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1 Title:

2 **Upregulation of Nrf2/HO-1 signaling and farnesoid X receptor and attenuation of**  
3 **oxidative stress and inflammation mediate the protective effect of sitagliptin against**  
4 **diabetic nephropathy in rats**

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## **Abstract:**

Diabetic nephropathy (DN) is a kidney complication associated with diabetes that can lead to renal failure. The dipeptidyl peptidase IV inhibitor sitagliptin (SITA) has shown potential therapeutic benefits for DN. This study investigated the effect of SITA on DN, focusing on its modulation of the farnesoid X receptor (FXR) and nuclear factor erythroid 2-related factor 2 (Nrf2)/heme oxygenase 1 (HO-1) signaling and its suppressive efficacy on inflammation and oxidative stress. Thirty-two male rats were divided into four groups: control, SITA-treated, diabetic, and SITA-treated diabetic rats. SITA was administered orally for 8 weeks to diabetic rats induced with streptozotocin, after which samples were collected for analysis. The results indicate that SITA effectively reduced hyperglycemia, weight loss, and kidney injury and fibrosis. SITA also decreased oxidative stress, inflammatory markers, and apoptosis, as demonstrated by reductions in kidney malondialdehyde (MDA), myeloperoxidase, nitric oxide (NO), nuclear factor-kappaB (NF- $\kappa$ B), interleukin (IL)-1 $\beta$ , inducible NO synthase (iNOS), tumor necrosis factor (TNF)- $\alpha$ , Bcl-2-associated X protein (Bax), and caspase-3. These protective effects were associated with Kelch-like ECH-associated protein (Keap)-1 inhibition, increased levels of Nrf2 and FXR, and enhanced antioxidant activity as well as Bcl-2 upregulation. *In silico* analysis showed the binding of SITA with FXR, NF- $\kappa$ B p65, iNOS, Keap-1, caspase-3, and HO-1. In conclusion, SITA mitigates DN by reducing hyperglycemia, inflammation, and oxidative stress, while enhancing antioxidant defenses, FXR and Nrf2/HO-1 signaling.

**Keywords:** Hyperglycemia; Dipeptidyl peptidase IV inhibitor; Nephropathy; Inflammation; Oxidative stress.

## **1. Introduction**

Diabetic kidney disease (DKD), also known as diabetic nephropathy (DN), is a chronic kidney disorder that emerges as a complication in diabetic patients [1]. Key risk factors for DKD

include prolonged periods of poorly controlled blood glucose (BG) levels and hypertension. Persistent hyperglycemia plays a central role in the onset and advancement of DN, despite the multifactorial nature of the disease [2]. Hyperglycemia not only provokes structural alterations in the kidney but also induces mechanical stress and hemodynamic changes in the glomeruli by activating several transcription factors and signaling molecules [2]. Chronic hyperglycemia can impact the glomerular basement membrane by causing non-enzymatic protein glycation, leading to stiffening of the efferent arterioles [1]. This alteration elevates the glomerular filtration rate and promotes ultrafiltration, consequently increasing intraglomerular pressure and triggering progressive hypertrophy of the glomeruli, which culminates in glomerulosclerosis [1]. The chronic microvascular effects of hyperglycemia involve tubuloglomerular atrophy, mesangial matrix expansion, interstitial fibrosis, and thickening of the basement membrane and arterioles. Approximately 40% of diabetic patients may eventually develop DN, making it the leading cause of end-stage renal disease [3, 4]. DKD arises approximately 10 years post-diagnosis in type 1 diabetes mellitus (T1DM) but can occur at various stages in individuals with type 2 diabetes [1].

Oxidative stress (OS) and inflammation are implicated in the complications of diabetes, including nephropathy and cardiomyopathy [5, 6]. Hyperglycemia-driven excess production of reactive oxygen species (ROS) and the resulting OS are recognized as key contributors to the pathophysiological mechanisms underlying diabetes-related vascular complications, including DKD [2, 7]. Elevated ROS levels can detrimentally affect cellular components, and this oxidative burden leads to lipid peroxidation (LPO), protein oxidation, DNA damage, and nuclear factor kappaB (NF- $\kappa$ B) activation, alongside an increase in pro-inflammatory mediators, all of which contribute to cellular injury [8]. OS and inflammation provoke cell death via apoptosis and both processes are therefore fundamental drivers in DN pathogenesis. Evidence from experimental models of DN reveals elevated OS markers, decreased antioxidant

80 defenses, activation of NF- $\kappa$ B, and upregulated inflammatory and apoptotic factors [5, 9].  
81 Clinically, hyperglycemia in diabetic patients has been correlated with an inflammatory  
82 response marked by enhanced macrophage infiltration [10]. Therefore, agents with combined  
83 antioxidant and anti-inflammatory properties hold promise for mitigating DN progression and  
84 its associated pathologies. In this context, the activation of nuclear factor erythroid 2-related  
85 factor 2 (Nrf2) and farnesoid X receptor (FXR) has recently been shown to effectively attenuate  
86 OS and inflammation in the kidney of diabetic and chlorpyrifos (CPF)-intoxicated rats [9, 11].  
87 In another recent study, the dual activation of FXR and Nrf2 prevented cholestatic liver injury  
88 [12]. Activation of Nrf2 represents a key approach to reducing OS and inflammation associated  
89 with various metabolic disorders [13]. Nrf2 is a transcription factor (TF) that controls the  
90 expression of antioxidant genes, such as heme oxygenase-1 (HO-1) under conditions of  
91 elevated ROS [14]. Nrf2 is found sequestered by Kelch-like ECH-associated protein 1 (Keap-  
92 1) in the cytoplasm, a binding that is dissociated in response to increased ROS [14]. The  
93 protective role of Nrf2 signaling in mitigating redox imbalance and inflammation in diabetes-  
94 related complications is well-documented [5, 14, 15]. FXR is expressed abundantly in the  
95 kidneys and in several tissues, and provides protective effects against metabolic dysregulation,  
96 and redox imbalance [16]. Studies indicate that FXR deficiency is associated with insulin  
97 resistance, hyperglycemia, and other metabolic abnormalities and accelerates the progression  
98 of DN. In contrast, its activation improves metabolic alterations and attenuates kidney injury  
99 by reducing inflammation, OS, and fibrosis [17-19].

100 Sitagliptin (SITA) is a dipeptidyl peptidase IV inhibitor and an oral anti-hyperglycemic drug.  
101 SITA enhances insulin secretion in a glucose-dependent manner and prolongs the half-life time  
102 of glucagon-like peptide-1 and other incretin hormones [20]. Its safety and efficacy both in  
103 mono- and combination therapies in adult diabetic patients have been suggested and it has also  
104 been shown to lower glycated hemoglobin levels [21, 22]. Besides its antidiabetic efficacy,

SITA has shown beneficial effects against diabetes complications and other disorders. Very recently, Li et al [23] reported the potential of SITA to prevent Parkinson's disease and other neurodegenerative disorders. Owing to its antiapoptotic and antifibrotic properties, the beneficial effects of SITA against COVID-19 has been suggested [24]. We have previously reported the protective efficacy of SITA against diabetic cardiomyopathy and investigated the involvement of JAK/STAT signaling [6]. The protective efficacy of SITA against kidney injury induced by chemotherapy [25, 26], and its antifibrosis effects in diabetic rats [27, 28] have also been reported. The role of FXR and Nrf2/HO-1 signaling in mediating the efficacy of SITA on DKD hasn't been explored yet.

This study aimed to investigate whether the attenuation of oxidative stress, inflammation, and renal damage by SITA in a diabetic rat model is linked to the upregulation of Nrf2/HO-1 signaling and FXR. Utilizing a combined *in vivo* experimental approach and *in silico* molecular docking analysis, we evaluated the hypothesis that SITA confers protection against DN by modulating Nrf2/HO-1 signaling and FXR and attenuating oxidative and inflammatory responses. By defining the involvement of Nrf2/HO-1 signaling and FXR, this research may offer an insight into the mechanism of action of SITA, potentially strengthening the rationale for its use in mitigating DN progression beyond glycemic control.

## **2. Materials and methods**

### **2.1. Animals and experimental design**

Thirty-two 10 week-old male Sprague-Dawley rats were housed at a standard temperature and humidity on a 12-h light/dark cycle in polypropylene cages (42.5 cm × 26.6 cm × 18.5 cm) enriched with corn cob. The rats (3 per cage) were given access to food and water *ad libitum*. The study was approved by the ethics committee of King Saud University (IRB: SE-19-155). Streptozotocin (STZ; Sigma, USA; Cat. no.: S0130) (55 mg/kg body weight) dissolved in cold citrate buffer (0.1 M, pH 4.5) was injected intraperitoneally (i.p.) to overnight fasted rats to

130 induce T1DM, while citrate buffer was given to the control rats. A commercial kit (Spinreact,  
131 Spain; Cat. no. 1001190) was used to measure BG after 72 h and animals with fasting BG  
132 values of at least 250 mg/dl were selected. Normal rats were allocated into Group I (Control)  
133 and Group II (SITA) that received 0.9% saline and 10 mg/kg SITA (Sigma, USA; Cat. no.  
134 SML3205) dissolved in 0.9% saline, respectively. Diabetic rats were allocated into Group III  
135 (STZ) and Group IV (STZ + SITA) that received 0.9% saline and 10 mg/kg SITA, respectively.  
136 Each group included 6 rats and SITA and 0.9% saline were supplemented daily for 8 weeks  
137 via oral gavage. For 24-h urine collection, rats were temporarily housed in individual metabolic  
138 cages equipped with wire-mesh floors and urine collection funnels.  
139 Following treatment, rats were fasted overnight, and blood was collected via cardiac puncture  
140 under ketamine/xylazine anesthesia. After immediate dissection, the kidneys were removed  
141 and weighed. Samples from the kidney were collected on RNALater (ThermoFisher, USA; Cat.  
142 no. AM7020) while others on 10% neutral-buffered formalin (NBF). Other samples were  
143 processed via homogenization (10% w/v) in Tris-HCl buffer (10 mM; pH 7.4), centrifuged,  
144 and the supernatant was stored at -80°C.

