Susceptibility to Anti–Glomerular Basement Membrane Disease and Goodpasture Syndrome Is Linked to MHC Class II Genes and the Emergence of T Cell–mediated Immunity in Mice

Raghuram Kalluri, Theodore M. Danoff, Hirokazu Okada, and Eric G. Neilson
Penn Center for Molecular Studies of Kidney Diseases, University of Pennsylvania, Philadelphia, Pennsylvania 19104

Abstract

We developed a new mouse model of human anti–glomerular basement membrane (GBM) disease to better characterize the genetic determinants of cell-mediated injury. While all major histocompatibility complex (MHC) haplotypes (H-2a, k, s, b, and d) immunized with α3 NC1 domains of type IV collagen produce anti–α3(IV) NC1 antibodies that cross-react with human Goodpasture [anti-GBM/anti–α3(IV) NC1] autoantibodies, only a few strains developed nephritis and lung hemorrhage associated with Goodpasture syndrome. Crescentic glomerulonephritis and lung hemorrhage were MHC-restricted in haplotypes H-2s, b, and d (Aβ/αα region in H-2k) and associated with the emergence of an IL-12/Th1-like T cell phenotype. Lymphocytes or anti–α3(IV) NC1 antibodies from nephritogenic strains transfer disease to syngeneic recipients. However, passive transfer of isogenic α3(IV) NC1 antibodies into −/− T cell receptor–deficient mice failed to produce nephritis. Finally, nephritis and its associated IL-12/Th1-like T cell response attenuate in disease-susceptible mice tolerized orally to α3(IV) collagen before immunization. Our findings suggest collectively, as a new mouse model of human anti–glomerular basement membrane disease to better characterize the genetic determinants of cell-mediated injury.

Introduction

The gene for human α3(IV) collagen is found in the q35–37 region of chromosome 2 (1), and its protein largely replaces α1(IV) and α2(IV) collagens in the specialized basement membranes of the kidney during organ development (2). Chain mutations in α3(IV) collagen produce autosomal recessive Alport syndrome (3), and humans occasionally develop autoimmune responses to the α3(IV) NC1 resulting in anti–glomerular basement membrane (GBM) nephritis or Goodpasture syndrome (GPS) (4, 5). Evidence of genetic susceptibility to human anti-GBM disease is limited to studies in twins (6, 7), and to an association with MHC haplotypes mapping to the class II HLA-D region (8, 9). Anti-GBM antibodies and T cells from Goodpasture patients recognize epitopes found at the distal termini of the recombinant α3 NC1 domain of type IV collagen (10–13).

Human anti-GBM antibodies eluted from nephritic kidney are thought to be etiologic for disease because they passively transfer a heterologous nephritis to primates (14). However, the nature of the passive experiments precluded allotype-matching of anti-GBM antibodies for recipients, and, therefore, was not a strong test for antibody sufficiency in an autologous environment. Likewise, it has not been feasible to adoptively transfer human lymphocytes reactive with α3(IV) collagen to assess cell-mediated immunity as an independent variable.

Animal models over the last few decades have attempted to approximate human anti-GBM disease (15, 16), the optimal binding conditions for antibody (17, 18), and the role of accessory molecules (19–21) or amplification mechanisms like the activation of complement (22). Unfortunately, these studies were performed before much was known about the specificity of the antibodies in human anti-GBM disease (3, 4) or in animal models that only mimic a portion of the disease seen in patients (4, 16). Furthermore, the tradition of calling anti-GBM nephritis an antibody-mediated disease has tempered greatly work evaluating the likely role of cellular immunity in its pathogenesis.

Immunization with crude GBM, or transfer of heterologous antibodies to crude GBM, produces a form of anti-GBM disease (15, 22) in monkeys, sheep, rabbits, and rats, but gives mixed results in mice (23–25). Initiation of nephritis with crude antigen, of course, does not reflect the more focused response typical of the human condition, that of antibodies directed to a single antigen like the α3(IV) NC1 domain (4, 26). Surprisingly, however, earlier attempts to produce aggressive disease with α3(IV) NC1 collagen in rabbits were only partially successful (27). Perhaps this was because the outbred rabbits did not carry MHC haplotypes fully favoring progressive injury.

The mouse model reported here is a significant advance over previous work because it comes much closer to the variable presentation of human anti-GBM disease, uses a single protein as antigen, and, with the availability of inbred and recombinant mice, lends itself to the segregation of important immunogenetic traits.

This work was presented initially in abstract form at the National Meeting of the American Society of Nephrology in November, 1995.

Address correspondence to Eric G. Neilson, M.D., C. Mahlon Kline Professor of Medicine, 700 Clinical Research Building, University of Pennsylvania, 415 Curie Boulevard, Philadelphia, PA 19104

Received for publication 21 May 1997 and accepted in revised form 11 September 1997.

J. Clin. Invest. © The American Society for Clinical Investigation, Inc
0021-9738/97/11/2263/13 $2.00
Volume 100, Number 9, November 1997, 2263–2275
http://www.jci.org

