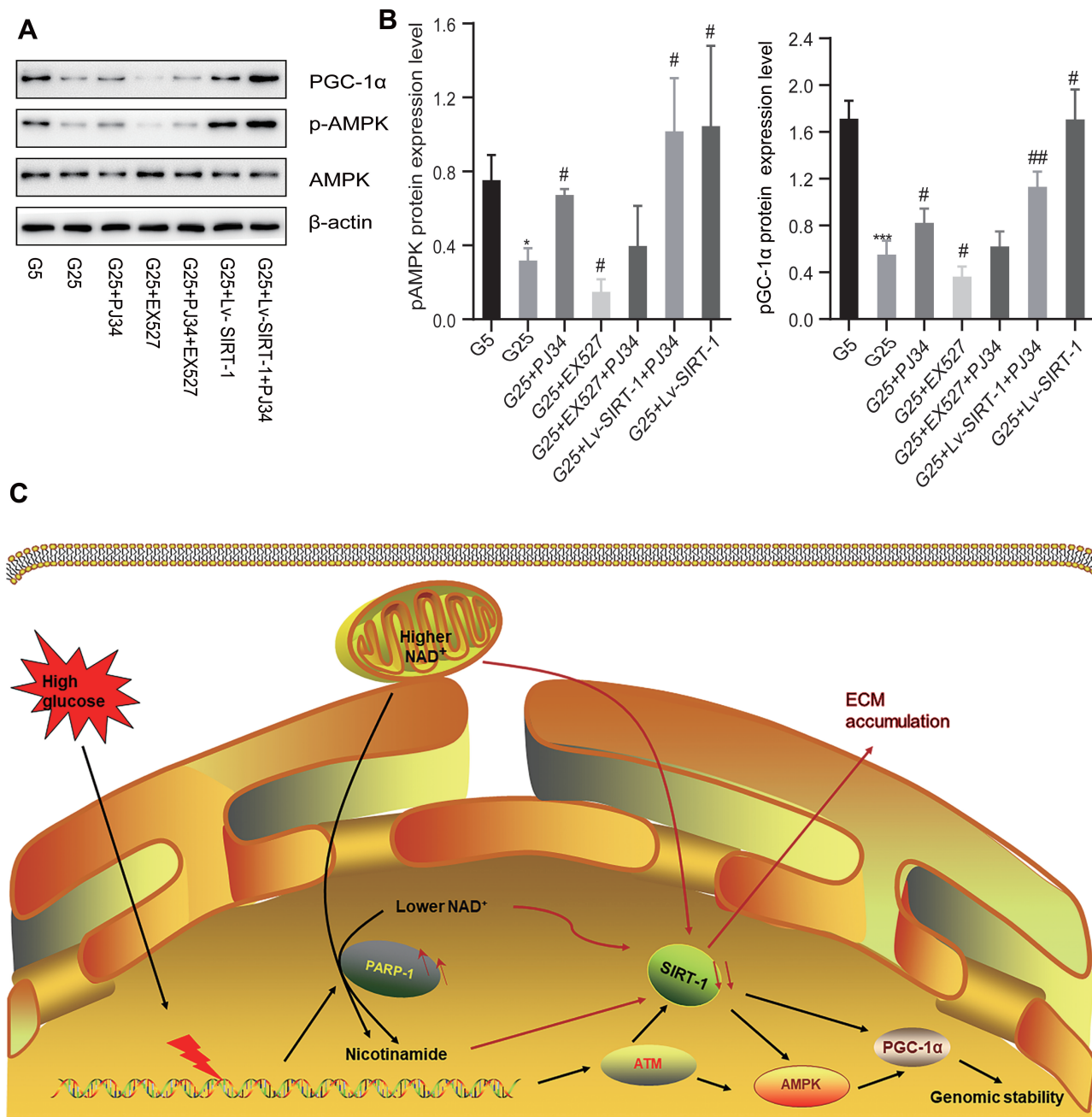


Figure 4 PARP-1 may have a crosslink with SIRT-1 via PGC-1α axis. **(A and B)** The expression of pAMPK and PGC-1α in MCs in high glucose condition or low glucose condition incubated with different reagents by Western blotting. Data of three experiments are presented as mean ± SD. *Compared with G5, $P < 0.05$; ***compared with G5, $P < 0.001$. #Compared with G25, $P < 0.05$, ##compared with G25, $P < 0.01$. **(C)** The possible pathway of the accumulation of ECM in DN patients: PARP-1-NAD⁺-SIRT-1-PGC-1α axis. High glucose can cause a strong stress to DNA replication, which then be transduced by DNA damage response sensors, such as PARP-1, ataxia telangiectasia mutated (ATM). Then SIRT-1 and AMPK propagate the signal via posttranslational modifications of PGC-1α and other proteins to maintain the integrity of genome. However, the overactivation of PARP-1 can reverse this protectable role, which subsequently contributes to the subcellular difference in NAD⁺ content between the nucleus and mitochondria and leads to the down-expression of SIRT-1 and finally causes the activation of ECM. The red arrow refers to the pathological phenomenon caused by excessive activation of PARP-1.

determined a significant connection of PARP-1 and ECM in the effect of fluorofenidone attenuating experimental cyclosporine A-induced kidney injuries through a variety of molecular biology experiments. And Xu et al⁶⁰ also illustrated that hyperhexosemia and diabetes cause an upregulation of ECM proteins at the transcriptional level in the

kidney via a signaling pathway mediated by PARP, which can be subsequently attenuated by PARP blockade. These findings suggest that PARP-1 attributes to the ECM accumulation in kidney. Moreover, previous studies indicated that there was a direct effect of SIRT-1 on PARP-1 regulation through the deacetylation of PARP-1 in many other

kidney-related diseases.^{20,61} However, the precise connection between SIRT-1 and PARP-1 in ECM accumulation in DN remains unknown.

