

Article **Metformin Attenuates Hyperglycaemia-Stimulated Pro-Fibrotic Gene Expression in Adventitial Fibroblasts via Inhibition of Discoidin Domain Receptor 2**

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Abstract: Molecular mechanisms underlying the diverse therapeutic effects of anti-diabetic metformin, beyond its anti-hyperglycaemic effects, remain largely unclear. Metformin is reported to reduce the long-term complications of diabetes, including cardiovascular fibrosis and remodelling. Our recent investigations show that Discoidin Domain Receptor 2 (DDR2), a Collagen receptor tyrosine kinase, has an obligate regulatory role in Collagen type I gene expression in cardiac and vascular adventitial fibroblasts, and that it may be a molecular link between arterial fibrosis and metabolic syndrome in rhesus monkeys. Using gene knockdown and overexpression approaches, the present study examined whether DDR2 is a target of metformin and whether, by targeting DDR2, it inhibits Fibronectin and Collagen type I expression in rat aortic adventitial fibroblasts exposed to hyperglycaemic conditions. Metformin was found to attenuate hyperglycaemia-induced increase in DDR2 mRNA and protein expression by inhibiting TGF-β1/SMAD2/3 signalling that mediates the stimulatory effect of hyperglycaemia on DDR2 expression. Metformin also inhibited DDR2 dependent expression of Fibronectin and Collagen type I, indicating that it regulates these matrix proteins via DDR2 inhibition. The findings identify DDR2, a mediator of cardiovascular remodelling, as a molecular target of metformin, thereby uncovering the molecular basis of its protective role in vascular fibrosis and possibly cardiac fibrosis associated with diabetic cardiomyopathy.

Keywords: metformin; Discoidin Domain Receptor 2; vascular adventitial fibroblasts; Fibronectin; Collagen type I; hyperglycaemia; reactive oxygen species

1. Introduction

Metformin is a widely used anti-diabetic drug whose beneficial effects far exceed its anti-hyperglycaemic action and are possibly mediated by mechanisms distinct from its metabolic actions such as inhibition of mitochondrial enzymes and consequent activation of AMP-activated protein kinase [\[1,](#page-10-0)[2\]](#page-10-1). Several studies point to its protective role in cancer, polycystic ovary syndrome, and neurological disorders [\[3–](#page-10-2)[6\]](#page-10-3). Metformin is reported to exert potent anti-fibrotic effects in the lung by modulating metabolic pathways, inhibiting TGF-β1 action, suppressing Collagen production, activating PPARγ signalling,

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and inducing lipogenic differentiation in myofibroblasts [\[7\]](#page-10-4). A number of clinical studies have shown that metformin reduces the incidence of cardiovascular events and associated mortality [\[8](#page-10-5)[,9\]](#page-10-6). It has been demonstrated that metformin mitigates cardiac fibrosis induced by pressure overload and inhibits Collagen synthesis in cardiac fibroblasts via inhibition of the TGF-β1/SMAD3 signalling pathway [\[10\]](#page-10-7). Another study showed that the drug inhibits inflammatory responses through suppression of the NF-κB inflammatory signalling pathway, improves vascular endothelial function, and inhibits cardiomyocyte apoptosis and myocardial fibrosis [\[9\]](#page-10-6). Notably, metformin protects against adverse fibrotic vascular remodelling [\[9\]](#page-10-6), which is a major complication of diabetes that contributes to end-organ damage. Notwithstanding these reports, the precise mechanisms underlying the diverse therapeutic effects of metformin remain largely elusive [\[2\]](#page-10-1). Admittedly, it is of immense scientific interest and clinical relevance to identify the molecular targets of metformin and delineate the mechanisms that underlie its widely acknowledged cardiovascular protective effects, especially its anti-fibrotic action.