1. Abbreviations used in this paper: ABM, alveolar basement membrane; GPS, Goodpasture syndrome; TcR, T cell receptor.

Murine Anti–Glomerular Basement Membrane Disease 2263
Methods

Immunoization of mice with α3(IV) NC1 domains. Inbred and recombinant mice SJL/J (H-2^k), C57BL/6J (B6; H-2^b), C57BL/10SnJ (B10; H-2^e), BALB/cJ (H-2^d), DBA/2J (H-2^d), A/J (H-2^s), B10.A/SvSnJ (H-2^s), AKR/J (H-2^s), CBA/J (H-2^s), A.TL/Sf-DvEg (H-2^s), B10.SJ/S (H-2^s), (BALB × A/J) F1, (BALB × DBA/2) F1, B10.BR/SvSnJ (H-2^s), and C57BL/d-TERb^b×Mon^b×Mom^b (H-2^d) T cell receptor (TCr)-deficient mice were purchased from The Jackson Laboratory (Bar Harbor, ME) at 4–6 wk of age. The TErrb^b×Mon^b×Mom^b TCr-deficient mice express neither αβ or γδ TCRs (28), and are designated TCr −/− in this report. The B10.S(9R) recombinants were obtained from Dr. Ronald Schwartz (National Institutes of Health, Bethesda, MD) through Contract Grant NO1-A0-52702, and CA21 (BALB/c-Igh^c; H-2^d) congenic mice were obtained from Dr. Roy Riblet (Molecular Biosystems, Inc., La Jolla, CA). (A/J × SJL) F1 and (BALB/c × CA21) F1 crosses were bred at the University of Pennsylvania. Four to six mice in each experimental group were immunized with 25 μg of α3(IV) NC1 collagen subcutaneously in CFA followed by a booster injection 3–4 wk later (29). Control mice were injected with CFA alone. Blood, kidneys, and lung tissues were collected from each mouse at time of killing. Tissues were either snap-frozen for antibody elution or immunolocalization, or fixed in 10% phosphate saline–buffered formalin for histopathology. Two mice from each group were placed randomly in metabolic cages for 16 h before time of killing to determine overnight urine protein excretion (29). A serum sample from each mouse was obtained for creatinine measurement (29) and anti-α3(IV) collagen antibodies by ELISA (26).

Preparation of the antigens. Bovine α3(IV) NC1 dimers were isolated from bovine testis (27, 30, 31). Bovine testis was digested with bacterial collagenase and the collagenase-solubilized material was resolved by gel filtration and reverse-phase HPLC, as described previously (27, 31). The isolated α3(IV) NC1 dimers were characterized by type IV collagen α chain–specific antibodies by ELISA and Western blot before use for the immunization of mice. Mouse NC1 hexamer was prepared as described for bovine and human NC1 hexamer (31).

Adoptive and passive transfer studies. For passive transfer experiments, 500 μl of undiluted serum from SJL mice with anti-GBM disease was injected intravenously into naive healthy SJL recipients. For adoptive transfer experiments, spleens and lymph nodes were removed from SJL mice immunized with α3(IV) NC1 collagen. Lymphocytes prepared from spleen and lymph nodes (32) were injected into naive SJL mice pretreated with cyclophosphamide (20 mg/kg) 48 h before the transfer. Each SJL recipient received 10^6 lymphocytes by intravenous injection. The control mice received lymphocytes from CFA-injected mice.

Histopathology and immunoﬂuorescence. Tissue fixed in 10% buffered formalin were sectioned and stained with periodic acid–Schiff (PAS) and/or hematoxylin and cosin. Frozen tissues were cryo-sectioned and stained with FITC-conjugated rabbit anti–mouse IgG, IgM, IgA, and C3 (Sigma Chemical Co., St. Louis, MO). All the sections were viewed blind in at least 10 fields using a microscope (Carl Zeiss, Inc., Thornwood, NY) with appropriate excitation filters (29). Tissues fixed in 10% buffered formalin were also stained with primary goat antibodies directed to II-12 and II-4 (R & D Systems, Inc., Minneapolis, MN) and developed with a secondary rabbit anti–goat antibody linked to horseradish peroxidase (33).

ELISA. For direct ELISA, the plates were coated in triplicate with 100 ng of antigen in 200 μl of coating buffer. The plates were incubated for 2 h at 37°C or overnight at room temperature. Upon coating, the plates were washed three times at intervals of 5 min with 0.15 M NaCl and 0.05% Tween 20 (washing buffer). After washing, the plates were blocked with 2% BSA in PBS for 30 min at 37°C. Upon blocking, the plates were washed again with the washing buffer and then incubated with primary antibody in appropriate dilution in PBS. The plates were incubated for 1 h at 37°C. After primary antibody incubation, the plates were washed again and subsequently incubated with the secondary antibody (anti-mouse IgG) conjugated to alkaline phosphatase at 1:5,000 dilution in PBS. The plates were incubated for 1 h at 37°C. Subsequently, the plates were washed again thoroughly, and substrate, disodium p-nitrophenyl phosphate (5 mg/ml), was added. After color development, the absorbance was measured using an ELISA autoreader at 405 nm. The native bovine and human recombinant antigens were prepared as described previously (26, 34).

Inhibition ELISA was performed as before (10, 11, 27, 34) with slight modification. The ELISA plates were coated with 100 ng of dissociated mouse NC1 hexamer in 6 M guanidine-HCl, 50 mM Tris-HCl, pH 7.5. Initially, dilution curves were obtained for the human Goodpasture and mouse antibodies. The plates were coated overnight at room temperature. Upon washing and blocking the plates, as described above for direct ELISA, saturating concentrations of Goodpasture or mouse antibodies were used in the assay. The plates were incubated overnight and developed as described above for direct ELISA.

Antibody elution from the tissue. Antibody elution from the kidneys and lungs was described previously (27). Briefly, the tissue was homogenized in the presence of protease inhibitors and washed thoroughly with distilled water. The resultant pellet was subjected to extraction with 0.1 M glycine, 0.5 M NaCl, pH 2.8. The extracted material was dialyzed extensively against 10 mM PBS and subsequently precipitated with 40% ammonium sulfate. The resultant pellet was solubilized in suitable volume of 10 mM PBS for future experiments.

