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Endothelin A receptor activation on mesangial cells initiates Alport glomerular disease

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Abstract

Recent work demonstrates that Alport glomerular disease is mediated through a biomechanical strain-sensitive activation of mesangial actin dynamics. This occurs through a Rac1/CDC42 cross-talk mechanism that results in the invasion of the sub-capillary spaces by mesangial filopodia. The filopodia deposit mesangial matrix proteins in the glomerular basement membrane, including laminin 211, which activates focal adhesion kinase in podocytes culminating in the up-regulation of pro-inflammatory cytokines and metalloproteinases. These events drive the progression of glomerulonephritis. Here we test whether endothelial cell-derived endothelin-1 is upregulated in Alport glomeruli, and further elevated by hypertension. Treatment of cultured mesangial cells with endothelin-1 activates the formation of drebrin-positive actin microspikes. These microspikes do not form when cells are treated with the endothelin A receptor antagonist sitaxentan, or under conditions of siRNA knockdown of endothelin A receptor mRNA. Treatment of Alport mice with sitaxentan results in delayed onset of proteinuria, normalized glomerular basement membrane morphology, inhibition of mesangial filopodial invasion of the glomerular capillaries, normalization of glomerular expression of metalloproteinases and pro-inflammatory cytokines, increased lifespan, and prevention of glomerulosclerosis and interstitial fibrosis. Thus endothelin A receptor activation on mesangial cells is a key event in initiation of Alport glomerular disease in this model.

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Disclosure Statement: This work was done as a collaboration between the Cosgrove Laboratory and Shiguang Liu's laboratory at Genzyme. Genzyme performed microarray analysis, analyzed serum samples for blood urea nitrogen levels, duplicated lifespan and physiologic measures (proteinuria and BUN) to independently verify our findings, and participated in the writing and review of the manuscript. There were no direct financial contributions to the Cosgrove laboratory (sponsored research support, licensing agreements or royalties).

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Keywords

Alport syndrome; glomerulonephritis; endothelin; actin dynamics

Introduction

The mature mammalian glomerular basement membrane (GBM) contains two separate networks of type IV collagen; a subendothelial network comprised of $\alpha 1(\text{IV})/\alpha 2(\text{IV})$ collagen heterotrimers, and a subepithelial network comprised of $\alpha 3(\text{IV})/\alpha 4(\text{IV})/\alpha 5(\text{IV})$ collagen heterotrimers (1). The latter contains more interchain disulfide cross-links, and thus would likely contribute a degree of tension resistance to the GBM (2). In Alport syndrome, mutations in the genes encoding any of the three chains comprising the sub-epithelial network results in the absence of all three chains, owing to the requirement of all three chains for assembly of the heterotrimers (3). The resulting Alport GBM is thinner and the type IV collagen network is comprised of only $\alpha 1(\text{IV})/\alpha 2(\text{IV})$ networks.

One consequence of a thinned and less crosslinked GBM collagen network is a presumed increase in its elasticity. Consistent with this, we previously showed that salt-induced hypertension resulted in elevated expression of pathologic genes in Alport glomeruli and accelerated progression of the disease (4). More recently we demonstrated that stretch mediated activation of the Rho GTPases Rac1 and CDC42 in Alport mesangial cells induced mesangial filopodial invasion of the glomerular capillary loops (5). The mesangial filopodia deposit laminin 211 in the GBM. This abnormal GBM laminin is known to progressively accumulate in the Alport GBM (6,7), but its cellular origin and its role in glomerular pathology, until recently, remained unknown. In a recent study it was shown that GBM laminin 211 activates focal adhesion kinase (FAK) on podocytes, which results in pathologic gene activation including those encoding matrix metalloproteinases (MMPs) and pro-inflammatory cytokines. Blocking FAK activation ameliorated Alport renal disease progression and improved renal function (8).

Here we show that biomechanical strain induces expression of endothelin-1 in endothelial cells from Alport mouse glomeruli. We further demonstrate that mesangial cells express endothelin A receptors ($\text{ET}_{\text{A}}\text{R}$), but not endothelin B receptors ($\text{ET}_{\text{B}}\text{R}$), unlike podocytes, which express predominantly $\text{ET}_{\text{B}}\text{R}$ with low levels of $\text{ET}_{\text{A}}\text{R}$. Specific blockade of $\text{ET}_{\text{A}}\text{R}$ in Alport mice by sitaxentan significantly delayed the onset of Alport glomerular disease, ameliorated mesangial filopodial invasion of the glomerular capillaries, prevented interstitial fibrosis, and improved renal function. Thus, strain-mediated induction of endothelin-1 activates $\text{ET}_{\text{A}}\text{R}$ on mesangial cells resulting in mesangial filopodial invasion of glomerular capillaries. The filopodia deposit mesangial proteins in the GBM. One of these mesangial proteins, laminin 211, activates expression of pro-inflammatory cytokines and MMPs in glomerular podocytes. Blocking this pathway by way of $\text{ET}_{\text{A}}\text{R}$ blockade has significant therapeutic benefits, and thus this mechanism reveals a new class of therapeutic targets that have the potential to delay the onset and slow the progression of Alport renal disease.

Results

Endothelin-1 is induced in Alport glomerular endothelium and further induced by hypertension

Preproteinuric C57Bl/6 X-linked Alport mice were made hypertensive by providing L-NAME salts in the drinking water from 4 to 7 weeks of age. Blood pressure was monitored twice weekly from 4 to 7 weeks of age using a CODA-2 tail cuff monitoring device. C57Bl/6 X-linked Alport mice were used because the CODA -2 system cannot accommodate measures in 2 week old mice, the age of pre-proteinuric 129 Sv autosomal Alport mice, due to their size. Figure 1A shows that animals given salt were consistently hypertensive relative to animals given water. At 7 weeks of age these animals are still pre-proteinuric (9). Cryosections were immunostained with antibodies specific for either endothelin-1 or CD31 (a marker for endothelial cells). Sections were stained simultaneously with the same antibody dilutions and imaged setting the gain of the confocal microscope using the hypertensive Alport sample immunostained for endothelin-1. Images for all remaining endothelin-1 immunostains were collected using this same gain setting, thus reflecting relative fluorescence to the hypertensive Alport sample. The lower panel of Figure 1B shows that the immunostaining intensity of ET-1 in glomeruli from hypertensive Alport mice is greater than that for normotensive Alport mice, which are greater than that for wild type mice. ET-1 co-localizes with CD31, indicating the cellular source of ET-1 is the endothelium. Western blot analysis shown in Figure 2A–B confirms significantly elevated expression of active ET-1 in Alport glomeruli compared to wild type. Significantly Elevated active endothelin was observed in the urine of 5 week old C57 Bl/6 XLAS mice, a biomarker for chronic kidney disease (10), by microplate ELISA (Figure 2C). Glomerular endothelin mRNA was not significantly elevated in either 129 Sv or the C56Bl/6 Alprot mouse models (Figure 2D).