## 145 2.2. Biochemical assays

146 Levels of creatinine and blood urea nitrogen (BUN) in serum and microalbumin in urine were  
147 assayed using Biodiagnostic (Egypt; Cat. no. CR 1250 and UR 2110) and Spinreact (Spain;  
148 Cat. no. 1107170) kits, respectively. Levels of reduced glutathione (GSH), malondialdehyde  
149 (MDA), and nitric oxide (NO) and activities of superoxide dismutase (SOD) and catalase were  
150 determined in the kidney homogenate supernatant using Biodiagnostic kits (Egypt; Cat. no.  
151 TA2511, MD2528, NO2533, SD2521, and CA2517, respectively). The activities of kidney  
152 HO-1 and myeloperoxidase (MPO) were measured according to Abraham et al. [29] and  
153 Krawisz et al. [30], respectively. NF- $\kappa$ B p65, TNF- $\alpha$  and IL-1 $\beta$  (ELabsience, China; Cat. no.  
154 E-EL-R0674, E-EL-R2856 and E-EL-R0012, respectively), and caspase-3 (Cusabio, China;

Cat. no. CSB-E08857r) were measured in the kidney homogenate supernatant. All assays were conducted strictly following the manufacturers' instructions.

### 2.3. Histopathology and immunohistochemistry (IHC):

Following fixation in 10% NBF for 24 h, the kidney samples were processed for embedding in paraffin. 5- $\mu$ m thick-sections were cut, stained with hematoxylin and eosin (H&E) and Masson's trichrome and examined using a light microscope (Olympus BX40, Olympus Corp., Japan). Other sections were processed for IHC staining to determine changes in Nrf2 and FXR. Briefly, the sections were dewaxed, rehydrated, and then treated with 0.05 M citrate buffer (pH 6.8) and 0.3% hydrogen peroxide (H<sub>2</sub>O<sub>2</sub>). After blocking, primary antibodies (Biospes, China; Cat. no. YPA1865 and YPA1581 for Nrf2 and FXR, respectively) were added overnight at 4°C, and the sections were washed and the secondary antibody (Biospes, China; Cat. no. BSA1031) was added. 3,3'-diaminobenzidine (Sigma, USA; Cat. no. D12384) in H<sub>2</sub>O<sub>2</sub> was used for color development and counterstaining was carried out using hematoxylin. ImageJ (NIH, USA) was used to measure intensity of the developed color (6/rat).

### 2.4. qRT-PCR

Changes in kidney Bax, Bcl-2, caspase-3, iNOS, Nrf2, FXR, NF- $\kappa$ B p65, and Keap-1 mRNA were assayed using qRT-PCR. RNA was isolated using Trizol (Invitrogen, ThermoFisher Scientific, Waltham, MA, USA, Cat. no. 15596026) and RNA samples with A260/A280 value  $\geq 1.8$  after purification were processed for cDNA synthesis via reverse transcription. cDNA was amplified using SYBR Green (ThermoFisher Scientific, USA; Cat. no. 4309155) and primer pairs in Suppl. Table I. The  $2^{-\Delta\Delta C_t}$  method [31] was employed for analysis using  $\beta$ -actin as a control.

### 2.5. *In silico* molecular docking

The affinity of SITA towards FXR (PDB ID: 7D42), NF- $\kappa$ B p65 (PDB ID: 5UO1), iNOS (PDB ID: 3EAI), caspase-3 (PDB ID: 1NME), HO-1 (PDB ID: 1DVE), and Keap-1 (PDB ID:



5CGJ) was investigated using PyRx virtual screening software (version 0.8) [32]. The target protein were prepared using Autodock Tools (ADT; v1.5.6) and PyMOL (v2.3.2) and LigPlot (v2.2.8) [33] were used for visualization of binding mode and protein-ligand interactions, respectively.

## 2.6. Statistical analysis

All the data were expressed as the mean  $\pm$  SEM. Statistics was performed using GraphPad Prism 8 software. The comparisons between the different groups were performed by one-way ANOVA, following with the post hoc Tukey's test. P values less than 0.05 were considered statistically significant.

## 3. Results

### 3.1. SITA ameliorates hyperglycemia and kidney damage in diabetic rats

Initial and final BG levels were markedly elevated in diabetic rats (Fig. 1A;  $P < 0.001$ ). The same animals exhibited significant decrease in body weight (BW) (Fig. 1B,C) and increased kidney weight (KW)/BW ratio (Fig. 1D) ( $P < 0.001$ ). Biochemical findings represented in Figures 2A-C revealed significant increase in creatinine, BUN, and microalbumin in diabetic animals ( $P < 0.001$ ). The findings in Fig. 3 showed degeneration of the epithelial lining of renal tubules, atrophied irregular renal corpuscle, and deposition of fibers in the interstitium in diabetic rats which also showed fiber deposition surrounding blood vessels. SITA effectively alleviated hyperglycemia, BW, KW/BW ratio (Fig. 1), kidney function markers (Fig. 2), and prevented kidney tissue damage (Fig. 3).

### 3.2. SITA mitigates kidney OS in diabetic rats

Diabetic rat kidney showed remarkable elevation in MDA (Fig. 4A) and MPO (Fig. 4B) along with declined GSH (Fig. 4C), SOD (Fig. 4D), and CAT (Fig. 4E) as compared to the control ( $P < 0.001$ ). SITA ameliorated MDA levels, MPO activity and antioxidants when supplemented to diabetic rats whereas had no effect on normal animals.

### 3.3. SITA attenuates kidney inflammation in diabetic rats

The effect of SITA on the inflammatory response in the kidney of diabetic rats was evaluated via assessment of changes in NF- $\kappa$ B p65 and pro-inflammatory mediators. Diabetes was associated with elevated NF- $\kappa$ B p65 levels (Fig. 5A,B) along with increased kidney TNF- $\alpha$  (Fig. 4C) and IL-1 $\beta$  (Fig. 4D). iNOS mRNA (Fig. 4E) and NO levels (Fig. 4F) were remarkably increased in the diabetic kidney when compared with the control group ( $P < 0.001$ ). Treatment with SITA effectively decreased NF- $\kappa$ B p65 and pro-inflammatory mediators (TNF- $\alpha$ , IL-1 $\beta$  and iNOS), and NO in diabetic rats ( $P < 0.001$ ).

*In silico* investigations revealed the affinity of SITA to bind NF- $\kappa$ B p65 and iNOS with binding energies -7.5 and -8.2 kcal/mol, respectively (Fig. 6 and Table 1). The complexes of SITA with NF- $\kappa$ B p65 and iNOS showed polar bonding with two amino acid residues of both proteins and hydrophobic interactions with five and three residues, respectively.

### 3.4. SITA prevents kidney apoptosis in diabetic rats

Kidney Bcl-2 mRNA was decreased in diabetic rats whereas Bax and caspase-3 were upregulated significantly as compared to control rats (Fig. 7A-D). SITA upregulated Bcl-2 and decreased Bax and caspase-3 in the diabetic rat kidney ( $P < 0.001$ ). Investigation of the binding of SITA with caspase-3 revealed 8 hydrophobic interactions and -5.3 kcal/mol binding energy (Fig. 7E and Table 1).

### 3.5. SITA upregulates Nrf2/HO-1 signaling and FXR in diabetic rats

Keap-1 (Fig. 8A) and Nrf2 (Fig. 8B) mRNA levels were significantly increased and decreased, respectively in the kidney of diabetic rats as compared to control rats ( $P < 0.001$ ). IHC staining of Nrf2 (Fig. 8C-D) and HO-1 activity (Fig. 8E) determination showed significant downregulation in diabetic rats as compared to control rats. SITA downregulated Keap-1 and boosted Nrf2 and HO-1 in the kidney of diabetic rats while had no effect on normal rats. The effect of SITA on Keap-1 and HO-1 was further investigated using *in silico* molecular docking

(Fig. 9 and Table 1). SITA exhibited -10.2 and -9.4 kcal/mol binding energies with Keap-1 and HO-1, respectively. SITA (Fig. 9A) and the Nrf2 activator RA839 (Fig. 9B, Suppl. Table II) showed 7 and 17 hydrophobic interactions and 5 and 3 polar bonding, respectively, with Keap-1. The complex of SITA with HO-1 showed polar bonding and hydrophobic interactions with one and ten residues, respectively (Fig. 9C and Table 1).

