Membranous nephropathy (MN) is the most common cause of nephrotic syndrome in adults, and one-third of patients develop end-stage renal disease (ESRD). Circulating autoantibodies against the podocyte surface antigens phospholipase A₂ receptor 1 (PLA₂R₁) and the recently identified thrombospondin type 1 domain–containing 7A (THSD7A) are assumed to cause the disease in the majority of patients. The pathogenicity of these antibodies, however, has not been directly proven. Here, we have reported the analysis and characterization of a male patient with THSD7A-associated MN who progressed to ESRD and subsequently underwent renal transplantation. Enhanced staining for THSD7A was observed in the kidney allograft, and detectable anti-THSD7A antibodies were present in the serum before and after transplantation, suggesting that these antibodies induced a recurrence of MN in the renal transplanted. In contrast to PLA₂R₁, THSD7A was expressed on both human and murine podocytes, enabling the evaluation of whether anti-THSD7A antibodies cause MN in mice. We demonstrated that human anti-THSD7A antibodies specifically bind to murine THSD7A on podocyte foot processes, induce proteinuria, and initiate a histopathological pattern that is typical of MN. Furthermore, anti-THSD7A antibodies induced marked cytoskeletal rearrangement in primary murine glomerular epithelial cells as well as in human embryonic kidney 293 cells. Our findings support a causative role of anti-THSD7A antibodies in the development of MN.

Introduction

Membranous nephropathy (MN) is an autoimmune disease that is histologically characterized by thickening of the glomerular basement membrane (GBM), granular staining for IgG, positivity for components of the complement system, and the presence of electron-dense deposits in the subepithelial space and within the GBM. Clinically, most patients present with high levels of proteinuria that usually exceed 3.5 grams per day, in conjunction with a nephrotic syndrome. The pathophysiology of MN has mainly been studied in the rat model of Heymann nephritis (1, 2). In passive Heymann nephritis, the transfer of sheep antibodies against the podocyte membrane protein megalin results in subepithelial immune complex formation (3, 4), activation of the complement system (5), and development of proteinuria. The concept that human MN is an antibody-mediated autoimmune disease has been supported by the discoveries of neutral endopeptidase (NEP) (6), phospholipase A₂ receptor 1 (PLA₂R₁) (7), and thrombospondin type 1 domain–containing 7A (THSD7A) (8) as podocyte membrane proteins serving as antigens in this disease. The current view is that PLA₂R₁ and THSD7A are targets for a malfunctioning immune system in 70% and 5% of adult cases, respectively, and that NEP is important in a small number of neonates with MN caused by alloimmunization due to the vertical transfer of antibodies from a genetically NEP-deficient mother (9). The findings that antibodies against PLA₂R₁ are present in patients with a recurrence of MN after renal transplantation (10, 11) and that anti-PLA₂R₁ antibody levels are associated with disease remission (12, 13) and progression (13–15) strongly support a pathogenic role of these antibodies in the development of MN. However, attempts to provide final proof that anti-PLA₂R₁ antibodies indeed cause MN have been hampered by the fact that PLA₂R₁ is not expressed on rodent podocytes. Given the recent observations that THSD7A, in contrast to PLA₂R₁, is expressed on both human and murine podocytes (16, 17), we investigated whether anti-THSD7A antibodies can cause MN in mice.

Results

Serum anti-THSD7A antibody positivity and recurrence of MN after renal transplantation. In 2004, a 57-year-old male was diagnosed with MN. An extensive clinical workup including medical history, prior medication, diagnostic tests for infections (hepatitis B and C, HIV, tuberculosis, treponema) and autoimmune diseases (lupus, rheumatoid arthritis, antineutrophil cytoplasmic antibody–associated [ANCA-associated] vasculitis, anti-GBM disease, sar-
however, the patient’s proteinuria increased to 3.4 grams per day, and a subsequent renal transplant biopsy showed a recurrence of MN, with normal staining for PLA2R1, enhanced staining for THSD7A, and positivity for complement C4d (Figure 1A and Supplemental Table 1). At that time, no donor-specific antibodies were detectable, and no evidence of a concomitant antibody-mediated rejection or another condition that might have led to a secondary MN was present. The patient’s serum creatinine levels increased shortly thereafter (Supplemental Figure 1), and anti-THSD7A antibodies were detectable, with a titer of 1:320 (Figure 1B). As proteinuria continued to rise, the patient received 2 doses of rituximab in addition to the continuous immunosuppression. Subsequently, the proteinuria decreased, but persisted at approximately 2.5 grams per day. Anti-THSD7A antibody levels also diminished, but remained positive at a level of 1:100 (Figure 1B). Taken together, this clinical case suggests a causative role of persisting circulating anti-THSD7A antibodies in the recurrence of MN in the transplanted kidney.

Human anti-THSD7A autoantibodies bind to mouse glomerular THSD7A. Serum samples from 2 patients with anti-THSD7A antibody-positive MN were taken for further studies (Supplemental Figure 2 and Supplemental Table 2). Our experiments demonstrated the ability of human anti-THSD7A antibodies to bind to the mouse antigen on podocyte foot processes in vitro and in vivo. Precisely, anti-THSD7A antibody–containing sera immunoprecipitated THSD7A from mouse glomeruli (Figure 2A) and bound linearly along the glomerular capillary wall in colocalization with nephrin in indirect immunofluorescence analyses of naive murine kidneys (Figure 2B). In vivo analyses further demonstrated binding of huIgG4 (huIgG4), the predominant anti-THSD7A IgG subclass in the patient’s serum, exclusively along the glomerular filtration barrier in colocalization with nephrin 2 hours after i.v. injection of anti-THSD7A antibody–containing serum into mice (Figure 2C).
merular podocytes (16), and the ability of human anti-THSD7A autoantibodies to bind to the murine antigen, we hypothesized that human anti-THSD7A antibodies cause MN in mice. To investigate the pathogenicity of anti-THSD7A antibodies, we injected BALB/c mice with a single i.p. dose of anti-THSD7A antibody-containing sera from patients with MN that were previously validated for the binding of mouse THSD7A (Figure 2). Thirteen mice received MN serum 1, seven mice received MN serum 2, and twelve mice received serum from a healthy individual. The glomerular binding of huIgG in mice injected with anti-THSD7A antibody-containing serum was initially found to be linear and on the subepithelial aspect of the GBM, as shown by its relation to the GBM-specific protein laminin (Figure 3A). By day 14, this linearity had changed toward a more granular pattern that became more prominent over time and persisted for at least 70 days. In contrast, mice injected with control serum failed to show binding of huIgG at the glomerular filtration barrier (Figure 3A). Notably, huIgG colocalized with THSD7A in mice that were exposed to sera from MN patients, suggesting antigen specificity of bound huIgG in subepithelial immune complexes (Figure 3B). In order to further investigate the specificity of bound huIgG in these mice, we eluted huIgG from frozen sections of kidney specimens 14 days and 70 days after injection of anti-THSD7A antibody–containing or control serum. HuIgG eluted from mice that received anti-THSD7A antibody–containing serum, but not from control mice, solely recognized a 250-kDa protein present in both human and mouse glomerular extracts (HGEs and MGEs) as well as recombinant mouse THSD7A, confirming that the huIgG that bound in mouse glomeruli was indeed specific for THSD7A (Figure 3C).