ET_AR activation on mesangial cells induce the formation of filopodia

Previous studies in rat have shown that ET_ARs are expressed on glomerular mesangial cells and podocytes, while ET_BRs are on podocytes and endothelial cells (11). We examined the relative expression of ETRs in the 129 Sv mouse. Figure 3 (upper panels) shows that glomerular ET_AR immunostaining co-localizes with the mesangial cell marker, integrin α 8, consistent with its expression in mesangial cells. In contrast, ET_BR immunostaining was observed in podocytes, identified with α -actinin-4. Western blots (Figure 3, lower panel) confirm that only the ET_AR is detected in cultured mesangial cells, while both ET_AR and ET_BR are detected in cultured podocytes (GECs). Glomerular outgrowths were used as a positive control for both receptors.

It has been previously shown that endothelin-1 activates CDC42/RAC1 in mesangial cells via ET_ARs (12,13), and we showed that CDC42/RAC1 crosstalk was linked to the activation of mesangial filopodial invasion of glomerular capillaries (5). A direct link between mesangial cell activation and filopodial formation has not been established. To address this, we pre-treated (or not) serum-starved cultured mesangial cells with sitaxentan and then stimulated the cells with endothelin-1. Cells were then dual labeled with anti-drebrin antibodies (drebrin stabilizes actin filaments in filopodia) and phalloidin. The results in

Figure 4A demonstrate that treatment of cultured mesangial cells with endothelin-1 induces the formation of drebrin-positive filopodial microspikes, and pretreatment of cells with the ET_AR antagonist Sitaxentan blocks their formation. We further analyzed lysates from cultured mesangial cells using these same conditions for the activation of CDC42 (GTP-CDC42). The results in Figure 4B show that endothelin-1 treatment significantly activates CDC42 in these cells, and that pretreatment with sitaxentan prevents this activation.

ET_AR blockade delays Alport glomerular disease initiation and ameliorates interstitial fibrosis in Alport mice

To determine whether ETR antagonism prevents mesangial filopodia formation *in vivo*, 129 Sv autosomal Alport mice were given sitaxentan once daily by oral gavage from 2 weeks to 7 weeks of age. Kidney cryosections were immunostained using antibodies laminin α 5 and integrin α 8 to determine the degree of mesangial filopodial invasion in the glomerular capillaries. Transmission electron microscopy was used to determine whether the ET_AR antagonist ameliorates GBM damage. The results in Figure 5A (arrowheads/insets) demonstrate that both vehicle-treated 129 Sv autosomal Alport mice (7 weeks of age) and vehicle-treated C57 Bl/6 XLAS mice (15 weeks of age) showed extensive invasion of glomerular capillaries by mesangial filopodia. Sitaxentan-treated mice showed marked reduction of mesangial filopodial invasion of glomerular capillaries, as evidenced by the relative lack of integrin α 8 immunopositivity in the glomerular capillary loops. Normalization of GBM ultrastructural morphology was also observed (Figure 5B and C) compared GBM morphology observed in vehicle-treated Alport mice. Lower magnification images of the GBM from which these examples were taken can be viewed in supplementary Figure 1. Sitaxentan treatment did not effect blood pressure (Figure 5C), suggesting its effects on Alport renal disease are ET_AR-mediated rather than non-specific.

The idea of mesangial filopodial invasion is new to Alport research. To clearly demonstrate this, we performed ultra-high resolution structured illumination microscopy (SIM) analysis on 7 week old Alport mouse glomeruli. Figure 6A shows that filopodia (labeled with anti-integrin α 8 antibodies) are visible (arrow heads), contiguous with the mesangial angles (asterisks), and exclusive of both the glomerular endothelium (labeled with anti-CD31 antibodies) and the GBM (labeled with anti-laminin α 5 antibodies). The mesangial angles lie between the GBM and the endothelium (14). We further show in Figure 6A that the podocyte pedicles (labeled with anti-CD2AP antibodies) do not overlap with the mesangial filopodia, with space between the two markers defining the glomerular basement membrane (GBM). This rules out ectopic activation of integrin α 8 by either podocytes or endothelial cells. Figure 6B and C shows quantitative analysis of mesangial filopodial invasion based on total red fluorescence in the capillaries. The graph (Figure 6C-left) shows quantitative analysis of total red fluorescence (integrin α 8 immunostaining) in the capillary region from 6 independent glomeruli (from three individual mice) per group. A significant elevation in capillary red fluorescence was observed in glomeruli from vehicle-treated Alport mice compared wild type, and normalization of capillary integrin α 8 immunostaining in the sitaxentan-treated mice. The images used for this analysis are in Supplementary Figure 2.

Sitaxentan treatment ameliorated interstitial fibrosis and leukocyte recruitment to the renal interstitium, as evidenced by anti-collagen I (fibrosis marker) and anti-CD45 (pan-leukocyte marker) immunostaining (Figure 7A). Quantitative analysis for both fibrosis scores and % sclerotic glomeruli were reduced to near wild type levels in sitaxentan-treated mice relative to untreated Alport mice (Figure 7B and C). Glomerular expression of MMP-9, -10, and -12 as well as proinflammatory cytokines IL-6 and MCP-1 were significantly reduced in sitaxentan-treated Alport mice relative to Alport mice (Figure 7D) to near wild type levels. Analysis of RNA from renal cortex of these same mice using a mouse fibrosis microarray confirmed amelioration of expression for large number of genes associated with fibrogenesis in sitaxentan-treated Alport mice compared to Alport mice given vehicle (Table 1).