Data represented in Fig 10 A-C showed significant downregulation of FXR mRNA and protein in the kidney of diabetic rats as compared to control rats ( $P < 0.001$ ). SITA upregulated FXR remarkably in diabetic rats whereas showed no effect on normal rats. Twelve and two hydrophobic and polar interactions, respectively, between SITA and FXR were shown *in silico* and the binding energy is -9.2 kcal/mol (Fig. 10D and Table 1). Twelve of the residues in SITA/FXR complex were noticed in the complex formed by the FXR activator tropifexor (Suppl. Fig. I, Suppl. Table II).

#### 4. Discussion

Nephropathy is one of the serious complications of diabetes characterized by functional and structural changes in the kidney [1]. Despite the progression in understanding the pathophysiology of DN, the condition is still undertreated due to the lack of effective sustainable treatments. The implication of OS and inflammation in DKD has been acknowledged and agents that attenuate these processes showed beneficial effects against the disease progression [9]. This study revealed that SITA protects the kidney against diabetes-induced injury by ameliorating hyperglycemia, OS, inflammation, and apoptosis, an effect that is associated with upregulation of FXR and Nrf2/HO-1 signaling.

Diabetic rats in this study exhibited hyperglycemia and abnormal renal function markers, including creatinine, BUN, and microalbumin. Hyperglycemia is a result of  $\beta$ -cells destruction and impaired insulin secretion because STZ is particularly toxic to these cells and promotes oxidative DNA damage [34]. Hyperglycemia was associated with a decrease in BW and

255 increase in KW/BW ratio. These findings align with previous studies showing BW loss and  
256 increased KW/BW ratio along with hyperglycemia and altered levels of kidney function  
257 markers in diabetic rodents [5, 9, 35]. Additionally, examination of the stained tissue revealed  
258 glomerular atrophy, irregular corpuscles, tubular epithelium degeneration and interstitial  
259 fibrosis as previously reported [9]. The progression of DN involves distinctive changes such  
260 as interstitial fibrosis, atrophic changes and renal dysfunction [3, 36], all have been observed  
261 in this study. BW loss observed in the diabetic rats is directly ascribed to the surplus utilization  
262 of lipids and proteins to provide energy in the lack of ability to use glucose [37]. The structural  
263 abnormalities, such as collagen deposition, could explain the changes in KW which could also  
264 be attributed to the upregulation of fibronectin, collagen, and growth factors [38]. SITA  
265 effectively ameliorated hyperglycemia, kidney hypertrophy and injury along with amelioration  
266 of creatinine, BUN and microalbumin. These findings highlight the nephroprotective efficacy  
267 of SITA which has been previously demonstrated in animals challenged with chemotherapy  
268 [25, 26]. In uninephrectomized rats challenged with doxorubicin, SITA administration for 6  
269 weeks ameliorated tubulointerstitial injury and fibrosis [25], and its administration for 14 days  
270 prevented cyclosporine-induced kidney injury and ameliorated serum urea and creatinine [26].  
271 In addition to its antidiabetic efficacy, the nephroprotective mechanism of SITA could be linked  
272 to its role in attenuating OS and inflammation as reported in animal models of chemotherapy-  
273 induced nephrotoxicity [25, 26]. In this study, markers of OS were evident through increased  
274 MDA, NO, and MPO activity, accompanied by reduced GSH and antioxidant enzyme  
275 activities. Inflammation in diabetic kidneys was characterized by elevated levels of NF- $\kappa$ B p65,  
276 iNOS, TNF- $\alpha$ , and IL-1 $\beta$ . The pathogenesis of DN is heavily influenced by OS and  
277 inflammatory mechanisms [2, 7, 39]. Persistent hyperglycemia leads to excessive production  
278 of ROS, driving LPO and NF- $\kappa$ B activation, which collectively contribute to cellular injury  
279 [8]. Upon activation by ROS, NF- $\kappa$ B leads to increased secretion of IL-1 $\beta$  and TNF- $\alpha$ , and

promotes iNOS, thereby explaining the rise in NO production. In our recent study [9], we have provided further evidence of inflammation in DN manifested by upregulation of CD68 which is an indicator of macrophage infiltration and has been demonstrated in renal autopsy samples from diabetic patients [40]. This interaction between OS and inflammation fosters apoptotic pathways. For instance, superoxide, a prominent ROS, can combine with NO to form the highly reactive oxidant peroxynitrite that damages DNA and induces cell death [41]. In this study, diabetic rats showed declined Bcl-2 and increased Bax and caspase-3 expression in the kidney. Inflammatory mediators and ROS stimulate pro-apoptotic Bax expression and disrupt mitochondrial membrane potential [42] and together with the impairment of mitochondrial respiratory function, Bax results in DNA damage and increases mitochondrial membrane permeability and cytochrome c release into the cytosol and subsequent activation of caspase-3 [43]. Activated caspase-3 promotes DNA fragmentation and cytoskeleton breakdown, culminating in apoptosis.

SITA demonstrated significant protective effects against OS, inflammation, and kidney injury. These protective actions were reflected by reduced levels of markers associated with oxidative and inflammatory stresses, including MDA, NO, MPO, NF- $\kappa$ B p65, pro-inflammatory mediators, Bax, and caspase-3, alongside an increase in antioxidant levels and Bcl-2 expression. Previous pre-clinical studies have pointed to the efficacy of SITA in mitigating OS and inflammation in the kidney and other tissues [6, 25, 26]. SITA afforded a protective role against myocardial oxidative damage in diabetic animals [6]. SITA mitigated LPO, IL-6 and IL-1 $\beta$ , suppressed JAK/STAT signaling and enhanced antioxidants in the heart of diabetic rats [6]. In a rat model of cyclosporine nephrotoxicity, SITA ameliorated MDA, GSH, SOD, CAT, Bax and TNF- $\alpha$  [26]. In a doxorubicin nephropathy model, SITA mitigated inflammation and ROS generation as shown by decreased IL-1 $\beta$  and the mRNA levels of NADPH oxidase subunits [25]. In a rat model of high-fat diet-induced T2DM, SITA ameliorated renal function

305 and prevented TGF- $\beta$ 1/Smad-mediated kidney fibrosis [27]. Similarly, SITA has been shown  
306 to attenuate inflammation and fibrosis by suppressing TGF- $\beta$ , collagen deposition, and TNF- $\alpha$   
307 in diabetic rats [28]. *In silico* findings added further support to the anti-inflammatory and anti-  
308 apoptosis efficacy of SITA. The results pinpointed the affinity of SITA to bind NF- $\kappa$ B p65  
309 which can lead to suppression of its transcriptional activity. Moreover, SITA exhibited affinity  
310 towards numerous amino acid residues in iNOS and caspase-3.

311 To investigate the nephroprotective mechanism of SITA, its influence on the Nrf2 and FXR  
312 was evaluated in the kidney of rats and an *in silico* investigation was also conducted. The  
313 diabetic rat kidneys exhibited upregulation of Keap-1 and a concurrent decrease in Nrf2. Nrf2,  
314 HO-1, and FXR, indicating suppression of the Nrf2/HO-1 pathway and FXR expression. The  
315 suppression of Nrf2 was observed in rats with DN as previously demonstrated [5, 9, 44]. These  
316 findings are also consistent with previous research demonstrating that hyperglycemia and other  
317 metabolic disturbances in diabetes and obesity correlate with reduced FXR levels, which in  
318 turn contribute to the progression of DN [9, 18, 19, 45, 46].

319 SITA downregulated Keap-1 and enhanced the expression of Nrf2, HO-1, and FXR, results  
320 directly associated with the alleviation of metabolic disturbances, OS and inflammation. Nrf2  
321 induces the transcription of antioxidant and cytoprotective enzymes, thereby reducing ROS  
322 and mitigating oxidative damage, and directly inhibits NF- $\kappa$ B [14, 47]. In this study, the  
323 increased expression of FXR was linked to improved BG levels and mitigation of inflammatory  
324 and oxidative stresses. FXR modulates BG regulation by influencing insulin release and  
325 sensitivity and gluconeogenesis [48-50]. Conversely, FXR deficiency leads to hyperglycemia,  
326 which is associated with decreased insulin release [49], with research on  $\beta$ -cells indicating that  
327 FXR modulates insulin secretion [51]. Activation of FXR has been shown to protect against  
328 mitochondrial damage and excess ROS associated with ischemia/reperfusion in murine kidney  
329 [52]. Nrf2 mediates the beneficial role of FXR against OS, with the silencing of Nrf2 nullifying

the beneficial consequences of FXR activation [52]. FXR activation also confers protection to the kidney in diabetes by reducing oxidative damage, fibrosis, and glomerulosclerosis [17]. Recently, Nrf2 and FXR have been shown to jointly mitigate liver injury caused by cholestasis [12]. Therefore, the combined action of Nrf2 and FXR likely contributes, at least in part, to the nephroprotective effects of SITA.

