Glycine increases glyoxalase-1 function by promoting nuclear factor erythroid 2-related factor 2 translocation into the nucleus of kidney cells of streptozotocin-induced diabetic rats

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Keywords

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ABSTRACT

Aims/Introduction: We have previously reported that glycine suppresses the advanced glycation end-products signaling pathway and mitigates subsequent oxidative stress in the kidneys of diabetic rats. In the present study, we investigated whether this beneficial effect was associated with upregulation of glyoxalase-1 (Glo1) and activation of the nuclear factor erythroid 2-related factor 2 (Nrf2).

Materials and Methods: Both healthy rats and streptozotocin-induced diabetic rats were administrated with glycine (1% added to the drinking water) for 12 weeks. The function of Glo1, messenger ribonucleic acid (mRNA) and protein expressions of Nrf2, and markers of oxidative status were measured in the kidneys. The mRNA expressions of other downstream signaling molecules of the Nrf2 pathway were also determined.

Results: The mRNA and protein expressions, as well as the activity of Glo1, were decreased in the kidneys of diabetic rats, accompanied by diminished glutathione levels. After glycine treatment, these parameters of Glo1 function were markedly increased. Compared with the control group, the levels of Nrf2 mRNA and protein in the total kidney lysis were both markedly elevated in the diabetic group and glycine-treated group. However, the nuclear translocation of Nrf2 was significantly increased in the glycine-treated group than in the diabetic group. In addition, the anti-oxidant capacity and the expressions of other downstream molecules of the Nrf2 signaling pathway were significantly increased after glycine treatment.

Conclusions: The present study shows that glycine might enhance the function of Glo1 and restore anti-oxidant defense by promoting the nuclear translocation of Nrf2, thus inhibiting advanced glycation end-products formation and protecting against renal oxidative stress.

INTRODUCTION

Oxidative stress plays a central role in the development of diabetic nephropathy^{1,2}. The advanced glycation end-products (AGEs) signaling pathway is one of the five major mechanisms of oxidative stress³. In the traditional view, AGEs

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mainly result from a slow, non-enzymatic reaction between glucose and large biomolecules⁴. Once formed, AGEs can activate the receptor for AGEs (RAGE), thus triggering a vicious loop of inflammation and oxidative stress^{3,5}, which eventually leads to renal damages. Recent findings suggest that methylglyoxal (MG), a highly-reactive byproduct of glycolysis, is much more potent than glucose in promoting the

formation of AGEs and in activating the AGE/RAGE signaling pathway 6,7 .

The glyoxalase system plays a crucial role in detoxifying MG. Using glutathione (GSH) as a cofactor, glyoxalase-1 (Glo1) and glyoxalase-2 efficiently catalyze the conversion of MG into lactate, thus reducing the precursor of AGEs⁸. Studies have shown that knockdown of Glo1, the rate-limiting enzyme of the glyoxalase system, can mimic diabetic nephropathy in healthy mice⁹; whereas upregulation of Glo1 suppresses the AGE/RAGE pathway, thus attenuating renal damage, such as mesangial matrix expansion¹⁰ and podocytes apoptosis¹¹. Therefore, the enhancement of Glo1 function might suppress the AGE/RAGE signaling pathway and the subsequent renal damage.

It was reported that Glo1 is a target gene of nuclear factor erythroid 2-related factor 2 (Nrf2), a transcription factor. Under basal conditions, the activation of Nrf2 is inhibited by Kelchlike ECH-associated protein 1, which binds to Nrf2 and forms a suppressive complex in the cytoplasm¹². Under oxidative stress, Nrf2 dissociates from the complex and translocates from the cytoplasm to the nucleus, where it binds to the anti-oxidant response element and promotes the transcription of numerous genes that encode anti-oxidant enzymes, including superoxide dismutase (SOD), catalase (CAT), glutathione peroxidase (GPx), heme oxygenase-1 (HO-1) and NAD(P)H (the reduced form of either nicotinamide adenine dinucleotide or nicotinamide adenine dinucleotide phosphate) quinone dehydrogenase 1 (NQO1)¹³. Nrf2 also regulates genes that encode GSH-related enzymes, such as glutathione peroxidase-1 (GPx-1), glutathione synthetase (GSS) and glutamate-cysteine ligase¹³. However, when oxidative stress persists, the nuclear translocation of Nrf2 is hampered. As a result, the anti-oxidant response of Nrf2 is impaired¹⁴, leading to uncontrolled oxidative stress in the kidneys^{15,16}. Therefore, activation of Nrf2 might be beneficial for increasing the function of Glo1, as well as ameliorating oxidative stress.