Sitaxentan-treated mice showed delayed *onset* of proteinuria, as well as significantly reduced proteinuria after the onset (Figure 8A), which is normally detected by 3 weeks of age in the 129 Sv autosomal Alport mouse model (6). This is consistent with the idea that ET_AR blockade delays disease initiation. Furthermore, BUN levels were reduced in sitaxentan-treated Alport mice relative to vehicle treated Alport mice (8B). Surprisingly, lifespan was increased only by approximately 20% (data not shown). This nominal improvement likely reflects the sitaxentan toxicity, as this class of compounds is notoriously toxic.

Discussion

In an earlier report we showed that hypertension accelerated the progression of Alport glomerular disease, suggesting a key role for biomechanical strain in the disease mechanism (4). This observation was extended with the discovery that mesangial processes invade the glomerular capillaries in a biomechanical strain-mediated Rac1/CDC42-activation mechanism (5). Importantly, the mesangial filopodia in the GBM are depositing mesangial matrix proteins, including laminin 211, which activates focal adhesion kinase in glomerular podocytes, resulting in the activation of genes encoding pro-inflammatory cytokines and metalloproteinases through an NF-kappaB-dependent signaling pathway, which drive the progression of GBM damage (8). The link between NF-kappaB activation in podocytes and the progression of Alport glomerular disease in the autosomal mouse model was recently strengthened by the observation that deletion of p53 in podocytes accelerated the progression of the disease by limiting podocyte expression of pro-inflammatory cytokines (15). The tumor suppressor p53 is a well-established negative regulator of NF-kappaB (16).

We surmised the mechanism underlying the activation of mesangial filopodia might reveal novel targets of therapeutic intervention aiming to arrest the *initiation* of these events. Here we show that induction of ET-1 expression in glomerular endothelial cells activates ET_ARs on mesangial cells culminating in CDC42 activation and the invasion of the sub-endothelial spaces of the glomerular capillaries by mesangial filopodia. These events constitute the earliest pathologic events yet identified, occurring before the onset of proteinuria (8, and data presented herein). *In vitro*, formation of drebrin-positive actin microspikes in cultured primary mesangial cells in response to ET-1 stimulation demonstrates that the pathway is intact and active. *In vivo*, ultra-high resolution structured illumination microscopy proves that integrin $\alpha 8$ immunopositivity is *in between*, and not overlapping with the GBM and the

endothelium, and contiguous with the mesangial angles, providing strong evidence that these filopodial projections do indeed originate from the mesangial cells. Proteinuria has been shown as causative of interstitial fibrosis in the kidney (17), and thus might be partially responsible for the reduction in interstitial fibrosis in sitaxentan-treated Alport mice.

It is likely that that multiple factors contribute to Alport glomerular disease initiation. Altered type IV collagen composition of the GBM might influence cell signaling. Ultra high resolution immunolocalization studies showed that the $\alpha 3(\text{IV})/\alpha 4(\text{IV})/\alpha 5$ network in wild type mice may not engage podocyte receptors. In Alport mice, however, the $\alpha 1(\text{IV})/\alpha 2(\text{IV})$ network is proximate enough to bind to collagen receptors on podocyte foot processes (18). The collagen-binding discoidin domain receptor 1 is present on the podocyte pedicles, and could bind the $\alpha 1(\text{IV})/\alpha 2(\text{IV})$ network resulting in receptor activation (19). Deletion of DDR1 in Alport mice resulted in improved renal function and a 50% increase in lifespan, supporting this notion (19).

Biomechanical stretching of glomerular podocytes activates angiotensin II receptors and increases expression of secreted protein acidic and rich in cysteine (SPARC) (20,21). The former is associated with apoptosis, and the latter is associated with accelerated renal injury in diabetic mice. Thus, biomechanical stretching of podocytes has been shown to have some direct maladaptive effects on podocyte biology, and thus could contribute to Alport glomerular disease.

ET-1 is secreted by podocytes in adriamycin-induced glomerulosclerosis, a model for focal segmental glomerular sclerosis, where it binds to ET_A Rs on glomerular endothelial cells, activating mitochondrial-targeted reactive oxygen species leading to podocyte loss (22). The fact that this mechanism is so different from the one described here demonstrates the diversity of ET-1 mediated influences on glomerular cells. This work suggests that release of endothelin-1 by endothelial cells in our model might promote similar pathology. Thus, the influences of ET_A R blockade on Alport glomerular pathology might involve direct contributions of all three glomerular cell types.

Biomechanical stretching may not be the only factor contributing to elevated expression of endothelin-1 in Alport glomerular endothelium. TGF- β 1 and TNF- α are induced in glomerular podocytes of Alport mice (23,24). Both cytokines have been shown to up-regulate expression of endothelin-1 in other systems (25,26). Sustained induction of endothelin-1 might be dominated early by a strain-dependent mechanism, and later by a cytokine-dependent mechanism.

In summary, this work confirms that ET-1 mediated activation of ET_A R on glomerular mesangial cells activates CDC42-dependent filopodial invasion of glomerular capillaries in Alport mice. These filopodia are the likely source of laminin 211, as well as other mesangial matrix proteins known to accumulate in the GBM of Alport humans, dogs and mice (6,7), and shown to underlie the FAK-dependent activation of NF-KappaB, which induces expression of pro-inflammatory cytokines and MMPs in podocytes (8). These events underlie Alport glomerular disease initiation, and reveal new therapeutic targets. Sitaxentan treatment increased lifespan approximately 20%, which is disappointing given the degree of

renal protection observed. This nominal improvement might reflect the toxicity of Sitaxentan, and suggests additional work needs to be done, perhaps using genetic modeling, to determine the full extent of renoprotection that might be achieved by inactivating this pathway.

Identification of elevated ET-1 levels in the urine of pre-proteinuric Alport mice suggests that ET-1 may be an early biomarker for Alport renal disease as it has proven to be for other models of CKD (10). The development of early biomarkers is currently a priority in the Alport field, as it may provide a sensitive means of monitoring response of emergent therapies (27). It would thus be useful to determine whether urinary ET-1 levels are elevated in pre-proteinuric Alport patients.

