ARTICLE

Deletion of the gene for adiponectin accelerates diabetic nephropathy in the $Ins2^{+/C96Y}$ mouse

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Abstract

Aims/hypothesis Diabetic nephropathy is one of the most common forms of chronic kidney disease. The role of adiponectin in the development of diabetic nephropathy has not been elucidated, and the aim of the present study was to investigate the hypothesis that deletion of the gene for adiponectin would accelerate diabetic nephropathy in the Akita mouse.

Methods We followed four groups of mice from 4 weeks to 16 weeks of age ($n \ge 10$ in each group): wild-type (WT) ($Ins2^{+/+}Adipoq^{+/+}$) mice; APN^{-/-} ($Ins2^{+/+}Adipoq^{-/-}$) mice; Akita ($Ins2^{+/C96Y}Adipoq^{+/+}$) mice and Akita/APN^{-/-} ($Ins2^{+/C96Y}Adipoq^{-/-}$) mice. The mice were then killed and diabetic kidney injury was assessed. In vitro experiments were performed in primary mesangial cells.

Results Mice from both diabetic groups exhibited increased glomerular adiponectin receptor 1 (adipoR1) expression, kidney hypertrophy, glomerular enlargement, increased

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albuminuria and tissue oxidative stress compared with the WT control. Deletion of the adiponectin gene had no effect on glycaemia. However, Akita/APN^{-/-} mice exhibited a greater extent of renal hypertrophy. In vitro, adiponectin attenuated high-glucose-induced phosphorylation of mammalian target of rapamycin (mTOR) and ribosomal protein S6 kinase (S6K). A higher level of fibrosis was observed in the tubulointerstitial and glomerular compartments of the Akita/APN^{-/-} mice and adiponectin was found to inhibit TGFβ-induced Smad2 and Smad3 phosphorylation in vitro. There was an exaggerated inflammatory response in the Akita/APN^{-/-} mice. Adiponectin also inhibited high-glucose-induced activation of nuclear factor κB (NFκB) in mesangial cells.

Conclusions/interpretation Our data suggest that adiponectin is an important determinant of the kidney response to high glucose in vivo and in vitro.

Keywords Adiponectin · Diabetic nephropathy · High glucose · Kidney hypertrophy

Abbreviations

Abbieviations				
adipoR1	Adiponectin receptor 1			
adipoR2	Adiponectin receptor 2			
AMPK	AMP-activated protein kinase			
ColIV	Collagen IV			
DHE	Dihydroethidium			
MCP-1	Monocyte chemotactic protein-1			
MDA	Malondialdehyde			
mTOR	Mammalian target of rapamycin			
ME _E B	Nuclear factor KB			

NFκB Nuclear factor κB NO Nitric oxide

PAS Periodic acid–Schiff ROS Reactive oxygen species



αSMA α Smooth muscle actin S6K Ribosomal protein S6 kinase UAE Urinary albumin excretion

WT Wild-type WT-1 Wilm's tumour-1

Introduction

Diabetic nephropathy is a major form of chronic kidney disease leading to end-stage renal failure [1–3]. Much effort has gone into studying mechanisms by which hyperglycaemia causes chronic kidney injury [4, 5], but precise roles for factors involved in the pathogenesis of diabetic nephropathy are yet to be completely elucidated. Recent studies have shown that adiponectin may protect against mechanisms implicated in diabetic injury, including vascular injury [6, 7]. The goal of the current study was to define the role of adiponectin in the development of diabetic kidney injury.

Adiponectin is a 30 kDa hormone secreted predominantly by adipose tissue and has insulin-augmenting effects [8]. It has an impact on several signalling cascades, including the AMP-activated protein kinase (AMPK), peroxisome proliferator-activated receptor α , mitogen-activated protein kinase, Akt and cAMP pathways [9–11], after engagement of the two major receptors, adiponectin receptor 1 (adipoR1) and 2 (adipoR2) [12]. Adiponectin can also bind with T-cadherin to influence cellular responses, especially in cardiac tissue [13].

Previous studies have demonstrated that adiponectin may protect against diabetic injury by improving insulin sensitivity: replenishing adiponectin in obese mice attenuated insulin resistance and hyperglycaemia [14]. Additionally, mutant mice lacking adiponectin developed greater endothelial cell dysfunction in response to a high-fat diet, while the presence of adiponectin suppressed endothelial expression of adhesion molecules in response to TNF α stimulation [15, 16].