Oral feeding of protein fraction containing α3(IV) NC1 domain. Mice were gavaged either with 65 or 700 μg of BSA or α3(IV) NC1 dimer–enriched type IV collagen (principally α3, but likely also containing small amounts of α1, α4, and α5) in sterile PBS distributed over 5 alternate day feedings for 10 d before immunization with the α3(IV) NC1 domains in adjuvant. The dosages and regimen of feeding were extrapolated from similar studies using other mouse models for autoimmune disease (35, 36). The mice were immunized with 25 μg of antigen in CFA and boosted 3 wk later with the same amount in incomplete Freund’s adjuvant. 2 mo later, the mice were evaluated for renal function, and their blood and tissue were harvested for serological analysis and histopathology.

Results

Mice in this study were immunized with the GPS antigen in adjuvant (CFA) made from purified bovine α3(IV) NC1 collagen (27, 30) and the resulting immune response produced anti-GBM antibodies cross-reactive with human antibodies from Goodpasture patients, lung hemorrhage, crescentic glomerulonephritis, and progressive renal failure in susceptible haplotypes.

Characterization of anti-GBM disease in mice. 8 different inbred strains of mice reflecting a diversity of MHC haplotypes were selected for immunization with α3(IV) NC1 domains (Table 1). 12 wk later, SJL mice (H-2^b) emerged as strong nephritic responders, with the highest urine protein, followed by the moderate responders, B6 (H-2^b), BALB/c (H-2^b), and DBA/2 (H-2^b). A/J (H-2^b), AKR (H-2^b), and CBA (H-2^b) were nonnephritic and had normal levels of urine protein. The levels of renal failure as measured by serum creatinine in all groups reflected the trends set by the urine protein data; the nonnephritic mice had normal levels.

All mice injected with α3(IV) NC1 domains developed high titers of circulating anti-α3(IV) NC1 antibodies regardless of whether they developed cellular infiltrates and renal injury, suggesting that the intensity of the antibody response is not a selective determinant of susceptibility to disease (Fig. 1A). The circulating, kidney-, and lung-bound antibodies from SJL mice were quite specific for native bovine and recombi-
nont human α3(IV) NC1 domain (Fig. 1, B and C). Antibodies from A/J mice also had some minor additional reactivity to α1 and α2(IV) NC1 domains of type IV collagen (data not shown).

Histopathological examination of kidneys from nephritic mice revealed severe crescentic glomerulonephritis with considerable tubulointerstitial disease (Table I and Fig. 2). SJL mice with anti-GBM disease had full to partial crescents in 45% of their glomeruli, compared with none in the nonresponder mice, and 20–22% in B6 and DBA/2 mice (Table I). In the SJL mice, most of the glomeruli revealed cellular infiltrates and shrinkage. Mononuclear infiltrates were evident in glomeruli and in the renal interstitium (Fig. 2 A). Lung sections from these mice also contained focal to massive hemorrhage with extravasation of red blood cells into alveolar air spaces (Fig. 2 B). Kidney and lung sections from A/J, AKR, and CBA mice appeared normal and comparable to CFA controls.

Cell-mediated susceptibility genes permissive for anti-GBM disease. Age-matched recombinant and congenic mice were used to characterize further the role of individual polymorphisms at class I and class II MHC loci for their effect on the expression of murine anti-GBM disease (Table II).

8 wk after immunization, A/J, A.TL, B10.A, and B10.BR mice did not have significant increases in urine protein or serum creatinine levels, compared with CFA-injected control mice (Table II). Histopathological evaluation of kidneys and lungs in these animals also revealed no tissue abnormalities. However, at the same time point, SJL, B6, B10, B10.S, B10.S(9R), and A.SW mice demonstrated an increase in urine protein (Table II). Serum creatinine was also elevated when analyzed 12 wk after initial immunization. Kidney and lung tissue revealed varying degrees of crescentic glomerulonephritis with tubulointerstitial disease and linear binding of IgG on the GBM.

The presence or absence of glomerulonephritis among B10, B10.S, B10.S(9R), B10.BR, and B10.A recombinant mice indicates that glomerular inflammation is linked to a subregion of the MHC; differences among SJL, A.SW, and A/J mice were confirmatory. Comparisons of BALB/c, A/J, B10.K, and B10.A mice suggest that the class I/D region of the mouse H-2 is not relevant. The differences in disease among SJL, A.SW,