Methods

Mice

129 Sv autosomal Alport mice were developed here (28). C57 Bl/6 X-linked Alport mice were developed by Rheault et al.(29) and were obtained from the Jackson Laboratories. All mice were on pure genetic backgrounds and maintained as in house colonies. All procedures involving animals were conducted in accordance of an approved IACUC protocol. Every effort was made to minimize usage as well as minimize any pain or distress.

Antibodies

Anti- α -actinin 4 was from Santa Cruz Biotechnology, Inc (Dallas, TX, USA, Cat #: SC-49333); anti-CD31 was from Thermo Fisher Scientific, Inc (Rockford, IL, USA, Cat #: MA1-40074); Anti-collagen 1 was from CedarLane (Cat# CL50151AP) at 1:200 and CD45 was from eBioscience (Cat# 14-0451) at 1:100. anti-Endothelin-1 was from Abbotec (San Diego, CA, USA, Cat #: 250633) and Pierce (Lifetechnologies, Cat# PA3.067); anti-Endothelin Receptor A was from Novus Biologicals (Littleton, CO, USA, Cat #: NB600-836); anti-Endothelin Receptor B was from EMD Millipore (Billerica, MA, USA, Cat #: AB3284); anti-Integrin α 8 was from R&D Systems, Inc (Minneapolis, MN, USA, Cat #: AF4076); anti- Laminin α 2 was from Sigma-Aldrich (St. Louis, MO, USA, Cat #: L-0663); anti-Laminin α 5 was a gift from Dr. Jeff Miner (Washington University, St. Louis, MO).

Immunofluorescence microscopy

Fresh frozen kidneys were sectioned at 8- μ m and acetone fixed. Sections were incubated overnight at 4°C in the appropriate primary antibody solutions. The dual stains of rat anti-CD31 antibody at 1:50 and rabbit anti-Endothelin-1 antibody at 1:300. Dual stain of goat anti- α Actinin 4 antibody at 1:100 and rabbit anti-Endothelin Receptor B antibody at 1:100 were diluted in 1 % BSA in 0.3% PBST (Triton X-100). Anti-Collagen 1 was used at 1:200 and anti-CD45 was used at 1:100. The dual stains of anti-Endothelin Receptor A at 1:100, goat anti-Integrin α 8 at 1:1000, rat anti-Laminin α 2 at 1:200, and rabbit anti-Laminin α 5 at 1:2000 were diluted in 0.3% PBST + 5% FBS. Triple stain of rat anti-CD31 at 1:50, goat anti-Integrin α 8 at 1:1000, and rabbit anti-Laminin α 5 at 1:2000 were diluted in 0.3% PBST + 5% FBS. Secondary antibodies were at 1:500 for 1 hour at room temperature.

Mean fluorescence analysis

Images were analyzed using ImageJ. Each capillary loop was measured for area, integrated density and mean gray value. Three regions next to the capillary loop were measured for background values. The corrected total fluorescence was calculated by taking the integrated density subtracted by the area multiplied by the mean fluorescence of the background measurements.

Confocal Microscopy

Confocal images captured using a Leica TCS SP8 MP confocal imaging system, using a 63x NA: 1.4 oil or 10x NA: 0.3 objective.

Structured Illumination Microscopy

SIM images were captured using a Zeiss ELYRA PS.1 Super-resolution microscope.

Real Time qRT-PCR

Real time qRT-PCR was performed as described previously (5). Samples were normalized to Mouse GAPDH Endogenous Control VIC[®] Probe (Applied Biosystems catalog #: 4352339E) which was run alongside IL-6 (Catalog # 4331182, ID# Mm00446190_m1), MCP-1 (Catalog # 4331182, ID# Mm00441242_m1), MMP-9 (Catalog # 4331182, ID# Mm00442991_m1), MMP-10 (Catalog # 4331182, ID# Mm00444630_m1), and MMP-12 (Catalog # 4331182, ID# Mm00500554_m1) TaqMan[®] Gene Expression Assay Probes (Applied Biosystems).

Sclerotic glomeruli analysis

Paraffin tissue sections were stained with Periodic acid-Schiff (PAS). Percentage of sclerotic glomeruli were recorded for each section blinded.

Fibrosis scoring

Slides stained with anti-collagen 1 and anti-CD45 (pan leukocyte) antibodies were scored blinded based on the approximate percentage of the total area of the renal cortex that was inundated with collagen 1 and leukocytes. A score of 1 represents 10% fibrosis.

Albuminuria and Blood Urea Nitrogen (BUN) measures

Urine Albumin measurements were determined by using Molecular Innovation's (Novi, MI) Mouse Albumin Antigen Assay (cat. # MSAKT) and normalized using BioAssay Systems (Hayward, CA) QuantiChrom Creatinine Assay Kit (DICT-500). BUN measures were performed using a Cobas Integra 400 plus analyzer (Roche) in accordance with the manufacturer's instructions.

ET-1 Elisa

75 μ L of urine was assayed using a Quantikine[®] Elisa (R&D Systems, Minneapolis, MN) for active Endothelin-1 as per manufacturer's recommendations and normalized to creatinine (above).

Primary Mesangial Cells (MES) were derived and characterized as previously described (30). For Cdc42 G-Lisa® Activation Assay (Cytoskeleton, Denver, CO.) 30 % confluent MES cultures were incubated with 1% FCS-containing media for 24 hours, followed by 0.05% BSA containing media overnight. 10 μ M Sitaxentan was added to cultures for one hour, followed by 250 nM Endothelin-1 (ET-1) (Syd Labs, Boston, MA.) for 5 minutes. Lysates were collected and assayed as per manufacturer's instructions. For **ET-1 induction of Filopodial formation**, MES were cultured and serum deprived as above, exposed to 10 μ M Sitaxentan for one hour, followed by 250 nM EDN-1 for 30 minutes. Slides were fixed in -20° acetone, perfused with 0.1% Triton/PBS and stained with 1:250 dilution of antibody specific for Drebrin (Gene Tex, Irvine CA) and phalloidin (Molecular Probes, Grand Island, NY).

Conditional Immortalized Glomerular Epithelial Cells (GEC's), previously derived and characterized (31), were grown under permissive conditions (10 % FCS, 10 U/ml -interferon at 33° C).