In this regard, several studies have shown that Discoidin Domain Receptor 2 (DDR2), a Collagen receptor tyrosine kinase, is a major regulator of a wide array of cellular processes such as extracellular matrix (ECM) turnover, proliferation, differentiation, migration, and survival in different cell types, including cardiac fibroblasts and vascular cells [\[11](#page-10-8)[–17\]](#page-11-0). Further, our recent studies provide robust evidence that DDR2 may be a molecular link between arterial fibrosis and metabolic syndrome in rhesus monkeys [\[18\]](#page-11-1). The predominant localisation of DDR2 in cells of mesenchymal origin and its indispensable role in Collagen type I production in cardiovascular cells make it a potential player and hence, a drug target in adverse cardiovascular tissue fibrosis and remodelling associated with pathological states such as diabetes and atherosclerosis.

The present study examined whether DDR2 is a target of metformin and whether, by targeting DDR2, the drug inhibits Fibronectin and Collagen type I expression in rat aortic adventitial fibroblasts exposed to hyperglycaemic conditions. We demonstrate for the first time that metformin prevents hyperglycaemia-induced increase in DDR2 expression in vascular adventitial fibroblasts by inhibiting the TGF-β1/SMAD2/3 signalling pathway, which enhances DDR2 expression in response to hyperglycaemia. Additionally, we also show that inhibition of DDR2 by metformin results in inhibition of Fibronectin and Collagen type I expression in these cells. The findings are important because they identify DDR2 as a molecular target of metformin, uncovering the molecular basis of its protective role in vascular fibrosis and possibly cardiac fibrosis associated with diabetic cardiomyopathy.

2. Results and Discussion

The incidence of diabetes mellitus is on the rise globally due to a variety of reasons, including increasing prevalence of obesity, genetic susceptibility, and aging [\[19\]](#page-11-2). Since conventional treatment modalities aiming at rigorous glycaemic control may not completely eliminate the cardiovascular risk burden, other strategies targeting critical pathogenetic mechanisms may offer hope. In this regard, it is pertinent to note that one of the hallmarks of vascular complications associated with diabetes is abnormal matrix deposition by vascular adventitial fibroblasts and vascular smooth muscle cells, leading to vascular fibrosis and remodelling of the ECM within the vascular wall [\[20\]](#page-11-3). This is integral to intimalmedial thickening, reduced lumen diameter, and compromised vascular resilience [\[9\]](#page-10-6). Hyperglycaemia-induced alterations in vascular wall structure may contribute to the pathogenesis of microvascular and macrovascular complications attributed to diabetes, resulting eventually in impairment of target organs such as the heart, kidneys, and brain. Targeting mechanisms that regulate ECM gene expression in vascular cells is clearly a promising option to prevent the cardiovascular complications of diabetes. This study focused on DDR2 as a target of metformin, examined the mechanism by which metformin inhibits DDR2, and in light of our earlier published studies, explored the consequences of DDR2 inhibition by metformin vis-a-vis matrix protein expression.

Exposure of adventitial fibroblasts to high glucose at 30 mM (HG) was found to cause a significant increase in Fibronectin (Figure [1A](#page-2-0)) and Collagen type I protein expression
(Figure 1B). HG also caused a significant increase in DDR2 mRNA and protein and protein and protein and protein (Figure [1B](#page-2-0)). HG also caused a significant increase in DDR2 mRNA and protein levels (Figure 2A, B). O (Figure [2A](#page-3-0),B). Our previous studies with mannitol at the same concentration had ruled μ out the possibility that these effects of HG may be osmolarity-related [\[18\]](#page-11-1). To ascertain $\frac{1}{100}$ whether DDR2 may have a role in HG-stimulated Fibronectin expression in adventitial
Glad described with DDR2 siRNA, and redescribed under the cells with deviating fibroblasts, the cells were transfected with DDR2 siRNA, as described under Methods, and
after confirming knockdown, Fibronectin expression was determined following expressure of after confirming knockdown, Fibronectin expression was determined following exposure of and community ancesation, I is following expression was determined following exposure of the cells to HG. DDR2 knockdown attenuated the stimulatory effect of HG on Fibronectin α expression (Figure [3\)](#page-3-1). In tandem with earlier observations [\[11–](#page-10-8)[14,](#page-10-9)[21\]](#page-11-4), the finding shows expression (rigino s). In dindem with cannot esservations [11,11], the finding shows conclusively that DDR2 has an indispensable regulatory role in the expression of Fibronectin Forefactively that be full that an induspendance regulatory recent the expression of Fibronectin and Collagen type 1 in both cardiac fibroblasts and vascular adventitial fibroblasts. The shared characteristic of a regulatory relationship between DDR2 and the two major ECM proteins in cardiac and vascular cells that express these proteins may be an important proteins may be an important of fibrosis in the cardiovascular system. Admittedly, any drug that targets DDR2 would impact the expression of these key matrix proteins in cardiovascular cells and mitigate the cardiovascular complications of diabetes, which affect the prognosis of ${\mathsf p}$ atients. Exposure of adventitial fibroblasts to high glucose at 30 mM (HG) was found to Exposure or adventural increases to figure grucose at 50 min (Fig) was found to cause