Adiponectin has also been found to influence kidney function. Sharma and colleagues identified a critical role of adiponectin in maintaining podocyte health that was dependent on inhibition of oxidative stress [17]. Pharmacological activation of AMPK, a major downstream signal mediator of adiponectin, was shown to suppress kidney H₂O₂ and monocyte chemotactic protein-1 (MCP-1) in mice on a high-fat diet [18]. However, the role of adiponectin in the development of mesangial expansion and glomerular hypertrophy has not been fully elucidated. We hypothesise that adiponectin attenuates hyperglycaemic kidney injury. To test this hypothesis, we crossed $Ins2^{+/-}Adipoq^{-/-}$ (APN---) mice with the type 1 diabetic $Ins2^{+/-}O6Y$ $Adipoq^{+/-}$ (Akita) mice to create a line of double-mutant $Ins2^{+/-}O6Y$ $Adipoq^{-/-}$ (Akita/APN---) mice and examined the pathological changes in the kidneys of these mice.

Methods

Animals Ins2^{+/+}Adipoq^{+/+} (wild-type [WT]) and Akita mice were purchased from The Jackson Laboratory (Bar Harbor, ME, USA). APN^{-/-} mice were generously donated by S. Verma (Department of Surgery, University of Toronto, Toronto, ON, Canada) and were crossed with the Akita mice to generate the Akita/APN^{-/-} mice. All mice were maintained on a C57BL/6 background, and were housed at the Division of Comparative Medicine at University of Toronto, with standard chow diet and free access to food and water. Male mice were followed from 4 to 16 weeks of age, with weekly blood glucose and body weight recordings. Twenty-four hour urine samples were obtained at 16 weeks of age and mice were then killed to extract plasma and kidney tissue. HbA_{1c} levels were measured at the time of killing. See electronic supplementary material (ESM) Methods for further details. All experiments were conducted under the guidelines of the University of Toronto Animal Care Committee.

Cells Human renal mesangial cells were purchased from Lonza (Clonetics, Lonza, Walkersville, MD, USA) and cultured as previously described [19]. Cells were maintained at 37°C in 5% CO₂–95% air and used at passages 4–7.

Histology and immunohistochemistry Mesangial matrix expansion and glomerular volume were measured on periodic acid–Schiff (PAS)-stained sections. Primary antibodies against adipoR1, adipoR2, α smooth muscle actin (α SMA), collagen IV (ColIV), F4/80 and Wilm's tumour-1 (WT-1) were used for immunohistochemistry. Positivity for adipoR1, ColIV and WT-1 staining was assessed in glomeruli, α SMA staining was assessed on whole-kidney sections excluding the large blood vessels and F4/80 staining was assessed on whole-kidney sections. Quantification of PAS-positive staining and immunohistochemistry was completed using the Aperio ImageScope software (Leica Microsystems, Concord, ON, Canada). See ESM Methods for further details.

ELISA and colorimetric assays ELISA kits were used to measure urinary levels of albumin, nitric oxide (NO), malondialdehyde (MDA) and nephrin, as well as the tissue level of MCP-1 in the kidney. Urinary albumin levels were used to calculate the 24 h urinary albumin excretion (UAE) rate. Levels of NO and MDA in urine were normalised to urinary creatinine concentration, while the level of MCP-1 in the kidney was normalised to total protein concentrations. See ESM Methods for further details.

Western blot Primary antibodies against the following proteins were used for immunoblot: phospho- and total mammalian target of rapamycin (mTOR); phospho- and total p70 ribosomal protein S6 kinase (S6K); phospho- and total



Smad2; phospho- and total Smad3 and β -actin. Mesangial cells were stimulated with high glucose (30 mmol/l, 1 h) or TGF- β (2 ng/ml, 1 h) with or without adiponectin (100 ng/ml, 30 min pre-incubation), then collected and lysed. Mannitol was used as an osmotic control for high-glucose experiments. Protein concentrations were determined by the Bradford method. Scion Image software (Scion, release beta 4.0.2, Frederick, MD, USA) was used for densitometry calculation. See ESM Methods for further details.

NFκB activity assay Mesangial cells were co-transfected with pNFκB-Luc plasmid and pRL-TK reporter vector, then treated with high glucose or TNF α (20 ng/ml), with or without adiponectin (100 ng/ml) or apocynin (100 μmol/l), for 24 h. Luciferase activity was determined with a spectrometer. Total protein measured with Bradford assay was used to normalise activity results. See ESM Methods for further details.