Western Blot(s) for EDNr (A), (B), ET-1 and β -Actin were performed on protein lysates from glomeruli, GEC's, MES, and outgrowths. 10 μ g samples were denatured/reduced with the addition of 6X SDS loading buffer containing fresh β -mercapoethanol at 65° for at least two hours. 10% SDS PAGE gels were run as described (31). A 1:2000 dilution of primary antibodies (for ET-1, 1:500) were incubated in 5 % non-fat dry milk in PBST (0.1 % Triton), overnight at 4° . For EDNr (A) a 1:10,000 dilution of Donkey anti- Sheep IgG Antibody [HRP] (NB7190 NOVUS) was incubated for 2 hours in 5% milk/PBST and then developed using ECL (32109 Pierce, Grand Island, NY). For EDNr (B) a 1:5000 dilution of Goat anti-Rabbit IgG Antibody [HRP] (057K4857 Sigma) was incubated for 2 hours in 1% BSA/ PBST and then developed using ECL 2 (80196 Pierce). For β -Actin a 1:10,000 dilution of Goat anti-Mouse IgG Antibody [HRP] (025K6105 Sigma) was used.

Glomerular isolation

Glomeruli were isolated by transcardiac perfusion of 4 μ M Dynabeads (Life Technologies) and glomeruli isolated using a magnet as described by Takemoto et al., (32).

Blood pressure monitoring

Blood pressures were measured using the CODA2 tail cuff system as previously described (4).

Microarray

Total RNA was isolated from renal cortex (n=5 per group) from wild type and Alport mice given 10 mg/kg sitaxentan or vehicle from 2 to 7 weeks of age. RNA was analyzed by real time RT-PCR using the Mouse Fibrosis PCR Array (Qiagen) according to the manufacturer's instructions.

Statistical analysis

Real time qRT-PCR data was analyzed using the one sample Student's t-test (controls were always set arbitrarily equal to 1). Mean fluorescence and % sclerotic glomeruli data were analyzed by two-tailed Student's t-test.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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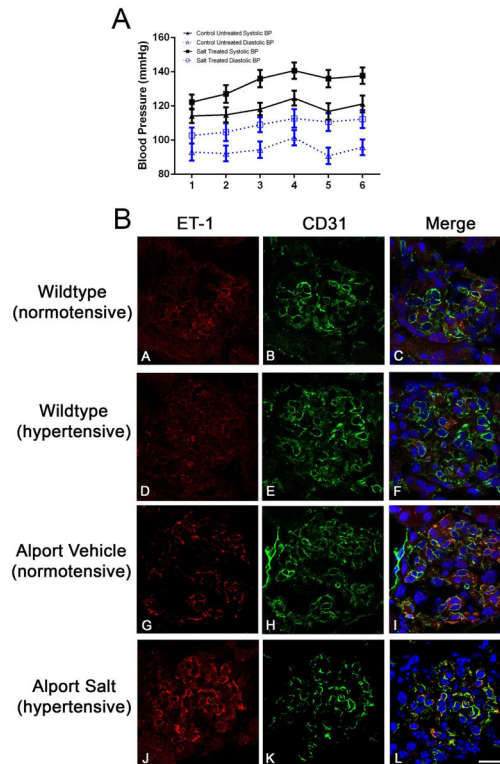


Figure 1.

1A: Induction of hypertension in Alport mice. C57Bl/6 X-linked Alport mice were made hypertensive by giving them the L-NAME salts in their drinking water from 4 to 7 weeks of age. Normotensive mice were given plain drinking water. Blood pressures were measured longitudinally using the CODA-2 tail cuff system. Numbers on the X-axis represent individual measurements which were performed twice weekly from 4 to 7 weeks of age. Both systolic and diastolic changes comparing normotensive and hypertensive mice were statistically significant ($P < 0.05$). **1B:** Hypertension induces endothelin-1 in Alport glomerular endothelial cells. Cryosections were dual immunostained with antibodies specific for either endothelin-1 or CD31 (a marker for endothelial cells). ET-1, endothelin-1. Elevated expression of endothelin-1 is clearly evident in glomeruli from hypertensive mice relative to normotensive mice (Compare G and J). This was not observed in wild type mice (compare A and D). Co-localization of endothelin-1 with CD31 demonstrates this induction is coming from the endothelial cell compartment. Bar=30 μ m.

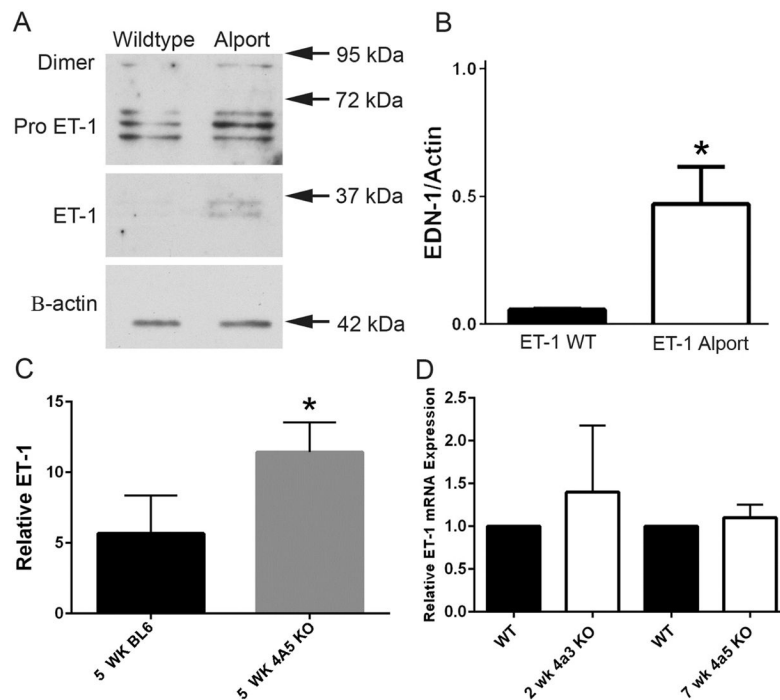


Figure 2.