Figure 2. Human anti-THSD7A antibodies bind to mouse THSD7A in vitro and in vivo. (A) Sera from 2 patients with anti-THSD7A antibody–positive MN, but not serum from a healthy individual, immunoprecipitated THSD7A from MGEs. (B) Indirect immunofluorescence analysis using serum from a patient with THSD7A–associated MN or control serum in frozen sections of mouse kidneys. Original magnification, ×1,000, 4× zoom (enlargement of the boxed area). (C) Immunofluorescence staining for huIgG4 and nephrin 2 hours after i.v. injection of 100 μl serum from a patient with THSD7A–associated MN or a from healthy individual into male BALB/c mice (weight, 30 g). Original magnification, ×1,000, 4× zoom (enlargement of the boxed area). (B and C) Asterisks indicate podocyte nuclei. Scale bars: 50 μm.
within the subepithelial aspect of the GBM, suggesting shedding of THSD7A to the subepithelial space in the mouse injected with anti-THSD7A antibody–containing serum (Figure 5A). THSD7A expression was also found on foot processes in the control mouse (Figure 5A), which was comparable in extent and localization to what was described previously (17). In immunohistochemical analyses, we found strong granular staining for hulgG along the GBM on day 70, closely resembling the morphology of human MN (Figure 5B). In line with these findings, ultrastructural studies of the same mouse demonstrated electron-dense deposits in a strictly subepithelial localization and some broadening of podocyte foot processes (Figure 5C). We also found regions with focal podocyte foot process effacement over electron-dense deposits (Supplemental Figure 5). None of these changes were seen in mice that received the control serum (Supplemental Figure 5).

Podocyte injury in MN is accompanied by oxidative stress and an upregulation of superoxide dismutase 2 (SOD2) (18) to detoxify ROS. In line with this, we found SOD2 expression to be enhanced in mouse podocytes 70 days after injection of anti-THSD7A antibody–containing serum (Figure 5D). Moreover, podocyte injury was also reflected by disruption of nephrin immune localization in these mice but not in controls (Figure 5E).

Human anti-THSD7A autoantibodies induce proteinuria in mice. We examined whether the transfer of human anti-THSD7A antibody–containing serum caused proteinuria in mice. We found that around day 3, mice injected with anti-THSD7A antibody–containing serum developed significant albuminuria. The albuminuria was further increased after 14 days and remained elevated during the 70-day observation period and was supposedly induced by the autologous reaction against the persistently bound hulgG, the glomerular deposition of mouse anti-hulgG, and the subsequent activation of complement (ref. 19 and Figure 5F). Of note, albuminuria on day 3 occurred prior to enhanced glomerular binding of mIgG and deposition of complement (Supplemental Figure 6).
Urinary albumin excretion in mice injected with serum from a healthy donor remained unchanged (Figure 5F).

Affinity-purified THSD7A-specific huIgG causes the histological features of MN and transient proteinuria. In order to investigate whether it is indeed the serum autoantibodies against THSD7A that are causative for MN, we affinity purified THSD7A-specific huIgG from anti-THSD7A antibody–containing serum using chromatographic columns containing beaded agarose coupled to DDK-purified recombinant THSD7A. Three rounds of purification were performed, with a total of six acid elutions, reducing the content of anti-THSD7A antibodies in the serum to nondetectable levels (Figure 6A). Subsequently, 4 mice each were injected with anti-THSD7A antibody–depleted serum or with affinity-purified anti-THSD7A antibodies from the pooled eluates. Morphological analyses demonstrated granular subepithelial deposition of purified anti-THSD7A antibodies, whereas solely mesangial huIgG deposition was observed in mice injected with the depleted serum (Figure 6B). Again, huIgG colocalized with THSD7A in immune deposits (Figure 6C) and induced a granular subepithelial mlG deposition (Figure 6D) only in mice injected with purified anti-THSD7A antibodies. Altogether, the subepithelial deposition of huIgG and mlG was less pronounced in mice injected with purified anti-THSD7A antibodies than in mice that received anti-THSD7A antibody–containing whole serum (Figure 3), a finding that was likely related to antibody loss during the purification procedure.

Clinically, injection of purified anti-THSD7A antibodies induced significant albuminuria after 3 days when compared with mice that received anti-THSD7A antibody–depleted serum (Figure 6E). After 3 weeks, proteinuria gradually decreased to the level seen in depleted serum–injected mice. No complement C3 deposition was observed in any of the mice (data not shown), suggesting that the amount of injected purified anti-THSD7A antibodies was not sufficient to induce an autologous phase with maintenance of proteinuria. Taken together, these experiments demonstrate that anti-THSD7A antibodies alone are sufficient to induce MN.

Anti-THSD7A autoantibodies cause cytoskeletal rearrangement in primary cultured glomerular epithelial and human embryonic kidney 293 cells. We next examined whether anti-THSD7A antibodies can directly affect the cellular architecture of THSD7A-expressing cells. Immortalized cultured murine or human podocyte cell lines do not express THSD7A at detectable protein levels (our unpublished data). We therefore generated primary glomerular epithelial cells (GECs) by culturing isolated murine glomeruli for 3 to 5 days on collagen I–coated plates. Western blot analysis showed THSD7A expression in GECs (Figure 7A). Exposure of GECs to affinity-purified anti-THSD7A antibodies resulted in binding of huIgG to the cell membrane. Cells with membrane-bound huIgG stained positive for the standard podocyte marker protein α-actinin-4, suggesting that these cells were indeed podocytes (Figure 7B). Cells expressing α-actinin-4 that were treated with serum depleted of anti-THSD7A antibodies or with huIgG that was purified from a healthy donor failed to bind huIgG (Figure 7B and Supplemental Figure 7). Importantly, cells that did not stain for α-actinin-4 also did not bind huIgG following incubation with affinity-purified anti-THSD7A antibodies (arrows in Figure 7B), indicating that the binding of THSD7A autoantibodies was specific for mouse primary podocytes.

GECs that exhibited huIgG binding following exposure to affinity-purified anti-THSD7A antibodies showed marked cytoskeletal rearrangement, with formation of stress fibers and strong enhancement of F-actin staining (Figure 7C). When quantifying F-actin OD, we found a 2.5-fold increase in cells treated with purified anti-THSD7A antibodies as compared with that detected in cells treated with depleted serum (Figure 7D). Similar results were obtained when anti-THSD7A antibody–containing whole serum was used for experimental procedures (Supplemental Figure 8).