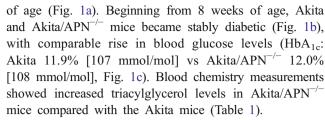
Dihydroethidium stain Mesangial cells cultured on glass cover slips were incubated for 18 h in normal glucose or high glucose with or without adiponectin. Cells were then washed with PBS and stained with dihydroethidium (DHE; 2 μmol/l, 1 h at 37°C in the dark). After washing, cells were mounted onto glass slides and visualised under a confocal microscope. At least 12 high-power fields were captured for semiquantitative scoring of fluorescent intensity. See ESM Methods for further details.

Lucigenin enhanced chemiluminescence assay NADPH oxidase activity was measured by lucigenin enhanced chemiluminescence assay as previously described [20]. Cultured mesangial cells were treated under the following conditions for 18 h before collection: normal glucose; high glucose with or without adiponectin (pretreatment for 1 h); specific antibodies against adipoR1 (10 μ g/ml, pretreatment for 1 h) and adipoR2 (10 μ g/ml, pretreatment for 1 h) and non-specific goat IgG. Mannitol was used as an osmotic control. Cell lysates were mixed with NADPH (100 μ mol/l) and lucigenin (5 μ mol/l). Peak light emission was recorded and normalised to the total protein concentration as determined by Bradford assay. See ESM Methods for further details.

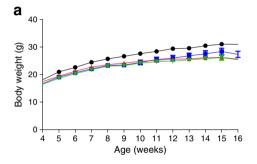
Statistical analysis Results are shown as mean \pm SE. GraphPad Prism 5 (GraphPad Software, La Jolla, CA, USA) was used for statistical analyses. For multiple-group comparison, one-way ANOVA with Bonferroni's post hoc test was performed and for two-group comparison, unpaired t test was performed. Statistical significance was set at $p \le 0.05$.

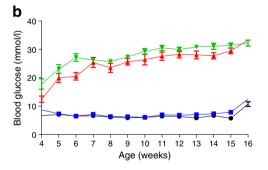
Results

Whole-animal data All four groups of experimental mice exhibited an increase in body weight between 4 and 16 weeks



Immunohistochemistry for adipoR1 showed the most intense staining in glomeruli, with positive signals in both podocytes and mesangia (Fig. 2a–d, i). Staining was increased in the glomeruli of diabetic mice (Fig. 2k). Deletion of the gene for adiponectin did not alter baseline glomerular adipoR1, but caused a trend towards more glomerular adipoR1-positive areas in diabetic mice. Positive staining for adipoR1 could also be identified in the proximal tubules. No





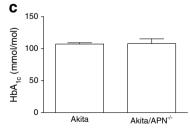


Fig. 1 (**a**, **b**) Changes in body weight (**a**) and blood glucose (**b**) levels were followed in the WT (circles), APN^{-/-} (squares), Akita (triangles) and Akita/APN^{-/-} (inverted triangles) mice weekly from 4 weeks to 16 weeks of age. (**c**) HbA_{1c} levels in diabetic mice were measured at the time of death. At least six mice were used in each experimental group at any given age for body weight and blood glucose assessments. For HbA_{1c}, *n*=3 for Akita mice, *n*=5 for Akita/APN^{-/-} mice. To convert values for HbA_{1c} in mmol/mol to %, use the equation: HbA_{1c} (%)=(0.09148×HbA_{1c} [mmol/mol])+2.152



Table 1 Whole-animal data

Variable	WT	APN ^{-/-}	Akita	Akita/APN ^{-/-}
Blood glucose (mmol/l)	8.0±0.60	10.3±0.62	30.7±0.67*	29.9±1.02*
Body weight (g)	29.4 ± 0.63	26.5±0.73*	26.5±0.44*	$23.7 \pm 0.60*^{\dagger}$
Kidney weight (g)	0.17 ± 0.003	0.18 ± 0.005	0.27±0.009*	$0.30 {\pm} 0.008*^{\dagger}$
Cholesterol (mmol/l)	3.0 ± 0.16	2.3±0.04*	2.2±0.12*	2.5±0.12*
Triacylglycerol (mmol/l)	0.85 ± 0.079	0.61 ± 0.040	1.1 ± 0.10	1.7±0.21* [†]
Urea (mmol/l)	17.4±1.49	16.1±1.13*	21.6±0.96*	21.4±0.78*
Creatinine (µmol/l)	17±1.5	16±1.1	22±1.0	21 ± 0.8
Sodium (mmol/l)	146±1.4	146±1.6	138±1.5*	141 ± 0.627
Potassium (mmol/l)	5.0±0.17	5.2±0.13	4.7±0.15	4.6±0.19
Chloride (mmol/l)	$116{\pm}0.8$	114 ± 0.6	$105 \pm 1.6*$	103±1.5*