Glycine is one of the predecessors of GSH, a major endogenous anti-oxidant in the human body. GSH is crucial for the detoxifying process catalyzed by Glo1, and depletion of GSH can suppress the function of Glo117-19 and induce significant MG accumulation²⁰. In individuals with either diabetes or metabolic syndrome, the levels of glycine and GSH are reduced as compared with healthy individuals, but this reduction can be reversed after oral glycine treatment²¹⁻²³. We have previously reported that oral supplement of glycine can inhibit the AGE/RAGE signaling pathway and the subsequent oxidative stress in the kidneys of diabetic rats^{24,25}, but whether this beneficial effect is associated with the upregulation of Glo1 still remains unknown. Glycine also increases the activity of anti-oxidant enzymes, such as SOD, CAT and GPx, under oxidative stress²⁶⁻²⁹, but the mechanism is unclear. Therefore, we speculate that the beneficial effects of glycine on suppressing the formation of AGEs and increasing anti-oxidant capacity might be related to the activation of Nrf2.

In the present study, we investigated the effect of oral glycine treatment on the function of Glo1 and the anti-oxidant defense in the kidneys of STZ-induced diabetic rats, and explored whether this effect is associated with Nrf2 activation.

METHODS

Experimental animals

All protocols and procedures of the present study were approved by the Ethics Committee for Animal Experimentation of the Faculty of Peking University First Hospital. All animal experiments were conducted following the national guidelines and the relevant national laws on the protection of animals. Six-week-old male Wistar rats were housed under a 12-h lightdark cycle. After 2 weeks of acclimation, all rats were randomly assigned either to an experimental group or to a healthy control group. The experimental group was intraperitoneally injected with a single dose of streptozotocin (STZ; 65 mg/kg of bodyweight; Sigma-Aldrich, St. Louis, MO, USA). Diabetes mellitus was confirmed after 2 weeks by measuring blood glucose levels (levels >16.7 mmol/L were confirmed as diabetic). The experimental group was then divided into a diabetes mellitus group (non-treated diabetic rats receiving normal drinking water, n = 8) and a DG group (diabetic rats receiving 1% (w/ v) glycine in drinking water ad libitum, as reported by others^{30–32}, n = 7). Likewise, the healthy control group was divided into two groups: a control group (healthy rats receiving normal drinking water, n = 8) and a CG group (healthy rats receiving 1% [w/v] glycine in drinking water ad libitum, n = 8). After 12 weeks, all rats were killed. Serum samples were collected to determine biochemical profiles by an Automatic Biochemical Analyzer (7600; Hitachi, Tokyo, Japan). Equal amounts of urine and serum samples were used to measure the levels of cystatin-C (CSB-E08385r; CUSABIO, Wuhan, China).

Measurement of renal Glo1 activity

The activity of Glo1 was measured according to the method established by Arai *et al.*³³. Samples of kidney cortex were washed in phosphate-buffered saline, and sonicated in 10 mmol/L sodium phosphate buffer (pH 7). The protein concentration of the supernatants was adjusted to the same level in all samples. In a 96-well UV plate, 125 μ L sodium phosphate buffer (100 mmol/L, pH 6.6), 25 μ L GSH (40 mmol/L), 25 μ L MG (40 mmol/L) and 70 μ L Millipore water were added into each well and incubated at 37°C for 15 min. After incubation, 5 μ L solubilized protein was added into each well. The absorbance at 240 nm (A240) was monitored every 2 min. The Glo1 activity was displayed as the change in A240 per minute per mg protein.

Quantitative real-time reverse transcription polymerase chain reaction

The method of real-time reverse transcription polymerase chain reaction has been previously reported²⁴. The primer sequences were as follows: Nrf2 GAATAAAGTTGCCGCTCAGAA (forward), AAGGTTTCCCATCCTCATCAC (reverse); Glo1 GAA