Figure 1. Metformin prevents HG-induced increase in Fibronectin and Collagen type I protein pression in vascular adventitial fibroblasts. Cells were pre-treated with metformin hydrochloride expression in vascular adventitial fibroblasts. Cells were pre-treated with metformin hydrochloride
(10 **h of HG and HG** and HG and HG and HG and HG and HG and (10 μM) for 45 min prior to HG supplementation. (**A**) Protein was isolated after 12 h of HG and subjected to western blot analysis for detection of Fibronectin, with β-Actin as loading control. **(B)** Protein was isolated after 12 h of HG and subjected to western blot analysis for detection of Collagen type I, with β-Actin as loading control. Data are expressed as mean \pm SEM (standard error of mean). The groups were analysed by one-way ANOVA ($p \leq 0.05$) and Tukey's multiple comparisons ($p \leq 0.05$). Significant comparisons along with their p values are depicted in the Figure. The data are representative of three independent experiments, $n = 3$.

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Figure 2. Metformin prevents HG-induced increase in DDR2 expression in vascular adventitial fibroblasts. Cells were pre-treated with metformin hydrochloride (10 μ M), 45 min prior to HG supplementation. (A) DDR2 mRNA levels were determined by Taqman real-time PCR at 6 h post-HG treatment, with β -Actin as loading control. (B) Protein was isolated after 12 h of HG treatment and expressed to overtext blot as classic for detection of DDP2, with θ , Astin as leading control. Det subjected to western blot analysis for detection of DDR2, with β -Actin as loading control. Data are expressed as mean \pm SEM (standard error of mean). The groups were analysed by one-way ANOVA $(p \le 0.05)$ and Tukey's multiple comparisons ($p \le 0.05$). Significant comparisons along with their p values are depicted in the Figure. The data are representative of three independent experiments, $n = 3$. b_{max} becomes the second with metroman hydrochloride (10 μ *M), 15 min pri* $\varphi \leq 0.05$ and Tukey's mample comparisons $\varphi \leq 0.05$. Significant comparisons along φ

Figure 3. DDR2 knockdown prevents HG-induced increase in Fibronectin expression in vascular adventitial fibroblasts. Cells were transiently transfected with DDR2 siRNA (5 pmol) (indicated by $\frac{1}{2}$ $(+)$ sign) or scrambled siRNA (5 pmol) (indicated by $(-)$ sign). Following exposure of the transfected cells to HG for 12 h, Fibronectin protein expression was examined by western blot analysis and normalized to β -Actin. 3,3'-diaminobenzidine (DAB) reagent was used for detection of β -Actin, adventitial fibroblasts. Cells were transiently transfected with DDR2 siRNA (5 pmol) (indicated by and brightness/contrast was adjusted uniformly to match with other blots. Data are expressed as mean \pm SEM (standard error of mean). The groups were analysed by one-way ANOVA ($p \leq 0.05$) and Tukey's multiple comparisons ($p \leq 0.05$). Significant comparisons along with their *p* values are depicted in the Figure. The data are representative of three independent experiments, $n = 3$.