A key finding of this study is the demonstration that the protective role of SITA against DN involves the concerted upregulation of both Nrf2/HO-1 signaling and FXR. The integrated approach, combining *in vivo* experimental validation with *in silico* molecular docking to predict the interactions of SITA with key targets, including NF- $\kappa$ B p65, iNOS, caspase-3, Keap-1, HO-1, and FXR, provides insights into its mechanism of renoprotection. However, certain limitations should be acknowledged. The study utilized a single dose and duration of SITA treatment; exploring dose-response relationships and longer-term effects would provide further insights. While the STZ-induced diabetic rat model is well-established for studying DN pathogenesis and drug effects, the obtained findings should be investigated in clinical settings.

## 5. Conclusion

This study provides new insights into the protective mechanisms of SITA against DKD. SITA alleviated hyperglycemia and kidney damage, fibrosis, OS, inflammation, and cell death. These protective effects were linked to the activation of FXR and Nrf2/HO-1 signaling, which led to increased antioxidant defenses and a reduced inflammatory response. Molecular docking studies suggested that SITA interacts with FXR, HO-1, Keap1, NF- $\kappa$ B, iNOS, and caspase-3, further supporting its therapeutic potential. In summary, SITA shows promise in mitigating DN by activating Nrf2/HO-1 signaling, enhancing FXR, and reducing OS, inflammation, and apoptosis. Future studies are recommended to further investigate other underlying mechanisms.

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Availability of data and materials

The manuscript and supplementary material contain all data supporting the reported results.

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520 **Tables:**

521 Table 1. Binding affinities and interaction of sitagliptin with different protein targets.

| Target             | Lowest binding energy (kcal/mol) | Polar bonds                            | Hydrophobic interactions                                                                                              |
|--------------------|----------------------------------|----------------------------------------|-----------------------------------------------------------------------------------------------------------------------|
| NF- $\kappa$ B p65 | -7.5                             | His58, Ser45                           | Lys56, Pro47, Gln220, Glu222, Thr52                                                                                   |
| iNOS               | -8.2                             | Tyr293, Gln265                         | Pro291, Ile285, Gln282                                                                                                |
| Caspase-3          | -5.3                             |                                        | Arg164, Glu124, Pro201, Tyr1975, Gly125, Cys264, Tyr195, Val266                                                       |
| Keap-1             | -10.2                            | Leu365, Val465, Cys513, Val418, Arg415 | Ala556, Gly462, Leu557, Gly 509, Val604, Gly364, Gly603                                                               |
| HO-1               | -9.4                             | Gly139                                 | Phe214, Gly143, Ser142, Ala28, Asn210, Leu138, His25, Arg136, Phe207, Tyr134, Ile352, His447, Phe329, Phe443, Arg331, |
| FXR                | -9.2                             | Tyr369, Ser332                         | Met265, Ile335, Gly343, Met328, Val325, Met290, Leu348                                                                |

522 NF- $\kappa$ B p65: nuclear factor-kappaB p65; iNOS: inducible nitric oxide synthase; Keap-1: Kelch-  
523 like ECH-associated protein 1; HO-1: heme oxygenase 1.

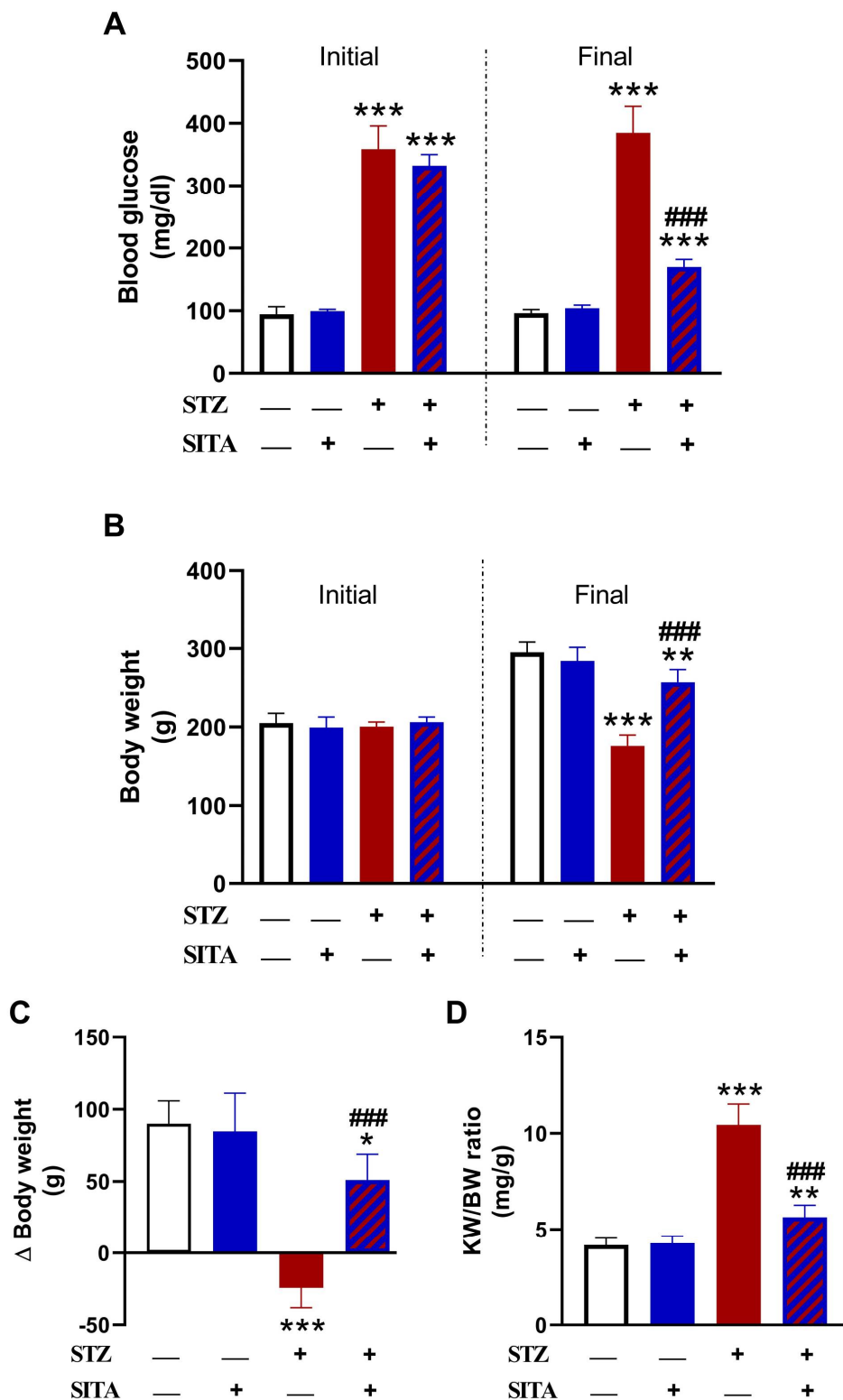


Fig. 1. Sitagliptin ameliorated blood glucose (BG) (A), body weight (BW) (B-C) and kidney weight (KW)/BW ratio (D) in diabetic rats. Data are mean  $\pm$  SEM, (n=8). \*P<0.05, \*\*P<0.01 and \*\*\*P<0.001 versus Control, and ###P<0.001 versus Diabetic.