### Table I. Mice Immunized to Produce Anti-GBM Disease

<table>
<thead>
<tr>
<th>Strain*</th>
<th>H-2</th>
<th>K</th>
<th>Aβ</th>
<th>Aα</th>
<th>Eβ</th>
<th>Eα</th>
<th>D</th>
<th>UP</th>
<th>sCr</th>
<th>α3(IV) Ab</th>
<th>cGN</th>
<th>TIN</th>
<th>IF-Kidney</th>
<th>IF-Lung</th>
</tr>
</thead>
<tbody>
<tr>
<td>SJL</td>
<td>s</td>
<td>s</td>
<td>s</td>
<td>s</td>
<td>–</td>
<td>–</td>
<td>s</td>
<td>16.5±6.3</td>
<td>4.20±1.6</td>
<td>+</td>
<td>45</td>
<td>4</td>
<td>+</td>
<td>–</td>
</tr>
<tr>
<td>BALB/c</td>
<td>d</td>
<td>d</td>
<td>d</td>
<td>d</td>
<td>d</td>
<td>d</td>
<td>d</td>
<td>6.70±1.3</td>
<td>3.00±1.1</td>
<td>+</td>
<td>12</td>
<td>3</td>
<td>+</td>
<td>–</td>
</tr>
<tr>
<td>DBA/2</td>
<td>d</td>
<td>d</td>
<td>d</td>
<td>d</td>
<td>d</td>
<td>d</td>
<td>d</td>
<td>2.11±1.1</td>
<td>3.20±1.4</td>
<td>+</td>
<td>22</td>
<td>2</td>
<td>+</td>
<td>–</td>
</tr>
<tr>
<td>B6</td>
<td>b</td>
<td>b</td>
<td>b</td>
<td>b</td>
<td>–</td>
<td>–</td>
<td>b</td>
<td>5.00±2.1</td>
<td>3.20±1.3</td>
<td>+</td>
<td>20</td>
<td>2</td>
<td>+</td>
<td>–</td>
</tr>
<tr>
<td>CBA</td>
<td>k</td>
<td>k</td>
<td>k</td>
<td>k</td>
<td>k</td>
<td>k</td>
<td>k</td>
<td>1.90±0.8</td>
<td>1.20±0.8</td>
<td>+</td>
<td>&lt; 0.5</td>
<td>0</td>
<td>+</td>
<td>–</td>
</tr>
<tr>
<td>A/J</td>
<td>a</td>
<td>k</td>
<td>k</td>
<td>k</td>
<td>k</td>
<td>k</td>
<td>k</td>
<td>1.20±0.5</td>
<td>0.65±0.4</td>
<td>+</td>
<td>&lt; 0.5</td>
<td>0</td>
<td>+</td>
<td>–</td>
</tr>
<tr>
<td>AKR</td>
<td>k</td>
<td>k</td>
<td>k</td>
<td>k</td>
<td>k</td>
<td>k</td>
<td>k</td>
<td>0.4±0.0</td>
<td>0.85±0.2</td>
<td>+</td>
<td>&lt; 0.5</td>
<td>0</td>
<td>+</td>
<td>–</td>
</tr>
<tr>
<td>Controls</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td>1.00±0.8</td>
<td>0.90±0.7</td>
<td>–</td>
<td>0</td>
<td>0</td>
<td>–</td>
<td>–</td>
</tr>
</tbody>
</table>

* n = 4–12 mice in each group; controls include 2 mice from each strain. mRNA for Eα in SJL and B6 mice is not transcribed, hence the Eα/Eβ heterodimer protein is also not expressed on cell surface. Protein and serum creatinine measurements were made separately for each group. UP, Urine protein (27); sCr, Serum creatinine (29); cGN, Crescentic glomerulonephritis, as measured by percent fully developed crescentic glomeruli. TIN, Tubulointerstitial nephritis, estimated by scale (29); 4 is highest and 0 is absent. IF, Immunofluorescence (27).

### Table II. MHC Recombinant Mice Immunized to Produce Anti-GBM Disease

<table>
<thead>
<tr>
<th>Strain*</th>
<th>H-2</th>
<th>K</th>
<th>Aβ</th>
<th>Aα</th>
<th>Eβ</th>
<th>Eα</th>
<th>D</th>
<th>UP</th>
<th>sCr</th>
<th>α3(IV) Ab</th>
<th>cGN</th>
<th>TIN</th>
<th>IF-Lung</th>
</tr>
</thead>
<tbody>
<tr>
<td>SJL</td>
<td>s</td>
<td>s</td>
<td>s</td>
<td>s</td>
<td>–</td>
<td>–</td>
<td>s</td>
<td>10.8±2.8</td>
<td>3.12±1.2</td>
<td>+</td>
<td>29</td>
<td>3</td>
<td>+</td>
</tr>
<tr>
<td>A.SW</td>
<td>s</td>
<td>s</td>
<td>s</td>
<td>s</td>
<td>–</td>
<td>–</td>
<td>s</td>
<td>9.00±1.7</td>
<td>2.51±1.4</td>
<td>+</td>
<td>16</td>
<td>2</td>
<td>+</td>
</tr>
<tr>
<td>A.TL</td>
<td>t</td>
<td>s</td>
<td>k</td>
<td>k</td>
<td>k</td>
<td>d</td>
<td>1.18±0.9</td>
<td>0.85±0.5</td>
<td>+</td>
<td>6</td>
<td>0</td>
<td>–</td>
<td>–</td>
</tr>
<tr>
<td>A/J</td>
<td>a</td>
<td>k</td>
<td>k</td>
<td>k</td>
<td>k</td>
<td>k</td>
<td>d</td>
<td>1.32±1.1</td>
<td>0.91±0.6</td>
<td>+</td>
<td>2</td>
<td>0</td>
<td>–</td>
</tr>
<tr>
<td>B10</td>
<td>b</td>
<td>b</td>
<td>b</td>
<td>b</td>
<td>–</td>
<td>–</td>
<td>b</td>
<td>4.90±1.3</td>
<td>1.85±0.7</td>
<td>+</td>
<td>19</td>
<td>1.5</td>
<td>+</td>
</tr>
<tr>
<td>B10.(9R)</td>
<td>t</td>
<td>s</td>
<td>s</td>
<td>s</td>
<td>s</td>
<td>k</td>
<td>d</td>
<td>7.30±1.5</td>
<td>1.71±0.1</td>
<td>+</td>
<td>10</td>
<td>1.5</td>
<td>+</td>
</tr>
<tr>
<td>B10.BR</td>
<td>k</td>
<td>k</td>
<td>k</td>
<td>k</td>
<td>k</td>
<td>k</td>
<td>d</td>
<td>1.50±0.1</td>
<td>0.61±0.2</td>
<td>+</td>
<td>3</td>
<td>1</td>
<td>–</td>
</tr>
<tr>
<td>B10.A</td>
<td>k</td>
<td>k</td>
<td>k</td>
<td>k</td>
<td>k</td>
<td>k</td>
<td>d</td>
<td>1.46±0.1</td>
<td>0.81±0.0</td>
<td>+</td>
<td>5</td>
<td>0</td>
<td>–</td>
</tr>
<tr>
<td>Controls</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td>1.40±1.1</td>
<td>0.80±0.7</td>
<td>–</td>
<td>0</td>
<td>0</td>
<td>–</td>
</tr>
</tbody>
</table>