Results are presented at mean±SE

Blood glucose was measured at 16 weeks of age before tissue harvest. Body weight and kidney weight (average of left and right kidney) were recorded at the time of killing. Concentrations of cholesterol, triacylglycerol, urea, creatinine, sodium, potassium and chloride were determined in plasma samples. At least 19 mice in each group were used to derive data for blood glucose, body weight and kidney weight and eight mice in each group were used for blood chemistry measurements

*p<0.05 vs WT; †p<0.05 vs Akita

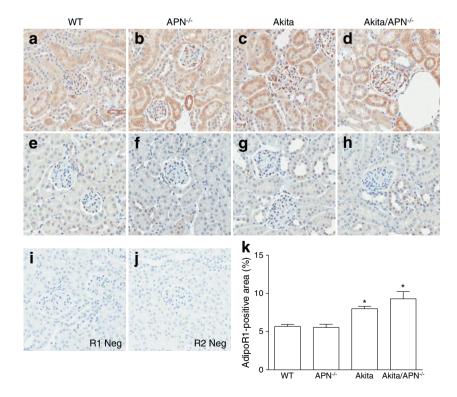
reliable staining for adipoR2 was detected in the kidney sections of our experimental mice (Fig. 2e-h, j).

Assessment of kidney hypertrophy Kidney weight was increased in the Akita mice and further in the Akita/APN^{-/-} mice (Fig. 3a, b). This was accompanied by even greater increases in kidney-to-body weight ratio (Fig. 3c). In agreement with increased kidney weight, mean glomerular volume was increased in the diabetic Akita and Akita/APN^{-/-} mice

compared with WT mice (Fig. 3d). The mean size of glomeruli was not significantly different between the two diabetic groups despite a higher mean glomerular volume in the Akita/APN^{-/-} mice.

To identify the mechanism(s) responsible for kidney hypertrophy we studied the effect of adiponectin on high-glucose-induced mTOR activation in mesangial cells. The dose of adiponectin was shown to be effective previously in cultured proximal tubular cells [20], and again verified in cultured

Fig. 2 Expression of adiponectin receptors in kidney. (a-h) Representative images for adipoR1 (a-d) and adipoR2 (e-h) immunohistochemical stainings are shown. (i, j) Sections incubated without primary antibodies were used as negative controls for adipoR1 (R1 Neg) and adipoR2 (R2 Neg). Magnification, ×200. (k) The percentage of adipoR1-positive area in glomeruli was computed using ImageScope software. Results are presented as mean \pm SE, $n \ge 9$ for each group. *p<0.05 vs WT mice





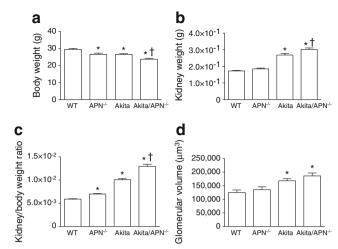


Fig. 3 Body weight (a), kidney weight (b), kidney/body weight ratio (c) and glomerular volume (d) were assessed in 16-week-old mice. Glomerular volumes were calculated from mean glomerular areas in PAS-stained sections. Data are shown as mean \pm SE. $n\geq$ 19 for body weight, kidney weight and kidney/body weight ratio in each experimental group; $n\geq$ 8 for glomerular volume in each experimental group. *p<0.05 vs WT mice; p<0.05 vs Akita mice

mesangial cells (ESM Fig. 1). Phosphorylation of mTOR and S6K was increased by high glucose and suppressed by co-incubation with adiponectin (Fig. 4). The high-glucose-induced mTOR phosphorylation diminished to baseline after 24 h, while a non-significant increase in S6K phosphorylation was observed (ESM Fig. 2). Adiponectin was able to suppress the phosphorylation of S6K after incubation for 24 h. Increased osmotic pressure yielded no change in activation of mTOR signalling (ESM Fig. 3).