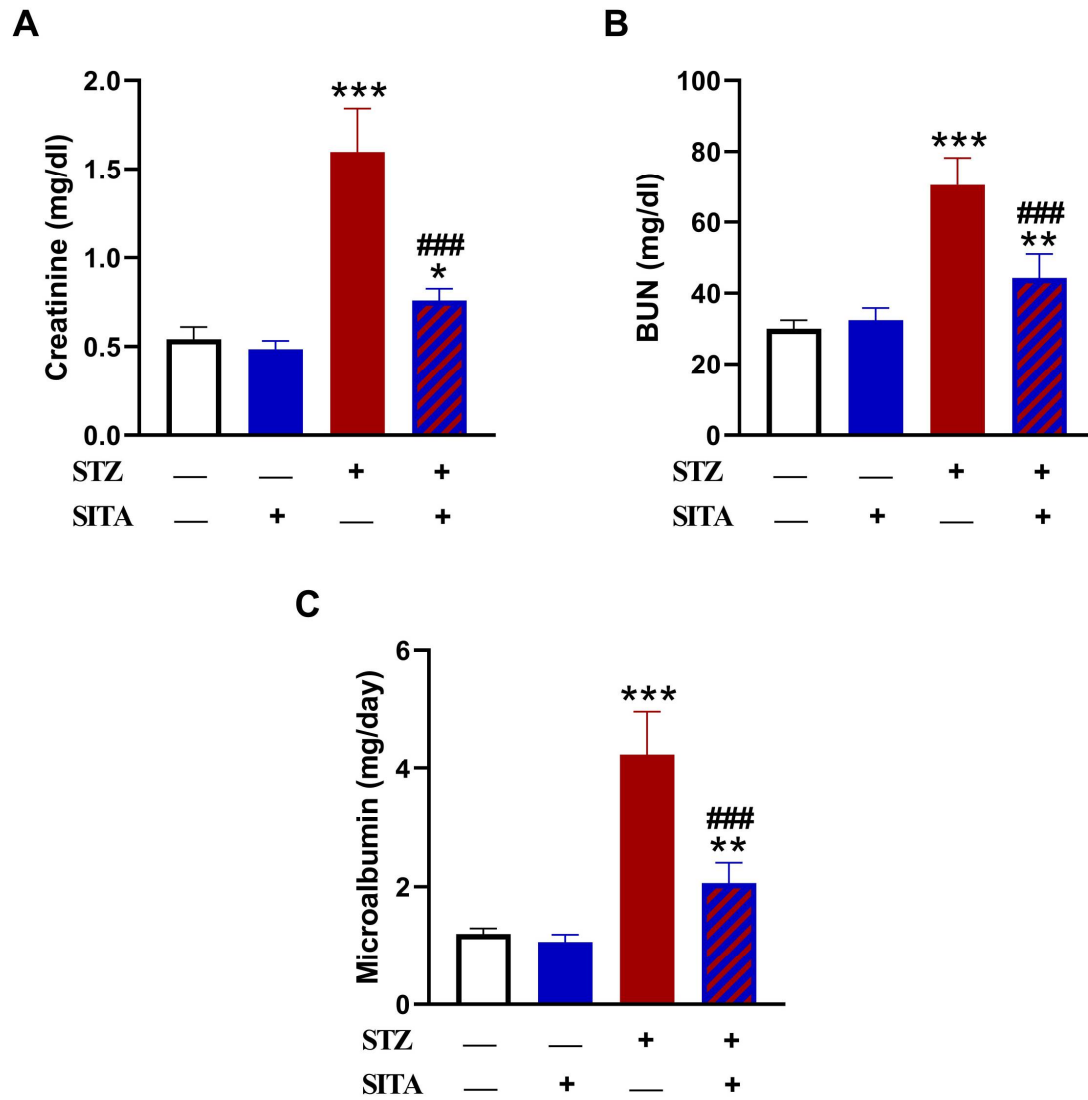


Fig. 2. Sitagliptin alleviated creatinine (A), blood urea nitrogen (BUN) (B) and microalbumin (C) in diabetic rats. Data are mean  $\pm$  SEM, ( $n=8$ ). \*  $P<0.05$ , \*\*  $P<0.01$  and \*\*\*  $P<0.001$  versus Control, and ###  $P<0.001$  versus Diabetic.



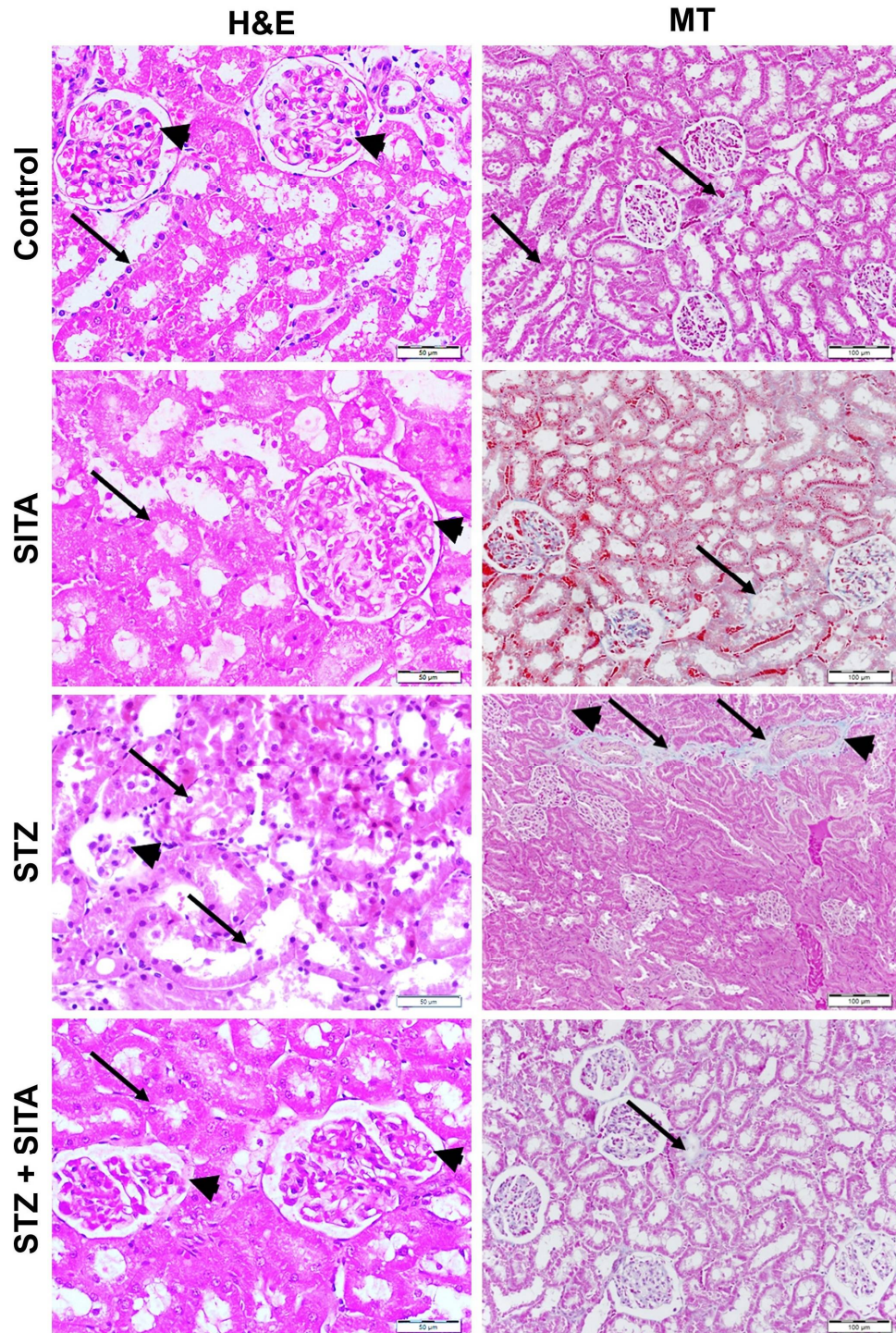


Fig. 3. Sitagliptin attenuated kidney injury in diabetic rats. H&E-stained sections from the control and SITA-supplemented rats showing normal renal corpuscle and glomeruli (arrowhead) and normal different types of tubules (arrow), diabetic rats showing irregular atrophied renal corpuscle (arrowhead), and degenerated tubular epithelium (arrows), and diabetic rats treated with SITA showing remarkable amelioration of the renal corpuscles degeneration (arrowhead) and normal tubules (arrow). (Scale bar = 50 µm). Masson's trichrome-stained sections from the control and SITA-supplemented animals showing normal distribution and amount of interstitial fibers (arrow), diabetic rats showing marked increase in fibrous tissue in the interstitium (arrow), and around blood vessels (arrowhead), and diabetic rats treated with SITA showing marked decrease in fibrotic deposition. (Scale bar = 100 µm).