* n = 4 mice in each group; controls include 2 mice from each strain. mRNA for Eα in SJL and B6 mice is not transcribed, hence the Eα/Eβ heterodimer protein is also not expressed on cell surface. Protein and serum creatinine measurements were made separately for each group. UP, Urine protein (27); sCr, Serum creatinine (29); cGN, Crescentic glomerulonephritis, as measured by percent fully developed crescentic glomeruli. TIN, Tubulointerstitial nephritis, estimated by scale (29); 4 is highest and 0 is absent. IF, Immunofluorescence (27).
and A.TL mice indicate further that the class I/K region is also not important. Therefore, it is quite likely that some Aβ/Aα haplotypes are linked to important susceptibility genes mediating cellular reactions in glomeruli and the interstitium, since the E-region gene products of MHC class II are not expressed in H-2^s,b haplotypes (37), and since A.SW, B10.S, and B10.S(9R) mice are susceptible to anti-GBM disease, whereas A.TL mice are not.

*Immunofluorescent staining for immunoglobulin in tissues.* Linear binding of IgG along the GBM and alveolar basement membrane (ABM) of immunized animals was observed by immunofluorescent staining on frozen kidney and lung sections (Fig. 2, C and D, respectively). Control SJL mice appeared normal, with no overt histopathological changes or specific IgG binding in the kidney or lung. IgA and IgM staining in the GBM and ABM was not noted in either the experimental or

---

**Figure 1.** Evaluation of humoral response in α3(IV) collagen–immunized mice. (A) Binding curves for circulating anti–α3(IV) NC1 collagen antibodies from four different inbred mice. The coating antigen is 100 ng bovine α3(IV) NC1 collagen (immunogen). (B) Specificity of circulating, kidney-, and lung-bound antibodies to the bovine α3(IV) NC1 collagen. (C) Specific binding of the same antibodies as in B to recombinant human type IV collagen NC1 domain. The antibodies bind specifically to the α3(IV) NC1 recombinant human antigen. Significant binding to other domains is not observed. The control antibodies do not reveal binding to any of the recombinant domains. In B and C, the antigen is coated at a concentration of 100 ng/well. The antibody dilution used in B and C was 1:200.
control mice (data not shown). C3 staining was observed as patchy immunofluorescence along the GBM of SJL mice with anti-GBM disease, as well as in A/J mice without glomerular pathology. B6 and DBA/2 mice also presented with similar immunofluorescent staining in the kidney as SJL mice, in spite of their less intense renal disease (data not shown).

Kidney sections from A/J, AKR, and CBA mice that did not develop cellular injury demonstrated the same linear bind-
Digestion to extract the NC1 domain of type IV collagen. 15 μg of the digest was analyzed by SDS-PAGE and immunoblotting with human Goodpasture antisera (31). These antibodies recognize specifically the α3(IV) NC1 domain (Goodpasture antigen). D and M. The dimers and monomers, respectively, of NC1 domain that are recognized by the Goodpasture antibodies (31). The molecular masses of the dimers and monomers are ~54 and 28 kD, respectively. Lanes 1 and 2 denote SJL, lanes 3 and 4 denote DBA/2, lanes 5 and 6 denote A/J, lanes 7 and 8 denote (A/J × SJL) F1. Lanes 1, 3, and 5 are kidney NC1, and lanes 2, 4, 6, and 8 are lung NC1. The two bands in the monomer region could represent two size isoforms of α3(IV) NC1 monomer (67).

Immunoblot analysis of Goodpasture antigen [α3(IV) collagen] in the kidney and lung tissue from mice. Lungs from eight naive mice of each inbred strain were used for bacterial collagenase digestion to extract the NC1 domain of type IV collagen. 15 μg of the digest was analyzed by SDS-PAGE and immunoblotting with human Goodpasture antisera (31). These antibodies recognize specifically the α3(IV) NC1 domain (Goodpasture antigen). D and M. The dimers and monomers, respectively, of NC1 domain that are recognized by the Goodpasture antibodies (31). The molecular masses of the dimers and monomers are ~54 and 28 kD, respectively. Lanes 1 and 2 denote SJL, lanes 3 and 4 denote DBA/2, lanes 5 and 6 denote A/J, lanes 7 and 8 denote (A/J × SJL) F1. Lanes 1, 3, and 5 are kidney NC1, and lanes 2, 4, 6, and 8 are lung NC1. The two bands in the monomer region could represent two size isoforms of α3(IV) NC1 monomer (67).

Immunoblot analysis of Goodpasture antigen [α3(IV) collagen] in the kidney and lung tissue from mice. Lungs from eight naive mice of each inbred strain were used for bacterial collagenase digestion to extract the NC1 domain of type IV collagen. 15 μg of the digest was analyzed by SDS-PAGE and immunoblotting with human Goodpasture antisera (31). These antibodies recognize specifically the α3(IV) NC1 domain (Goodpasture antigen). D and M. The dimers and monomers, respectively, of NC1 domain that are recognized by the Goodpasture antibodies (31). The molecular masses of the dimers and monomers are ~54 and 28 kD, respectively. Lanes 1 and 2 denote SJL, lanes 3 and 4 denote DBA/2, lanes 5 and 6 denote A/J, lanes 7 and 8 denote (A/J × SJL) F1. Lanes 1, 3, and 5 are kidney NC1, and lanes 2, 4, 6, and 8 are lung NC1. The two bands in the monomer region could represent two size isoforms of α3(IV) NC1 monomer (67).