Concurrent treatment of vascular adventitial fibroblasts with metformin at $10 \mu M$ was found to significantly reduce HG-induced increase in Fibronectin and Collagen type I expression in these cells (Figure [1A](#page-2-0),B). At this concentration, metformin did not affect cell viability, which is consistent with what has been previously reported [\[22\]](#page-11-5). Moreover, plasma concentrations of metformin in patients on standard clinical doses of the drug $(1.5-2$ g per day) are reported to be within 5 to 30 μ M [\[23\]](#page-11-6). Notably, metformin also reduced DDR2 expression in HG-treated cells (Figure [2A](#page-3-0),B), indicating that it may regulate Fibronectin and Collagen type I via a DDR2-dependent mechanism in vascular adventitial fibroblasts. Both Collagen type I and Fibronectin are importantly involved in the regulation of cardiovascular pathophysiology. Although earlier studies focused predominantly on Collagen type I, Fibronectin is now implicated in the pathogenesis of pulmonary and vascular fibrosis. Inhibition of Fibronectin as a strategy to diminish fibrosis and improve cardiac function has been demonstrated in an animal model of heart failure [\[24\]](#page-11-7). Polymerized Fibronectin influences the deposition, maturation, and stability of other ECM proteins, including Collagen type I, which makes it an important regulator of ECM [\[24\]](#page-11-7). Therefore, targeted regulation of Collagen type I and Fibronectin expression in cardiovascular cells or a common regulator of these two important ECM proteins would provide a novel therapeutic approach to prevent or reduce cardiovascular fibrosis. The inhibitory action of metformin on DDR2 reported here needs to be viewed in light of these observations.

Next, we examined how metformin may inhibit DDR2 expression, focusing on the role of Transforming Growth Factor-β 1 (TGF-β 1) that is reported to be involved in many pathologies, including cardiovascular diseases in which it is associated with fibrosis, epithelial-to-mesenchymal transition, and inflammation [\[25](#page-11-8)[,26\]](#page-11-9). In the present study, hyperglycaemia was found to enhance TGF-β1 mRNA and protein expression in vascular adventitial fibroblasts (Figure [4A](#page-5-0),B). Inhibition of TGF-β1 using TGF-β1 siRNA (Figure [5A](#page-6-0)) and SMAD2/3 inhibition using PD169316 or SIS3 (Figure [5B](#page-6-0)) abolished the HG-induced increase in Fibronectin expression in these cells. Interestingly, while TGF-β1 siRNA abolished HG-induced increase in DDR2 expression (Figure [5C](#page-6-0)), consistent with the previously reported regulatory role of TGF-β1 in DDR2 expression [\[18\]](#page-11-1), overexpression of DDR2 in TGF-β1-silenced cells restored Fibronectin (Figure [5C](#page-6-0)), which showed that DDR2 acts downstream of TGF-β1 to regulate Fibronectin expression. Notably, metformin was found to abolish HG-stimulated increase in TGF-β1 mRNA and protein expression (Figure [4A](#page-5-0),B), indicating that suppression of $TGF-\beta1/SMAD2/3$ signalling underlies the negative regulatory effect of the drug on DDR2. Figure [6](#page-7-0) provides a schematic representation of the plausible molecular events by which metformin inhibits hyperglycaemia-induced profibrotic gene expression in vascular adventitial fibroblasts via TGF-β1 and DDR2 inhibition.

While discussing the action of metformin in the context of hyperglycaemia, it is pertinent to appreciate the reported link between hyperglycaemia and reactive oxygen species (ROS), whose role in mediating both physiological and pathophysiological signal transduction profoundly impacts gene expression in diverse cellular contexts. Many studies report that high glucose concentrations promote increased ROS generation, leading to oxidative stress and consequent cellular changes that underlie the complications of diabetes, including cardiovascular diseases [\[27\]](#page-11-10). High glucose concentrations are also reported to enhance intracellular synthesis of Angiotensin II (Ang II) in cardiac myocytes, vascular smooth muscle cells, and cardiac fibroblasts [\[28\]](#page-11-11). Moreover, in cardiac fibroblasts, Ang II acts via ROS to enhance DDR2 expression via the redox-sensitive transcription factor, NF-κB [\[11\]](#page-10-8). Remarkably, several studies report the inhibitory effect of metformin on ROS and oxidative stress [\[29\]](#page-11-12). It is therefore possible that metformin may negatively impact ROS signalling to inhibit DDR2 and DDR2-dependent downstream events.