Focal adhesions are highly organized multiprotein assemblies that mediate signaling between the actin cytoskeleton and the extracellular matrix (20). Paxillin is a vital member of focal adhesions (21) and connects integrin signaling with cytoskeletal remodeling (22). In order to investigate whether the changes in cytoskeletal organization were accompanied by altered focal adhesions, we performed staining for phosphorylated paxillin (p-paxillin). We found an increase in focal adhesions with positive p-paxillin staining in GECs that were exposed to purified anti-THSD7A antibodies, but not in GECs treated with depleted serum (Figure 7C). Moreover, p-paxillin was shifted from the leading edges of cells toward cell centers, according to the formation of stress fibers in

**Figure 4.** Mouse IgG and complement C3 are deposited in the glomeruli of mice that were injected with anti-THSD7A–containing serum. (**A** and **B**) Immunofluorescence staining (paraffin sections) for mlG and laminin (**A** and complement component C3 and laminin (**B** on day 70. Scale bars: 10 μm.
cells with bound huIgG, but not in cells devoid of huIgG binding (Figure 7C, arrows).

Furthermore, GECs exhibited an elongated morphology following exposure to anti-THSD7A antibody–containing serum (Supplemental Figure 9), and some detached from the culture plates and were lost to the medium (data not shown). Since GECs are a heterogeneous population of glomerular cells, we repeated the experiments in human embryonic kidney 293 (HEK293) cells for adequate quantification of the observed effects. HEK293 cells transiently transfected with human THSD7A or empty vector (Figure 8A and Supplemental Figure 10) were incubated with medium containing 2% of the anti-THSD7A antibody–containing or control serum. When incubated with anti-THSD7A antibody–containing serum, binding of huIgG on the cell membrane and predominantly at cell-cell junctions of THSD7A-expressing HEK293 cells and colocalization of huIgG with THSD7A were detected (Supplemental Figure 10). Following exposure to anti-THSD7A antibody–containing serum, confluent THSD7A-expressing cells significantly detached from their culture plates (Supplemental Figure 11A) and were lost to the culture medium (Figure 8B). FACS-based analyses demonstrated that detached cells were THSD7A positive and had huIgG bound to the cell membrane (Supplemental Figure 11B). Additionally, detached cells with bound huIgG were alive, as demonstrated by exclusion of ViViD staining in THSD7A-expressing cells (Supplemental Figure 11C). THSD7A-expressing HEK293 cells did not detach from the culture plates when incubated with control serum, serum from patients with PLA2R1-associated MN, or serum from patients with MN that had neither anti-THSD7A nor anti-PLA2R1 autoantibodies (Figure 8B). In contrast, cells significantly detached when incubated with affinity-purified anti-THSD7A antibodies, but not with anti-THSD7A–depleted serum or purified control IgG (Figure 8C).
In line with these observations, incubation with anti-THSD7A antibody-containing serum and purified anti-THSD7A antibodies resulted in the cytoskeletal rearrangement of THSD7A-expressing cells that was characterized by strong enhancement of cortical F-actin fibers (Figure 8D and Supplemental Figure 12). Taken together, these results suggest a direct influence of anti-THSD7A antibodies on cytoskeletal organization of THSD7A-expressing GECs and HEK293 cells.

Discussion
A number of glomerular proteins have been identified as potential targets for autoimmunity in MN in recent years. These molecules include the podocyte membrane proteins NEP (6) and PLA2R1 (7). Additionally, other intracellular antigens such as SOD2, ubiquitin C-terminal hydrolase L1 (UCHL1), and aldose reductase have been described (18). However, the question of whether autoantibodies against all these antigens are actually causing the morphological and clinical picture of MN has intrigued researchers in the field for years (23). The recent discovery of THSD7A as another endogenously expressed podocyte antigen in human MN (8) and the fact that THSD7A is expressed in mouse podocytes (16, 17) with more than 90% sequence homology to the human molecule have given rise to experimental approaches in the investigation of MN pathogenesis.

In light of our observation of a patient who presented with a recurrence of THSD7A-associated MN in a renal transplant in
anti-THSD7A antibodies to THSD7A resulted in the onset of proteinuria, establishing the causal relationship between podocyte-directed autoantibodies and the development of MN. That this concept is also operative in human MN was first shown in 2002, when Debiec et al. discovered autoantibodies against NEP to be associated with a rare form of newborn alloimmune MN (6). They could also demonstrate that injection of whole IgG isolated from the mother (who gave birth to a child with antenatal NEP-associated MN) into rabbits induced proteinuria and glomerular immune complexes (6). In our experiments, both whole serum and affinity-purified THSD7A-specific IgG induced proteinuria within 3 days after administration to mice. In the presence of circulating anti-THSD7A antibodies, we hypothesized that anti-THSD7A antibodies could be causative for MN in humans. We therefore initiated studies in mice, transferring anti-THSD7A antibodies from 2 patients with THSD7A-associated MN. Our data demonstrate that human anti-THSD7A antibodies can bind to the murine antigen in vivo and in vitro. Moreover, the administration of both anti-THSD7A antibody–containing serum and isolated THSD7A-specific IgG to mice led to the development of proteinuria and a histomorphological pattern of MN. Injected antibodies thereby reached the glomerulus, bound to their target antigen, and induced subepithelial immune deposits containing THSD7A and anti-THSD7A antibodies. The binding of anti-THSD7A antibodies to THSD7A resulted in the onset of proteinuria, establishing the causal relationship between podocyte-directed autoantibodies and the development of MN.

That this concept is also operative in human MN was first shown in 2002, when Debiec et al. discovered autoantibodies against NEP to be associated with a rare form of newborn alloimmune MN (6). They could also demonstrate that injection of whole IgG isolated from the mother (who gave birth to a child with antenatal NEP-associated MN) into rabbits induced proteinuria and glomerular immune complexes (6). In our experiments, both whole serum and affinity-purified THSD7A-specific IgG induced proteinuria within 3 days after administration to mice. In the ani-