Figure 3. Immunoblot analysis of Goodpasture antigen [α3(IV) collagen] in the kidney and lung tissue from mice. Lungs from eight naive mice of each inbred strain were used for bacterial collagenase digestion to extract the NC1 domain of type IV collagen. 15 μg of the digest was analyzed by SDS-PAGE and immunoblotting with human Goodpasture antisera (31). These antibodies recognize specifically the α3(IV) NC1 domain (Goodpasture antigen). D and M. The dimers and monomers, respectively, of NC1 domain that are recognized by the Goodpasture antibodies (31). The molecular masses of the dimers and monomers are ~54 and 28 kD, respectively. Lanes 1 and 2 denote SJL, lanes 3 and 4 denote DBA/2, lanes 5 and 6 denote A/J, lanes 7 and 8 denote (A/J × SJL) F1. Lanes 1, 3, and 5 are kidney NC1, and lanes 2, 4, 6, and 8 are lung NC1. The two bands in the monomer region could represent two size isoforms of α3(IV) NC1 monomer (67).

Figure 4. Epitope recognition of murine and human anti–α3(IV) NC1 collagen antibodies. Binding curves for the human Goodpasture (GP) antibodies, circulating, kidney-, and lung-bound, from SJL mice immunized with bovine α3(IV) NC1 collagen were obtained using denatured mouse NC1 hexamer as the coating antigen in direct ELISA. The 50 and 100% saturating human Goodpasture antibody concentrations were used as inhibitor to binding of the mouse antibodies to mouse antigen. (A) Inhibition curves from such experiments. The assay is developed using anti–mouse IgG conjugated to alkaline phosphatase.

(B) The reciprocal experiment, in which mouse circulating anti–α3(IV) NC1 collagen antibodies are used as inhibitor to binding of the human Goodpasture antibodies to mouse antigen. The assay is developed using anti–human IgG conjugated to alkaline phosphatase. In B, the 50 and 100% saturating concentrations for mouse circulating antibodies are 1:600 and 1:25, respectively. The concentration of denatured mouse NC1 hexamer used to coat the wells in all the experiments is 150 ng.
day 7 in susceptible SJL mice, and persisted in high titer until the end of the study (data not shown). Circulating anti-α3(IV) NC1 antibodies from SJL mice with nephritis stained GBM from human kidney by indirect immunofluorescence (data not shown).

Circulating SJL mouse antisera or antibody eluted from mouse lungs and kidneys were used to compete with human Goodpasture antibody for recognition of dissociated mouse NC1 hexamer. Antibodies eluted from SJL kidney, lung, or the circulation all exhibited > 60% inhibition of human Goodpasture antibodies to the mouse hexamer (Fig. 4 A). Conversely, human Goodpasture antibodies inhibited the binding of antibody eluted from mouse kidney by 65%, and that from lung by 42%, but little inhibition was observed with circulating antibodies (Fig. 4 B). These latter results suggest that anti-α3(IV) NC1 antibodies bound to tissue share some target epitopes with human antibodies.

Passive transfer of anti–α3(IV) NC1 antibodies. Antisera against anti-α3(IV) NC1 domains were harvested and pooled from SJL mice immunized to produce anti-GBM disease. Anti-GBM or control antibodies were passively transferred into naïve syngeneic recipients. Urine protein for all the experimental and control mice was monitored on day 3 after transfer. All the experimental SJL mice showed slightly elevated urine protein compared with vehicle-injected controls. On day 5, an experimental SJL mouse was killed, and the tissue was analyzed. The day 5 kidney exhibited some mild inflammation within the glomeruli, with minimal interstitial changes; lung histopathology was normal (data not shown). Immunofluorescent studies revealed linear IgG binding to the GBM (Fig. 5 C). These results suggest that anti-GBM disease can be passively transferred to naïve SJL mice by administration of anti-α3(IV) NC1 collagen antibodies.
Since passive transfer of antibodies into a naive recipient of a susceptible strain produced nephritis, we decided to evaluate the nephritogenic capacity of anti-α3(IV) NC1 antibodies transferred from a nonsusceptible strain into allotypically compatible recipients. However, when anti-α3(IV) antibodies from immune SJL and A/J mice were passively transferred into (A/J × SJL) F1 recipients, neither produced an expected nephritis. The F1 crosses were then immunized to produce anti-GBM disease, and while robust titers of anti-α3(IV) antibodies appeared (data not shown), there was still no nephritis.

MHC-identical F1 crosses between disease-susceptible BALB/c and congenic CA21 mice (BALB.1gh) were next generated and used as recipients to test the nephritogenicity of passively transferred anti-α3(IV) collagen antibodies from disease-susceptible BALB/c and nonsusceptible A/J mice. The development of antiallotype antibodies against the A/J (Igh) allotype was avoided by using congenic CA21 (BALB.1gh) mice in the cross (39, 40). 3 wk after the passive transfer of immune sera from BALB/c or A/J mice into the (BALB/c × CA21) F1 recipients, we observed that both sets of antibodies produced anti-GBM disease. Proteinuria in F1 recipients injected with BALB/c or A/J anti-α3(IV) antibodies was much higher than in controls: 7.45±2.23 (BALB/c nephritic sera) vs. 6.67±2.53 (A/J nephritic sera) vs. 0.8±0.31 mg/16 h (CFA control sera). Histology revealed hypercellularity in the glomeruli with occasional crescents (20% of the glomeruli at 3 wk) and early tubulointerstitial infiltrates in F1 mice receiving either BALB/c or A/J sera (data not shown). The kidneys from control mice were histologically normal.