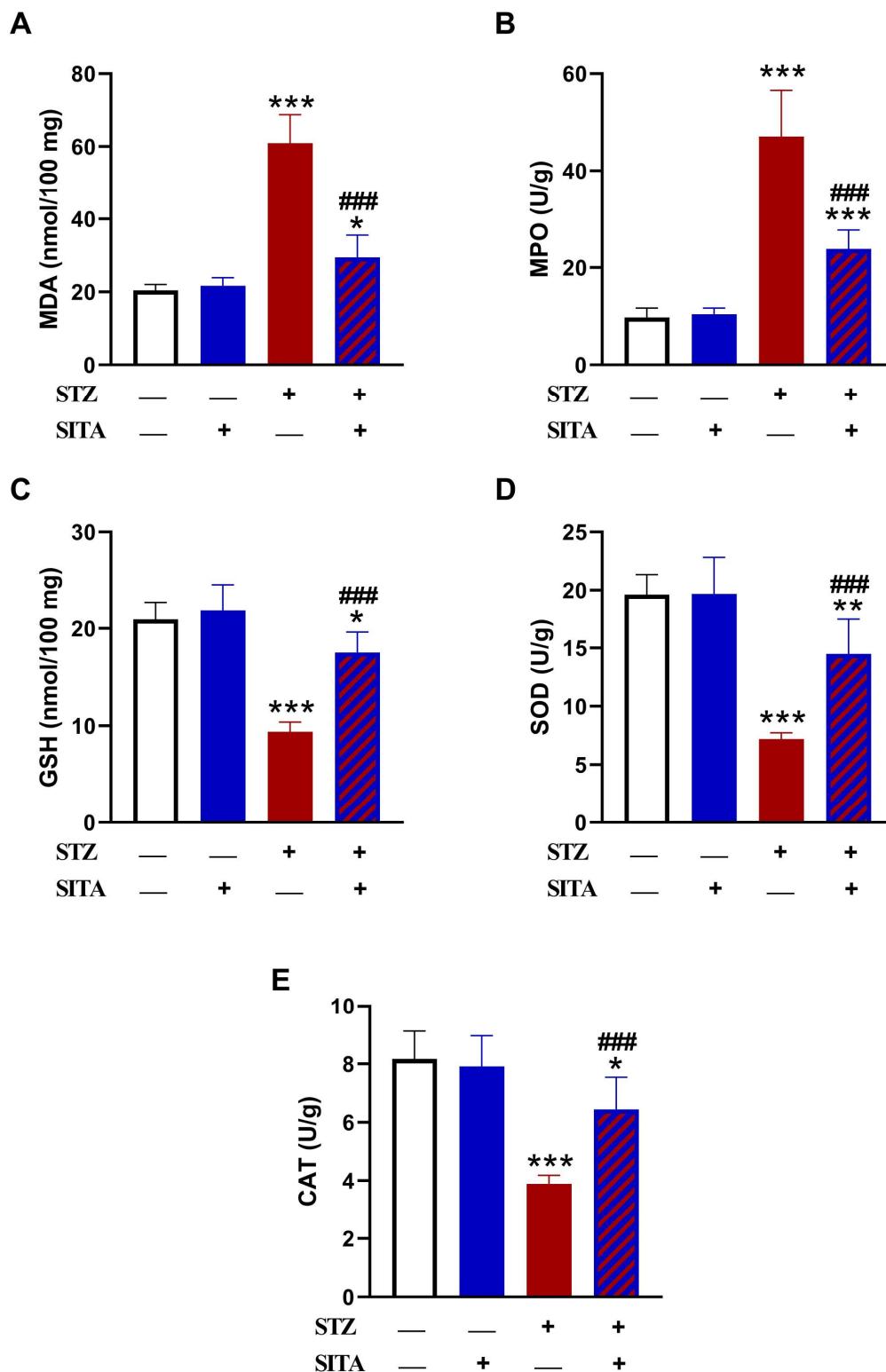


Fig. 4. Sitagliptin prevented oxidative stress in diabetic rats. SITA ameliorated renal malondialdehyde (MDA) (A), and myeloperoxidase (MPO) (B), and increased reduced glutathione (GSH) (C), superoxide dismutase (SOD) (D) and catalase (E) in diabetic rats. Data are mean  $\pm$  SEM, ( $n=8$ ). \* $P<0.05$ , \*\* $P<0.01$  and \*\*\* $P<0.001$  versus Control, and ### $P<0.001$  versus Diabetic.



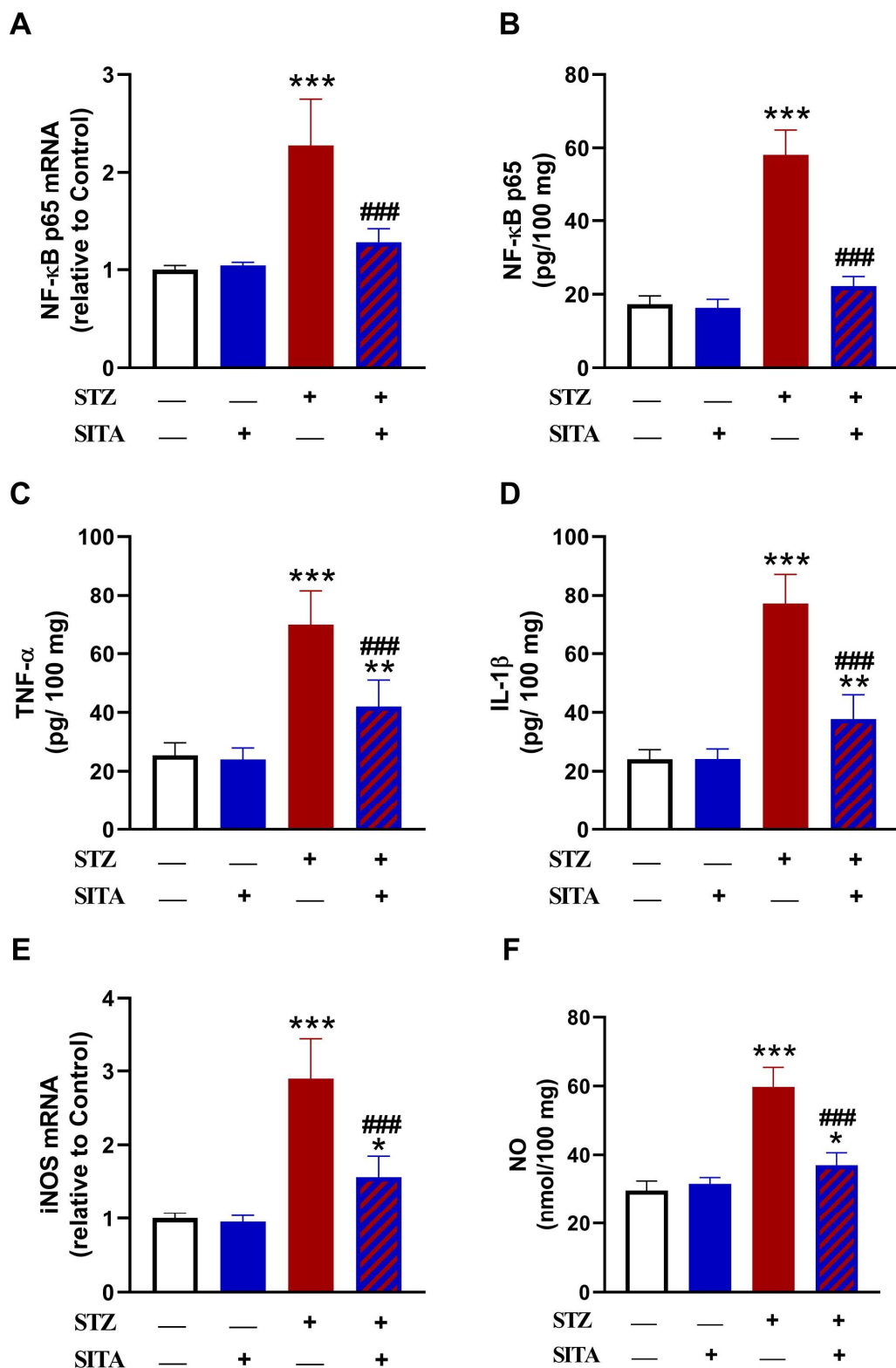


Fig. 5. Sitagliptin downregulated kidney nuclear factor-kappaB (NF- $\kappa$ B) p65 mRNA (A) and protein (B), tumor necrosis factor (TNF)- $\alpha$  (C) and interleukin (IL)-1 $\beta$  (D), inducible nitric oxide synthase (iNOS) mRNA (E), and nitric oxide (NO) levels (F) in diabetic rats. Data are mean  $\pm$  SEM, (n=8). \* P<0.05, \*\* P<0.01 and \*\*\* P<0.001 versus Control, and ### P<0.001 versus Diabetic.

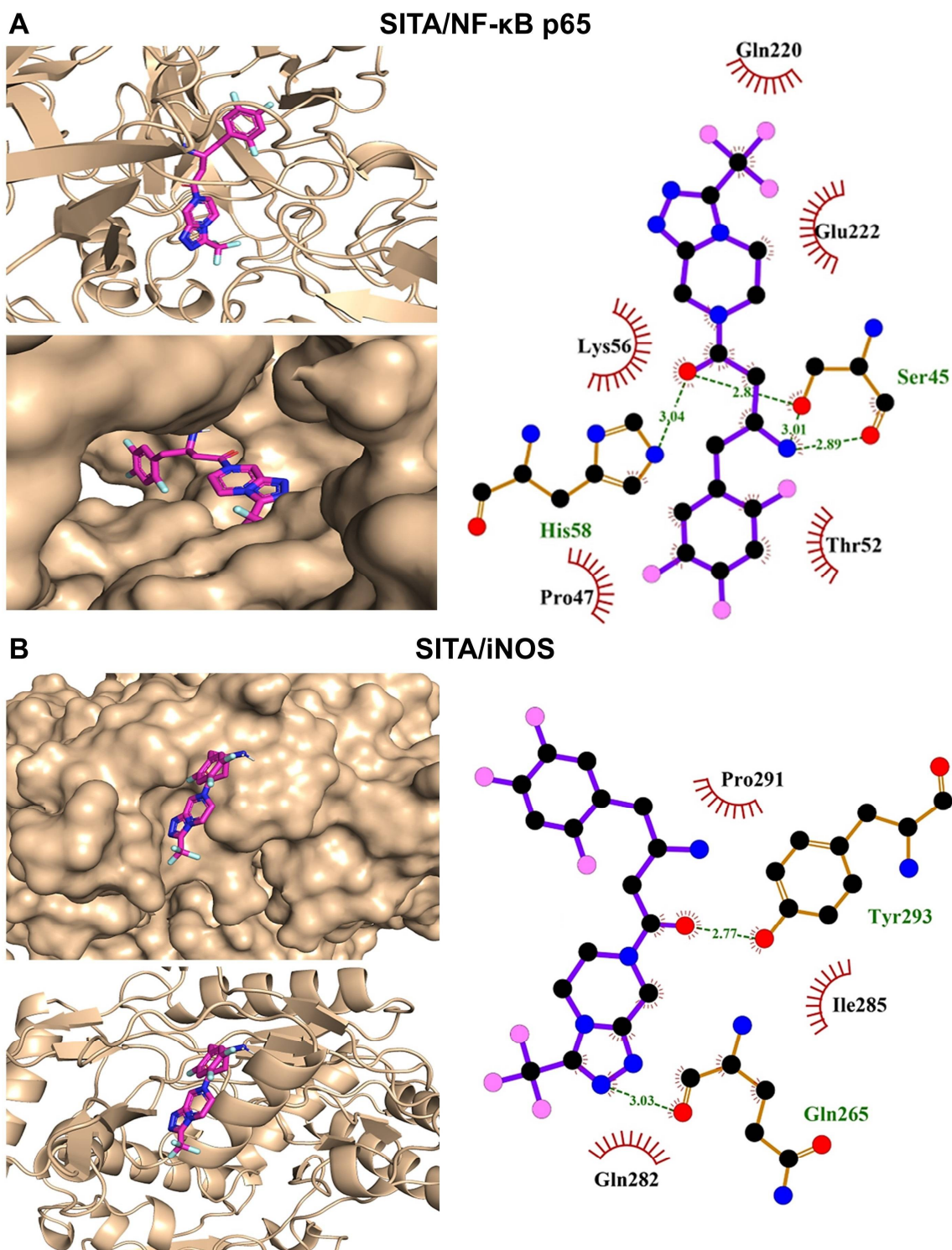


Fig. 6. Molecular docking shows the interaction between sitagliptin and nuclear factor-kappaB (NF- $\kappa$ B) p65 (A) and inducible nitric oxide synthase (iNOS) (B).