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**Figure 7. Anti-THSD7A antibodies cause cytoskeletal rearrangement in THSD7A-expressing primary cultured GECs.** (A) Western blot analysis shows expression of THSD7A in GECs. Rec. THSD7A, recombinant human THSD7A. (B) Immunofluorescence staining for hulgG and α-actinin-4 in GECs following a 40-minute exposure to affinity-purified anti-THSD7A antibodies or to serum that was depleted of anti-THSD7A antibodies. Arrows indicate cells that did not express α-actinin-4 and that showed no hulgG binding. Asterisks indicate the glomerulus from which the cells grew. Scale bars: 50 μm. Enlargements of boxed areas are shown in the far right upper and lower panels (original magnification, ×1,000, 4× zoom). (C) Immunofluorescence staining for hulgG, F-actin (phalloidin), and p-paxillin in GECs following a 40-minute exposure to affinity-purified anti-THSD7A antibodies or to serum that was depleted of anti-THSD7A antibodies. Arrows indicate a cell that had no hulgG bound to the membrane and that did not exhibit enhanced F-actin staining. Asterisks indicate the glomerulus from which the cells grew. Scale bars: 50 μm. (D) Quantification of stress fiber formation after treatment of GECs with purified anti-THSD7A antibodies or serum depleted of anti-THSD7A antibodies. A total of 30 images from 3 independent experiments were analyzed. Data indicate F-actin OD of individually circled cells normalized to the control condition (depleted serum). Error bars represent the mean ± SEM. ***P < 0.001, by 2-tailed, nonparametric Mann-Whitney U test.
and the development of proteinuria (5, 24). Other studies suggest that proteinuria can develop in the absence of certain complement factors (25, 26). We therefore analyzed mouse kidneys from our experiments for the presence of complement components in the subepithelial immune deposits. In the early phase of the disease, we could not detect C3 or C5b-9 in mice that received whole serum or in mice that received purified anti-THSD7A antibodies alone. Later in the course, marked C3 deposits were found in the mice injected with whole serum, but were absent in the animals receiving anti-THSD7A antibodies alone. Thus, it is possible that at early time points, complement activation is not vital to the initiation of podocyte injury and proteinuria in this model, but that at later time points, when larger amounts of autologous mouse anti-

Figure 8. Anti-THSD7A antibodies induce cell detachment and cytoskeletal rearrangement in THSD7A-expressing HEK293 cells. (A) Western blot analysis shows reactivity of a specific anti-THSD7A antibody in HEK293 cells that were transfected with THSD7A cDNA or empty vector. (B and C) Quantification of detached cells by measurement of total protein concentration in cell pellets after centrifugation of cell culture supernatants following a 60-minute incubation with (B) anti-THSD7A antibody-containing or control serum (results are from at least 5 independent experiments with 2 anti-THSD7 antibody–positive sera, 5 anti-PLA2R1 antibody–positive sera, 5 sera with no antibodies against THSD7A or PLA2R1, and 1 control serum) or with (C) affinity-purified anti-THSD7A antibodies, depleted serum, and affinity-purified IgG from a healthy individual (results are from 5 independent experiments with antibodies purified from 1 patient, the corresponding depleted serum, and purified control IgG). Data indicate the fold change in protein content in the cell culture supernatant in comparison with the cells that were transfected with empty vector and treated with serum from a healthy donor (B) or with purified control IgG (C). Error bars indicate the mean ± SEM. ***P < 0.001, by 1-way ANOVA with Bonferroni’s post test; statistical significance is shown for all conditions versus THSD7A-transfected cells that were treated with anti-THSD7A antibody–containing serum (B) or affinity-purified anti-THSD7A antibodies (C). (D) Immunofluorescence staining for huIgG and F-actin (phalloidin) in empty vector– and THSD7A-transfected HEK293 cells after a 30-minute exposure to anti-THSD7A antibody–containing serum or control serum. Arrows show accentuated cortical F-actin rings; asterisks indicate detaching cells. Scale bars: 20 μm.

In contrast, proteinuria in animals that received the purified anti-THSD7A antibodies was transient, which we cannot sufficiently explain at this point. However, some issues may be relevant. There is considerable evidence from studies in passive Heymann nephritis that proteinuria in this model depends on activation of the complement system and that the complement membrane attack complex C5b-9 is responsible for podocyte injury and the development of proteinuria (5, 24). Other studies suggest that proteinuria can develop in the absence of certain complement factors (25, 26). We therefore analyzed mouse kidneys from our experiments for the presence of complement components in the subepithelial immune deposits. In the early phase of the disease, we could not detect C3 or C5b-9 in mice that received whole serum or in mice that received purified anti-THSD7A antibodies alone. Later in the course, marked C3 deposits were found in the mice injected with whole serum, but were absent in the animals receiving anti-THSD7A antibodies alone. Thus, it is possible that at early time points, complement activation is not vital to the initiation of podocyte injury and proteinuria in this model, but that at later time points, when larger amounts of autologous mouse anti-
bodies bind to huIgG, complement gets activated and maintains the disease. Also, the acid-eluted heterologous huIgG alone might be incapable of inducing or insufficient to induce the mouse complement system, with complement activation in this case entirely depending on the presence of large amounts of mIgG. Several studies describe direct effects of antibodies on molecular processes that are potentially relevant in the pathogenesis of MN. In the rat model of Heymann nephritis, anti-megalin antibodies directly affect podocyte function by preventing the uptake of apolipoproteins through megalin, resulting in apolipoprotein deposition in immune deposits (27). In alloimmune antenatal MN, different anti-NEP antibody subclasses have different potencies in inhibiting enzymatic NEP activity in vitro (9). Moreover, in a case of IgG4-related disease that was associated with MN, it was demonstrated that the binding of circulating IgG4 anti-carbonic anhydrase II antibodies to the membrane of cultured podocytes induced a podocyte stress reaction with externalization of SOD2 (28). Given the surprising result that mice developed significant proteinuria in vivo as well, leading to cell damage and proteinuria.

In summary, the present study describes a patient with THSD7A-associated MN, whose rapid recurrence of MN after kidney transplantation in the presence of anti-THSD7A antibodies suggests a causative role of these antibodies in disease onset. We translated this clinical observation into animal studies and demonstrated that anti-THSD7A autoantibodies from patients with THSD7A-associated MN bound to podocyte-expressed THSD7A on foot processes in mice and induced proteinuria and the morphologic hallmarks of MN. The glomerular binding of huIgG was thereby THSD7A specific, demonstrating that it is in fact this antigen-antibody interaction that induced MN pathology in mice. We believe that our study uncovers the causative role of autoantibodies in the development of MN and supports the pathogenic concept of this disease.

Methods

Patients. The diagnosis of MN was made by renal biopsy. For animal experiments, sera were taken from 2 patients with THSD7A-associated MN and from healthy controls. All patients were without previous immunosuppressive therapy at the time of serum collection. Sera were investigated for anti-PLA1, anti-THSD7A antibodies by means of ELISA (29). Anti-THSD7A antibodies were detected using Western blotting, and titers were quantified by indirect immunofluorescence testing (IFT) as described previously for PLA1 (30) (test was provided by EUROIMMUN AG, Lübeck, Germany).

Immunohistochemical analyses. Paraffin sections (1-µm) of renal biopsies from a 57-year-old patient with MN and a healthy control patient were deparaffinized and rehydrated. Antigen retrieval was obtained for THSD7A by boiling in Dako antigen retrieval buffer, pH 9 (15 minutes at 98°C); for C4d by boiling in citrate buffer, pH 6.0 (30 minutes at 98°C); and for huIgG by protease XXIV treatment (5 µg/ml at 37°C for 15 minutes; Sigma-Aldrich). Nonspecific binding was blocked with 5% horse serum (Vector Laboratories) with 0.05% Triton X-100 (Sigma-Aldrich) in PBS for 30 minutes at room temperature (RT) prior to incubation at 4°C overnight with rabbit anti-THSD7A (1:400; Atlas; catalog HPA009293); anti-C4d (1:200; BIOZOL; catalog BI-RC4D); or anti-huIgG (1:200, Cy2-huIgG H+L; Dianova; catalog 709-225-149) in blocking buffer. Staining was visualized with a ZytochemPlus AP Polymer Kit (ZytoVision Systems) according to the manufacturer’s instructions. Nuclei were counterstained with hemalaun, and sections were mounted with gum arabic (Sigma-Aldrich). Negative controls were performed by omitting primary antibodies. Stainings were evaluated with an Axioskop using the Axiosvision software (both from Zeiss).