Adoptive transfer studies in disease-susceptible SJL mice. Donor SJL mice were immunized with α3(IV) NC1 collagen. 18 d later, spleen and lymph node cells were harvested for transfer (32). 10^7 immune cells were administered intravenously to low-dose cyclophosphamide-pretreated syngeneic recipients. Low-dose cyclophosphamide pretreatment was used because initial cell transfers without pretreatment did not produce disease (data not shown); cyclophosphamide is known to improve adaptation of transferred cells by reducing native regulatory T cell populations in naive recipient mice (41). Recipient mice were monitored for elevation in urine protein and serum creatinine levels. Control mice were injected with cells from adjuvant control mice. These cohorts were then killed after 4 mo. The kidneys and lungs were evaluated for immune cell infiltrates and pathology associated with disease. Kidneys from the experimental animals revealed moderate crescentic glomerulonephritis, with tubulointerstitial disease and linear staining for mouse IgG on the GBM and tubular basement membrane (Fig. 5, E and F). The lung histopathology was normal (data not shown). Control mice did not exhibit abnormal pathology in the kidney and lungs (Fig. 5 D). The lungs from the experimental group revealed insignificant changes compared with control mice within the time frame of the experiment (data not shown). These results suggest that immune cells also transfer nephritis adaptively. As in the passive transfer experiments above, whether a longer period of observation would have produced more damage to the lungs, or what the special role of adjuvant in accelerating inflammation might be, were not determined in this set of experiments.

We next performed two additional experiments to assess further the role of T cells in the expression of anti-GBM disease. In the first experiment, Tcr−/− mice (28) were immunized with α3(IV) NC1 collagen and, when compared with Tcr+/+ controls, did not produce anti-GBM antibodies (data not shown) or disease (Fig. 6, A and C); these Tcr−/− mice also did not develop proteinuria (0.54±0.12 mg/16 h) when compared with wild-type mice (5.43±1.22 mg/16 h). In a second experiment, anti-α3(IV) NC1 antibodies generated in disease-susceptible C57BL/6 mice were passively transferred into naïve syngeneic Tcr+/+ or Tcr−/− recipients. After 3 wk, kidneys from Tcr−/− recipients contained anti-GBM antibody but did not develop disease when compared with Tcr+/+ mice (Fig. 6, B and D); these Tcr−/− mice also did not develop proteinuria (0.44±0.11 mg/16 h) when compared with wild-type mice (3.78±1.01 mg/16 h).

Assessment of the nephritogenic T cell phenotype. In the first experiment, antibodies reactive to α3(IV) NC1 domains were IgG-isotyped in a panel of SJL, A/J, A.SW, and (A/J × SJL) F1 mice immunized to produce disease. All mice demonstrated significant levels of the IgG1/Th2-like isotype, whereas only SJL and A.SW nephritic mice showed significant titers for IgG2a/Th1-like isotype (Fig. 7). The same pattern of isotype IgG was observed in antibodies eluted from the kidneys of SJL and A/J mice (data not shown). Anti-α3(IV) NC1 IgG2b and IgG3 antibodies were not detectable in any of the immunized strains.

Kidney tissue from susceptible SJL and nonsusceptible A/J mice were further stained by immunohistochemistry for the presence of IL-12 and IL-4. Cortical kidney tissue from SJL mice with 12 wk of disease stained for IL-12 principally in glomerular crescents (Fig. 8 A), but not the tufts themselves, and in tubular casts (Fig. 8 B). Rare tubular cells stained for IL-12 (not shown); presumably, the IL-12 in the casts derived from release into Bowman’s space. A/J kidneys did not stain for IL-12 (Fig. 8 C), and neither stained positive for IL-4 (Fig. 8 D).

Figure 7. Isotyping of anti-α3(IV) NC1–specific IgG repertoire. Direct ELISA was performed to analyze the α3(IV) NC1–reactive IgG repertoire in SJL, A/J, A.SW, and (A/J × SJL) F1 mice. Isotype-specific anti-mouse IgG antibodies were used to develop this analysis (Sigma Chemical Co.). Denatured mouse α3(IV) NC1 hexamer (100 ng) was used as the antigen. The primary antibody dilutions for each of the mice were 1:200.
8 D shows SJL). We also stained for IL-10 and IFN-γ, and the findings were identical to those observed for IL-4 and IL-12, respectively (data not shown). These results along with the antibody isotyping suggest a connection between the presence of a Th1-like response (42) and susceptibility to anti-GBM nephritis.

**Oral tolerance studies in susceptible mice.** SJL mice were fed one of two doses (65 or 700 μg) of either type IV collagen containing α3(IV) NC1 dimers or BSA before immunization with the antigen, to determine if oral feeding could attenuate tissue inflammation. Two different doses were used because of reports that low and high doses of tolerogen induced tolerance by different mechanisms (43, 44). 2 mo later, they were evaluated for urine protein and serum creatinine. The BSA-fed or unfed mice immunized with antigen revealed increases in urine protein and serum creatinine levels (Fig. 9). Histological examination of kidneys from these mice demonstrated crescentic glomerulonephritis with interstitial infiltrates. However, mice immunized with antigen after being prefed NC1 domains (both 700- and 65-μg doses) showed a substantial decrease in numbers of crescents, and complete amelioration of the interstitial disease (Fig. 9); the effect in mice fed with NC1 domains at high doses was similar in those fed low doses. Our results are consistent with those observed in other antigen systems (35, 36, 43, 45–47). IgG-isotyping experiments using α3(IV) NC1 domains as the target antigen revealed a decrease in the titers of IgG2a antibodies in mice prefed α3(IV) NC1 domains in the type IV collagen diet compared with BSA-fed and unfed mice. All groups had robust levels of anti-α3(IV) IgG1 antibodies (Fig. 9). IL-12 expression was observed in all the unfed and BSA-fed mice immunized to produce disease (Fig. 10); mice prefed NC1 domains had notably less IL-12 expression in the kidney. These results suggest that oral tolerance to the α3(IV)...
NC1 antigen in murine anti-GBM disease attenuates IL-12/Th1-like T cell responses.