Detection of renal fibrosis Kidney pathology was first examined at the microscopic structural level by PAS stain. The UAE rate was used for functional assessment (Fig. 5). Akita mice showed mesangial expansion (as quantified by PAS-positive area in glomerular profiles; Fig. 5a–d) that was exacerbated by

the loss of the adiponectin gene (Fig. 5e). A similar pattern was seen for albuminuria (Fig. 5f). However the difference in UAE between the Akita and the Akita/APN $^{-/-}$ mice did not reach statistical significance by one-way ANOVA analysis of the four experimental groups, possibly due to the limited time of observation in the current study. Fibrosis within the tubulointerstitial and glomerular compartments was further assessed by immunohistochemical staining for αSMA (Fig. 6a–d) and ColIV (Fig. 6e–h), respectively. Similar trends emerged for quantification of both αSMA -positive and glomerular ColIV-positive areas, which were significantly increased in Akita/APN $^{-/-}$ mice compared with WT mice (Fig. 6i, j). Furthermore, for ColIV, the difference was significantly greater when comparing Akita/APN $^{-/-}$ mice with Akita mice.

To investigate the molecular mechanism(s) responsible for the role of adiponectin in fibrosis, mesangial cells were treated with TGF- $\beta1$ with or without adiponectin. Phosphorylation of Smad2 and Smad3 was strongly stimulated by TGF- $\beta1$ treatment, and adiponectin partially blocked this activation (Fig. 7). Treatment with adiponectin did not affect the baseline Smad2 and Smad3 activity (ESM Fig. 4).

Pathological changes in inflammation Infiltrating macrophages based on F4/80 staining (Fig. 8a–d) were significantly increased only in Akita/APN^{-/-} mice compared with controls; Akita mice showed a trend towards increased macrophage infiltration when compared with WT mice (Fig. 8e). In agreement with F4/80 immunohistochemistry, tissue MCP-1 content was significantly higher only in the kidneys of Akita/APN^{-/-} mice when compared with the WT mice, with a similar trend of increase in the Akita mice (Fig. 8f).

 $NF\kappa B$ is a central mediator of inflammation and hence its activation was examined. Cultured mesangial cells were transfected with an NF κB -sensitive luciferase reporter construct and treated with high glucose with or without

Fig. 4 Adiponectin prevented high-glucose-induced activation of the mTOR signalling pathway in human mesangial cells. Cells were treated with the respective medium and collected for immunoblotting against phosphoand total mTOR (a, c) and S6K (b, d). NG, normal glucose (5.6 mmol/l); HG, high glucose (30 mmol/l); HA, high glucose with adiponectin (100 ng/ml). Results are presented as mean \pm SE, n=3 in each group. *p<0.05 vs normal glucose: $^{\dagger}p$ <0.05 vs high glucose

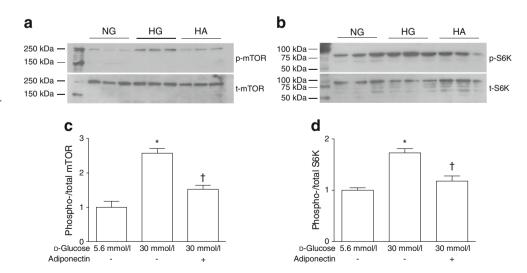
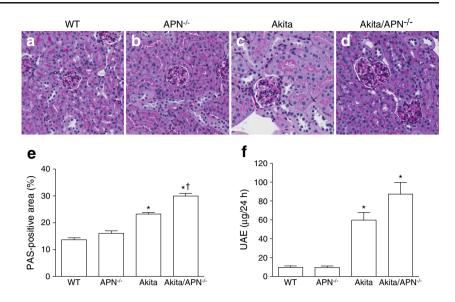




Fig. 5 Structural and functional changes in kidneys of experimental mice. (a–d) Representative images for PAS staining of kidney sections. Magnification, $\times 200$. (e) Quantification of PAS-positive area in glomeruli. (f) Quantification of UAE rates in 16-week-old mice. Results are presented as mean \pm SE. For PAS-positive area, n=10 in each group; for UAE, $n\geq 6$ for each experimental group. *p<0.05 vs WT mice; $^{\dagger}p<0.05$ vs Akita mice

