

RESEARCH PAPER

Effects of xanthine oxidase inhibition with febuxostat on the development of nephropathy in experimental type 2 diabetes

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BACKGROUND AND PURPOSE

Elevated serum uric acid (UA) is a risk factor for the development of kidney disease. Inhibitors of xanthine oxidase (XO_i), an enzyme involved in UA synthesis, have protective effects at early stages of experimental diabetic nephropathy (DN). However, long-term effects of XO_i in models of DN remain to be determined.

EXPERIMENTAL APPROACH

The development of albuminuria, renal structure and molecular markers of DN were studied in type 2 diabetic Zucker obese (ZO) rats treated for 18 weeks with the XO_i febuxostat and compared with vehicle-treated ZO rats, ZO rats treated with enalapril or a combination of both agents, and lean Zucker rats without metabolic defects.

RESULTS

Febuxostat normalized serum UA and attenuated the development of albuminuria, renal structural changes, with no significant effects on BP, metabolic control or systemic markers of oxidative stress (OS). Most of these actions were comparable with those of enalapril. Combination treatment induced marked decreases in BP and was more effective in ameliorating structural changes, expression of profibrotic genes and systemic OS than either monotherapy. Febuxostat attenuated renal protein expression of TGF- β , CTGF, collagen 4, mesenchymal markers (FSP1 and vimentin) and a tissue marker of OS nitrotyrosine. Moreover, febuxostat attenuated TGF- β - and S100B-induced increased expression of fibrogenic molecules in renal tubular cells *in vitro* in UA-free media in an Akt kinase-dependent manner.

CONCLUSIONS AND IMPLICATIONS

Febuxostat is protective and enhances the actions of enalapril in experimental DN. Multiple mechanisms might be involved, such as a reduction of UA, renal OS and inhibition of profibrotic signalling.

Abbreviations

8-epi PGF_{2 α} , 8-isoprostane; ACEI, ACE inhibitor; α -SMA, α -smooth muscle actin; BG, blood glucose; CKD, chronic kidney disease; Coll4, collagen IV; CTGF, connective tissue growth factor; DM2, type 2 diabetes; ECM, extracellular matrix; ENL, enalapril; FBX, febuxostat; FSP1/S100A4, fibroblast-specific protein 1; GFR, glomerular filtration rate; GSS, glomerulosclerosis score; HBA1c, glycosylated haemoglobin; IHC, immunohistochemistry; MS, metabolic syndrome; NT, nitrotyrosine; PAI-I, plasminogen activator inhibitor-I; RAS, renin–angiotensin system; RAGE, receptor for advanced glycosylation end products; SAG, severely affected glomeruli; TBARS, thiobarbituric acid reactive substances; TIF, tubulointerstitial fibrosis; TIFS, tubulointerstitial fibrosis score; UA, uric acid; U_{alb}V, 24 h urinary albumin excretion; XO, xanthine oxidase; XO_i, xanthine oxidase inhibition/inhibitor; ZL, Zucker lean; ZO, Zucker obese

Tables of Links

TARGETS	
Other protein targets^a	Enzymes^b
RAGE	Akt (PKB)
	Angiotensin-converting enzyme
	Xanthine dehydrogenase/oxidase

LIGANDS	
CTGF	TGF-β1
Enalapril	Uric acid
Febuxostat	Xanthine
Fibronectin	

These Tables list key protein targets and ligands in this article which are hyperlinked to corresponding entries in <http://www.guidetopharmacology.org>, the common portal for data from the IUPHAR/BPS Guide to PHARMACOLOGY (Southan *et al.*, 2016) and are permanently archived in the Concise Guide to PHARMACOLOGY 2015/16 (^{a,b}Alexander *et al.*, 2015a,b).

Introduction

In addition to its emerging roles in the pathophysiology of cardiovascular disease, metabolic syndrome (MS) and type 2 diabetes (DM2) (Cirillo *et al.*, 2006; Johnson *et al.*, 2013a), increases in serum concentrations of uric acid (UA) have been identified as a risk factor for the development and progression of chronic kidney disease (CKD) including nephropathy in both types of diabetes mellitus (Ficociello *et al.*, 2010; Hovind *et al.*, 2009; Jalal *et al.*, 2010; Zoppini *et al.*, 2012).

UA is generated from xanthine by xanthine oxidase (XO) as a final step in the metabolism of endogenous and exogenous purines. The XO inhibitor (XOi) allopurinol has been used for the treatment of hyperuricaemia and prevention of gout for decades. Reflecting the evidence showing hyperuricaemia as a risk factor for CKD (Ficociello *et al.*, 2010; Hovind *et al.*, 2009; Jalal *et al.*, 2010; Zoppini *et al.*, 2012), recently, we have investigated the effects of an XOi as a new nephroprotective treatment. The nephroprotective potential of XOi has been shown in various models of CKD (Kang *et al.*, 2002; Kosugi *et al.*, 2009; Omori *et al.*, 2012; Ryu *et al.*, 2013; Sanchez-Lozada *et al.*, 2008b; Sanchez-Lozada *et al.*, 2008c). Yet studies exploring the effects of XOi on the development of nephropathy in insulin-resistant states and DM have been sparse and focused mainly on early stages of this disorder (Kosugi *et al.*, 2009; Sanchez-Lozada *et al.*, 2008b). It remains to be determined whether favourable effects of XOi will translate into long-term kidney protection in diabetes.

In parallel, some studies exploring the nephroprotective potential of XOi in patients with kidney disease have indicated that this treatment can be beneficial as it slows down the progression of CKD (Goicoechea *et al.*, 2010; Siu *et al.*, 2006). However, recent meta-analysis of interventional studies with allopurinol (Bose *et al.*, 2014) has shown that the data on effects of UA-lowering therapy on renal outcomes are still insufficient, and it remains unclear whether long-term XOi has beneficial effects on proteinuria, loss of glomerular filtration rate (GFR) and progression to end-stage renal disease (Bose *et al.*, 2014).

In this context, more studies are needed to elucidate the potential of XOi in the treatment of DN. To address this issue, we studied the long-term effects of the more recently introduced XOi febuxostat on the development and progression of proteinuria as well as on structural and molecular markers of DN *in vivo* in experimental DM2. In addition, we

conducted experiments exploring effects of febuxostat on molecular markers of fibrosis and fibrogenic signalling in renal cells stimulated *in vitro* by the components of diabetic milieu.

The results indicate that treatment with febuxostat has long-term nephroprotective effects in experimental DM2 and enhances the protective effects of renin–angiotensin system (RAS) inhibitors in the diabetic kidney. Multiple mechanisms might contribute to the beneficial actions of febuxostat in the diabetic kidney, including those related to a reduction in serum UA levels and attenuation of renal peroxynitrite formation. Moreover, febuxostat possesses antifibrotic effects *in vitro*, which are independent of changes in extracellular UA levels and attributable to inhibition of profibrotic signalling via the kinase Akt.

Methods

In vivo studies

Rat model of DM2 and nephropathy. The studies were performed in male Zucker obese (ZO, fa/fa; Charles River, Wilmington, MA, USA) rats. The evolution of metabolic changes due to an autosomal recessive mutation of the fa-gene, encoding the leptin receptor, resulting in hyperphagia, obesity, and hyperlipidaemia, IR, hyperinsulinaemia, impaired glucose tolerance, and slowly progressive increases in blood glucose (BG) levels (Ionescu *et al.*, 1985; Kasiske *et al.*, 1992; Phillips *et al.*, 1996); as well as renal structural changes and molecular markers of nephropathy in ZO rats have been well characterized (Chander *et al.*, 2004; Coimbra *et al.*, 2000; Figarola *et al.*, 2008; Kasiske *et al.*, 1985). Age-matched Zucker lean (ZL) rats without metabolic defects served as controls.

Animal welfare. The animals were housed with a light-dark cycle of 12 h each, and with free access to food (standard chow) and water. All experiments were carried out with the approval of and in accordance with the regulations of the Portland VA Medical Center IACUC Committee and are reported in compliance with the ARRIVE guidelines (Kilkenny *et al.*, 2010; McGrath & Lilley, 2015).

Study design. The treatment was initiated in 10-week-old ZO rats. The rats were randomly selected to receive the following

treatments: febuxostat [ZO-FBX, $n = 12$, $5 \text{ mg}\cdot\text{kg}^{-1}$ body wt (bwt) in drinking water after dissolution by sonication in 0.05 N NaOH ; Sanchez-Lozada *et al.*, 2008b; 2008c] or drinking water vehicle alone (ZO-VE, $n = 12$), supplemented with $0.6 \text{ mg}\cdot\text{kg}^{-1}$ of NaCl to maintain a salt concentration corresponding to that of the febuxostat-containing water. An additional group of ZO rats received 'late' intervention (ZO-FBXlate, $n = 12$) starting at week 10 of the follow up (20 weeks of age). The ZO-FBX and ZO-VE rats were compared with animals treated with the ACE inhibitor (ACEI) enalapril (ZO-ENL, $n = 12$, $10 \text{ mg}\cdot\text{kg}^{-1}\cdot\text{day}^{-1}$), an established treatment for DN. Furthermore, the effects of the combination of febuxostat ($5 \text{ mg}\cdot\text{kg}^{-1}$) and enalapril ($10 \text{ mg}\cdot\text{kg}^{-1}\cdot\text{day}^{-1}$) were also evaluated in ZO animals (ZO-FBX/ENL, $n = 12$). The results of structural and molecular analyses were also compared with vehicle-treated age-matched ZL rats (ZL-VE, $n = 10$).