Animal care. WT male BALB/c mice (10- to 14-weeks of age) were bred in the animal facility at the University Medical Center Hamburg-Eppendorf. Animals had free access to water and standard animal chow.

Animal experiments. The pathogenicity of anti-THSD7A antibodies was evaluated in 2 experimental setups. In the first setup, anti-THSD7A antibody–positive sera from 2 patients with THSD7A-associated MN were used for disease induction, while serum from a healthy donor served as a negative control. In the second approach, affinity-purified anti-THSD7A antibodies and the remaining serum that was depleted of anti-THSD7A antibodies were used. Prior to injection, human sera were concentrated 2-fold using Amicon Ultra-15 centrifugal filters with a molecular cut-off of 100 kDa (EMD Millipore). Sera were then decomplexed by heating for 30 minutes at 56°C in a water bath. Debris was removed by centrifugation at 20,817 g for 15 minutes. As the sera from the 2 nephrotic patients contained subnormal levels of total proteins, the huIgG serum levels were quantified by SDS-PAGE and adjusted to equal levels. BALB/c mice were injected i.v. with 100 µl of adjusted sera for analysis after 2 hours and i.p. with 900 µl of adjusted sera for disease induction. Development of proteinuria was monitored using metabolic cages every 3 to 4 days for 2 weeks and then weekly. The histological images presented in the figures represent analyses of mice that were sacrificed at different time points (2 animals after 3 days, 3 animals after 7 days, 3 animals after 14 days, and 9 animals after 70 days). For the second experimental setup, anti-THSD7A antibodies were purified from 10 ml serum from a patient with THSD7A-associated MN and concentrated using Amicon Ultra-15 centrifugal filters with a molecular cut-off of 100 kDa to a final volume of 1 ml. Four male BALB/c mice were then i.v. injected with 250 µl affinity-purified anti-THSD7A antibodies. The remaining 8 ml of depleted serum was concentrated using Amicon Ultra-15 centrifugal filters with a molecular cut-off of 100 kDa to a final volume of 4 ml. Male BALB/c mice were then i.p. injected with 1 ml of depleted serum. Development of proteinuria was monitored as described above.

Immunofluorescence analyses. For immunolocalization of nephrin (guinea pig pAB, 1:100; Acris; catalog BP5030); laminin (rabbit pAB, 1:1,000; Sigma-Aldrich; catalog L9393); huIgG (Cy2 huIgG H+L, 1:200; Dianova; catalog 709-225-149); murine IgG (H+L Cy2 mIgG, 1:400; Dianova; catalog 715-225-151); complement C3 (FITC goat pAB, 1:100; Cappel; catalog 55500); or SOD2 (rabbit pAB, 1:100; Acris; catalog AP03023PU-S), 2-µm paraffin sections of normal or experimental mouse kidneys were deparaffinized and rehydrated with water. Antigen retrieval was obtained by boiling in citrate buffer, pH 9. Immunofluorescence analyses were performed as described previously for PLA1 (30) (test was provided by EUROIMMUN AG, Lübeck, Germany).
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6.1 (both 30 minutes at a constant temperature of 98°C) or by digestion
with protease XXIV (5 μg/ml; Sigma-Aldrich) for 15 minutes at 37°C.
Unspecific binding was blocked with 5% horse serum (Vector Labora-
tories) with 0.05% Triton X-100 (Sigma-Aldrich) in PBS for 30 min-
utes at RT prior to incubation at 4°C overnight with primary antibodies
in blocking buffer. Staining was visualized with fluorochrome-conju-
gated secondary antibodies (1:400; all affinity purified from Jackson
Immunoresearch Laboratories) for 30 minutes RT in 5% horse serum
with 0.05% Triton X-100. Nuclei were counterstained with DRAQ5
(1:1,000; Thermo Scientific; catalog 62252).

For indirect immunofluorescence using anti-THSD7A antibody-
positive sera or healthy control sera, 5-μm cryosections were fixed
with ice-cold 100% acetone for 10 minutes at −20°C. Unspecific
binding was blocked with 5% normal horse serum containing 0.05%
Triton X-100 for 30 minutes at RT. Sera were diluted at 1:250 and
incubated overnight at 4°C in blocking buffer concomitantly with anti-
nephrin antibody (1:100). Autoantibody binding was visualized using
Cy2 anti-hulG, and nephrin was visualized using Cy3 anti-guinea pig
(1:200; Dianova; catalog 706-165-148) for 30 minutes in blocking buf-
erg. Nuclei were counterstained with DRAQ5 (1:1,000).

For visualization of autoantibody binding following i.v. injection
of sera, 5-μm cryosections were fixed with ice-cold 100% acetone for
10 minutes at −20°C. Unspecific binding was blocked with 5% nor-
mal horse serum containing 0.05% Triton X-100 for 30 minutes at RT.
Autoantibody binding was visualized using hulgG4 (mouse mAb,
1:1,000; Binding Site; catalog MC011) incubated overnight at 4°C in
blocking buffer concomitantly with anti-nephrin antibody (1:100).
Bound autoantibodies were visualized using Cy2 anti-mouse IgG and
nephrin using Cy3 anti-guinea pig for 30 minutes in blocking buffer.
Nuclei were counterstained with DRAQ5 (1:1,000).

All negative controls were performed by omitting primary anti-
body solutions. Stainings were evaluated with a confocal LSM 510 meta
microscope and LSM software (both from Zeiss).

Generation of antibodies against native human THSD7A. A rabbit
antisera (K1405) recognizing native human and mouse THSD7A
was produced by cDNA immunization essentially as described pre-
viously (31). An expression construct encoding full-length human
THSD7A was conjugated to 1-μm gold particles (Bio-Rad Laborato-
ries). These were ballistically injected into a rabbit at the antibody core
unit of the University Medical Center Hamburg-Eppendorf. The rabbit
received 4 immunizations in 3- to 6-week intervals, each with 12 shots
of plasmid-conjugated gold particles (1 μg DNA/mg gold per shot).
Serum was obtained 3 weeks after the last DNA immunization. The
specificity of the antisera for native human THSD7A was verified
using immunofluorescence staining of Chinese hamster ovary (CHO)
cells with serial dilutions of the antisera 24 hours after transient
cotransfection of cells with expression constructs for nuclear GFP
and full-length THSD7A and by Western blotting with native human
THSD7A from glomerular extracts, recombinant human THSD7A,
and recombinant mouse THSD7A. Bound antibodies were detected
with phycoerythrin-conjugated anti-rabbit IgG (Dianova). Serum
K1405 was used at a dilution of 1:400 for staining of frozen kidney
sections and transfected HEK293 cells (Figure 3B, Figure 6C, and Sup-
plemental Figures 4 and 10). For FACS analyses, K1405 anti-THSD7A
antibody was labeled with AF647 using the Alexa Fluor 647 antibody
labeling kit according to the manufacturer’s instructions and was used
at a dilution of 1:400 (Supplemental Figure 11, B and C).