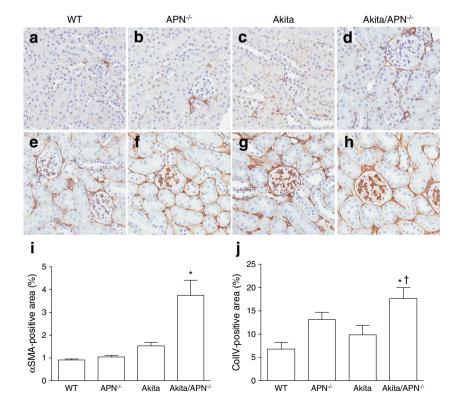


adiponectin. High glucose strongly activated NF κ B in mesangial cells, and adiponectin reversed this effect (Fig. 9a). Reactive oxygen species (ROS) have been implicated in mediating many injurious effects of diabetes. In support of this, co-treatment with the antioxidant apocynin inhibited the high-glucose-induced NF κ B activation (Fig. 9a). To test whether the inhibitory effect of adiponectin on NF κ B activation was stimulus specific in cultured mesangial cells, the potential interaction between adiponectin and TNF α , a known

activator of NF κ B signalling, was investigated. Incubation with adiponectin caused a small, yet significant, reduction in TNF α -induced NF κ B activation (Fig. 9b).

Evaluation of oxidative stress To further study the role of adiponectin in oxidative stress in the kidney in response to high glucose, the excretion of MDA and NO was quantified in 24 h urine samples. Diabetic mice exhibited higher levels of urinary NO (Fig. 10a) and MDA (Fig. 10b). Despite a higher

Fig. 6 (a–h) Immunohistochemical staining for α SMA (a–d) and ColIV (e–h) on renal sections. Magnification, $\times 200$. (i, j) Quantification of α SMA-positive (i) and glomerular ColIV-positive areas (j). Results are presented as mean \pm SE, $n\geq 6$ for each experimental group. *p<0.05 vs WT mice; $^{\dagger}p<0.05$ vs Akita mice





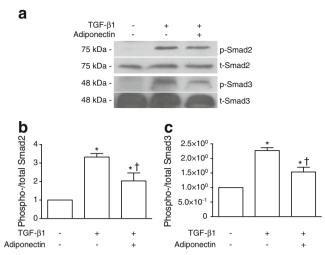


Fig. 7 Adiponectin partially inhibited phosphorylation of Smad2 and Smad3 following TGF- β 1 stimulation in mesangial cells. Representative images of immunoblot (a) and the densitometry for Smad2 (b) and Smad3 (c) are shown. Results are presented as mean \pm SE. n=4 for Smad2 immunoblot; n=3 for Smad3 immunoblot. *p<0.05 vs control; $^{\dagger}p$ <0.05 vs TGF- β 1 treatment

mean value, the levels of MDA and NO in Akita/APN^{-/-} mice failed to reach statistical significance when compared with the levels in Akita mice.

The direct effect of adiponectin on cellular oxidative stress in response to high glucose was then studied in mesangial cells, with DHE stain as a marker for superoxide generation. High glucose raised ROS levels in mesangial cells, and coincubation with adiponectin reduced this effect (Fig. 11a–d). In addition, adiponectin was able to inhibit the elevation in NADPH oxidase activity in mesangial cells exposed to high glucose (Fig. 11e). Stimulation of NADPH oxidase activity was not seen when mannitol was added as an osmotic control (ESM Fig. 5). As shown in vivo, adipoR1 appeared to be the

dominant receptor for adiponectin in the kidney in our model (Fig. 2). Similarly in vitro, the suppression of high-glucose-induced NADPH oxidase activation by adiponectin was primarily mediated by adipoR1 in cultured mesangial cells, as demonstrated by the effect of blocking antibodies (ESM Fig. 6).

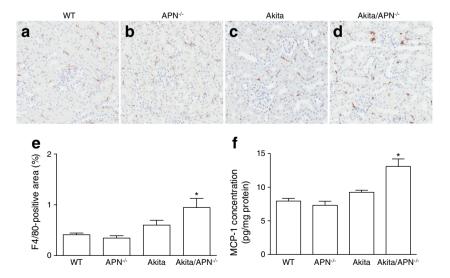
Since it has been reported that the antioxidative effect of adiponectin could also contribute to a protective effect on podocytes [17], we evaluated podocyte numbers by quantifying WT-1-positive cells and urinary nephrin excretion (ESM Fig. 7). Neither the number of podocytes nor urinary nephrin excretion was significantly altered (ESM Fig. 7b, c), suggesting that pathological changes in podocytes were limited in our experimental mice at this stage of disease development.