The treatment in ZO-FBX, ZO-ENL and combination groups was maintained for 18 weeks and 8 weeks in the late FBX treatment group. BG and awake systolic BP (SBP) (tail plethysmography) (Anderson *et al.*, 2010) were measured at 6 week intervals together with 24 h urinary albumin excretion (U_{albV}), using metabolic cages. Following the last measurements in metabolic cages, the animals were anaesthetized and blood samples were obtained from the abdominal aorta for biochemical analyses [glycosylated haemoglobin (HBA1c), serum triglyceride and UA concentrations, and markers of systemic oxidative stress]. The kidneys were then harvested, partitioned and processed for histological evaluation, immunohistochemistry (IHC), western blotting and mRNA expression studies targeting molecular markers of DN and key signalling events involved in the pathophysiology of the disorder.

Renal structural analysis. Structural analyses were performed as previously described (Komers *et al.*, 2011; Raij *et al.*, 1984). Glomerulosclerosis score (GSS) was determined on periodic acid Schiff-stained paraffin sections using a scale ranging from 0 to 4 for normal (0), 1 = 25% sclerosis, 2 = 50% sclerosis, 3 = 75% sclerosis and 4 = 100% sclerosis. On average, 200 glomeruli were evaluated per rat. Glomeruli with scores of 3 and 4 were considered to be severely affected. The proportion of severely affected glomeruli (SAG) per section was determined for each kidney. The measurements of tubulointerstitial fibrosis score (TIFS) were performed on Masson's Trichrome-stained sections. The 100-square grid was applied at low magnification ($50\times$) on consecutive microscopic cortical fields. The number of squares containing stained fibrous tissue or atrophic tubules was recorded for each field and averaged for each kidney. Eighteen fields were evaluated for each kidney. All structural analyses were conducted in a blind manner on unidentified sections.

In vitro studies

Cell culture. Rat proximal tubule epithelial cells (NRK-52E; American Tissue Culture Collection, Rockville, MD, USA) were maintained in DMEM (Life Technologies, Carlsbad, CA, USA) supplemented to 25 mM glucose with 10% fetal bovine serum.

Treatments. Serum-deprived NRK were cultured and stimulated with TGF- β ($2 \text{ ng}\cdot\text{mL}^{-1}$; R&D Systems, Minneapolis, MA, USA) or with the receptor for advanced glycosylation end-products (RAGE) ligand S100B ($10 \text{ }\mu\text{g}\cdot\text{mL}^{-1}$, Calbiochem-Millipore, Billerica, MA, USA) for prespecified periods of time (details provided in figure legends) alone or after 30 min pretreatment with (100 nM). The febuxostat concentration used in these experiments is based on published concentrations far exceeding K_i values for purified bovine milk XO, indicating inhibition of both the oxidized and reduced forms of XO, but no significant effects on the activities of enzymes of purine and pyrimidine metabolism (Takano *et al.*, 2005). After completion of study periods, the cells were collected, homogenized and processed for further analyses by immunoblotting. All *in vitro* studies were performed at least in triplicate.

General methods

Immunoblotting and IHC. The kidney cortical samples or cells were homogenized, processed and analysed by immunoblotting as previously described (Komers *et al.*, 2006; Komers *et al.*, 2011), using primary antibodies raised against TGF- β (1:600; Cell Signaling, Beverly, MA, USA), connective tissue growth factor (CTGF, 1:800; Santa Cruz, Santa Cruz, CA, USA), collagen IV (Coll4, 1:1500; Abcam, Cambridge, MA, USA), fibronectin (1:1000; Sigma, St. Louis, MO, USA), fibroblast-specific protein 1 (FSP1/S100A4, 1:2500; Abcam), α -smooth muscle actin (α -SMA, 1:500; Abcam), vimentin (1:600; Santa Cruz), phospho-Smad3 (1:500; Cell Signaling), total Smad3 (Cell Signaling), phospho Ser⁴⁷³ Akt (1:500; Cell Signaling), total Akt (Cell Signaling) and nitrotyrosine (NT, 1:500; Santa Cruz) followed by HRP-conjugated secondary antibodies (Pierce, Rockford, IL, USA). Visualization, loading control and image analysis were accomplished as described previously (Komers *et al.*, 2006). The same antibodies directed against FSP1, vimentin, α -SMA and nitrotyrosine were used for IHC, as well as anti-ED-1 antibody (1:100; Santa Cruz) on formalin-fixed, paraffin-embedded kidney sections and processed as previously described (Komers *et al.*, 2006). ED-1 positive cells, as a marker of macrophage infiltration, were counted in 10 consecutive microscopic fields ($\times 100$) and data presented as a mean \pm SEM per field.

Isolation of total RNA, synthesis of cDNA and quantitative real-time PCR. RNA was extracted from frozen tissue fragments using trizol reagent (Invitrogen, Carlsbad, CA, USA) as per the manufacturer's protocol. To synthesize cDNA, 400 ng of trizol-extracted total RNA was reverse transcribed with 200 units MMLV reverse transcriptase (Invitrogen) and oligo-DT primers in 20 μL reactions. RT reactions were diluted 100-fold with water prior to use in qPCR (Komers *et al.*, 2013). The PCR reactions were done in 384-well plates 2.0 μL final volume (1.0 μL cDNA template and 1.0 μL master mix, Bioline, Taunton, MA, USA) on ABI7900HT Real Time PCR system (40 cycles, annealing temperature 60°C for all primers) (ABI Biotechnology, Eldersburg, MD, USA). All PCR reactions were run in quadruplicates. mRNA levels are represented as ratios to β -actin transcript calculated as differences from their respective average cycle threshold (Ct) values, deltaCt. The

gene-specific primers Coll4, fw, are as follows: CACTATGAA-AACCGTAAAGTGCCTTA, rev: GCAAACAGAGGCCAACGAA; Fibronectin, fwd: GGTCTGGGATCAAAGGGAAA, rev: CGCT-CTTGGGTACTACCATT; CTGF, fw: GACAGCTTGTGGCAAG-TGAA, rev: TCGATGGTGTGGAGTTT; plasminogen activator inhibitor-I (PAI-I), fw: GCTCCTGGTCAACCACCTTA, rev: TGGAGATGTAACGGATGCAG; FSP1, fw: AGGACAGACGA-AGCTGCATT, rev: CTCACAGCCAACATGGAAGA; vimentin, fw: ACTTCTCAGCACCACGATGA, rev: CTTTTGGGGGTGT-CAGTTGT; β -actin, fw: CTCTGAACCCTAAGGCCAAC, rev: GACCAGAGGCATACAGGGAC.

Measurements of UA and xanthine. Serum UA and xanthine concentrations were measured by HPLC as previously described (Cooper *et al.*, 2006) using HPLC column Synergi 4u Hydro-RP 80A, 250 × 4.6 mm and guard column SecurityGuard Guard Cartridge from Phenomenex (Torrance, CA, USA). Standards and samples were diluted 1:5 with 47 mM KH_2PO_4 , filtered through Centrifree YM-30 filters and injected (50 μL) onto an HPLC column.

Biochemical methods. Serum creatinine concentrations were measured by spectrophotometric assay (Biovision, Mountain View, CA, USA). Urinary albumin concentrations were determined using the Nephurat kit (Exocell, Philadelphia, PA, USA). HBA1c was determined by affinity column chromatography (Glyco-Gel B; Pierce Chemical). Thiobarbituric acid reactive substances (TBARS; Cayman Chemical, Ann Arbor, MI, USA) (Yagi, 1998) and 8-isoprostane (8-epi $\text{PGF}_{2\alpha}$; Cayman Chemical) (Morrow *et al.*, 1995) concentrations were analysed by ELISA according to the manufacturer's instructions. Blood glucose was measured using a reflectance meter (One Touch II; Lifescan, Milpetas, CA, USA).

Statistical analysis. The data and statistical analysis comply with the recommendations on experimental design and analysis in pharmacology (Curtis *et al.*, 2015). Data are expressed as means \pm SEM and analysed by ANOVA with Newman-Keuls multiple comparison *post test*. Statistical significance was defined as $P < 0.05$. The *post hoc* test was

performed only if $P < 0.05$, and there was no significant inhomogeneity of variance. Urinary albumin excretion was not normally distributed and therefore analysed after logarithmic transformation and expressed as geometric means \times/\div tolerance factor.

Results

Physical and metabolic parameters in Zucker lean (ZA) and ZO rats

As summarized in Table 1, all ZO rats demonstrated increases in body weight and renal hypertrophy, moderate hyperglycaemia and increases in HBA1c and serum triglyceride concentrations as compared with lean animals. None of these parameters were influenced by treatment with febuxostat, enalapril or their combination. Treatment with febuxostat normalized UA in all treated groups and resulted in reciprocal increase in xanthine concentrations. In ZO-ENL, UA and xanthine concentrations remained comparable with those in ZO-VE rats.

Effects of febuxostat, enalapril and their combination on BP, $U_{\text{alb}}V$, renal structural parameters and gene expression

At baseline, there were no differences in SBP among the groups of rats. Figure 1 shows mean treatment-induced changes in SBP from baseline in individual groups of rats. At week 12, SBP responses in enalapril-treated animals (both in monotherapy and in combination with febuxostat) were different from ZO-VE. At the end of the follow up, SBP responses in ZO-ENL were different from ZO-VE and ZO-FBXlate, whereas ZO rats receiving febuxostat and enalapril combination demonstrated different responses as compared with other groups of rats except the ZO-ENL counterparts. The means of SBP in individual groups at baseline and at subsequent study time points are presented in the Supporting Information Fig. S1.