To elucidate this point, we measured the expression of PARP-1 and SIRT-1, as well as FN in kidney tissue of db/db mice and wild-type mice. The increase in PARP-1 activity and the reduction expression of SIRT-1 were detected, coupled with increased FN expression in db/db mice kidney compared with wildtype mice kidney. PARP-1 inhibitor treatment by PJ34 reversed these changes, suggesting that there may exist a potential relationship between PARP-1 and SIRT-1 and down-expression of PARP-1 could exert a vital protective role in the diabetic mice.

We further tested the respective effects of an overexpression or decline of SIRT-1 on PARP-1 and an inhibitor of PARP-1 on SIRT-1 in hyperglycemia-activated MCs. Hyperglycemia induced the accumulation of ECM in MCs and increased the expression of FN and PARP-1, followed by the down-expression of SIRT-1. The reduction in PARP-1 activity resulting from PARP-1 inhibition by PJ34 in the hyperglycemia mice was followed by increasing SIRT-1 and FN expression in MCs. And PARP-1 levels were significantly declined in MCs treated with overexpression SIRT-1 or PJ34 and increased after down-expression SIRT-1 via the treatment of EX257. These findings indicate that SIRT-1 and PARP-1 have an interplay in hyperglycemia-activated MCs.

In addition, PARP-1 can be modulated by PGC-1 α signaling.^{14,39,40} Wadman et al⁴⁰ reported that activation of SIRT-1-PGC-1 α signaling pathway can protect cardiomyocytes from suffering diabetes through downregulating PARP-1. In this study, we found that hyperglycemia-induced MC activation suppressed SIRT-1 activity along with declining PGC-1 α signaling, both of which were reversed by PARP-1 inhibitor PJ34. The possible mechanism employed by PJ34 is the inhibition of AMPK-PGC-1 α signaling by declining SIRT-1 deacetylation. To prove this hypothesis, MCs were pretreated with the Lv-SIRT-1 or inhibitor EX257. MCs treated with Lv-SIRT-1 showed enhanced PGC-1 α and higher p-AMPK levels compared to PJ34 group, while a completely opposite effect was seen with inhibitor EX257 treatment. Therefore, PARP-1 acts as an SIRT-1 activity opponent to inhibit the deacetylation of PGC-1 α and enhances DNA breaks in hyperglycemia-activated MCs.

To date, the role of PARP-1 and SIRT-1 in DN remains still controversial. In this experiment, an obvious increase of PARP-1 and a dramatic decrease of SIRT-1 were observed in hyperglycemia-activated MCs and db/db mice followed by the

imbalance of AMPK/PGC-1 α pathway, suggesting it reasonable to consider PARP-1 and SIRT-1 as adaptive targets to combat kidney damage caused by hyperglycemia in the future. The treatment of regulating AMPK/PGC-1 α pathway could protect the kidney from damage caused by hyperglycemia and possibly prevents stress-induced PARP-1 activation.

Conclusion

In summary, we elucidated the possibility of a potential relationship between PARP-1 and SIRT-1 at the animal and cell levels, respectively. Our study suggests that PARP-1 involves in the downregulation of SIRT-1 impeding renal function in db/db mice. And the reduction of PARP-1 could protect MCs from hyperglycemia-induced MC activation ameliorating renal ECM accumulation via SIRT-1-AMPK-PGC-1 α pathway. These findings provide the experimental basis to further investigate the therapeutic target to DN through this intricate connection between PARP-1 and SIRT-1.

Abbreviations

DN, diabetic nephropathy; ECM, extracellular matrix; PARP-1, poly (ADP-ribose) polymerase 1; SIRT-1, sirtuin 1; AMPK, AMP-activated protein kinase; PGC-1 α , peroxisome proliferator-activated receptor gamma coactivator 1- α ; HG, high glucose; FN, fibronectin; DKD, diabetic kidney disease; ESRD, end-stage renal disease; NAD⁺, nicotinamide adenine dinucleotide; PAR, ADP-ribose polymers; CR, caloric restriction; NF- κ B, nuclear factor kappa B; DMEM, Dulbecco's Modified Eagle Medium; FBS, fetal bovine serum; MOI, multiplicity of infection.

Data Sharing Statement

The datasets used and analyzed during the current study are available from the corresponding author on reasonable request.

Ethics Approval

All applicable international, national, and/or institutional guidelines for the care and use of animals were followed. All animal experiments were guided in accordance with the guidelines of the National Institutes of Health, eighth edition, 2011. All experiments involving ethics have been ethically approved by the Department of Science and Technology of the First Affiliated Hospital of Nanchang University, and the ethics number is 2020 Medical Research Ethics No. 116.

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Author Contributions

All authors made substantial contributions to conception and design, acquisition of data, or analysis and interpretation of data; took part in drafting the article or revising it critically for important intellectual content; agreed to submit to the current journal; gave final approval of the version to be published; and agree to be accountable for all aspects of the work. Hengmei Zhu and Zhi Fang contributed to the work equally. Everyone participated in the final approval of the manuscript.

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Disclosure

The authors report no conflicts of interest in this work.

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