Electron microscopy. Electron microscopic analyses were performed
on kidneys that were fixed in 4% buffered paraformaldehyde. Tissue
was postfixed with 1% osmium in 0.1 M sodium cacodylate buffer, stained
with 1% uranyl acetate, and embedded in epoxy resin (Serva). Ultrathin
sections were cut (Ultramicrotome; Reichert Jung) and contrasted with
uranyl acetate in methanol, followed by lead citrate. Micrographs were
generated with a transmission electron microscope (JEM 1010; JEOL).

Immunogold electron microscopy. Fixation for immunogold labeling:
The fixation procedure and preparation of Bouin’s solution for immuno-
gold staining were modified according to Somogyi and Takagi (32). Ani-
mals were transcardially perfused with 0.1 M phosphate buffer (PB), pH
7.4, at 4°C. The fixative consisted of 500 ml 0.2 M PB, pH 7.4, 150 ml sat-
urated picric acid (Bouin’s solution: 0.044 M picric acid, 5% [v/v] acidic
acid, and 10% [v/v] formaldehyde in distilled water); 346.8 ml of 4% para-
formaldehyde solution; and 3.2 ml of 25% glutaraldehyde at a final
pH of 7.4. The final concentrations were 4% paraformaldehyde, 0.08%
glutaraldehyde, and 0.15% picric acid. Kidneys were removed after the
perfusion procedure and postfixed in 100 ml of 0.1 M PB including 4%
paraformaldehyde and 0.15% picric acid for 24 hours at 4°C. Vibratome
sections (Leica VT 1000S) of fixed tissue were made at 60-μm thick-
ness for areas of interest, and the slices were placed into cryoprotection
solution containing 0.1 M PB, pH 7.4, 30% sucrose, 1% polyvinylpyrroli-
done, and 30% ethyleneglycol. Pre-embedding immunogold protocol:
To remove the cryoprotection solution, the 60-μm slices were washed
twice in 0.1 M sodium phosphate buffer (PBS, pH 7.4) for 20 minutes,
followed by washing of the slices in 0.1 M PBS with 10% sucrose for
30 minutes and 0.1 M PBS with 20% sucrose for another 30 minutes.
Tissue was transferred to a drop of PBS with 20% sucrose and frozen
briefly with nitrogen and then thawed to RT. Subsequently, the tissue
was washed 3 times in PBS, pH 7.4, for 10 minutes. Blocking reagent
solution (BRS) was prepared with 100 ml PBS; 800 mg BSA (Sigma-
Aldrich; catalog A7638-5G); 250 μl 40% cold water fish skin gelatin
(AURION; CFWSG 900.033); and 162.5 mg sodium acid. To increase
the specificity of the primary antibody reaction, BRS supplemented with
2% normal goat serum (NGS) (Sigma-Aldrich) was used for 1 hour for
the blocking reaction. The primary antibody was dissolved in BRS over-
night at 4°C. After 24 hours, the tissue was washed in PBS 3 times for
5 minutes each time and incubated with 12 nm gold-conjugated anti-
body solution (1:50) in BRS for 2 hours. Subsequently, the tissue was
washed 3 times in PBS, pH 7.4, for 5 minutes each time, followed by
1% glutaraldehyde solution in 0.1 M PB for 10 minutes. The tissue was
dissolved in 1% OsO4 in distilled water at 4°C for 10 minutes, followed
by dehydration in an ethanol series. Subsequently, the probes were
transferred twice into propylene oxide for 5 minutes and embedded in
a mixture of epon and propylene oxide (4:1) overnight. Tissue samples
were cut with an ultrathin microtome (Ultracut; Reichert Jung), and thin
sections were inspected in a Phillips CM 100 electron microscope. Pic-
tures were taken using iTEM 5.2 software (Quemesa, Olympus).
The primary antibodies used were: human anti-THSD7A antibody—contain-
ing or control sera (1:100) and rabbit anti-THSD7A (1:100; Atlas; catalog
HPA000923) in blocking reagent solution. The following gold-conju-
gated secondary antibodies were used: 12 nm gold-AffiniPure goat anti-
hulG (H+L) antibody (Jackson ImmunoResearch; catalog 109-205-
088) and 12 nm gold-AffiniPure donkey anti-rabbit IgG (H+L) antibody
(Jackson ImmunoResearch; catalog 711-205-152).

Immunoblotting. Protein samples were prepared for Western blot
analysis by addition of 5× Laemmli buffer (1.5 M Tris-HCl, pH 6.8,
the secondary antibody (1:40,000; HRP goat anti-mouse albumin). Diluted urine was incubated for 60 minutes at RT. Following the washes, RT with 50 mM Tris, 0.14 M NaCl, and 1% BSA, pH 8.0, and rewashed, the plates were blocked for 30 minutes at 0.5% dry milk in PBS-T and HRP-conjugated secondary antibodies in PBS-T alone. Membranes were washed 3 times for 5 minutes in PBS-T after incubation with primary and secondary antibodies.

If serum was used as the primary antibody, a dilution of 1:100 was used. For specific detection of THSD7A, we used a rabbit polyclonal antibody (1:1,000; Atlas). The secondary antibodies used were: HRP-conjugated goat anti-mouse IgG, goat anti-huIgG, and goat anti-rabbit IgG (all 1:20,000 and all from SouthernBiotech; catalogs 1010-05, 2040-05, and 4010-05, respectively).

### Immunoprecipitation

For the isolation of glomeruli from BALB/c mice, mice were perfused with Dynabeads (Dynal) as described previously (33). MGEs (50 μg) were mixed with 5 times the volume of 20 mM Tris, pH 7.4, 140 mM NaCl, 1 mM CaCl₂, and 5 μl serum from a patient with THSD7A-associated MN or from a healthy donor. Samples were incubated 2 hours at 4°C on a rotator. Next, 20 μl Protein G Affinity Matrix (GeneScript) was added, and samples were incubated overnight at 4°C on a rotator. Samples were then centrifuged for 1 minute at low speed, and the supernatant was discarded. The pellet was washed 3 times with 20 mM Tris, pH 7.4, 140 mM NaCl, 0.8 mM CaCl₂, 0.02% SDS, 0.2% NP-40, and 0.1% Na-DOC and subsequently resuspended in Laemmli buffer containing β-mercaptoethanol, loaded on polyacrylamide gels, and electrophoresed. Immunoprecipitates were detected with the rabbit polyclonal anti-THSD7A antibody (Atlas), as described above.