GATGACGAGACGCAGAGTTAC (forward), CAGGATCTT GAACGAACGCCAGAC (reverse); CAT ACACTTTGACAG AGAGCGGA (forward), TTTCACTGCAAACCCACGAG (reverse); SOD 2 CTGCTGGGGATTGATGTGTG (forward), CTACAAAACACCCACCACGG (reverse); HO-1 CTGGTGAT GGCCTCCTTGTA (forward), GATGAGTACCTCCCACCT CG (reverse); NQO1 CCTGATTGTATTGGCCCACG (forward), CTCAGGCGGCCTTCCTTATA (reverse); GPx-1 GAC CGACCCCAAGTACATCA (forward), GCAGGGCTTCTAT ATCGGGT (reverse); glutamate-cysteine ligase catalytic subunit GACCATCGACACGGAGATCT (forward), GAGTTTGGAG GAGGAGGCTT (reverse); glutamate-cysteine ligase modifier subunit CACAACTCAGGGGCCTTGTA (forward), CGCCAG GGAGGTACTCAAAC (reverse); GSS CGTGGTGCTACTGA TTGCTC (forward), AACAGCCTTCGGTTTTGGTC (reverse); β-actin ATCTGGCACCACACCTTC (forward), AGCCAGGT CCAGACGCA (reverse). Gene expression was normalized to β -actin and calculated using the $2^{-\Delta\Delta CT}$ method.

Western blot

To obtain total protein samples, frozen samples of kidney cortex were homogenized in ice-cold radioimmunoprecipitation assay lysis buffer (P0013; Beyotime, Beijing, China) with 1% protease inhibitors cocktail and phenylmethane sulfonyl fluoride. The protein concentration of the supernatants was determined by the bicinchoninic acid assay kit and adjusted to the same level in all samples. The separation of nuclear and cytosol protein was carried out according to the manufacturer's instructions (KGP1100; KeyGEN BioTECH, Beijing China).

The method of western blot has been previously reported elsewhere²⁴. The membrane was incubated overnight with a primary mouse anti-Nrf2 antibody (1:1,000 ab89443; Abcam, Cambridge, UK) or mouse anti-Glo1 antibody (1:3,000, MA1-13029; Invitrogen, Waltham, MA, USA) at 4°C. After washing, the membrane was incubated with a secondary antibody (1:5,000 HRP-coupled anti-mouse antibody, ZB-2305; ZSGB-BIO, Beijing, China) for 1 h at room temperature. The membrane was probed for β -actin (1:3,000, TA-09; ZSGB-BIO) as a loading control for total protein or cytoplasmic protein. Histone was used as a loading control for nuclear protein (1:3,000, ab11079; Abcam). The bands were visualized using the ECL Western Blotting Substrate (Thermo Fisher, Hudson, NY, USA), and quantified with Image-Pro Plus 6.0 software (Media Cybernetics, Rockville, MD, USA).

Measurement of renal anti-oxidant capacity

Briefly, renal cortex samples were lysed in ice-cold HEPES buffer and centrifuged at 17,000 g for 20 min at 4°C. The supernatants were collected and the protein concentrations of the samples were adjusted to the same level. Equal amounts of the protein samples were subjected to the measurement of total GSH levels (A061-1; Jiancheng, Nanjing, China), ratio of reduced GSH/oxidized GSH (GSSG; S0053 GSH kit; Beyotime), malondialdehyde levels (S0131; Beyotime), SOD activity (706002; Cayman Chemicals, Ann Arbor, MI, USA), CAT activity (707002; Cayman Chemicals), 8-hydroxy-2'-deoxyguanosine (8-OHdG) levels (CSB-E10526r; CUSABIO), protein carbonyl levels (STA-310, OxiSelectTM protein carbonyl ELISA kit; Cellbiolabs, San Diego, CA, USA) and 8-iso-prostaglandin F2 α (8-iso-PGF₂ α) levels (ab133043; Abcam) according to the manufacturers' instructions.

Statistical analysis

SPSS 20.0 (SPSS Inc., Chicago, IL, USA) was applied to carry out statistical analysis of the data. Quantitative data are shown as the mean \pm standard error of the mean (for normally distributed data) or medians and interquartile ranges (for non-normally distributed data). Differences between groups were assessed using one-way analysis of variance (ANOVA) for normally distributed data, followed by Tukey's post-hoc test. In addition, the Kruskal–Wallis test was used for non-normally distributed data. A *P*-value <0.05 was considered statistically significant.