As shown in Figure 2, $U_{\text{alb}}V$ progressively increased in ZO-VE rats. This increase in $U_{\text{alb}}V$ was attenuated by all

Table 1

Physical and metabolic parameters after completion of the study

Group	bwt (g)	RKW (g)	RKW/bwt ($\text{g} \cdot 100 \text{g}^{-1} \text{bwt}$)	BG (mM)	HBA1c (%)	TG (mM)	UA (μM)	Xanthine (μM)
ZL-VE	389 \pm 10	1.2 \pm 0.1	0.32 \pm 0.01	4.0 \pm 0.3	3.2 \pm 0.1	5.2 \pm 1.2	29 \pm 1	16 \pm 1
ZO-VE	459 \pm 13 ^a	2.0 \pm 0.1 ^a	0.44 \pm 0.02 ^a	16.2 \pm 1.5 ^a	4.9 \pm 0.2 ^a	26.6 \pm 2.4 ^a	58 \pm 12 ^{ab}	15 \pm 1
ZO-FBX	458 \pm 23 ^a	1.9 \pm 0.1 ^a	0.42 \pm 0.02 ^a	17.7 \pm 1.1 ^a	4.9 \pm 0.2 ^a	29.5 \pm 1.2 ^b	31 \pm 3	48 \pm 7 ^c
ZO-FBXlate	440 \pm 15 ^a	2.0 \pm 0.2 ^a	0.46 \pm 0.03 ^a	18.0 \pm 1.0 ^a	5.0 \pm 0.2 ^a	26.2 \pm 1.0 ^a	25 \pm 2	49 \pm 3 ^c
ZO-ENL	454 \pm 22 ^a	1.8 \pm 0.1 ^a	0.40 \pm 0.02 ^a	17.5 \pm 1.4 ^a	5.2 \pm 0.2 ^a	29.2 \pm 2.0 ^a	63 \pm 4 ^{ab}	19 \pm 1
ZO-FBX + ENL	464 \pm 14 ^a	2.0 \pm 0.1 ^a	0.44 \pm 0.03 ^a	16.0 \pm 1.6 ^a	5.2 \pm 0.2 ^a	25.5 \pm 1.5 ^a	29 \pm 2	45 \pm 4 ^c

There were 10 animals in the ZL-VE group and 12 animals in each ZO group.

RKW, right kidney wt; LKW, left kidney wt; TG, triglycerides.

^a $P < 0.05$ versus ZL-VE.

^b $P < 0.05$ versus FBX-treated rats (ZO-FBX, ZO-FBXlate, ZO-FBX + ENL).

^c $P < 0.05$ versus rats not receiving FBX.

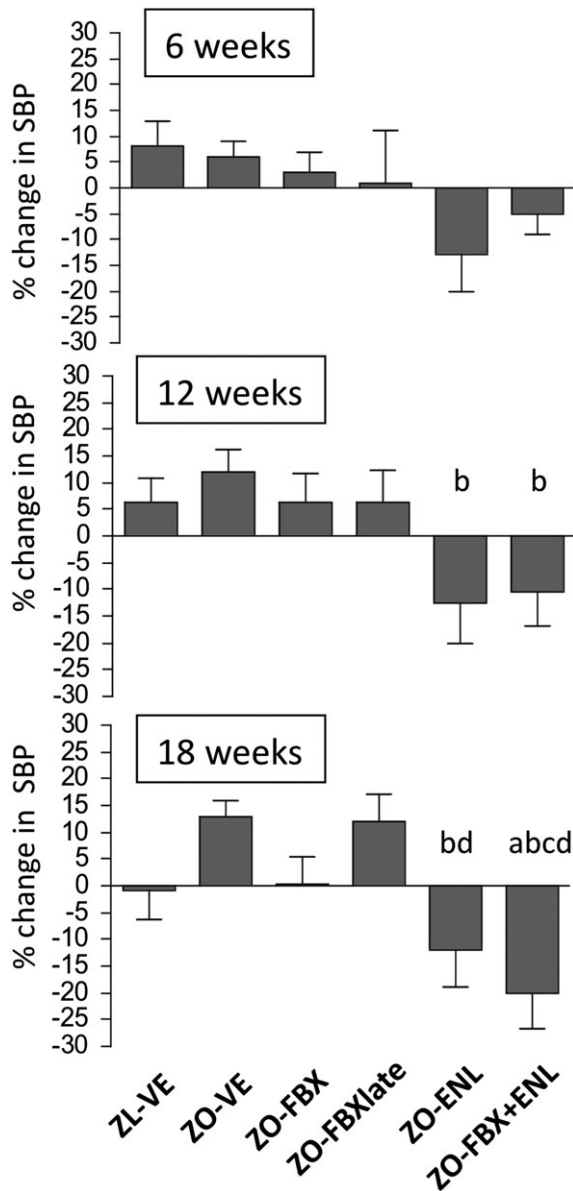


Figure 1

Treatment-induced changes in SBP. Graphs show % changes in SBP in individual groups of rats at different study time points as compared with baseline. There were 10 animals in ZL-VE group and 12 animals in each ZO group. ^a $P < 0.05$ versus ZL-VE; ^b $P < 0.05$ versus ZO-VE; ^c $P < 0.05$ versus ZO-FBX; ^d $P < 0.05$ versus ZO-FBXlate.

treatments including the febuxostat, albeit not when administered later in the course of the disease. There were no statistically significant differences in $U_{alb}V$ between the ZO-FBX, ZO-ENL and the ZO-FBX + ENL groups of rats. Serum creatinine concentrations, determined at the end of the follow up, were elevated in ZO-VE rats as compared with lean controls (ZL-VE: 78 ± 9 vs. ZO-VE: 143 ± 9 μM , $P < 0.05$). All treatments resulted in a decrease in S-Cr, albeit only the rats receiving FBX + ENL combination displayed a significant S-Cr reduction compared with untreated ZO rats (ZO-FBX: 116 ± 10 , ZO-FBXlate: 119 ± 13 , ZO-ENL: 110 ± 13 , ZO-FBX + ENL: 90 ± 6 ; $P < 0.05$, ZO-FBX + ENL vs. ZO-VE)

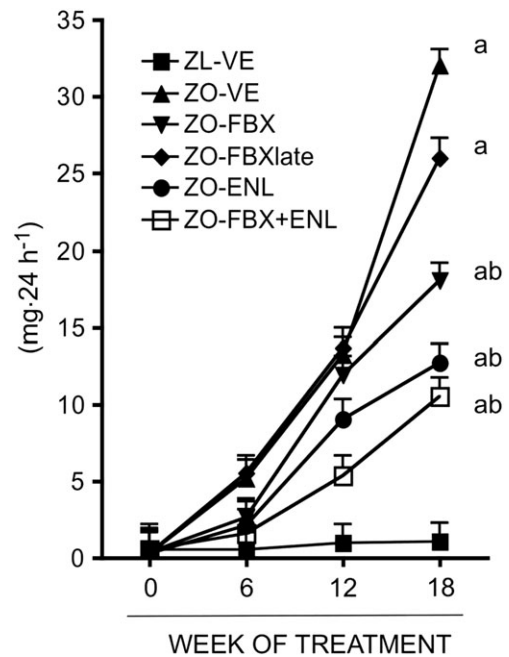


Figure 2

Effect of febuxostat, enalapril and their combination on urinary albumin excretion. Data are expressed as geometric means \times/\pm tolerance factor. Statistically significant differences between the groups were detected at week 18. There were 10 animals in ZL-VE group and 12 animals in each ZO group. ^a $P < 0.05$ versus ZL-VE; ^b $P < 0.05$ versus ZO-VE.

Compared with their lean counterparts, ZO-VE developed advanced GS and TIF, as determined by increases in GSS and TIFS as well as higher proportion of SAG (Figure 3). Treatment with febuxostat attenuated both the development of GS (GSS: $P < 0.05$; SAG: $P < 0.05$ vs. ZO-VE) and TIF ($P < 0.05$). The delayed treatment with febuxostat was also effective at reducing GS and TIF scores, albeit the proportion of SAG in ZO-FBXlate was not statistically lower as compared with that in ZO-VE rats. The monotherapy with enalapril also significantly ameliorated the development of GS and TIF in ZO rats ($P < 0.05$ vs. ZO-VE; Figure 3). Some of these effects of enalapril were more prominent as compared with those of febuxostat-treated animals (GSS: $P < 0.05$ vs. ZO-FBX, $P < 0.05$ vs. ZO-FBXlate; SAG: $P < 0.05$ vs. ZO-FBXlate; TIFS: $P < 0.05$ vs. ZO-FBX, $P < 0.05$ vs. ZO-FBXlate). The combination of both agents had similar beneficial effects on GS characteristics and TIF as the monotherapy with enalapril (Figure 3), being more prominent than in febuxostat-treated animals. However, the TIFS was, in ZO-FBX + ENL rats, lower than that in all other groups of ZO rats including the ZO-ENL and was not different from that in ZL rats.