ET-1 protein expression is elevated in glomeruli **and** urine from Alport mice relative to age/strain-matched wild type mice. **Panel A.** Glomeruli were isolated from 2 week old (pre-proteinuric) 129 Sv Alport mice and wild type mice. Lysates were analyzed by western blots and probed with anti-ET-1 antibodies. **Panel B.** Active ET-1 was quantified on quadruplicate blots by densitometry, demonstrating significantly elevated ET-1 in Alport versus wild type glomeruli. **Panel C.** Urine from 5 week C57 Bl/6 wild type and X-linked Alport mice was analyzed for active ET-1 using a microplate ELISA assay. **Panel D.** Glomerular RNA analysis by real time RT-PCR shows that ET-1 mRNA levels are not significantly elevated in either the pre-proteinuric 129 Sv autosomal Alport model or the C57 Bl/6 X-linked Alport model. * $P < 0.05$

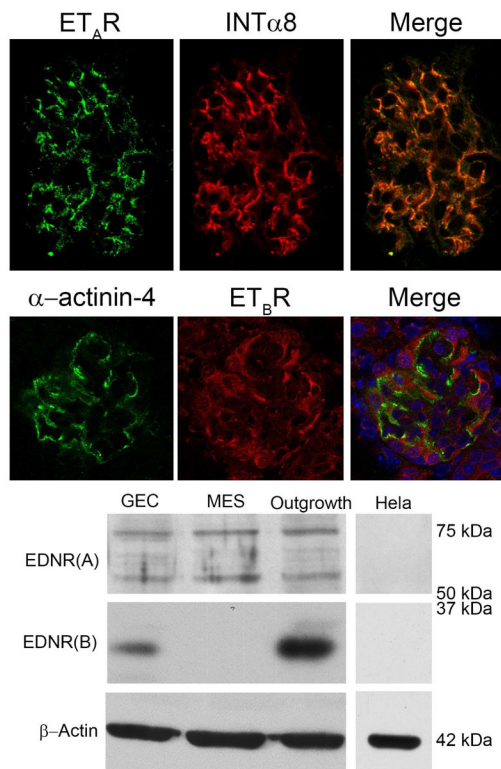


Figure 3.

Endothelin A receptor is the primary endothelin receptor on mouse mesangial cells. **Upper panel:** Kidney cryosections from wild type mice were dual stained with antibodies specific for the endothelin A receptor and the mesangial cell marker integrin α 8. Staining in a single glomerulus is shown. The merged image shows that the endothelin A receptor staining is primarily in the mesangial cell compartment. Bar=30 μ m. The **lower panel** shows that Endothelin B receptors are principally expressed on podocytes, consistent with earlier reports (10). Alpha-actinin-4 is used as a podocyte marker. Panel B. Western blots from cultured mesangial cells (MES) and podocytes (GEC) confirm that ET_AR is robustly expressed on mesangial cells, while ET_BR is not detected. Cultured podocytes and glomerular outgrowths (cultured for 24 hours after glomerular isolation, used as a positive control for both endothelin receptors) express both ET_AR and ET_BR. HeLa cell extracts were used as a negative control, and expressed neither ET_ARs or ET_BRs.

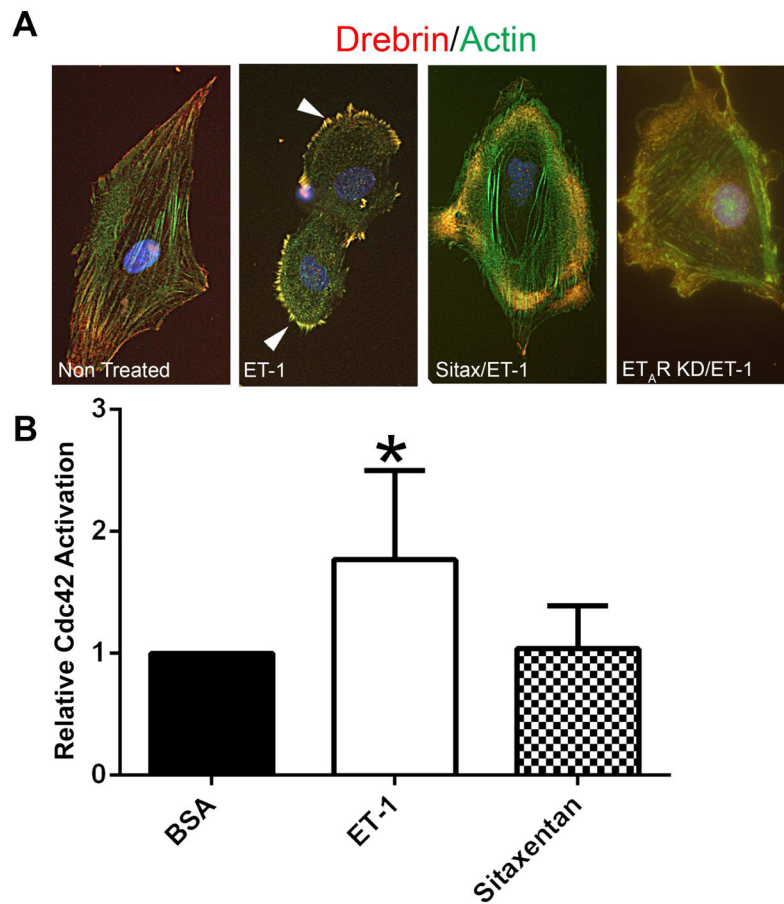


Figure 4. Treatment of mesangial cells with endothelin-1 activates CDC42 and induces the formation of drebrin-positive actin microspikes; microspikes and CDC42 activation are inhibited by pre-treatment of cells with Sitaxentan. Panel A: Cultured mesangial cells were serum starved, pretreated for 1 hour with or without Sitaxentan, treated for 30 minutes with endothelin-1, fixed with acetone, and dual stained with anti-drebrin antibodies (red) and phalloidin (green). Drebrin-positive microspikes (filopodia, denoted by arrowheads) are highly abundant on the endothelin-treated cells, but not detected when the cells are pre-treated with Sitaxentan or in ET_AR knockdown cells (control). Bar=12μm. Panel B: Cells were treated the same way as in panel A, then lysates prepared and assayed by ELISA for GTP-CDC42. Endothelin treatment significantly activates CDC42 in the cultured mesangial cells, and its activation is inhibited by pre-treatment of cells with Sitaxentan.