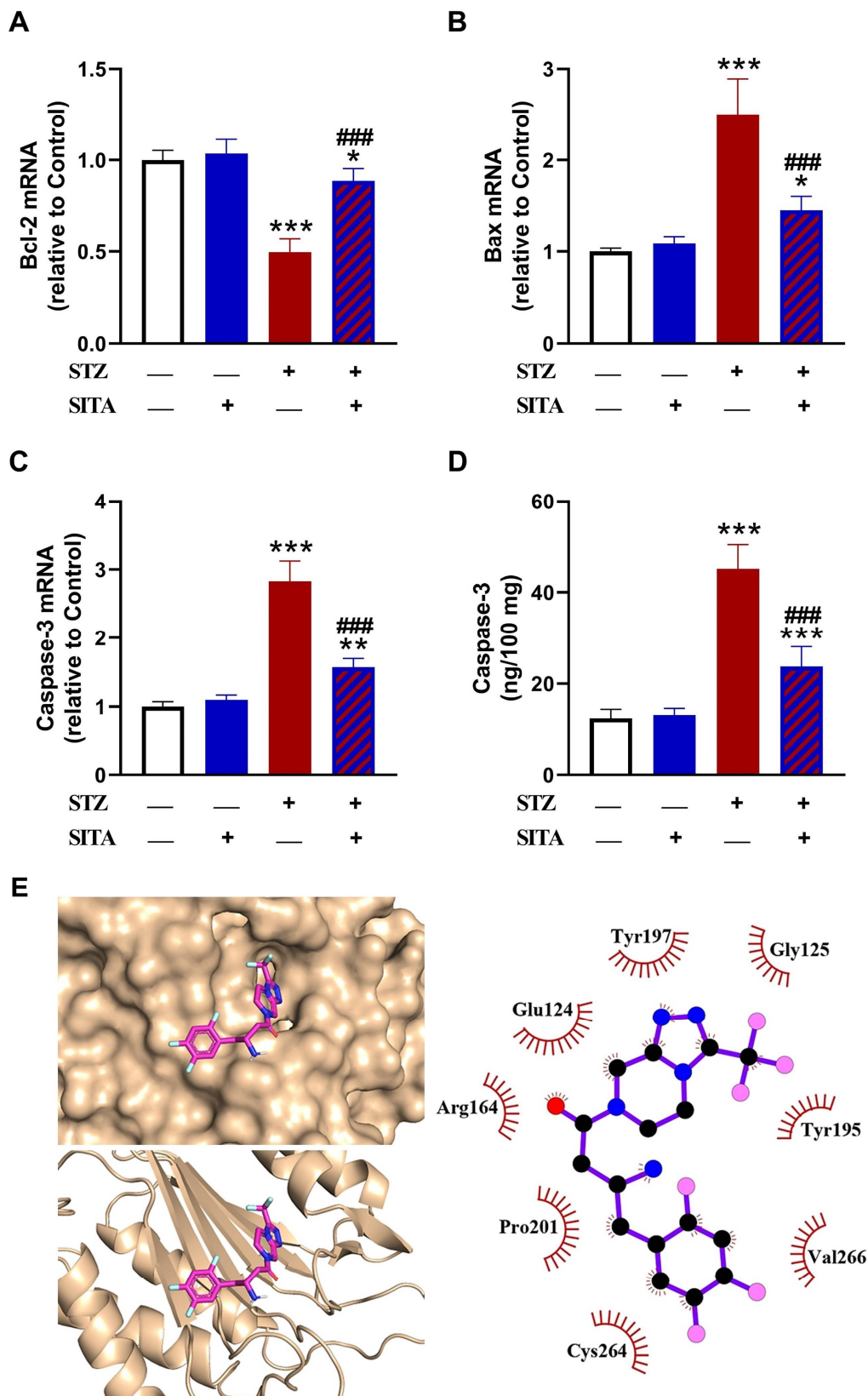


Fig. 7. Sitagliptin upregulated kidney B-cell lymphoma 2 (Bcl-2) (A) and suppressed Bcl-2-associated X protein (Bax) (B) and caspase-3 (C) mRNA, and caspase-3 protein levels (D) in diabetic rats. Data are mean  $\pm$  SEM, (n=8). \*P<0.05, \*\*P<0.01 and \*\*\*P<0.001 versus Control, and \*\*\*\*P<0.001 versus Diabetic. (E) Molecular docking shows the interaction between sitagliptin and caspase-3.

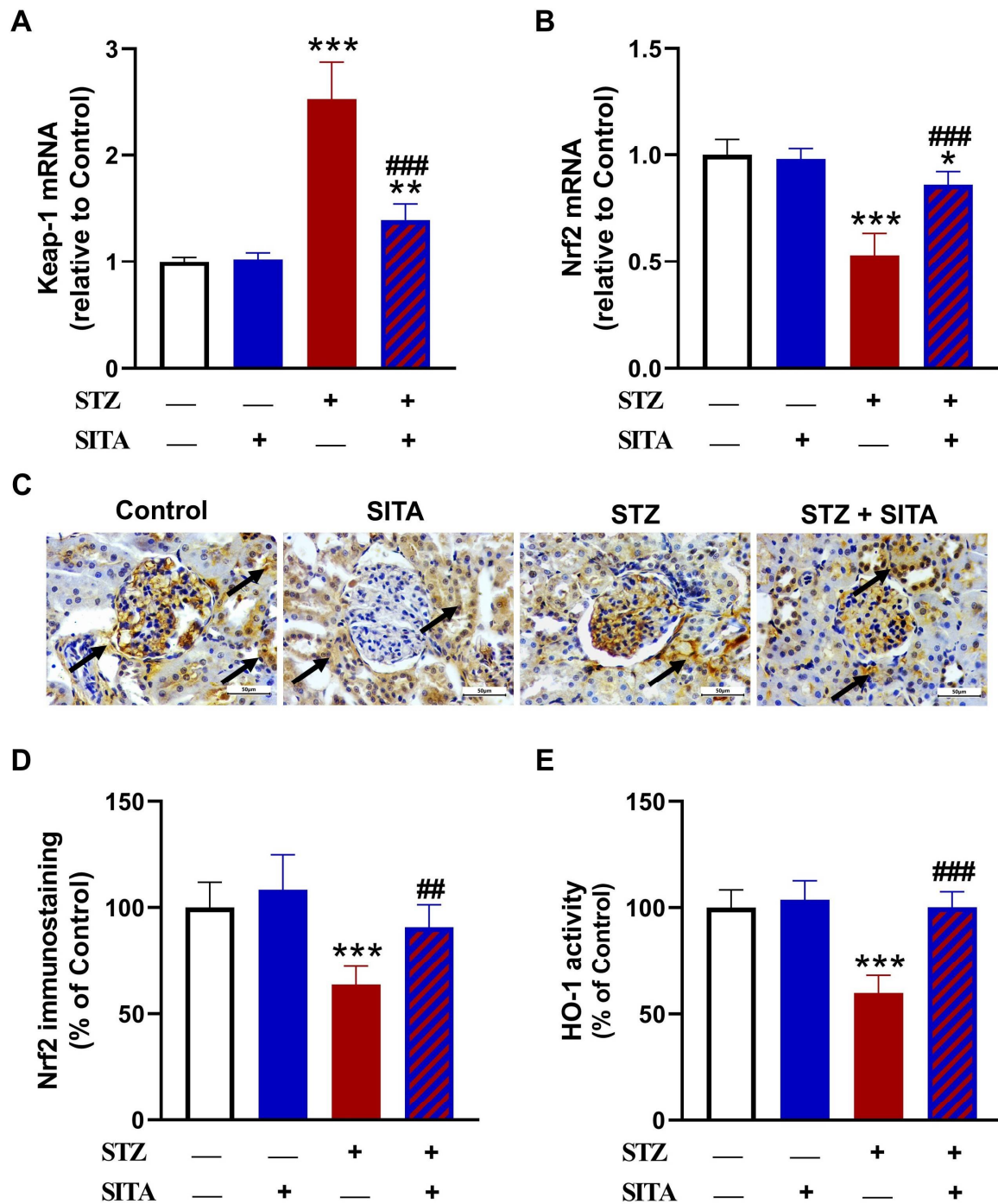


Fig. 8. Sitagliptin enhanced nuclear factor erythroid 2-related factor 2 (Nrf2)/heme oxygenase 1 (HO-1) signaling in diabetic rat. Sitagliptin decreased kidney Keap-1 mRNA (A) and upregulated Nrf2 mRNA (B), Nrf2 protein (C-D) and HO-1 activity (E). Data are mean  $\pm$  SEM, ( $n=8$ ). \* $P<0.05$ , \*\* $P<0.01$  and \*\*\* $P<0.001$  versus Control. ## $P<0.01$  and ### $P<0.001$  versus Diabetic.