In conclusion, this study expands our understanding of the mode of action of metformin, highlighting a novel mechanism by which this drug of choice in the treatment of diabetes may prevent excessive matrix accumulation in the vessel wall and protect against vascular fibrosis and dysfunction. The inhibitory effect of metformin on TGF-β1-dependent DDR2 expression and consequent downregulation of DDR2 signalling is an important

finding with immense clinical implications because of the characteristic presence of DDR2 in cardiac fibroblasts and vascular cells, and its critical role in the regulation of these cells. The obligate role of DDR2 in Collagen type I and Fibronectin expression in cardiac and vascular adventitial fibroblasts and its association with arterial fibrosis establish its role as an important determinant of fibrosis in the cardiovascular system [\[11,](#page-10-8)[12,](#page-10-10)[18,](#page-11-1)[21\]](#page-11-4). DDR2 also regulates the phenotypic transformation of Angiotensin II-stimulated cardiac fibroblasts [\[12\]](#page-10-10) and is a positive regulator of cardiac fibroblast proliferation [\[30\]](#page-11-13). It confers apoptosis resistance on cardiac fibroblasts [\[30\]](#page-11-13), which may underlie the role of these cells in adverse myocardial fibrosis following injury. Importantly, DDR2 regulates the expression of the Ang II receptor, AT1, in cardiac fibroblasts [\[21\]](#page-11-4), which would impact the pleiotropic actions of Ang II on cardiovascular cells. In this context, it is important to note that metformin is also reported to inhibit all of these processes that are under the regulatory control of DDR2. Thus, in different experimental models, metformin has been shown to inhibit the phenotypic transformation of Ang II-stimulated cardiac fibroblasts [\[31\]](#page-11-14) and negatively regulate the expression of Collagen type I, Fibronectin [\[32\]](#page-11-15), and the AT1 receptor [\[33\]](#page-11-16), and inhibit cardiac fibrosis via inhibition of the TGF-β1/SMAD3 signalling pathway [\[10,](#page-10-7)[34\]](#page-11-17). What is, however, not clear from these studies is whether a single critical regulator of multiple pathways could mediate the cardiovascular protective effects of metformin. Pending in vivo and clinical studies, it seems very likely that the manifold cardiovascular benefits of metformin action are attributable, at least in part, to the suppression of DDR2 signalling.

Figure 4. Metformin prevents HG-induced increase in TGF-β1 mRNA and protein expression in **Figure 4.** Metformin prevents HG-induced increase in TGF-β1 mRNA and protein expression in vascular adventitial fibroblasts. Cells were pre-treated with metformin hydrochloride (10 μ M), 45 min min prior to HG supplementation. (**A**) TGF-β1 mRNA levels were determined by Taqman real-time PCR at 6 h post-HG treatment, with β-Actin as loading control. (**B**) Protein was isolated after 12 h of prior to HG supplementation. (**A**) TGF-β1 mRNA levels were determined by Taqman real-time PCR at 6 h post-HG treatment, with β-Actin as loading control. (**B**) Protein was isolated after 12 h of HG and subjected to western blot analysis for detection of TGF-β1, with β-Actin as loading control. Data are expressed as mean \pm SEM (standard error of mean). The groups were analysed by one-way ANOVA ($p \leq 0.05$) and Tukey's multiple comparisons ($p \leq 0.05$). Significant comparisons along with their *p* values are depicted in the Figure. The data are representative of three independent experiments, $n = 3$.