Consistent with histological findings, this study shows that ZO-VE rats displayed increased mRNA expression of the selected genes associated with TIF (Coll4, fibronectin, PAI-I, FSP1 and vimentin) (Table 2), albeit the increase in CTGF mRNA did not reach statistical significance. Treatment with febuxostat resulted in lower FSP1 mRNA expression as compared with untreated ZO-VE. This effect was also apparent in rats with delayed febuxostat treatment. A reduction in

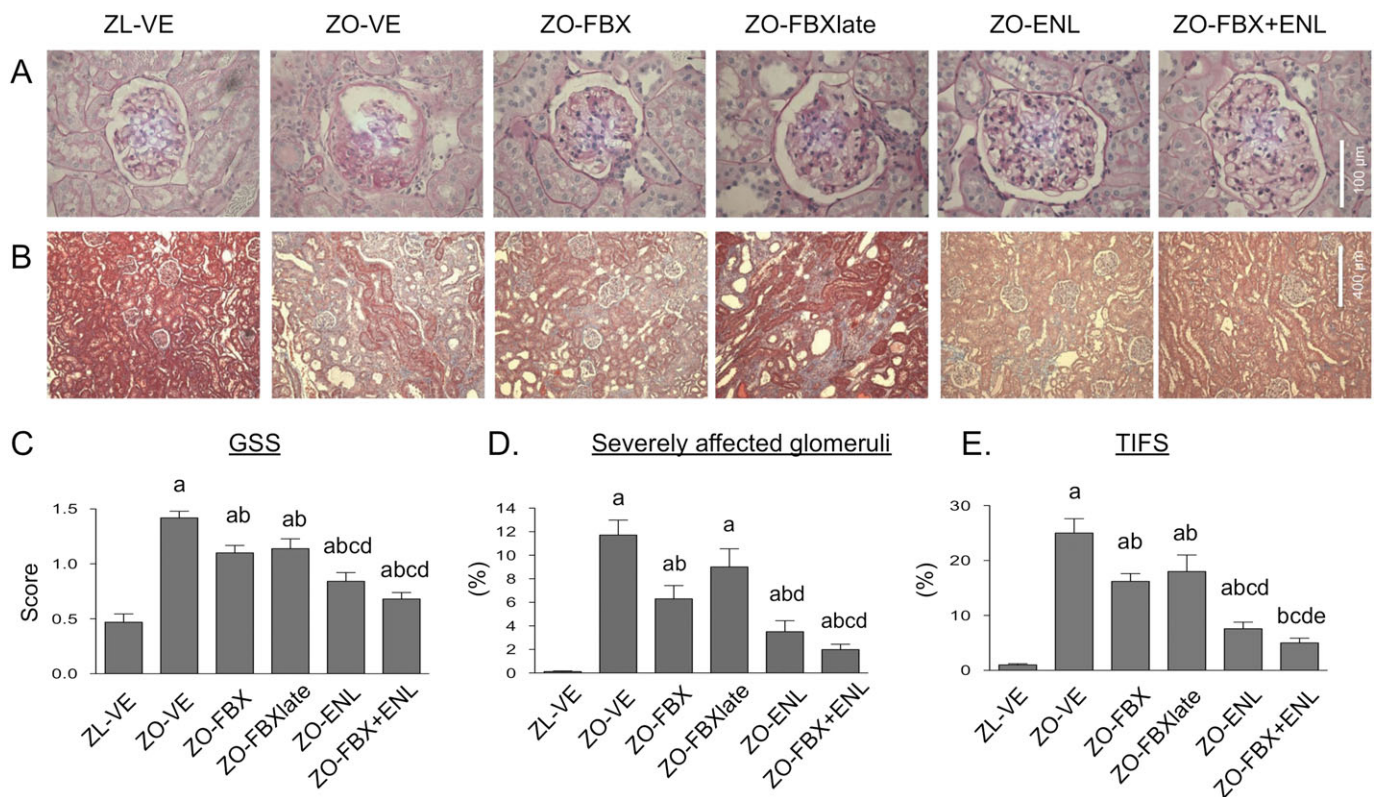


Figure 3

Analysis of GSS, % of severely affected glomeruli (SAG) and TIFS. (A) Representative microphotographs of glomeruli stained with periodic acid Schiff with different degrees of glomerular sclerosis (GS) and (B) interstitial regions stained with Trichrome showing areas of various degrees of interstitial fibrosis and tubular atrophy in individual groups of rats. (C) Bar graph presentation of quantitative evaluation of GSS, (D) % of SAG and (E) TIFS. There were 10 animals in ZL-VE group and 12 animals in each ZO group. ^a*P* < 0.05 versus ZL-VE; ^b*P* < 0.05 versus ZO-VE; ^c*P* < 0.05 versus ZO-FBX; ^d*P* < 0.05 versus ZO-FBXlate; ^e*P* < 0.05 versus ZO-ENL.

FSP1 mRNA was also detected in ZO-ENL, in addition to a reduction in PAI-I. Vimentin mRNA was lower in febuxostat-treated animals as compared with ZO-ENL. Importantly, the combination treatment was effective in reducing mRNA

expression of all selected profibrotic genes as compared with ZO-VE and in some cases (fibronectin, CTGF, PAI-I and vimentin) compared with febuxostat or enalapril monotherapies (Table 2).

Table 2

Expression of profibrotic genes in ZL and ZO rats

Gene	Rat group					
	ZL-VE	ZO-VE	ZO-FBX	ZO-FBXlate	ZO-ENL	ZO-FBX + ENL
<i>Coll4</i>	0.305 ± 0.037	0.503 ± 0.068 ^a	0.351 ± 0.036	0.425 ± 0.042	0.373 ± 0.025	0.300 ± 0.050 ^b
<i>Fibronectin</i>	0.026 ± 0.003	0.074 ± 0.006 ^a	0.065 ± 0.008 ^a	0.077 ± 0.007 ^a	0.059 ± 0.009 ^a	0.044 ± 0.002 ^{a b d}
<i>CTGF</i>	0.383 ± 0.034	0.490 ± 0.054	0.389 ± 0.033	0.406 ± 0.044	0.529 ± 0.055	0.192 ± 0.034 ^{b e}
<i>PAI-I</i>	0.009 ± 0.002	0.027 ± 0.005 ^a	0.024 ± 0.003 ^a	0.028 ± 0.004 ^a	0.016 ± 0.002 ^{b c d}	0.008 ± 0.001 ^{b c d}
<i>FSP1</i>	0.083 ± 0.018	0.185 ± 0.035 ^b	0.104 ± 0.011 ^d	0.137 ± 0.014 ^c	0.122 ± 0.008 ^c	0.095 ± 0.008 ^d
<i>Vimentin</i>	0.239 ± 0.057	0.605 ± 0.094 ^a	0.397 ± 0.054 ^{d e}	0.783 ± 0.108 ^a	0.632 ± 0.057 ^a	0.376 ± 0.025 ^{b d e}

mRNA level of a given gene in each sample was normalized to β-actin transcript. Explanation of abbreviations is in the text. There were 10 animals in ZL-VE group and 12 animals in each ZO group.

^a*P* < 0.05 versus ZL-VE.

^b*P* < 0.05 versus ZO-VE.

^c*P* < 0.05 versus ZO-FBX.

^d*P* < 0.05 versus ZO-FBXlate; ^e*P* < 0.05 versus ZO-ENL.

Impact of febuxostat on renal protein expression and localization of markers and mediators of diabetic glomerulosclerosis and TIF

We next determined the impact of febuxostat on renal protein expression of established mediators and markers of GS and TIF. Moreover, these analyses gave us an opportunity to compare the effects of febuxostat with those of enalapril and determine possible differences between the two interventions.

Immunoblot analysis of profibrotic factors and extracellular matrix (ECM) proteins showed increases in renal abundance of TGF- β , CTGF and Coll4 (Figure 4) in ZO-VE as compared with lean controls. Increases in these parameters were reduced to a similar extent by both febuxostat and enalapril treatments. Further analyses determined renal cortical abundance and localization of mesenchymal markers that are being expressed during the development of TIF (Kalluri, 2009; Ren and Duffield, 2013). As shown in Figure 5, vimentin and FSP1 (also known as S100A4) were elevated in ZO-VE as compared with ZL-VE rats. These changes were ameliorated in both febuxostat- and enalapril-treated animals. Vimentin immunoreactivity in ZL-VE was localized only in glomeruli and renal vessels. In addition to glomerular localization of vimentin, observed in all groups of rats, ZO-VE displayed abundant tubulointerstitial immunoreactivity of this protein, which was less apparent in both febuxostat- and enalapril-treated diabetic kidneys (Figure 5). FSP1

immunoreactivity was barely detectable in ZL-VE kidney sections. In contrast, ZO-VE rats demonstrated accumulations of FSP1-positive cells in the cortical interstitium (Figure 5), frequently associated with areas of tubular atrophy and surrounding fibrosis. Both treatments were associated with lower number of vimentin- and FSP1-positive cells in tubulointerstitial compartment. There were no differences in renal cortical α -SMA abundance between the ZL and ZO rats when analysed by immunoblotting. Yet IHC showed that in addition to vascular structures, vehicle-treated ZO rats displayed tubulointerstitial immunoreactivity of α -SMA. This phenomenon was not detected in ZO-FBX or ZO-ENL animals, suggesting the effects of treatments in non-vascular compartments.