### Antibody elution

Sixty to seventy 20-μm sections from frozen renal tissue were resuspended in 1 ml sterile PBS and centrifuged at low speed for 1 minute. The pellet was then washed 3 times in 1 ml PBS. After the last wash, the pellet was resuspended in 150 μl of 25 mM citrate, pH 3.2, and left on ice for 20 minutes, with intermediate shaking. Subsequently, the sample was centrifuged again, and the supernatant was added to 150 μl of 1 M Tris, pH 8. The pellet was resuspended in 150 μl of 25 mM citrate, pH 2.5, and left on ice for 20 minutes, with intermediate shaking. The sample was then centrifuged for 5 minutes at 20,817 g at 4°C. Next, the supernatant was added to the first elution and diluted to 2 ml using 0.05% dry milk in PBS-T. The resulting sample was used as the primary antibody in immunoblot analyses, as described above.

### Albumin ELISA

Urine albumin content was quantified using a commercially available ELISA system (Bethyl) according to the manufacturer’s instructions. Briefly, 96-well plates were coated 1:100 with goat anti-mouse albumin in binding buffer (0.05 M carbonate-bicarbonate, pH 9.6) for 60 minutes at RT. After washes in 50 mM Tris, 0.14 M NaCl, and 0.05% Tween-20, pH 8.0, the plates were blocked for 30 minutes at RT with 50 mM Tris, 0.14 M NaCl, and 1% BSA, pH 8.0, and rewashed. Diluted urine was incubated for 60 minutes at RT. Following the washes, the secondary antibody (1:40,000; HRP goat anti-mouse albumin) was applied for 60 minutes at RT. After the washes, enzyme substrate included in the kit was added, and the color development was stopped after 5 minutes with 2 M sulphuric acid. Extinction was measured at 450 nm in an ELISA plate reader (BioTek; ELX808). The urinary albumin concentration was calculated according to the formula for absorption: (A – D)/(1 + (x/C))² + D, where A and D are values from the standard curve. Regression values for the standard curve were calculated to assess the accuracy of the measured values. Standard curves with r values greater than 0.9950 were used. The urinary albumin values were standardized against urinary creatinine values of the same sample.

### Purification of human THSD7A and anti-THSD7A antibodies

Human DDK-tagged THSD7A (huTHSD7A) was expressed in HEK293 cells as described above. Forty-eight hours after transfection, cells were scraped in 50 mM Tris, pH 7.4, 150 mM NaCl, 1 mM EDTA, 1% Triton X-100, and protease inhibitor cocktail. Cells were sonicated shortly, incubated 1 hour at 4°C with slow rotation, and then centrifuged for 10 minutes at 20,000 g. A 2-ml chromatographic column was rinsed several times with PBS before addition of 1 ml anti-Flag M2 affinity gel (Sigma-Aldrich). The resin was equilibrated with a total of 15 ml PBS containing 0.2 mg/ml HSA and 1% Triton X-100. Subsequently, the cell lysate containing DDK-tagged huTHSD7A was loaded on the column. The flow-through was collected and loaded again. The column was then washed with 10 ml PBS containing 0.2 mg/ml HSA and 1% Triton X-100. Next, bound huTHSD7A was eluted in 3 steps with 2.25 ml each of 100 mM glycine, pH 3.5, containing 0.2 mg/ml HSA and 1% Triton X-100. In order to change the buffer, eluates were loaded onto PD-10 columns (GE Healthcare), the flow-through was discarded, and the columns were rinsed with 3 ml PBS containing 0.2 mg/ml HSA and 1% Triton X-100. Purification was verified using Western blot analysis, and purified THSD7A was quantified by Coomassie blue staining. Since approximately 95% of huTHSD7A was in the first 2 eluates, the eluates were pooled and concentrated using Amicon Ultra-15 centrifugal filters with a molecular cut-off of 100 kDa (EMD Millipore).

For purification of anti-THSD7A antibodies from serum from a patient with THSD7A-associated MN, we coupled purified huTHSD7A protein to beaded agarose using the AminoLink Plus Immobilization Kit (Thermo Scientific) according to the manufacturer’s instructions, with some modifications. Briefly, 250 μl purified huTHSD7A was mixed with 1 ml coupling buffer and loaded on a chromatography column. After addition of 20 μl sodium cyanoborohydride solution, the column was stored overnight at 4°C with gentle agitation. Subsequently, the column was washed twice with 1 ml quenching buffer, 4 times with 1 ml wash solution, and 3 times with 1 ml storage buffer. The column was then rinsed twice with PBS, and 10 ml serum containing anti-THSD7A antibodies was loaded onto the column. The flow-through was taken and loaded again. The column was then washed with 10 ml PBS containing 2% mouse serum. Elution was performed 2 times with 2.25 ml of 100 mM glycine, pH 3.5. Subsequently, the column was washed again with 10 ml PBS containing 2% mouse serum, and the flow-through from the first round was loaded again 2 times with subsequent elution. This procedure was repeated 3 times, resulting in 1 final flow-through devoid of anti-THSD7A antibodies (herein referred to as depleted serum) and a total of 6 eluates (herein referred to as E1–E6) containing the pulled anti-THSD7A antibodies. The eluate buffer was changed to PBS only using PD-10 columns as described above. Aliquots were taken at all steps of the experiment for Coomassie blue staining and Western blot analysis. Finally, eluates
were pooled and concentrated using Amicon Ultra-15 centrifugal filters with a molecular cut-off of 100 kDa (EMD Millipore). The reactivity of purified anti-THSD7A antibodies with THSD7A was tested using different dilutions in the Western blot analysis, compared with whole serum reactivity, and adjusted to equal amounts.