Figure 5. DDR2 acts downstream of TGF-β1 to regulate Fibronectin expression. (**A**) Cells were **Figure 5.** DDR2 acts downstream of TGF-β1 to regulate Fibronectin expression. (**A**) Cells were transient of *DDF-* acts downsteam of 1 cm^o 1 and 2 (5 pm/ 1 and 2 and 2 cmo 6 pmol) (in discrete by (-) signals and 2 (5 pm/ 1 and 2 a transiently transfected with TGF-β1 siRNA 1 and 2 (5 pmol) (indicated by $(+)$ sign) or scrambled siRNA (5 pmol) (indicated by (−) sign) prior to treatment with HG for 12 h. Protein was isolated and subjected to western blot analysis for detection of Fibronectin, with β-Actin as loading control. Validation of TGF-β1 knockdown is also shown. (**B**) Sub-confluent, quiescent cultures of vascular

adventitial fibroblasts in M199 were pre-treated with SMAD2 inhibitor, PD169316, or SMAD3 inhibitor, SIS3, for 1 h and subsequently with HG for 12 h. Following treatment with HG, protein was isolated and subjected to western blot analysis for detection of Fibronectin, with β-Actin as loading control. 3,3'-diaminobenzidine (DAB) reagent was used for detection of β-Actin and brightness/contrast was adjusted uniformly to match with other blots. (**C**) Cells were co-transfected with DDR2 cDNA plasmid (1μg) (indicated by (+) sign) or empty plasmid (1μg) (indicated by (−) sign) and TGF-β1 $\frac{1}{2}$ siRNA 1 and 2 (5 pmol) (indicated by (+) sign) or scrambled siRNA (5 pmol) (indicated by (−) sign) (lipofectamine 2000-complexed) for 8 h in OptiMEM. After a recovery period of 12 h, the cells were subjected to HG for 12 h. Following treatment with HG, protein was isolated and subjected to western blot analysis for detection of Fibronectin, with β-Actin as loading control. Validation of DDR2 OE and TGF-β1 knockdown are also shown. 3,3'-diaminobenzidine (DAB) reagent was used for detection of β-Actin and brightness/contrast was adjusted uniformly to match with other blots. Data are expressed as mean \pm SEM (standard error of mean). The groups were analysed by one-way ANOVA ($p \leq 0.05$) and Tukey's multiple comparisons ($p \leq 0.05$). Significant comparisons along with their *p* values are depicted in the Figure. The data are representative of three independent experiments, $n = 3$. μ (in potectation exponentially for 8 h in OptiMEM. After a recovery period of 12 h, the cens were $\frac{1}{2}$

Figure 6. Schematic depiction of plausible mechanisms underlying the protective effect of metmin against hyperglycaemia-induced vascular fibrosis. Metformin inhibits TGF-β1/SMAD2/3 sig-formin against hyperglycaemia-induced vascular fibrosis. Metformin inhibits TGF-β1/SMAD2/3 signalling to inhibit DDR2-dependent pro-fibrotic gene expression in HG-stimulated vascular adven-nalling to inhibit DDR2-dependent pro-fibrotic gene expression in HG-stimulated vascular adventitial fibroblasts.

In conclusion, this study expands our understanding of the mode of action of met-**3. Materials and Methods**

PD169316 (Cat No. 513030) and SIS3 (Cat No. 566405) were from Calbiochem (San Diego, USA). Collagenase type II (Cat No. LS0004194) and elastase (lyophilised) (Cat No. LS006363) were from Worthington (Columbus, USA). Glucose (Cat No. G8270), metformin hydrochloride (Cat No. 1396309), SB431542 (Cat No. 616461), M199 (Cat No. M2520), and TGF-β1 (Cat No. T7039) were obtained from Sigma-Aldrich (St. Louis,