Effect of febuxostat on systemic and tissue markers of oxidative stress and renal inflammation

Serum levels of TBARS (Yagi, 1998) and 8-epi PGF $_{2\alpha}$ (Morrow *et al.*, 1995), which reflect peroxynitrite formation (Ferraro *et al.*, 2003; White *et al.*, 1994) and have been linked to enhanced XO activity (Gao *et al.*, 2008; Khadour *et al.*, 2002), were elevated in ZO-VE rats and remained unchanged by monotherapies with febuxostat and enalapril (Figure 6A, B). By contrast, treatment with FBX + ENL combination near-normalized serum concentrations of these markers. NT has been used as a marker of tissue nitrosylated products and a

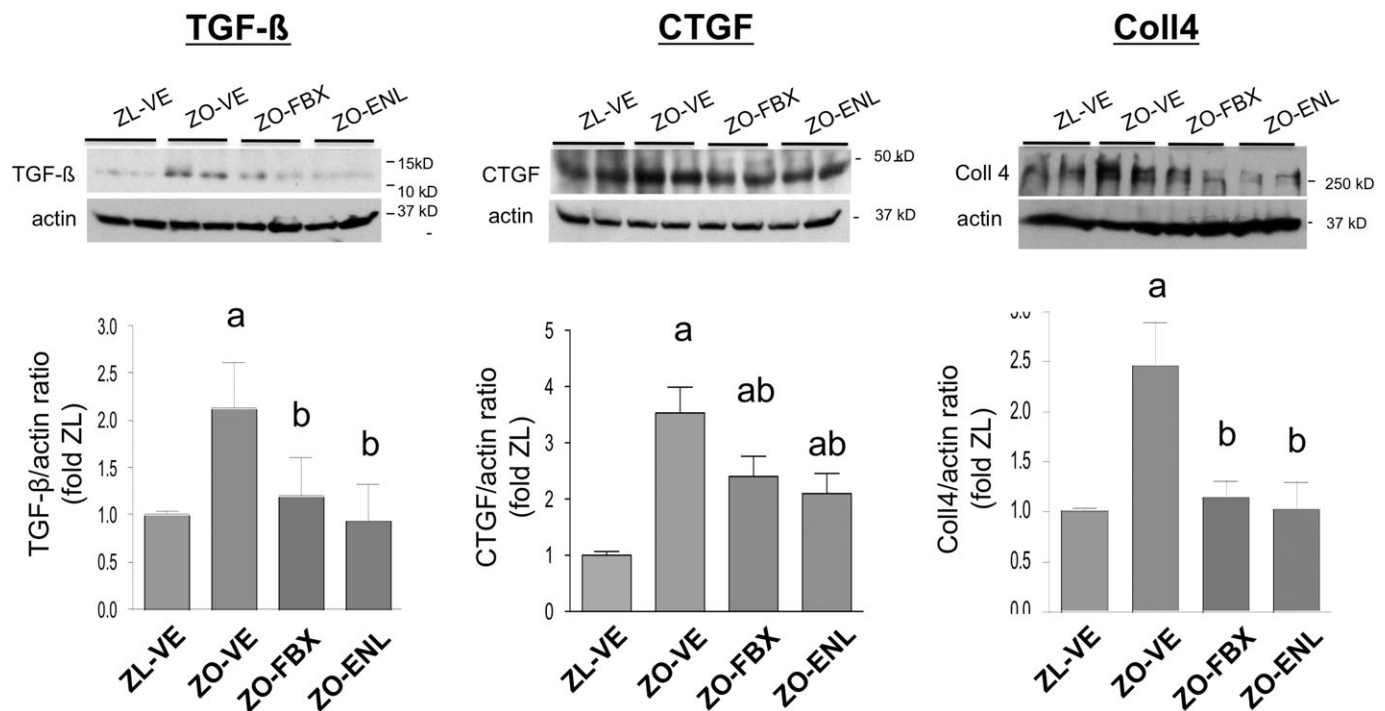


Figure 4

Renal protein expression of TGF- β , CTGF and Coll4. TGF- β (A), CTGF (B) and Coll4 (C) protein expression was analysed by immunoblotting in renal cortical homogenates. Renal homogenates from randomly selected six animals from each group were used. The upper insets show representative blots in ZL-VE (lanes 1–2) and ZO-VE (lanes 3–4) rats and in ZO-FBX (lanes 5–6) or ZO-ENL (lanes 7–8) rats. The graphs show densitometric analysis of western blots. Febuxostat and enalapril were both effective in reducing diabetes-induced increases in renal TGF- β and CTGF expression and reduced abundance of Coll4. The data are presented as mean \pm SEM of protein/actin ratios. ^a $P < 0.05$ versus ZL-VE; ^b $P < 0.05$ versus ZO-VE.

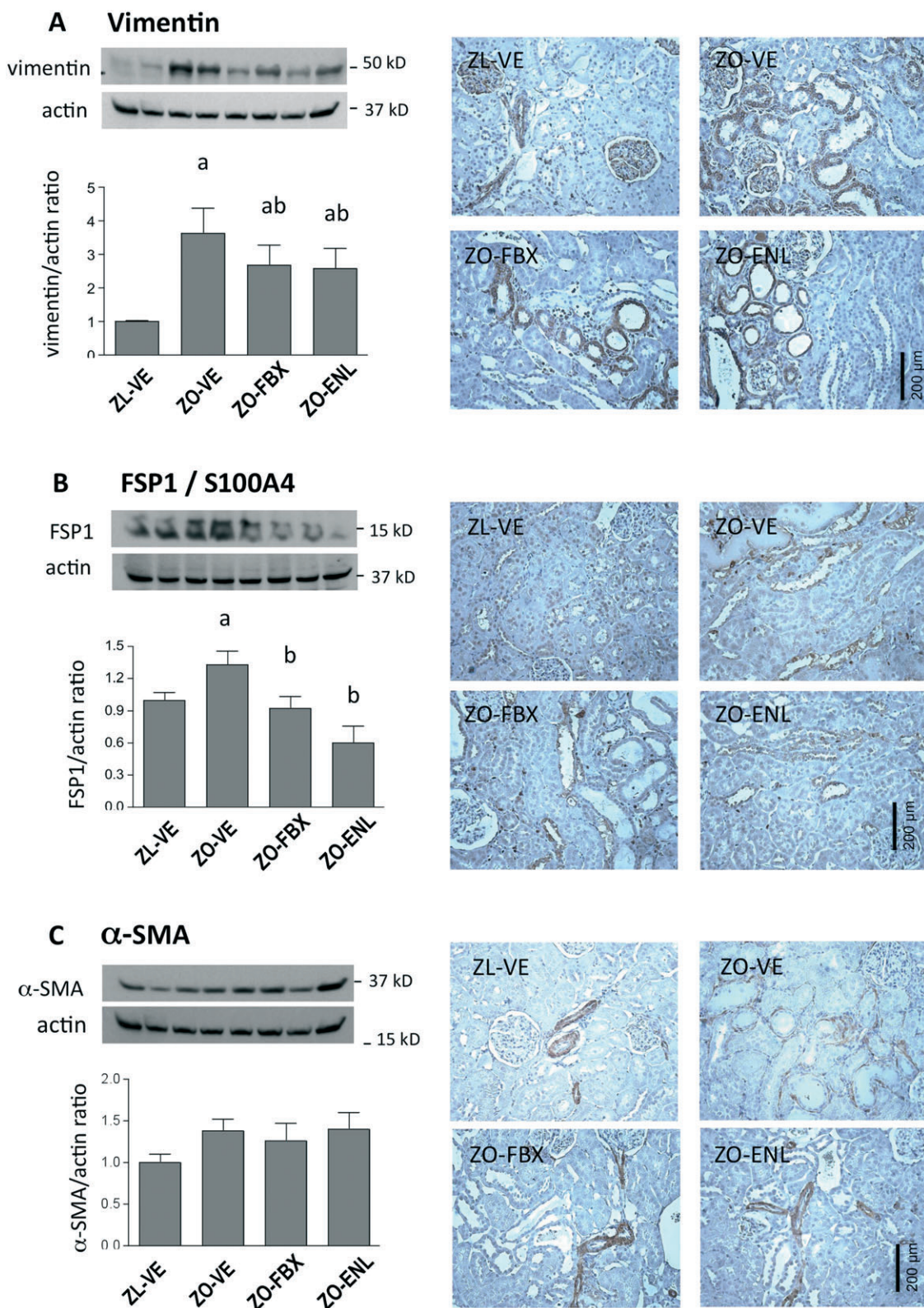


Figure 5

Immunohistochemical expression and localization of mesenchymal markers in renal cortex. Vimentin (A), FSP1/S100A4 (B) and α -SMA (C) protein expression was analysed by immunoblotting in renal cortical homogenates ($n = 6$ each group). The graphs show densitometric analysis of western blots. The upper insets show representative blots of vimentin FSP1 and α -SMA in ZL-VE (lanes 1-2) and ZO-VE (lanes 3-4) rats and in ZO-FBX (lanes 5-6) or ZO-ENL (lanes 7-8) rats. Data are presented as protein/actin ratios. Right panels show corresponding representative microphotographs ($\times 200$) of immunohistochemical localization of these proteins. ^a $P < 0.05$ versus ZL-VE; ^b $P < 0.05$ versus ZO-VE.