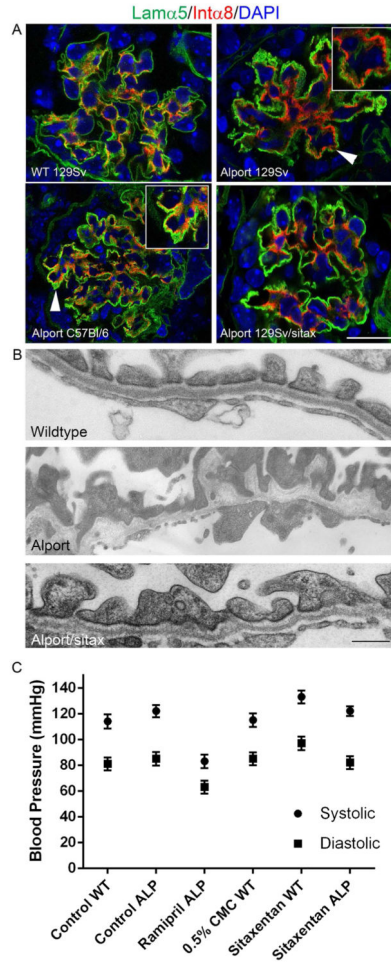


Figure 5. Endothelin A receptor blockade prevents mesangial process invasion of glomerular capillaries and ameliorates GBM damage in Alport mice. 129 Sv Alport animals were treated with the endothelin A receptor specific blocking agent Sitaxentan from 2 weeks to 7 weeks of age. Dual staining demonstrates absence of integrin $\alpha 8$ immunostaining in the glomerular capillaries, which are dual stained with anti-laminin $\alpha 5$ antibodies (**Panel A**). Arrows denote integrin $\alpha 8$ immunopositivity in the capillary loops of the glomeruli from untreated 129 Sv Autosomal Alport mice and C57 Bl/6 X-linked Alport mice, and the relative absence of integrin $\alpha 8$ immunopositivity in the sitaxentan-treated mice. Sitaxentan ameliorates GBM dysmorphology and (**Panel B**), largely normalizing the irregular thickening and thinning observed for the GBM of 7 week old Alport mice. Bar=500nm. Sitaxentan treatment does not affect blood pressure (**Panel C**). Wild type and Alport mice were treated with nothing (control), vehicle (CMC), ramipril (positive control for an agent known to effect blood pressure), or sitaxentan for 1 week and blood pressures measured (5 measures per condition) using a tail cuff method (CODA 2). While blood pressure was lower in the ramipril-treated mice, sitaxentan treatment had no effect on blood pressure.

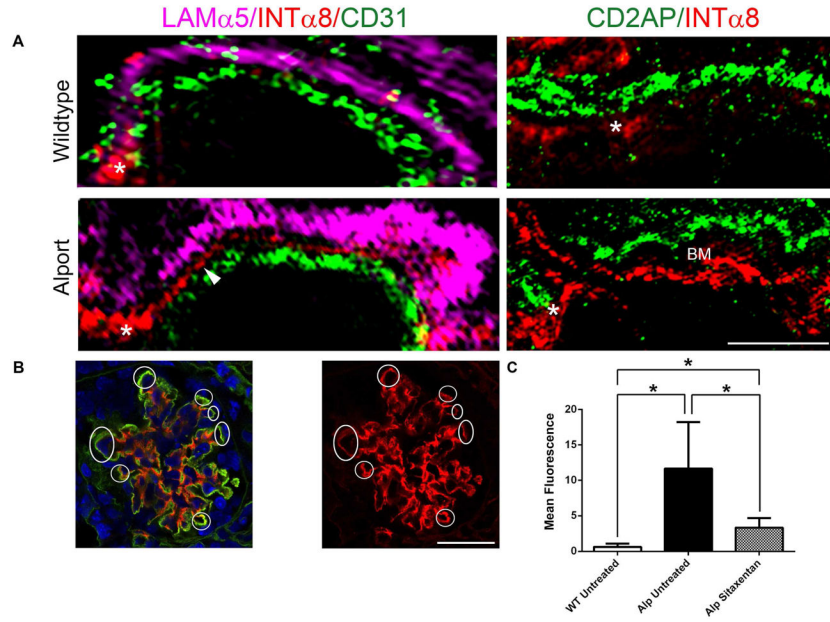


Figure 6.
Panel A (left panels) Ultra-high resolution Structured Illumination Microscopy (SIM) of Alport mouse glomeruli reveals that mesangial filopodia (arrowhead) emanate from the mesangial angles (asterisk), and flow between the glomerular endothelial cells, labeled using anti-CD31 antibodies, and the GBM, labeled using anti-laminin α 5 antibodies. These processes are not observed in wild type glomeruli. **Panel A (right panels)** SIM analysis of dual staining with the podocyte pedicle marker CD2AP and integrin α 8 demonstrate non-overlapping staining in the alport capillary loops with a space in between the labeled area where the basement membrane (BM) is located. Bar=2.5 μ m. **Panel B.** Integrin α 8 immunolabeling in the GBM demonstrates extensive filopodial invasion (circled) in 7 week Alport glomeruli that is prevented by treatment of animals with Sitaxentan. Bar=15 μ m. **Panel C.** Sitaxentan treatment significantly reduces integrin α 8 immunopositivity in the glomerular capillary loops. At least 6 glomeruli from at least 3 independent animals per group were analyzed by quantifying red fluorescence in circumferential capillary loops (denoted in Panel B as circled regions) using NIH image J software. Mesangial angles were excluded.