RESULTS

Effect of glycine on bodyweight, glucose control and biochemical profiles

Compared with the control group, the bodyweight in the diabetes mellitus group decreased significantly (Figure 1a). After glycine treatment, the bodyweight seemed unaltered. The blood glucose levels in the diabetes mellitus group increased sharply compared with the control group after the induction of diabetes mellitus. However, treatment of glycine did not affect the glucose levels (Figure 1b). Likewise, the levels of hemoglobin A1c in the diabetes mellitus group were higher than in the control group, but were not significantly changed after glycine treatment (Table 1). No marked alterations were observed in the healthy control rats treated with glycine (the CG group).

The serum creatinine levels in the diabetes mellitus group were not altered compared with those in the control group. However, after glycine treatment, the serum creatinine levels were significantly lower than in the diabetes mellitus group (P < 0.05). The levels of urea and cvstatin-C in the serum increased significantly in the diabetes mellitus group than in the control group (P < 0.05). After glycine treatment, the level of urea significantly was decreased (P < 0.05). The levels of serum cystatin-C tended to be reduced, but this reduction did not reach statistical significance. The levels of urinary cystatin-C in the diabetes mellitus group were increased compared with the control group (P < 0.05), but were significantly decreased after glycine treatment (P < 0.05). The levels of aspartate transaminase, alanine transaminase and triglyceride in the diabetes mellitus group were significantly higher than in the control group (P < 0.05), but seemed unaffected by glycine treatment. No marked changes in other parameters of lipid profiles were detected among the four groups.

Glycine increases renal Glo1 function

To assess the function of Glo1, we measured the messenger ribonucleic acid (mRNA) and protein expressions, as well as



Figure 1 | (a) Bodyweight and (b) blood glucose levels. ****P* < 0.001 compared with the control group. CG, healthy rats receiving tap water with 1% glycine added; Control, healthy rats receiving normal tap water; DG, diabetic rats receiving tap water with 1% glycine added; DM, diabetic rats receiving normal tap water.

Table 1 | Hemoglobin A1c levels and serum biochemical profiles after 12 weeks of treatment

Group	Control	CG	DM	DG
HbA1c (%)	4.338 ± 0.47	4.363 ± 0.30	9.587 ± 0.69*	9.429 ± 0.62*
CREA (µmol/L)	42.5 ± 2.26	39.38 ± 2.16	40.88 ± 2.69	34.14 ± 2.14**
UREA (µmol/L)	8.71 ± 1.06	7.40 ± 1.31	$20.49 \pm 2.29^{*}$	12.3 ± 1.89**
Serum cystatin-C (ng/mL)	5.85 ± 0.41	6.68 ± 0.68	10.75 ± 1.61*	7.30 ± 0.68
Urinary cystatin-C (ng/mL)	3.00 ± 0.29	3.30 ± 0.43	8.84 ± 1.19*	3.75 ± 0.85**
ALT (mmol/L)	38.0 ± 4.96	36.2 ± 3.97	276.5 ± 10.91*	200.0 ± 12.72*
AST (mmol/L)	207.75 ± 15.77	109.5 ± 10.50	488.13 ± 16.61*	401.43 ± 18.86*
TG (mmol/L)	0.41 ± 0.37	0.47 ± 0.44	2.3 ± 1*	2.08 ± 1.18*
TCHO (mmol/L)	2.29 ± 0.67	2.09 ± 0.69	2.55 ± 0.69	2.46 ± 0.81
HDL (mmol/L)	0.65 ± 0.30	0.65 ± 0.36	0.89 ± 0.40	1.05 ± 0.40
LDL (mmol/L)	0.33 ± 0.33	0.22 ± 0.22	0.41 ± 0.42	0.42 ± 0.51

Data are expressed as the mean \pm standard error of the mean. *P < 0.05 compared with the control group; **P < 0.05 compared with the diabetic rats receiving normal tap water (DM) group. ALT, alanine transaminase; AST, aspartate transaminase; CG, healthy rats receiving tap water with 1% glycine added; Control, healthy rats receiving normal tap water; CREA: creatinine; DG, diabetic rats receiving tap water with 1% glycine added; HbA1c, hemoglobin A1c; HDL, high-density lipoprotein; LDL, low-density lipoprotein; TCHO, total cholesterol; TG, triglyceride.

the activity of Glo1 in the kidneys. As GSH serves as the cofactor of Glo1, the levels of renal GSH were determined. A decrease of total GSH level in the diabetes mellitus group was observed compared with the control group (P < 0.05; Figure 2a), but this reduction was reversed by glycine supplement (P < 0.001). Although the total GSH levels in the CG group tended to be higher than in the control group, no statistical significance was observed. The levels of reduced GSH were decreased in the kidneys of diabetic rats as compared with the control group (P < 0.001; Figure 2b), but were increased by glycine treatment (P < 0.01). Conversely, the levels of GSSG were increased in the diabetes mellitus group (P < 0.05; Figure 2c), but were decreased in the DG group (P < 0.01). Consequently, the ratio of reduced GSH/GSSG was significantly increased in the DG group as compared with the diabetes mellitus group (P < 0.01; Figure 2d).