**Cell culture, cell transfection, and in vitro experiments.** HEK293 cells were cultured in DMEM supplemented with 10% FCS, 100 U/ml penicillin, and 100 mg/ml streptomycin (all from Invitrogen) at 37°C and 5% CO₂. cDNAs coding for human and murine THSD7A cloned into the pCMV6 entry vector containing a DDK tag were purchased from OriGene. HEK293 cells were transiently transfected using the calcium phosphate method. For each 10-cm dish of cultured cells, we used 15 μg plasmid DNA mixed with 36 μl of 2 M CaCl₂ and diluted with sterile water up to a volume of 300 μl. This solution was gently mixed with an equal volume of 2× HEPES-buffered saline (HBS) (275 mM NaCl, 55 mM HEPES, pH 7.0) and incubated for 30 minutes at RT. The resulting solution was added drop-wise to the cells. The medium was changed to serum-free medium 24 hours after transfection. For quantification and FACS analyses of detached cells, cell culture medium was collected 60 minutes after treatment and centrifuged at 500 g for 5 minutes to sediment the cell pellets. The cell pellets were resuspended in PBS and centrifuged again. For quantification of detached cells, cell pellets were then lysed in 50 mM Tris, pH 7.4, 150 mM NaCl, 0.1% SDS, 1% NP-40, and 0.5% Na-DOC supplemented with protease inhibitor cocktail as described above. Protein concentration was measured using the Pierce BCA protein assay kit according to the manufacturer’s instructions. Cell detachment experiments were conducted 10 times, each time with 2% of both anti-THSD7A antibody-positive sera, MN 1 and MN 2, and serum from a healthy donor. Experiments were conducted independently 3 times with 5 different sera from patients with anti-PLA2R1 antibody–positive MN and 5 different sera from patients with anti-THSD7A and anti-PLA2R1 antibody-negative MN. Additionally, experiments were performed 5 times with affinity-purified anti-THSD7A antibodies, serum depleted of anti-THSD7A antibodies (2%), and affinity-purified IgG from a healthy donor. The amount of purified anti-THSD7A antibody was adapted to the anti-THSD7A antibody–containing whole serum by analysis of reactivity in Western blot analysis. For FACS analyses, cells were resuspended in MACS buffer (Miltenyi Biotec) and stained with AF647 anti-THSD7A (1:400) and with Cy2 anti-huIgG (1:200) for 20 minutes at 4°C. For the detection of cell death, the LIVE/DEAD kit (Thermo Fisher Scientific) was used according to the manufacturer’s instructions. Experiments were performed using a BD LSR II Flow Cytometer and analyzed with FlowJo software. FACS experiments were conducted 3 independent times with MN serum 1.

**Phase-contrast micrographs** were taken with an Axiosvert 40 microscope using Axiovision software (both from Zeiss). For immunofluorescence analyses, HEK293 cells were fixed with ice-cold 100% methanol for 10 minutes at −20°C or with 4% paraformaldehyde for 8 minutes at RT. Staining was performed as described for the frozen tissue sections, using Cy2 anti-huIgG (1:200; Dianova) and K405 anti-THSD7A antibody (1:400) for the visualization of THSD7A expression and autoantibody binding to HEK293 cells, and phalloidin-AF568 (1:200; Molecular Probes; catalog A12380) for visualization of cytoskeletal rearrangement.

**GEC culture.** For the isolation of glomeruli, mice were perfused with Dynabeads (Dynal) as described previously (33, 34). Decapsulated glomeruli were plated on collagen 1-coated 10-cm culture plates (Biocoat) in RPMI 1640 supplemented with 10% FCS, 1 mmol/l HEPES, 1 mmol/l sodium pyruvate, 100 U/ml penicillin, and 100 mg/ml streptomycin at 38°C and 5% CO₂. Three to five days after the initial glomerular isolation, primary culture GECs were challenged with 1:50 (2%) anti-THSD7A antibody–containing patient’s serum or serum from a healthy donor. Experiments were conducted independently 3 times with 2 anti-THSD7A antibody–containing sera (MN 1 and MN 2) and 2 different control sera. Moreover, cells were treated with affinity-purified anti-THSD7A antibodies, serum depleted of anti-THSD7A antibodies, and affinity-purified IgG from a healthy donor. These experiments were conducted independently 2 times (with 10 for each condition). Immunofluorescence analyses were performed 40–60 minutes after serum challenge. For immunofluorescence analyses, GECs were fixed with 4% paraformaldehyde for 8 minutes at RT. Staining was performed as described for the frozen tissue sections. The following primary antibodies were used: Cy2-coupled anti-huIgG (1:100; Dianova); α-actinin-4 (1:200; Immunoglob; catalog 0042-05); p-paxillin (Tyr118) (1:400; Thermo Fischer Scientific; catalog 44-722G); and phalloidin-AF568 (1:200; Molecular Probes).

**F-actin quantification** was performed using ImageJ software (NIH). For analysis, grayscale images from 5 independent experiments (with at least 4 per condition) were used at a 1,000-fold magnification. Individual cell areas were circled, and optical density was analyzed.

**Statistics.** Data are expressed as the mean ± SEM. For statistical analysis for comparison of mouse treatment groups, we performed a 2-tailed, nonparametric Mann-Whitney U test to enable robust conclusions on effect significance in the event of departures from normality associated with small sample sizes. The number of animals used for the experiments was limited by the availability of serum from patients with THSD7A-associated MN. Therefore, no statistical method was used to predetermine sample size. Replicates used were biological replicates, which were measured using different samples derived from distinct mice. All animals were WT mice and were blindly assigned to the experimental groups. No inclusion or exclusion criteria were defined for animals, and no animals were excluded from the analysis. Examiners were not blinded, except for those conducting the following in vitro analyses: serum treatment of GECs and transfected HEK293 cells for all experiments, phase-contrast microscopy of nondetached GECs and HEK293 cells after treatment, culture medium withdrawal for protein quantification of detached HEK293 cells and FACS analyses, protein quantification of detached cells, sample preparation for Western blot analyses, and all histological stainings. A P value of less than 0.05 was considered significant.

**Study approval.** All patients provided written, informed consent, and the study was approved by the local ethics committee and conducted according to Declaration of Helsinki guidelines. All animal experiments were performed according to national and institutional animal care and ethics guidelines and were approved by the veterinary agency and the local animal care committee of Hamburg, Germany (BSG Amt für Gesundheit und Verbraucherschutz, Lebensmittelsicherheit und Veterinärwesen).

**Author contributions** NMT conducted animal experiments, performed protein and antibody purification and in vitro studies, analyzed the data, and wrote the first draft of the manuscript. EH acquired and analyzed patients’ clinical data and measured autoantibody titers. ATR per-
formed FACS analyses. LF and GR performed immunogold electron microscopic studies. UH performed histologic analyses of human biopsy samples and conventional electron microscopy. JG, FB, and KB provided reagents and acquired clinical data. GZ provided reagents. FKN generated a rabbit antibody for the detection of THSD7A. GL designed some of the protein biochemical experiments. CMS designed and conducted animal experiments, generated GECs, performed confocal microscopy, and revised the manuscript. RAKS designed the study and finalized the manuscript.

Acknowledgments

These studies were supported by grants from the Deutsche Forschungsgemeinschaft, KFO 228 (STA-193/9-2) and the Sonderforschungsbereich 1192 (projects B1, B2, and B3). We would like to thank Wolfgang Schlumberger of EUROIMMUN AG (Lübeck, Germany) for providing the indirect immunofluorescence test (IFT), which allowed the measurement of THSD7A serum antibody titers in patients. We thank Ursula Knießler and Barbara Holstermann for performing the electron microscopic studies. We thank Eugene Kinzler, Samaneh Liagos, and Daniela Bergleiter for following the patients and collecting sera. We also thank Marlies Sachs, Marion Nissen, Silke Dehde, and Olga Ilin for their excellent technical assistance.

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