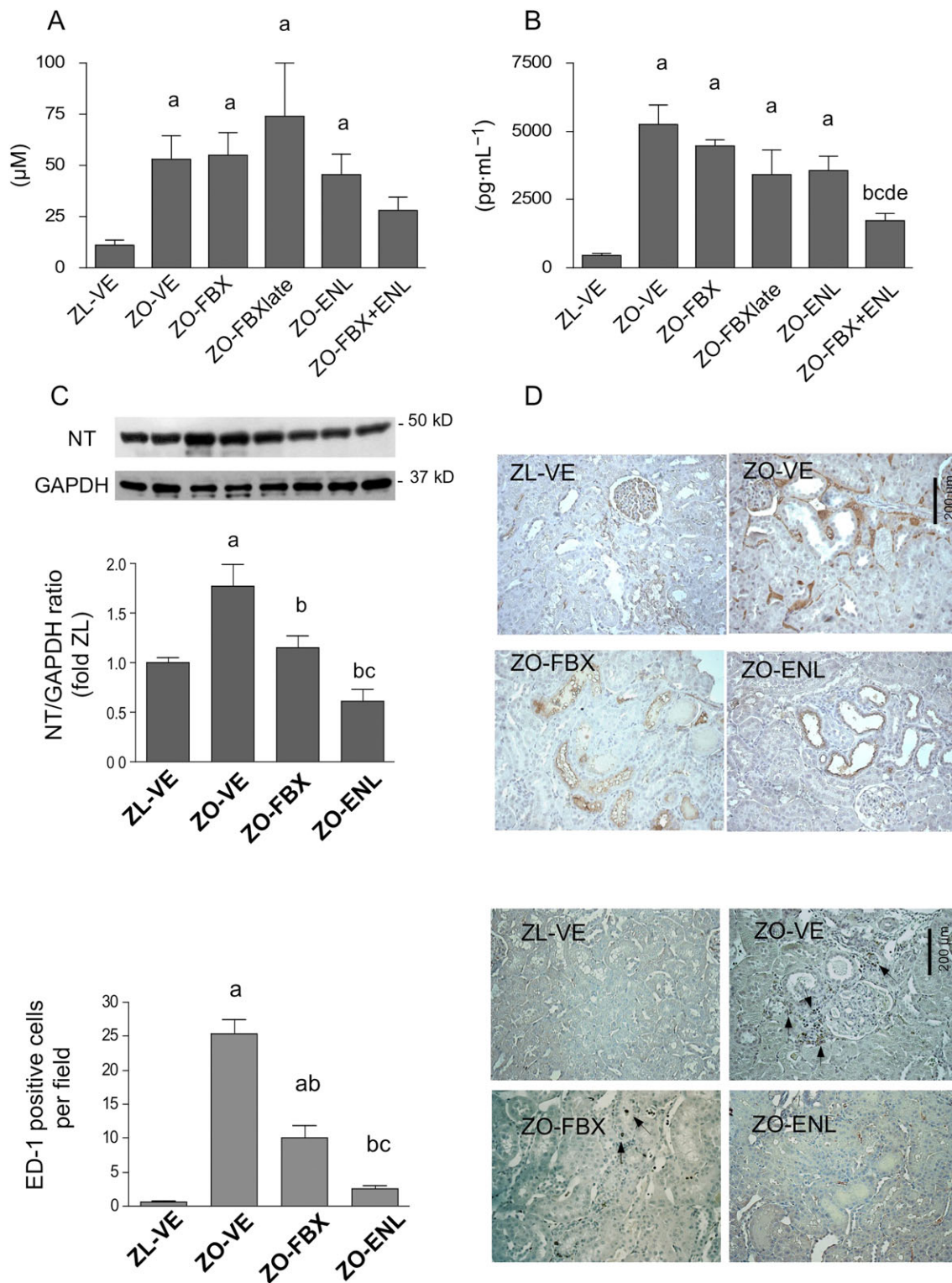


Figure 6

Systemic and renal markers of oxidative stress. TBARS (A) and 8-epi PGF_{2α} (B) were measured in serum as markers of the systemic oxidative stress by ELISA. Immunoblotting of renal cortical homogenates was used to determine abundance of NT (C) as a marker of protein nitrosylation in kidney tissue ($n = 6$ each group). The upper insets show representative blots of NT and a loading control GAPDH in ZL-VE (lanes 1–2) and ZO-VE (lanes 3–4) rats and in ZO-FBX (lanes 5–6) and ZO-ENL (lanes 7–8) groups. The graphs show densitometric analysis of western blots. Data are presented as protein/GAPDH ratios. (D) Renal immunohistochemical localization of NT in these groups of rats. (E and F) Mean numbers of ED-1 positive cells (arrows) per microscopic field and representative images in the same groups of rats. ^a $P < 0.05$ versus ZL-VE; ^b $P < 0.05$ versus ZO-VE; ^c $P < 0.05$ versus ZO-FBX; ^d $P < 0.05$ versus ZO-FBXlate; ^e $P < 0.05$ versus ZO-ENL.

fingerprint of local production of peroxynitrite (Chander *et al.*, 2004; Xia and Zweier, 1997), which could be in part attributable to actions of XO (Bagi *et al.*, 2002; Zweier *et al.*, 1994). Immunoblotting showed increased abundance of NT in ZO-VE (Figure 6C), which was significantly reduced by treatment with both febuxostat and enalapril, the latter being more effective than febuxostat. In all groups of rats, NT immunoreactivity was detected in some glomeruli and occasional endothelia of intertubular capillaries. However, tubular and interstitial NT immunoreactivity was detectable only in ZO-VE and to a lesser degree in ZO-FBX rats, whereas ZO-ENL displayed a pattern akin to lean rats (Figure 6D).

Number of ED-1-positive cells, determined as a measure of macrophage infiltration, was markedly elevated in ZO-VE rats and reduced by both febuxostat and enalapril treatment (Figure 6E, F). Treatment with enalapril was more effective than febuxostat in reducing macrophage infiltration.

Effect of febuxostat on profibrotic signalling *in vitro*

Further studies were conducted to evaluate whether the antifibrotic actions of febuxostat detected *in vivo* could be reproduced *in vitro* in a more controlled environment. Rat renal tubular cells (NRK) were stimulated with TGF- β or the RAGE ligand S100B with and without pretreatment with febuxostat. As depicted in Figure 7, febuxostat attenuated TGF- β - and S100B-induced increases in CTGF, fibronectin and α -SMA protein expression.

As shown in Figure 8A, pretreatment with febuxostat had no impact on TGF- β -induced Ser^{423/425} phosphorylation of its major signalling effector, Smad3. Analysis of renal cortical homogenates (Figure 8B) was in accord with these *in vitro* findings; ZO-VE rats displayed increased Smad3 phosphorylation as compared with lean counterparts, which was normalized by treatment with enalapril, but not with febuxostat.

There were no differences between the groups in total Smad3 abundance.

Further experiments focused on possible effects of febuxostat on Smad-independent pathways of TGF- β signalling. Akt has been implicated in this process (Lan and Du, 2014). Stimulation of NRK cells with TGF- β induced rapid Ser⁴⁷³ Akt phosphorylation in NRK cells, which was attenuated by pretreatment with febuxostat (Figure 9A). Since RAGE stimulation has been also shown to activate Akt (Arcuri *et al.*, 2005; Leclerc *et al.*, 2007), we also evaluated whether the effects of febuxostat can be observed in cells stimulated with S100B. Similar to TGF- β -stimulated cells, febuxostat attenuated S100B-induced Akt (Figure 9B). Parallel analysis of renal cortical homogenates from ZL and ZO rats confirmed DM-induced Akt phosphorylation and its reduction by treatment with febuxostat. This effect was not, however, observed in ZO rats treated with enalapril (Figure 9C).

Discussion

The treatment with febuxostat attenuated the development of proteinuria and renal structural changes in ZO rats with DM2. These effects of febuxostat were independent of major risk factors for initiation and progression of DN, such as metabolic as well as BP control. Consistent with beneficial effects in renal architecture, febuxostat reduced renal expression of some prosclerotic/profibrotic genes and attenuated the abundance of protein markers and mediators of renal fibrosis and GS.

Only a few studies have thus far evaluated the effects of XO_i in models of MS and DM2. Sanchez-Lozada *et al.*, (2008b) reported that the treatment with febuxostat decreased BP, glomerular capillary pressure and afferent arteriolar hypertrophy in insulin-resistant fructose-fed rats, findings consistent with nephroprotective effects. Renal structural

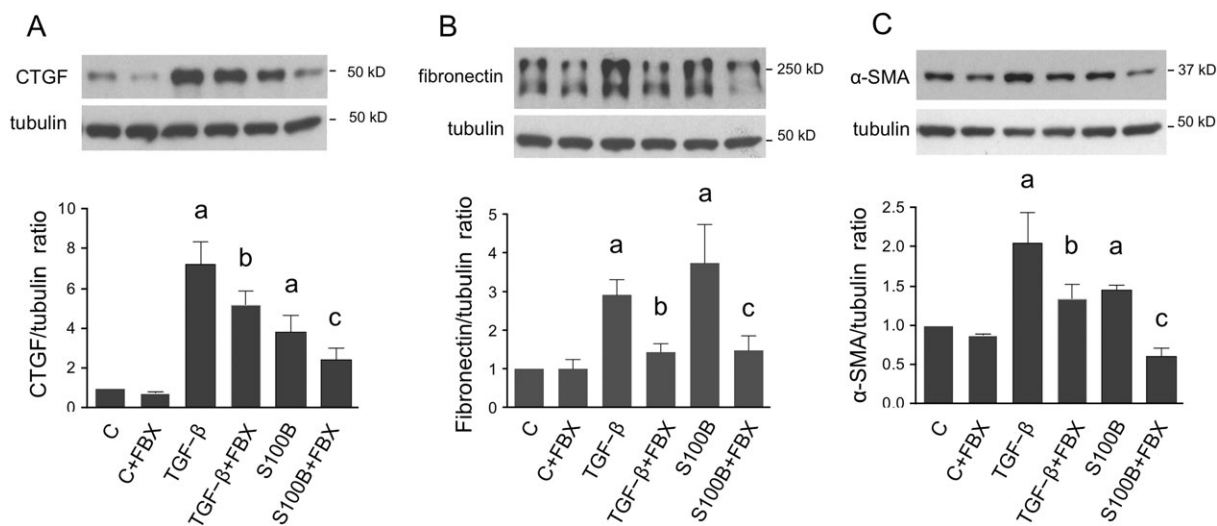


Figure 7

Effect of FBX on markers of fibrosis in renal tubular cells *in vitro*. Serum-deprived NRK cells cultured in UA-free media were stimulated with TGF- β (2 ng·mL⁻¹) or with RAGE ligand S100B (10 μ g·mL⁻¹) for 48 h alone or after 30 min pretreatment with FBX (100 nM) and compared with unstimulated cells with (C-FBX) or without FBX treatment as time controls (designed as C). CTGF (A), fibronectin (B) and α -SMA (C) were analyzed by western blot. The order of lanes corresponds to conditions in bar graphs. ^a*P* < 0.05 versus C; ^b*P* < 0.05 versus TGF- β ; ^c*P* < 0.05 versus S100B.