MO, USA). Lipofectamine 2000 (Cat No. 11668027) was from Invitrogen (Carlsbad, CA, USA). The Chemiluminescence western blot detection reagent (Cat No. 34095) and Pierce™ DAB Substrate Kit (Cat No. 34002) was from Thermo Fisher Scientific (Waltham, MA, USA). DDR2 (Cat No. 4390771, assay ID: s189847), TGF-β1 siRNA (Cat No. AM16708, assay ID: 199140 and 199141), and control siRNAs (Cat No. 4390844) were from Ambion (Foster City, CA, USA). Taqman RT-PCR probes for DDR2 (Cat No. 4351372, assay ID: Rn01490282_m1, FAM-labelled), TGF-β1 (Cat No. 4351372, assay ID: Rn00572010_m1, FAM-labelled), β-Actin (Cat No.4448489, assay ID: Rn00667869_m1, VIC-labelled), and 18s rRNA (Cat No. 4331182, assay ID: Rn03928990_g1, VIC-labelled) were from Thermo Fisher Scientific (Waltham, MA, USA). The rat DDR2/CD167b Gene ORF cDNA clone expression plasmid (Cat No. RG80341-G) was obtained from Sino Biologicals (Beijing, China). Opti-MEM (Cat No. 31985070) and fetal bovine serum (FBS) (Cat No. 10-082-147) were from GIBCO (Waltham, MA, USA). All cell culture ware were purchased from BD Falcon (Corning, NY, USA). Primary antibodies against DDR2 (Cat No. 12133S) and TGF-β1 (Cat No. 3711) were obtained from Cell Signalling Technology (Danvers, MA, USA). Primary antibodies against Fibronectin (Cat No. sc9068) and Collagen type I (Cat No. sc293182) were from Santa Cruz Biotechnology (Dallas, TX, USA). Loading control β-Actin (Cat No. A2228) antibody was obtained from Sigma-Aldrich. All antibodies were used after dilution (1:1000). XBT x-ray film was from Carestream (Rochester, NY, USA).

4. Isolation of Rat Vascular Adventitial Fibroblasts

Vascular adventitial fibroblasts were isolated from young adult male Sprague-Dawley rats following protocols previously standardised by us [\[18\]](#page-11-1). Aorta was removed from the left subclavian origin, washed in Hank's Balanced Salt Solution (HBSS), and then transferred to a 100 mm dish with fresh HBSS in a sterile hood. Fat was removed gently under a microscope and incubated in a 35 mm dish containing 2 mL Digestion Medium I for 30 min at 37 °C. The tissue was then placed in fresh HBSS, cut open, separated into the outermost adventitial layer after scraping out the endothelium, and incubated in Digestion Medium II for 90 min at 37 °C. After incubation, an equal volume of Medium-199 (M199) containing 10% FBS was added to the dish and mixed well for proper dissociation. Cells were collected by filtering through a cell strainer $(70 \mu m)$ and centrifuged at 1500 rpm for 5 min. The cell pellet was collected and resuspended in 0.5 mL of M199 containing 10% FBS and seeded on a 12-well plate. At 24 h after isolation, the supernatant containing unattached cells was discarded and the cells were incubated with M199 containing 10% FBS in a CO_2 incubator at 37 °C. Cultures of adventitial fibroblasts were characterised by immunocytochemistry and checked for cross-contamination. Adventitial fibroblasts stained positive for DDR2 and negative for Desmin, as shown earlier [\[18\]](#page-11-1). Cells from passages 3–4 were used for the experiments.

5. High Glucose and Metformin Treatment

Vascular adventitial fibroblasts were grown in Dulbecco's Modified Eagle Medium (DMEM) (5.5 mM glucose, normal medium) for 24 h prior to the experiments. D-glucose was then supplemented to achieve a final concentration of 30 mM D-glucose (high glucose (HG); hyperglycaemic media) and the cells were incubated for 6 h (for mRNA expression) and 12 h (for protein expression). In the metformin group, metformin hydrochloride was added to the normal medium for a final concentration of 10 μ M, 45 min prior to high glucose supplementation.

6. Western Blot Analysis

Sub-confluent cultures of adventitial fibroblasts were subjected to the indicated treatments for 12 h, following which relative protein expression was determined by western blot analysis by standard protocols, using β-Actin as the loading control. Enhanced chemiluminescence reagent was used to detect the proteins. Where indicated in the Figure legends, 3,3'-diaminobenzidine (DAB) reagent was used for detection of β-Actin and

brightness/contrast was adjusted uniformly to match with other blots. Protein expression was quantified by densitometric analysis of scanned films using ImageJ software [\[35\]](#page-11-18). Full blots of the representative western blot images are provided as Supplementary Files.