Discussion

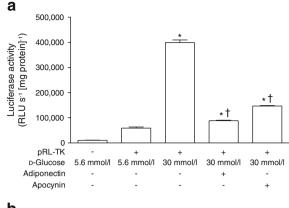
Adiponectin has been characterised as an insulin-sensitising adipokine that promotes glucose utilisation and lipid metabolism. In pancreatic beta cells, adiponectin prevents lipid-induced apoptosis by activating sphingosine-1-phosphate generation [21], as well as beta cell death from combined IL-1 and IFN- γ stimulation [22]. In the cardiovascular system, overexpression of adiponectin protects against atherosclerosis, while adiponectin deficiency exacerbates cardiac ischaemia—reperfusion injury [23, 24]. The administration of adiponectin as well as deletion of the gene for adiponectin have been reported to attenuate ischaemia—reperfusion-induced inflammation and apoptosis in the kidney [25, 26].

Sharma and colleagues were the first to establish a critical role for adiponectin in podocyte health [17]. These authors showed that adiponectin-null mice developed albuminuria and podocyte foot process effacement. Urinary measurements of oxidative stress and albuminuria were exacerbated by

Fig. 8 (a–d) Immunohistochemical staining for F4/80 on kidney sections (a–d) and the corresponding quantification (e) are shown. Magnification $\times 200$. (f) Measurement of tissue MCP-1 concentration in kidney. Results are presented as mean \pm SE. For F4/80, $n{\ge}8$ for each group; for MCP-1 measurement, $n{\ge}4$ for each group. *p<0.05 vs WT







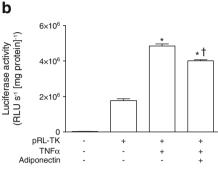


Fig. 9 (a) High-glucose-induced NFkB activation in mesangial cells was inhibited by adiponectin. Co-incubation with apocynin also suppressed the NFkB activity after high-glucose treatment. (b) Adiponectin induced a modest reduction in NFκB activity after TNFα stimulation. Vectortransfected cells cultured in regular medium were used as controls. Results are shown as mean \pm SE, n=3 in each group. *p<0.05 vs controls; p < 0.05 vs high glucose (30 mmol/l) or TNF α treatment. RLU, relative light units

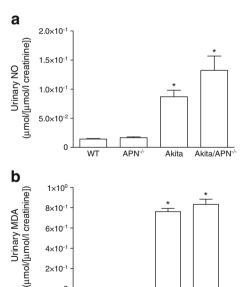


Fig. 10 Concentration of urinary oxidative stress markers, NO (a) and MDA (b), were quantified by colorimetric assays and normalised by urinary creatinine levels. Results are shown as mean \pm SE, $n \ge 6$ for each experimental group. *p<0.05 vs WT mice

APN-/

Akita Akita/APN

WT

2×10-

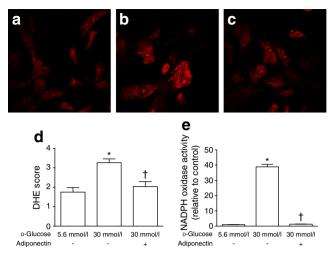


Fig. 11 Adiponectin reversed the increase in high-glucose-induced oxidative stress in mesangial cells. (a-c) Representative images of DHE stains in cells treated with normal glucose (5.6 mmol/l, a), high glucose (30 mmol/l, b) and high glucose with adiponectin (c). Magnification, ×400. (d) Semiquantitative score for DHE intensity. (e) NADPH oxidase activity measurement in mesangial cells. Results are presented as mean± SE. At least 12 images were used for DHE score; for NADPH oxidase activity assay, $n \ge 4$ for each group. *p < 0.05 vs normal glucose; †p < 0.05vs high glucose

streptozotocin-induced diabetes mellitus. They demonstrated that adiponectin activated AMPK in cultured podocytes, maintained cell-cell contact and prevented high-glucoseinduced changes in podocyte monolayer permeability. Adiponectin also prevented high-glucose-induced increases in NADPH oxidase 4 (Nox4) expression in cultured podocytes. Given these observations, we sought to more fully characterise the effect of deletion of the gene for adiponectin on the diabetic kidney. We also sought to extend our understanding of the impact of adiponectin on signal transduction in the mesangial cell. We selected the Akita mouse for our in vivo studies because it spontaneously develops profound but stable hyperglycaemia uncomplicated by the hormonal changes that accompany obesity. Our findings suggest that adiponectin plays a broad protective role in kidney responses to high glucose.