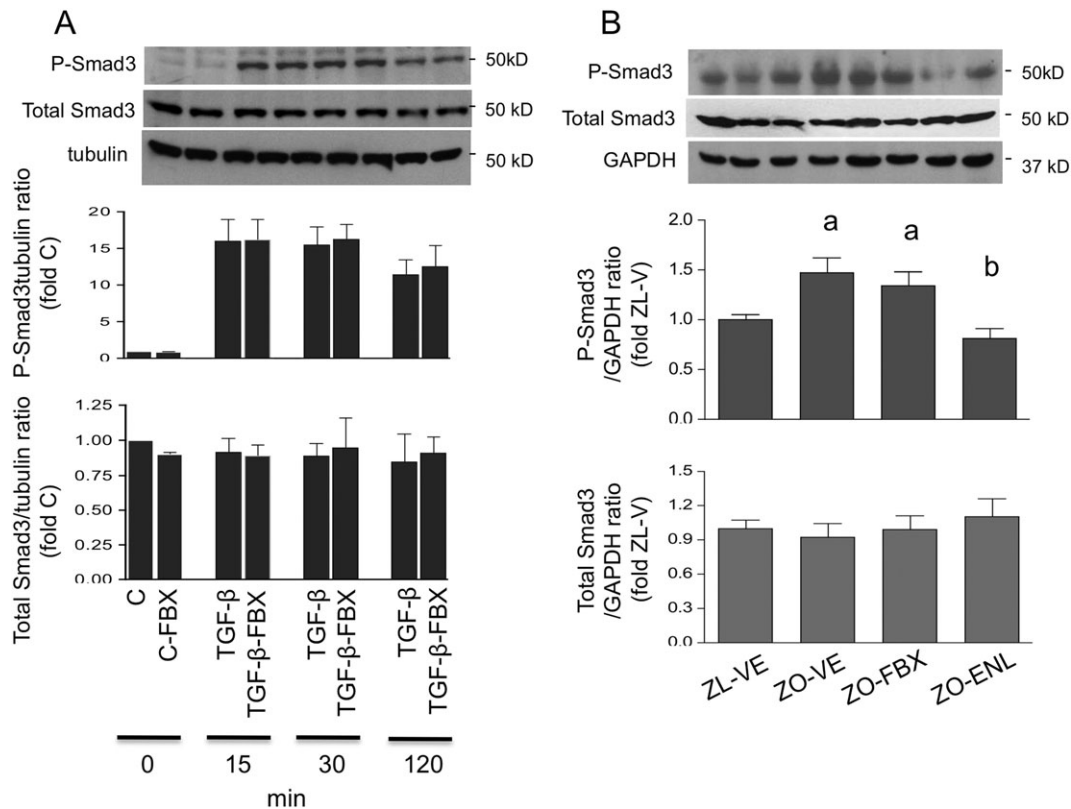


Figure 8

Effect of febuxostat on TGF- β -induced Smad3 phosphorylation. (A) Smad3 Ser^{423/425} phosphorylation (P-Smad3) and total Smad3 were determined by immunoblotting in serum-deprived NRK cells stimulated with TGF- β for 15, 30 and 120 min in the presence or absence of FBX and compared with unstimulated controls. Upper insets show representative blots of Smad3 and tubulin used as the loading control. The order of lanes corresponds to conditions in bar graphs. (B) The same technique was used for analysis of renal cortical P-Smad3 and total Smad3 protein expression in ZL and ZO rats ($n = 6$ each group). The upper insets show representative Smad3 blots and GAPDH loading control in ZL-VE (lanes 1–2) and ZO-VE (lanes 3–4) rats and in ZO-FBX (lanes 5–6) or ZO-ENL (lanes 7–8) rats. The graphs show densitometric analysis of western blots. ^a $P < 0.05$ versus ZL-VE; ^b $P < 0.05$ versus ZO-VE.

analysis or measurements of proteinuria were not performed in that study. In another report, allopurinol reduced albuminuria in db/db mice (Kosugi *et al.*, 2009) together with renal immunohistochemical reactivity of TGF- β , osteopontin and collagen III. In addition, allopurinol prevented the development of tubular dilation in db/db mice, used as a marker of tubular injury. Our present observations are in accord with these reports but extend these findings to show that a spectrum of beneficial effects of XOis found previously at early stages of nephropathy in MS and/or DM2 can be translated into long-term nephroprotection.

These complex actions of an XOis interfering with the development of TIF and GS on both the structural and molecular levels have not been thus far described in experimental DM. An effect on fibrogenesis is particularly important since the development of TIF is the major driving force of a progressive decline in GFR at the later stages of kidney diseases (Diamond and Anderson, 1990). Previous studies have indicated antifibrotic actions of XOis in non-diabetic context. In rats, oxonic acid-induced hyperuricaemia stimulated expression of mesenchymal markers in renal tubular cells after a relatively short period of time (6–7 weeks) (Ryu *et al.*, 2013). These changes were ameliorated by the treatment with

allopurinol, which also led to a significant reduction in serum UA levels. Similarly, allopurinol ameliorated TIF in rats with unilateral ureteral obstruction (Omori *et al.*, 2012) and in the rat remnant kidney model (Sanchez-Lozada *et al.*, 2008c).

Hyperuricaemia can trigger spectrum mechanisms, which are relevant for the pathophysiology of DN, such as haemodynamic changes leading to increased glomerular pressure and structural changes in afferent arterioles (Sanchez-Lozada *et al.*, 2008a), endothelial dysfunction with reduced bioavailability of NO (Gersch *et al.*, 2008; Hu *et al.*, 2009; Sanchez-Lozada *et al.*, 2008a), activation of RAS (Corry *et al.*, 2008; Mazzali *et al.*, 2002) and a variety of pro-oxidant, inflammatory and fibrogenic actions (Hu *et al.*, 2012; Ryu *et al.*, 2013; Sanchez-Lozada *et al.*, 2008a; Yang *et al.*, 2010). Consequently, as an XOis, febuxostat is most likely to exert its protective effects by lowering UA. Indeed, protective effects of febuxostat in the kidney were paralleled by normalization of UA levels in ZO rats without affecting established driving forces of DN such as hyperglycaemia or hyperlipidaemia.

Yet in addition to its role in UA synthesis, XO is an enzyme with dual function that contributes to the generation of ROS (Zweier *et al.*, 1994) and secondarily to peroxynitrite