The mRNA and protein expression of renal Glo1 were lower in the diabetes mellitus group than in the control group (P < 0.05; Figure 2e,f). After glycine treatment, the mRNA and protein expression of Glo1 were markedly increased (P < 0.001and P < 0.05, respectively). The activity of renal Glo1 was decreased in the diabetes mellitus group compared with the control group (P < 0.05; Figure 2g), but was increased in the DG group (P < 0.05). No obvious changes in the protein expression and activity of Glo1 in the CG group were observed compared with the control group.

Glycine promotes the translocation of Nrf2 from the cytoplasm into the nucleus

Both mRNA and protein expression of Nrf2 in the total protein of kidney samples were increased significantly in the diabetes mellitus group compared with those in the control group



Figure 2 | The assessments of glyoxalase-1 (Glo1) function in the kidneys. (a) The levels of total glutathione (GSH) in the kidneys. (b) The levels of reduced GSH in the kidneys. (c) Levels of oxidized GSH (GSSG) in the kidneys. (d) The ratio of reduced GSH/GSSG in the kidneys. (e) The messenger ribonucleic acid expression of renal Glo1. (f) The protein expression of renal Glo1. (g) The activity of renal Glo1. *P < 0.05; ***P < 0.001 compared with the control group. *P < 0.05; ***P < 0.001; *##P < 0.001 compared with the diabetic rats receiving normal tap water; CG, healthy rats receiving tap water with 1% glycine added; DG, diabetic rats receiving tap water with 1% glycine added.

(P < 0.05; Figure 3a,b), but remained unaltered after glycine treatment. However, the protein expression of Nrf2 in the cytoplasm of kidney samples was significantly higher in the diabetes mellitus group than in the DG group (P < 0.001; Figure 3c). Conversely, the protein expression of Nrf2 in the nucleus of kidney samples was significantly higher in the DG group than in the diabetes mellitus group (P < 0.001; Figure 3d). The mRNA expressions of other downstream signaling molecules in the Nrf2 pathway were detected in the kidneys. The mRNA expressions of CAT, SOD2, HO-1 and NQO1 in the diabetes mellitus group were not markedly changed as compared with the control group. However, the mRNA expressions of these genes were significantly increased in the DG group as compared with the diabetes mellitus group (P < 0.05, P < 0.05, P < 0.05 and P < 0.01, respectively; Figure 4a-d). The mRNA expression of GPx-1 was decreased in the diabetes mellitus group as compared with the control group (P < 0.05; Figure 4e), but was significantly increased after glycine treatment (P < 0.001). The mRNA expressions of GSS and glutamate-cysteine ligase catalytic subunit were lower in the diabetes mellitus group than in the control group (P < 0.01; Figure 4f,g), but were not altered after glycine treatment. No statistically significant changes were detected in the mRNA expression of glutamate-cysteine ligase modifier subunit (Figure 4h).

Glycine restores the impaired redox defense

The activities of CAT and SOD in the diabetes mellitus group were lower than in the control group (P < 0.05 and P < 0.01, respectively; Figure 5a,b), but were restored after glycine treatment (P < 0.001 and P < 0.01, respectively). The levels of renal 8-iso-PGF₂ α , 8-OHdG, malondialdehyde and protein carbonyls were increased in the diabetes mellitus group compared with the control group (P < 0.05, P < 0.01, P < 0.05 and P < 0.05, respectively; Figure 5c–f), but were significantly decreased after glycine treatment, except for 8-iso-PGF₂ α (P > 0.05, P < 0.01, P < 0.05, P < 0.05, P < 0.01, P < 0.05, P < 0.05, P < 0.01, P < 0.05, P < 0.05, P < 0.01, P < 0.05, P

DISCUSSION

The present study shows that oral glycine treatment increases the expression and activity of renal Glo1 by promoting the nuclear translocation of Nrf2 and increasing GSH synthesis, thus ameliorating oxidative stress and renal injury in diabetic rats.