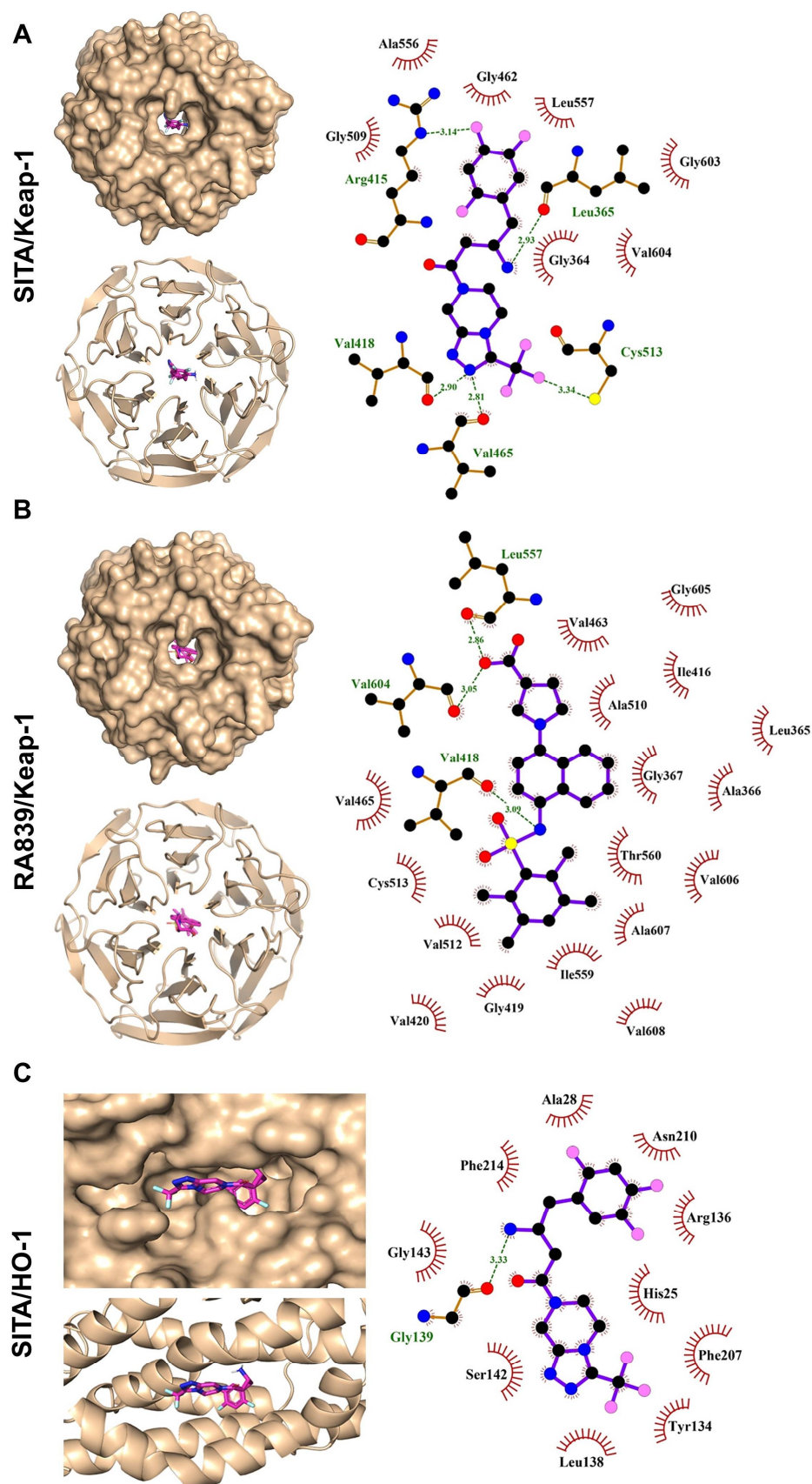


Fig. 9. Molecular docking shows the binding modes of sitagliptin (A) and RA839 (B) with Kelch-like ECH-associated protein 1 (Keap-1) and sitagliptin with heme oxygenase 1 (HO-1) (C).

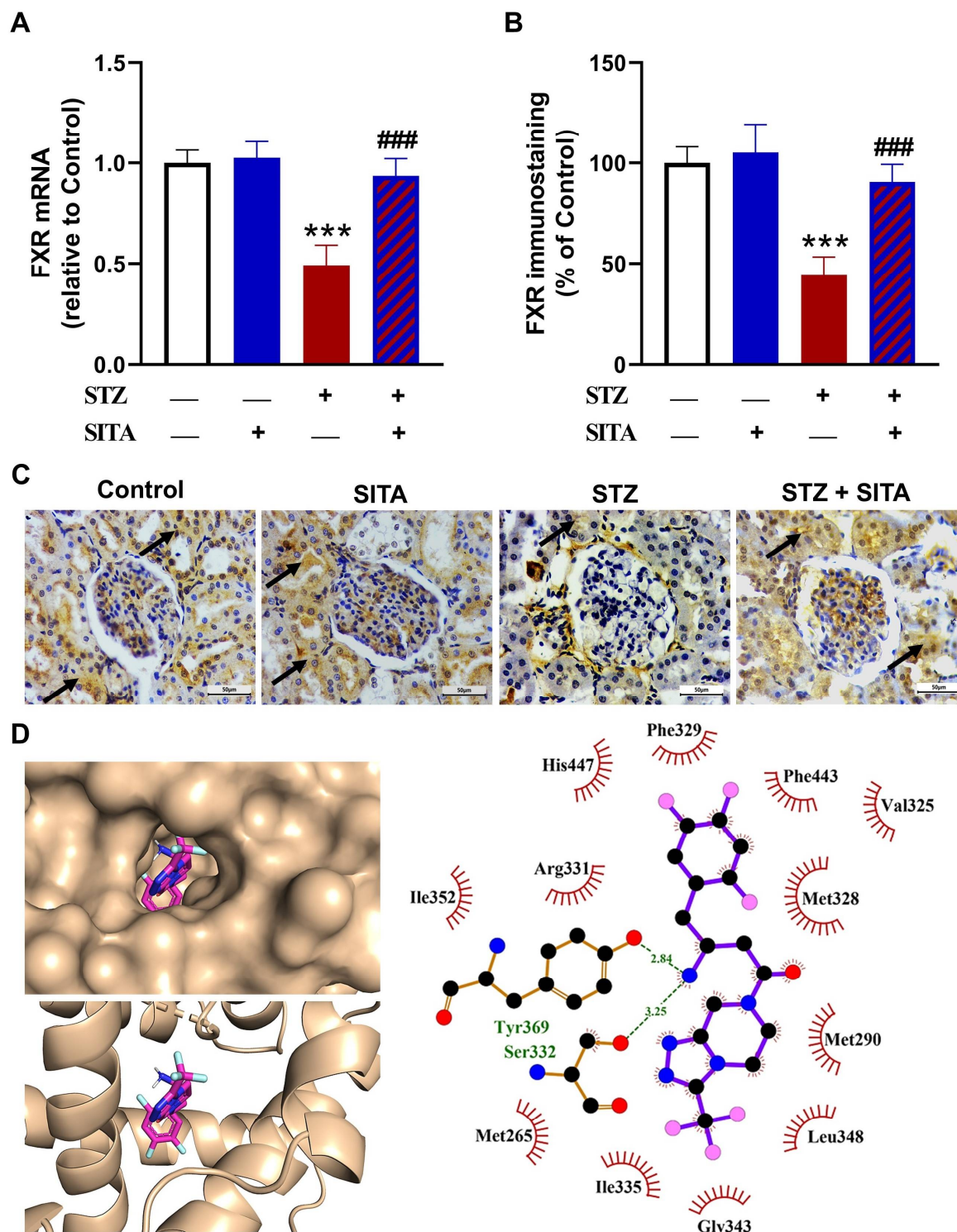


Fig. 10. Sitagliptin upregulated kidney farnesoid X receptor (FXR) in diabetic rat. Sitagliptin increased FXR mRNA (A) and protein (B) in the diabetic kidney. Data are mean  $\pm$  SEM, ( $n=8$ ). \*\*\* $P<0.001$  versus Control, and ### $P<0.001$  versus Diabetic. (D) Molecular docking showing the binding of sitagliptin with FXR.