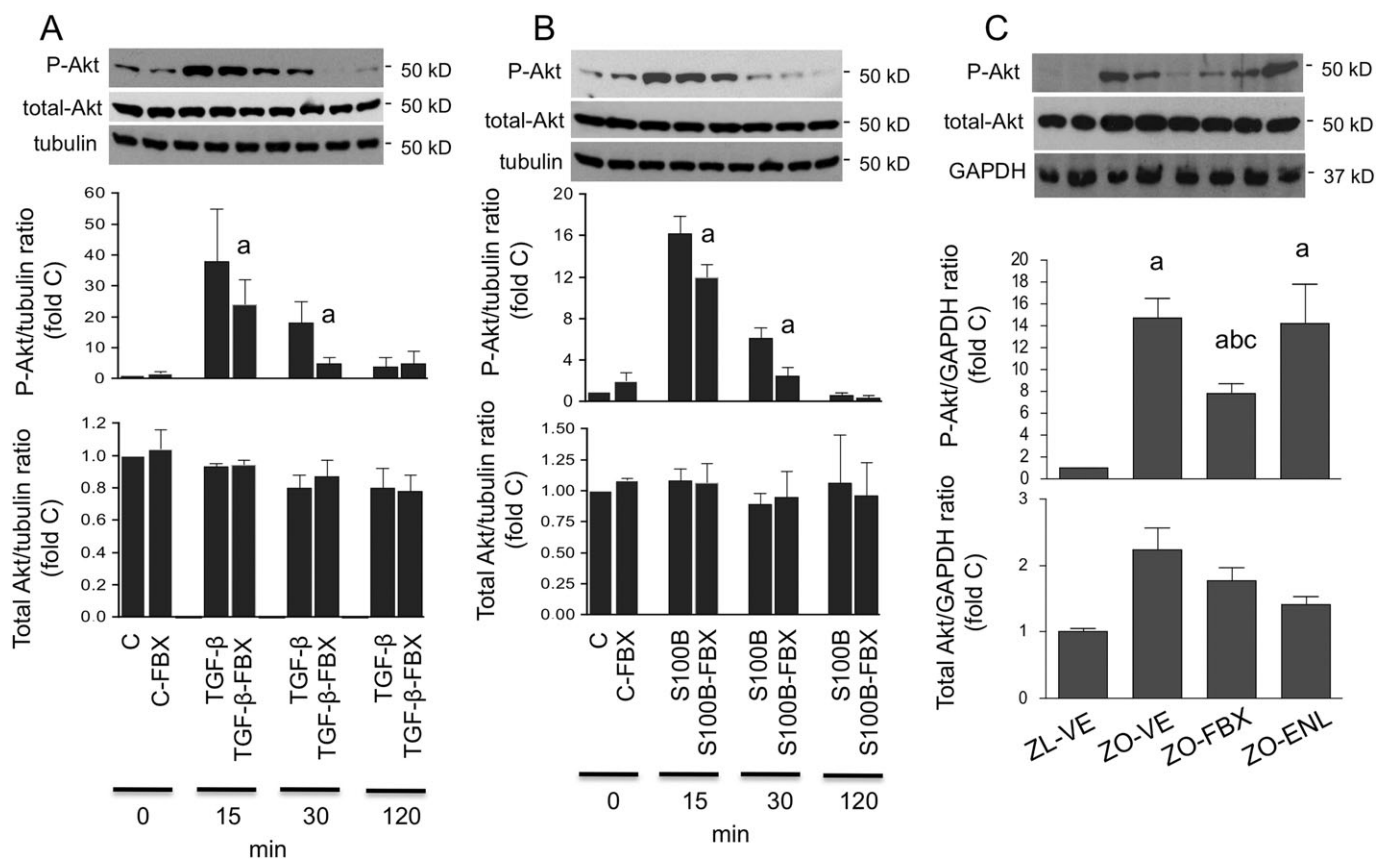


Figure 9

Effect of febuxostat on Akt phosphorylation. Serum-deprived NRK cells were stimulated with TGF- β (A) or S100B (B) for 15, 30 and 120 min in the presence or absence of FBX and compared with unstimulated cells (abbreviated as C). Ser⁴⁷³ phosphorylation (P-Akt) and total Akt were determined by immunoblotting. Upper insets show representative blots of Akt and tubulin loading control. The order of lanes corresponds to conditions in bar graphs. ^a $P < 0.05$ versus TGF- β - or S100B-stimulated counterparts without FBX treatment. (C) The same technique was used for analysis of renal cortical P-Akt and total Akt protein expression in ZL and ZO rats ($n = 6$ each group). The upper insets show representative Akt blots and GAPDH loading control in ZL-VE (lanes 1–2) and ZO-VE (lanes 3–4) rats and in ZO-FBX (lanes 5–6) or ZO-ENL (lanes 7–8) rats. The graphs show densitometric analysis of western blots. ^a $P < 0.05$ versus ZL-VE; ^b $P < 0.05$ versus ZO-VE; ^c $P < 0.05$ versus ZO-ENL.

formation (Bagi *et al.*, 2002), one of the most powerful reactive species produced by the reaction of NO and superoxide radicals. Consequently, XO actions may contribute to renal pathophysiology via increases of the systemic or local nitrosative stress.

Treatment with febuxostat partially ameliorated the increases in renal NT abundance observed in ZO rats, in accord with the postulated impact of the drug on enhanced peroxynitrite formation in the kidney. A reduction in renal NT has been previously linked to nephroprotective actions of a variety of interventions, including XO_i, in different models of kidney disease (DeRubertis *et al.*, 2004; Omori *et al.*, 2012; Onozato *et al.*, 2002). In contrast, previously described beneficial renal effects of allopurinol in db/db mice were not associated with reductions of renal NT and other indicators of oxidative stress (Kang *et al.*, 2002; Kosugi *et al.*, 2009; Omori *et al.*, 2012; Ryu *et al.*, 2013; Sanchez-Lozada *et al.*, 2008b; 2008c). Longer treatment period, as applied in the present study, may be required for detection of this effect of XO_i in the diabetic kidney, or as suggested by some studies in the non-renal context, febuxostat may be more effective than allopurinol in its antioxidant efficacy (Malik *et al.*,

2011; Tausche *et al.*, 2014). Unlike tissue NT, treatment with febuxostat did not influence systemic markers of ROS generation, suggesting that XO is not the major source of the systemic production of ROS at least in this model of DM2. Yet some antioxidant potential could be inferred from its ability to enhance the antioxidant actions of an ACEI, discussed below. Altogether, a reduction of nitrosative stress could, in addition to UA lowering, contribute to the protective actions of febuxostat in the kidney. Adding complexity to this matter, hyperuricaemia *per se* has been implicated in oxidative stress, leading to early functional and molecular changes in the rat kidney (Sanchez-Lozada *et al.*, 2008a), and UA lowering could further enhance the antioxidant effects of the XO_i.

We compared the nephroprotective potential of febuxostat with the ACEI enalapril, one of the established treatments for DN. As expected, monotherapy with enalapril had beneficial effects in ZO rat kidneys, and the data indicate that enalapril is more protective than febuxostat in diabetic kidney considering its more prominent impact on TIF, NT abundance and inflammatory infiltration. Yet other effects of febuxostat on major molecular markers of DN and fibrosis were comparable with those of enalapril. Importantly, the

best therapeutic effect was observed in ZO rats treated with a combination of both agents as the kidney function was best preserved in this group and TIFS was reduced to a greater degree than achieved by both monotherapies. Superior effects of the combination were also apparent when we analysed the renal mRNA expression of profibrotic genes, up-regulated in diabetic kidneys, and system markers of oxidative stress. Altogether, these observations suggest potentially clinically relevant additive nephroprotective effects of the combination as compared with both monotherapies. Of note, SBP was at the end of the follow up the lowest in the combination group, which could have contributed to superior nephroprotection. However, as shown in Figure 1, the effect of combined treatment on SBP was driven mainly by enalapril.

As the antifibrotic effects of febuxostat were prominent in this study, we embarked on *in vitro* studies in renal tubular cells to further evaluate these actions in a more controlled environment. In accord with *in vivo* observations, febuxostat attenuated TGF- β -stimulated production of CTGF, fibronectin and α -SMA also in NRK cells. The results were reproduced in cells exposed to the RAGE ligand S100B selected as another component of the diabetic milieu. These findings suggest that febuxostat interferes with general mechanisms of fibroproduction in renal cells. Of note, these experiments were conducted in the UA-free environment, suggesting that febuxostat can exert beneficial effects independently of changes in extracellular UA levels. We cannot exclude that these effects are mediated by suppression of intracellular UA. The importance of the reduction of intracellular UA has been previously emphasized (Johnson *et al.*, 2013b).

Interestingly, febuxostat did not attenuate TGF- β -stimulated Smad3 phosphorylation in NRK cells. This lack of effect on Smad3 was then also detected in renal cortical homogenates from ZO-FBX rats contrasting with the *in vivo* effect of enalapril. Akt has been implicated in Smad-independent TGF- β signalling (Lan and Du, 2014; Park *et al.*, 2013), being involved in a number of pathophysiological processes in the diabetic kidney (Zdychova and Komers, 2005), including growth and fibrosis (Mariappan *et al.*, 2007). Our previous studies in ZO rats (Zdychova *et al.*, 2008) as well as studies by others in models of DM2 (Feliars *et al.*, 2001) have shown enhanced renal Akt activity. In addition, Akt operates downstream of RAGE and can be stimulated with S100B (Arcuri *et al.*, 2005; Leclerc *et al.*, 2007). In contrast to Smad3, febuxostat attenuated TGF- β -induced Akt phosphorylation in NRK cells. Similar to TGF- β stimulation, S100B led to marked Akt activation in NRK cells, which was attenuated by the pretreatment with febuxostat. This effect of febuxostat on Akt was in parallel observed in renal tissue *in vivo* in rats and contrasted with the lack of effect of enalapril. These data suggest another mechanism of febuxostat's action mediated via inhibition of renal Akt activation. Disparate effects on Smad3 and Akt activation represent the major difference between the actions of febuxostat and enalapril detected in this study and may help explain additive effects of both agents in the diabetic kidney.

Several caveats of this study should be pointed out. Quantitative analyses of protein/mRNA expression were performed in renal cortical homogenates and consequently do not provide renal cell- or compartment-specific information.

Moreover, beneficial effects of febuxostat on GS and proteinuria and possibly in renal endothelia combine molecular actions with a haemodynamic component (Sanchez-Lozada *et al.*, 2008d) and require appropriate study design involving measurements of glomerular haemodynamics.

In summary, treatment with febuxostat has long-term nephroprotective effects in experimental type 2 DN. Moreover, febuxostat enhances the protective effects of RAS inhibitors. Multiple mechanisms might contribute to the beneficial actions of febuxostat in the diabetic kidney, including those related to a reduction in serum UA levels and attenuation of renal peroxynitrite formation. Moreover, febuxostat possesses antifibrotic effects *in vitro* independently of changes in extracellular UA levels and attributable to inhibition of profibrotic signalling via Akt. The findings in this study strongly support the rationale for a large clinical trial evaluating XO_i in DN. New information in this field could be expected after the completion of the ongoing PERL trial (Maahs *et al.*, 2013) in type 1 diabetic patients with nephropathy.

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Author contributions

R.K. conceived the studies, planned the experiments, analysed the data, prepared the manuscript and performed image analyses. B.X. performed cell studies, rtPCR, western blotting, IHC and HPLC. T.T.O. was responsible for animal care and *in vivo* measurements as well as for selected western blot analyses and ELISA. J.S. contributed selected western blot measurements and IHC and to data analysis preparation of the MS.

Conflict of interest

The authors declare no conflicts of interest.

Declaration of transparency and scientific rigour

This Declaration acknowledges that this paper adheres to the principles for transparent reporting and scientific rigour of preclinical research recommended by funding agencies, publishers and other organisations engaged with supporting research.

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Supporting Information

Additional Supporting Information may be found in the online version of this article at the publisher's web-site:

<http://dx.doi.org/10.1111/bph.13527>

Figure S1 Effect of FBX, ENL and their combination on SBP. ^a $P < 0.05$ versus ZL-VE; ^b $P < 0.05$ versus ZO-VE; ^c $P < 0.05$ versus ZO-FBX and/or ZO-FBXlate; ^d $P < 0.05$ versus ZO-ENL.