First we found that deletion of the adiponectin gene did not have a significant effect on weight gain or blood glucose in diabetic Akita mice. We next examined the effect of hyperglycaemia on adiponectin receptors. Based on our immunohistochemical analyses, adipoR1 appeared to be the dominant receptor. AdipoR1 was expressed in glomeruli, the primary target of diabetic injury, and its expression was increased in diabetic mice. In cultured mesangial cells, adipoR1 also predominantly mediated the inhibitory effect of adiponectin on high-glucose-induced NADPH oxidase activation. These results are consistent with previous findings of adiponectin receptor expression patterns in tissues and in kidney cell culture [12, 17, 20]. Interestingly, in studies using streptozotocin-induced diabetic rats and type 2 diabetic db/

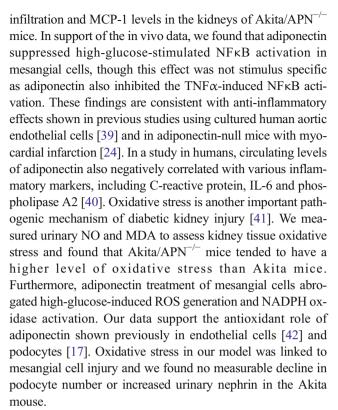


db mice, the expression of adipoR1 mRNA and protein was decreased in whole renal cortical tissue [27, 28]. These differences may be a consequence of the specific disease models but also suggest that adipoR1 expression may be regulated differentially in the glomerular and tubulointerstitial compartments of the kidney.

Although deletion of the adiponectin gene had no effect on blood glucose levels, we found that deletion led to exaggerated kidney hypertrophy in diabetic mice. Mesangial cells are important targets in diabetes [29], and we elected to study the effect of adiponectin on signal transduction pathways linked to cell hypertrophy, specifically the activation of mTOR by high glucose [30]. Previous studies in vascular smooth muscle cells [31] and in mesangial cells treated with platelet-derived growth factor [32] have demonstrated an interaction between adiponectin and this signalling pathway via AMPK. In accordance with these observations, our findings in mesangial cells showed that adiponectin attenuated high-glucose-induced mTOR and S6K activation, suggesting that mTOR pathway activation potentially mediates the exaggerated renal hypertrophic effect we observed in the diabetic Akita/APN^{-/-} mice.

In addition to hypertrophy, long-standing hyperglycaemia is associated with fibrosis in the glomerular and tubulointerstitial compartments of the kidney. We found that glomerular matrix expansion was significantly increased in Akita/APN^{-/-} mice compared with Akita mice, and there was a trend towards higher UAE. Consistent with the findings of renal hypertrophy and the structural and functional changes, the accumulation of aSMA and ColIV increased in the kidneys of Akita/APN^{-/-} mice compared with WT mice. Our findings suggest that lack of adiponectin could accelerate fibrotic changes in the diabetic kidney. To better understand the effect of adiponectin on high-glucose-induced kidney fibrosis, we studied the effect of adiponectin on TGF-β signal transduction, because TGF-β has been linked to the expression of both α SMA and ColIV in the diabetic kidney [33, 34]. We observed that adiponectin reduced the phosphorylation of Smad2 and Smad3 in mesangial cells following TGF-ß stimulation. Our results are in agreement with previous findings that adiponectin-null mice were susceptible to kidney fibrosis after subtotal nephrectomy and that administration of adiponectin ameliorated fibrosis in streptozotocin-induced diabetic rats [35, 36]. In contrast, Yang and colleagues discovered that adiponectin-null mice displayed less accumulation of matrix proteins in the kidney after both unilateral ureteral obstruction and renal ischaemia-reperfusion injury [37]. Adiponectin was able to stimulate monocyte-fibroblast differentiation in their model and thus promote renal fibrosis. Therefore, the mechanism by which adiponectin affects fibrotic changes in the kidney may be stimulus- and cell-typespecific.

Inflammation is a critical contributor to diabetic kidney injury [38]. We observed a marked increase in macrophage



The current study has some important limitations. First, although several groups have used a 16 week time point for analyses of diabetic nephropathy, the changes in kidney structure and function are relatively modest on the C57BL/6 background [43–45]. This may have limited our ability to detect a significant difference in the UAE. Second, although WT-1 and nephrin excretion have been commonly used to assess podocyte injury [46–48], the lack of changes in these markers does not exclude a role for the podocyte in the phenotypical changes that we observed in our mice.

In summary, we found that deletion of the adiponectin gene exacerbated diabetes-induced kidney hypertrophy, fibrosis and inflammation. In vitro, adiponectin attenuated high-glucose-induced mTOR, TGF- β and NF κ B signalling in glomerular mesangial cells. Together, this work supports the hypothesis that adiponectin is a protective factor in the kidney. Augmenting adiponectin signal transduction in the kidney may be an important treatment strategy in diabetic nephropathy.

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