We have previously reported that oral glycine treatment inhibited the formation of AGEs in the kidneys of STZ-induced diabetic rats, thus suppressing the expression of RAGE, Nox4 and the subsequent renal oxidative stress^{24,25}. As a result, renal damage, such as increased urinary albumin/creatinine rate, mesangial expansion and thickening of glomerular basement membrane, was decreased after glycine treatment^{24,25}. Besides our previous findings, Bahmani *et al.*³⁰ reported that oral



Figure 3 | The (a) messenger ribonucleic acid expression, (b) protein expression in the total tissue lysis, (c) cytoplasm and (d) nucleus of nuclear factor erythroid 2-related factor 2 (Nrf2) in the kidneys. **P < 0.01; ***P < 0.001 compared with the control group. ##P < 0.001 compared with the diabetic group without treatment (DM). CG, control rats receiving 1% oral glycine treatment; Control, healthy control group; DG, diabetic rats receiving 1% oral glycine treatment; PCR, polymerase chain reaction.

glycine treatment in diabetic rats significantly lowers plasma AGEs levels, thus delaying the development of cataracts. It was also hypothesized that glycine might work as a scavenger of intracellular glucose, and thereby protect the protein in the lens from excessive glycation³⁴. In addition, the high solubility of glycine might prevent the formation and precipitation of AGEs, as raised by Alvarado-Vásquez *et al.*³¹. Despite these findings, the upstream mechanism whereby glycine reduces the formation of AGEs still remains unclear.

The glyoxalase system plays a vital role in degrading MG, a highly-reactive metabolite that accelerates the formation of AGEs³⁵. As the rate-limiting enzyme of the glyoxalase system, Glo1 efficiently detoxifies MG, thus suppressing AGEs formation^{12,36,37}. It was reported that in non-diabetic mice, knockdown of Glo1 increased AGEs formation and accelerated oxidative stress to the extent observed in the diabetic mice, causing morphological changes, as seen in diabetic nephropathy⁹. However, this damage could be completely prevented by



Figure 4 | The messenger ribonucleic acid expression of (a) catalase (CAT), (b) superoxide dismutase 2 (SOD2), (c) heme oxygenase-1 (HO-1), (d) reduced nicotinamide adenine dinucleotide (phosphate) quinone dehydrogenase 1 (NQO1), (e) glutathione peroxidase-1 (GPx-1), (f) glutathione synthetase (GSS), (g) glutamate-cysteine ligase catalytic subunit (GCLC) and (h) glutamate-cysteine ligase modifier subunit (GCLM) in the kidneys. *P < 0.05; **P < 0.01 compared with the control group. #P < 0.05; ##P < 0.01; ##P < 0.01 compared with the diabetic group without treatment (DM). CG, control rats receiving 1% oral glycine treatment; Control, healthy control group; DG, diabetic rats receiving 1% oral glycine treatment; PCR, polymerase chain reaction.

overexpression of Glo1^{9,38–40}. Hence, in order to protect against renal damage induced by AGEs, it is of great importance to enhance the function of Glo1. In the present study, the mRNA and protein expressions, as well as the activity of renal Glo1, in the diabetes mellitus group were decreased compared with those in the control group. After glycine treatment, these parameters of Glo1 function were significantly increased. This result indicated that the decreased formation of AGEs in the glycine-treated rats observed in our previous study could be explained, at least in part, by the enhancement of Glo1 function.

It was reported that Glo1 is transcriptionally regulated by Nrf2¹², and activators of Nrf2 can promote mRNA and protein expressions of Glo1, and increase Glo1 activity^{10,41}. Recently, the role of anti-oxidants in activating Nrf2 has sparkled intensive attention. Anti-oxidants, such as resveratrol, quercetin and mangiferin, exert protective effects against the formation of AGEs and dicarbonyl stress by promoting nuclear translocation

of Nrf2, thus increasing Glo1 expression⁴¹⁻⁴⁵. In the present study, glycine treatment significantly enhanced the nuclear translocation of Nrf2, and increased the mRNA expression of Nrf2-targeted genes, such as Glo1, CAT, SOD2, HO-1, NQO1 and GPx-1, suggesting that the Nrf2 pathway was activated. The activities of SOD and CAT were also increased after glycine treatment, which is consistent with previous studies^{21,23}, but the mechanism remains unclear. One speculation is that glycine increases the level of GSH, which has been reported to increase SOD activity by abolishing reactive oxygen speciesinduced inactivation⁴⁶. This could also be explained by the increased mRNA expression of CAT and SOD2 after glycine treatment observed in the present study. In addition, glycine treatment decreased the levels of oxidative markers, such as 8-OHdG, malondialdehyde and protein carbonyls, indicating that the increased oxidative stress in the kidneys was suppressed. The renal levels of 8-iso-PGF2a in diabetic rats were not significantly reduced after glycine treatment, which might