7. Real-Time Polymerase Chain Reaction Analysis

Sub-confluent cultures of adventitial fibroblasts were subjected to the indicated treatments and total RNA was isolated using TRI reagent, following the manufacturer's instructions. After DNase I treatment, 2 µg of total RNA was reverse transcribed to cDNA with random primers and M-MLV reverse transcriptase. TaqMan quantitative RT-PCR analysis was performed using the ABI prism 7500 Sequence Detection System (Applied Biosystems, CA, USA) with specific FAM-labelled probes. PCR reactions were carried out over 40 cycles, as per the manufacturer's instructions. DDR2 and TGF-β1 mRNA expression levels were normalised to β-actin and 18s rRNA, respectively.

8. RNA Interference

The gene knockdown protocol by RNA interference was as reported by us earlier (18). Briefly, cells were seeded on 12-well plates at 8×10^4 cells/well. After 24 h, the cells were incubated in Opti-MEM with Ambion pre-designed Silencer Select siRNA [5 pmol DDR2, 5 pmol TGF-β1 (siRNA 1 and 2) or scrambled siRNA, control] and Lipofectamine 2000 $(2 \mu L)$ for 19 h. After an additional incubation in M199 with 10% FBS for 12 h, the cells were treated with high glucose for the indicated duration.

9. Overexpression of DDR2

Strong constitutive expression of DDR2 was achieved, as previously described by us [\[18\]](#page-11-1), under a pCMV promoter, purchased as a cDNA clone from Sinobiologicals, China (Rat DDR2/CD167b Gene ORF cDNA). The DDR2 clone size was verified by PCR amplification of the primers in the kit. The size of the plasmid was analysed by single site restriction digestion at the Mlu-1 site. DDR2 overexpression plasmid cocktail was prepared with 1 μ g plasmid and 3 μ L Lipofectamine 2000 in 100 μ L Opti-MEM, and added to the cultures for incubation in M199 + 10% FBS for 8 h.

The transfection procedures did not affect cell viability.

10. Statistical Analysis

GraphPad Prism (version 9.4.1(681)) software was used for the illustration of graphs and for statistics. The groups were analysed by one-way ANOVA and $p \leq 0.05$ was considered significant. Pair-wise post-hoc testing for multiple comparisons between groups was by Tukey's method and $p \leq 0.05$ was considered significant. The data presented are mean \pm SEM (standard error of mean) of three independent experiments.

Supplementary Materials: The following supporting information can be downloaded at: [https:](https://www.mdpi.com/article/10.3390/ijms24010585/s1) [//www.mdpi.com/article/10.3390/ijms24010585/s1.](https://www.mdpi.com/article/10.3390/ijms24010585/s1)

Author Contributions: A.S.T., S.K., H.V., M.G.U., E.G.L. and M.W. conceived the study; A.S.T., S.K., H.V. and M.G.U. designed the study; A.S.T., H.V. and M.G.U. performed the experiments; A.S.T. and S.K. analysed the data; A.S.T. and S.K. interpreted the results of the experiments; A.S.T. prepared the figures; A.S.T. and S.K. drafted the manuscript. All authors have read and agreed to the published version of the manuscript.

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Institutional Review Board Statement: The study on rats was approved by the Institutional Animal Ethics Committee of Sree Chitra Tirunal Institute for Medical Sciences and Technology, Trivandrum (B form No: SCT/IAEC-233/AUGUST/2017/94 and SCT/IAEC-268/FEBRUARY/2018/95).

Informed Consent Statement: Not applicable.

Data Availability Statement: The datasets generated during and/or analysed during the current study are available from the corresponding author on request.

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Abbreviations

DDR2: Discoidin Domain Receptor 2, TGF-β1: Transforming Growth Factor-β1, SMAD2/3: SMAD Family Member 2/3, PPARγ: Peroxisome proliferation-activated receptor gamma, NF-κB: Nuclear factor-kappa B, Ang-II: Angiotensin II, AT1: Angiotensin II receptor type 1, ROS: Reactive Oxygen Species.

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