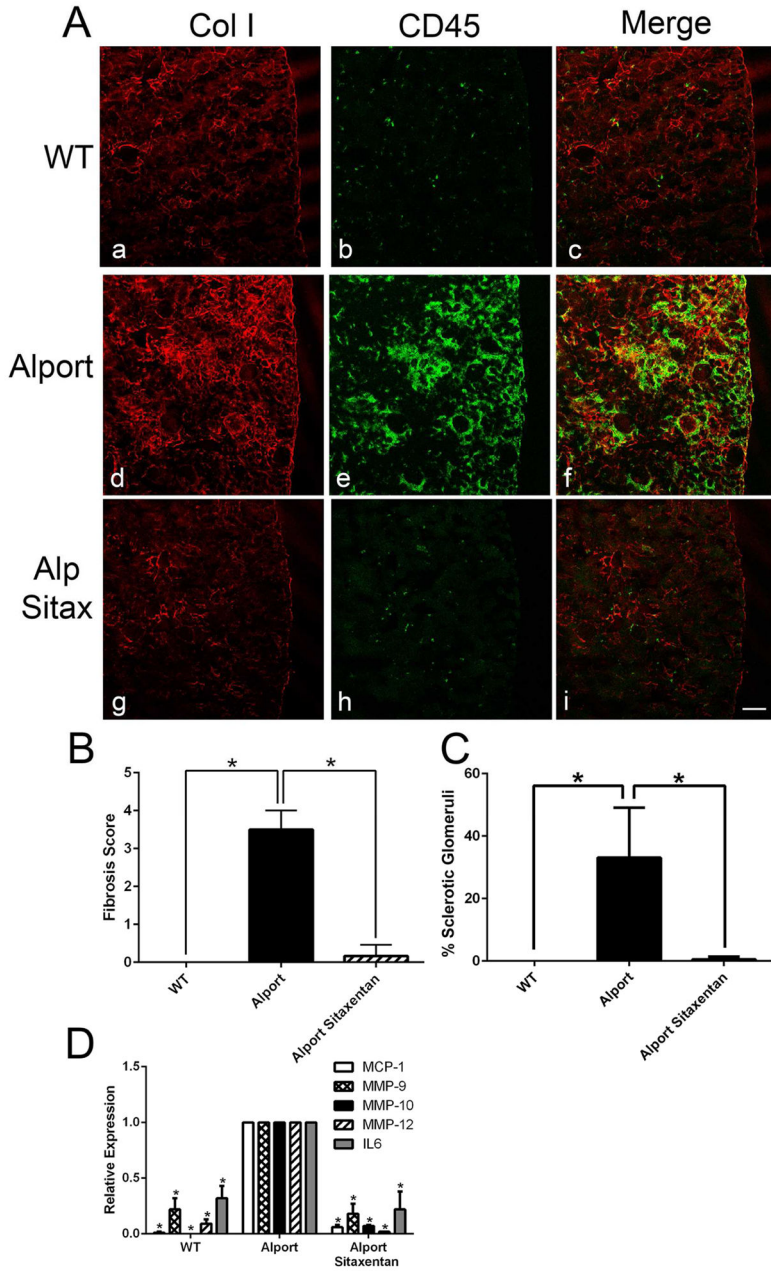


Figure 7.

Panel A Sitaxentan treatment ameliorates interstitial fibrosis and monocyte infiltration in Alport mouse kidneys. Cryosections from Sitaxentan-treated and vehicle treated wild type and Alport mice were immunostained with antibodies specific for collagen I (a marker for fibrosis) and CD45 (a pan-leukocyte marker). Note the complete absence of fibrosis and interstitial leukocytes in the treated mice. Bar=50µM. **Panel B.** Sitaxentan treatment ameliorates Alport interstitial fibrosis. Fibrosis scores were determined by blinded quantification of the relative cortical area with dense collagen I immunostaining (five animals per group). A score of 1 is 10%; a score of 4 is 40%. **Panel C.** Sitaxentan treatment ameliorates glomerulosclerosis in Alport mice. Sclerotic glomeruli were counted (blindly)

and plotted as percent relative to the total number of glomeruli (five animals per group).

Panel D. Sitaxentan treatment reduces mRNA expression of MMP-9, MMP-10, MMP-12, MCP-1, and IL-6 in glomeruli from Alport mice. Glomerular RNA from Sitaxentan-treated and vehicle-treated mice was analyzed by real time RT-PCR for the indicated transcripts. MMP, matrix metalloproteinase; MCP-1, monocyte chemoattractant protein-1; IL-6, interleukin 6. *P<0.001

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Table 1

Expression profiling of renal cortex mRNA from wild type, Alport, and sitaxentan-treated Alport mice using a mouse fibrosis microarray.

		Fold Change		
		Alport vs WT	Sitax Alp vs WT	Sitax Alp vs Alp
	Itgb8	0.5	1.2	2.4
	Itgb3	4.9	1.9	0.5
	Itga2	6.1	3.5	0.7
	Col1a2	8.6	3.0	0.4
	Col3a1	16.5	4.7	0.3
	Mmp1a	0.4	0.8	1.8
	Plat	1.8	1.9	1.1
Extracellular Matrix and Cell Adhesion	Timp2	1.9	1.7	0.9
	Serpinh1	1.9	1.2	0.7
	Lox	6.2	3.8	0.7
	Mmp2	6.6	3.2	0.6
	Mmp14	6.8	2.4	0.5
	Mmp9	10.5	3.2	0.4
	Mmp3	10.6	2.4	0.3
	Serpine1	23.7	10.7	0.6
	Timp1	525.4	113.1	0.3
Growth Factors	Egf	0.2	1.0	5.9
	Vegfa	0.6	1.2	1.8
	Pdgfa	1.7	1.4	0.8
	Pdgfb	2.3	1.1	0.6
	Hgf	4.3	3.4	0.8
Inflammatory	Il1b	9.0	2.5	0.4
	Tnf	18.6	4.7	0.4
	Ccr2	19.5	4.4	0.2
	Ctgf	1.8	3.3	1.9
Pro-Fibrotic	Snai1	2.2	1.7	0.8
	Grem1	10.0	4.2	0.5
	Ccl3	10.4	3.6	0.3
	Ccl11	11.3	3.8	0.3
	Ccl12	104.4	19.6	0.2
	Bmp7	0.4	0.9	2.5
	Tgfb2	1.6	1.2	0.8
TGF β Superfamily	Tgfb1	1.7	1.6	0.9
	Cebpb	1.8	1.9	1.0
	Nfkb1	1.9	1.5	0.8
	Tgif1	2.2	1.5	0.8
	Dcn	2.6	3.7	1.4

		Fold Change		
		Alport vs WT	Sitax Alp vs WT	Sitax Alp vs Alp
	Tgfb3	3.0	1.6	0.6
	Tgfb1	3.3	1.6	0.6
	Myc	3.4	2.1	0.7
	Thbs2	6.7	2.3	0.4
Apoptosis	Fasl	16.7	3.4	0.2

= 2-Tailed t-test (p<0.05)

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