Figure 5 | Estimation of anti-oxidant capacity in the kidneys of the four groups. (a) The activity of renal catalase. (b) The activity of renal superoxide dismutase. (c) The levels of renal 8-iso-prostaglandin F2 α (8-iso-PGF₂ α). (d) The levels of renal 8-hydroxy-2'-deoxyguanosine (8-OHdG). (e) The levels of renal malondialdehyde (MDA). (f) The levels of protein carbonyl in the kidneys. *P < 0.05; **P < 0.01; compared with the control group. "P < 0.05; **P < 0.01; compared with the control group. "P < 0.05; **P < 0.01; control rats receiving 1% oral glycine treatment; Control, healthy control group; DG, diabetic rats receiving 1% oral glycine treatment.

be explained by the limit of the method^{47,48} and the possible increase in arachidonic acid levels after glycine treatment⁴⁹. Consistent with the ameliorated oxidative stress, renal damage, such as mesangial expansion and microalbumin excretion, was reduced²⁵. The renoprotective effect of glycine was also supported by the decreased serum creatinine and urea levels, as well as the decreased urinary cystatin-C levels in the present study. It was reported that renal tubules are more prone to be injured by oxidative stress⁵⁰, and our previous study showed that glycine can specifically suppress oxidative stress in the tubular area rather than the glomerular area in the kidneys²⁴. In the present study, the levels of urinary Cystatin-C were significantly decreased in the DG group, indicating that glycine treatment is more effective in protecting against tubular damage.

It was reported that activators of Nrf2, such as sulforaphane and cinnamic aldehyde, could significantly attenuate hyperglycemia⁵¹. However, in the present study, the treatment of glycine did not exert such a beneficial effect in diabetic rats. Although there is not total agreement about the effect of glycine on hyperglycemia in diabetic models^{21,24,30,32,52}, the unchanged glucose levels after glycine treatment in the present study indicated that the beneficial effects of glycine might be independent from the alleviation in hyperglycemia.

GSH is a crucial cofactor of the glyoxalase system^{19,20}. The activity of Glo1 is proportional to the levels of GSH, and depletion of GSH can suppress the function of Glo1¹⁷⁻¹⁹ and induce significant MG accumulation²⁰. Glycine is one of the predecessors of GSH. It was suggested that glycine is rate-limiting for GSH synthesis⁵³. Under diabetes or metabolic syndrome, the level of GSH is reduced, but this reduction can be reversed by glycine treatment²¹⁻²³. Our previous study showed that the levels of glycine in both serum and kidneys were decreased in diabetic rats, but were restored after oral glycine treatment²⁴. In the present study, the levels of total GSH and the ratio of GSH/GSSG in the kidneys of the diabetes mellitus group were significantly lower than the control group, but were significantly elevated after glycine treatment, which is in accordance with previous studies. However, the mRNA expressions of the enzymes that are involved in GSH synthesis, such as glutamate-cysteine ligase catalytic subunit, glutamate-cysteine ligase modifier subunit and GSS, were not significantly altered. Therefore, we speculate that the supplementation of glycine might increase the content of GSH by sustaining the availability of the substrate in the kidneys.

Taken together, the present study showed that glycine treatment increases the expression and activity of renal Glo1 by promoting the nuclear translocation of Nrf2 and increasing GSH synthesis, thus ameliorating oxidative stress and renal injury in diabetic rats. The enhanced function of Glo1 might, to some extent, account for the suppressive effects of glycine on the renal AGE/RAGE signaling pathway observed in our previous study. However, the favorable effects in the present study are limited to our model of diabetic rats, and the optimal serum levels of glycine in humans have not reached total agreement. Although further investigation is required, the role of glycine in attenuating glycation and restoring anti-oxidant defense might provide new insights into the therapeutic options of diabetic nephropathy.

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DISCLOSURE

The authors declare no conflict of interest.

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