Discussion

Immunization of disease-susceptible mice with α3(IV) NC1 collagen produces lung hemorrhage, crescentic glomerulonephritis, interstitial renal disease, and progressive renal failure. Tissue-eluted antibodies bind specifically to α3(IV) NC1 collagen, and these antibody repertoires overlap in specificity with human Goodpasture antibodies. Nephritic antibodies from SJL mice bind to the human and mouse GBM in a linear fashion, suggesting corecognition of antigens in similar distribution. Human Goodpasture antibody also blocks significantly the binding of nephritic mouse anti-α3(IV) antibody to mouse hexamer. This was observed best with mouse antibodies eluted from kidney and lung, but not at all with circulating antibody.

This seemingly curious finding with circulating antibody is likely explained by the differences in how these sets of antibodies were generated. Human Goodpasture antibodies arise spontaneously and are directed to cryptic epitopes buried in associated hexamer that escape tolerance (10). Circulating antibodies from immunized mice recognize the same cryptic α3(IV) NC1 epitopes as the human antibodies and antibodies eluted from mouse lung and kidney, but also contain species that recognize other disease-irrelevant α3(IV) epitopes probably created by exposure of the immunogen to biochemical purification and the process of adjuvant immunization. These latter antibodies would not be expected to inhibit human Goodpasture antisera. In spite of this understandable quirk, there appears to be appropriate commonality in the recognition specificity of tissue-bound antibodies between humans and mice with disease.

Most recent investigations suggest that the principal specificity of human anti-GBM antibodies is towards α3(IV) NC1 domain of type IV collagen (26). Antibodies from nephritic mice in these experiments share this human specificity and, therefore, are capable of transferring disease to syngeneic naive recipients. Earlier passive transfer experiments performed in sheep and primates (14, 48) also demonstrated the capacity of heterologous antibodies to crude GBM to transfer nephritis. Whether this nephritis is the same as that in humans or in our mouse model is unclear. The target antigens for those heterologous anti-GBM antibodies are not known. Additionally, inflammatory mediators stimulated by the antiallotype effect of passively transferring heterologous antibodies were not taken into consideration when assigning pathogenicity to primary antibodies expressed in an autologous environment.
We also identified several genetic factors pertaining to the location, initiation, and intensity of anti-GBM disease. Human patients with anti-GBM disease and GPS are phenotypically diverse. Some have disease limited to the kidney (4, 26), like DBA/2 mice, some have aggressive disease in both organs (4, 26), like SJL and to a lesser extent BALB/c, B6, and B10 mice, and some have antibodies without much inflammation (22, 49), like A/J, AKR, and CBA mice. Some patients have disease limited to the lung; however, no mice to date express this phenotype.

The class II MHC locus is a polymorphic region for immune response genes, and seems to be an important modulator of the immune response to α3(IV) NC1 domains. Human alleles mapping to HLA-DR (Eβ/Eα in mice) and -DQ (Aβ/Aα in mice) are associated with the inflammatory response of GPS in humans (8, 9). Limited sample size and the linkage disequilibrium between certain DR and DQ alleles in human anti-GBM nephritis have made it difficult to attribute susceptibility to one region more than the other (9). In this study, we were able to link susceptibility to anti-GBM disease in mice to the Aβ/Aα region in H-2; this suggests that the HLA-DQ region should be studied more closely in humans with anti-GBM nephritis. Of course, non-MHC genes also map in or near the genomic location of class II. However, the non-MHC loci in this region are usually considered unlikely candidates for susceptibility because they do not express enough polymorphisms to explain multiple strain variations.

Curiously, the accessibility of α3(IV) collagen to anti-GBM antibodies in lung basement membrane also maps to MHC class II. Western blots of lung tissue in vitro from A/J, SJL, A.SW, and (A/J × SJL) F1 mice indicate that similar amounts of α3(IV) collagen are present in all lung tissues in spite of the failure of lung from A/J and (A/J × SJL) F1 crosses to bind antibody in vivo. These findings suggest that epitope accessibility in the lung is polymorphic. How prevalent such polymorphisms are in humans, and whether this MHC gene effect is a modification of structure or bears subtly on the expression of the immune repertoire, are not yet clear.

MHC class II genes are important in the initiation of helper T cell activation (50, 51). The role of T cells in human anti-GBM disease is now beginning to be studied, and reports to date have simply identified α3(IV) NC1 autoantigenic CD4+ and CD8+ T cells in patients with disease (12, 13). Earlier experiments in chickens demonstrated the ability of immune cells reactive to a mixture of antigens in solubilized GBM to transfer glomerulonephritis to naive recipients, even upon removal of the bursa (52). Anti-GBM disease in our experiments can also be transferred adoptively by lymphocytes. Additionally, TCR−−/− mice do not develop anti-GBM disease after immunization with α3(IV) NC1 domains compared with wild-type controls. To address further the significance of the role of T cells in mediating inflammation, we transferred syngeneic anti-α3(IV) antibodies passively from TCR +/+ disease-susceptible mice into TCR −/− or +/- recipients. After 3 wk, anti-GBM disease was observed only in +/- mice. These results lend further credence to the notion that nephritogenic T cells in disease-susceptible mice are of pivotal importance in the expression of inflammation in anti-GBM disease. Since TCR deficiency should not affect natural killer cells or the expression of antibody-dependent cell-mediated cytotoxicity, we assume that antibody binding can further educate nephritogenic T cell repertoires in disease-susceptible haplotypes.

Helper T cells coordinate the preference for emerging effector cell or antibody repertoires probably as a net balance of Th1 and Th2 response, respectively (42, 51, 53, 54). Production of IgG1 is part of an IL-4/Th2-like response in mice, and IgG2a is part of an IL-12/Th1-like response; these cytokines and antibody isotypes are not the only ones involved in this selection but they are reasonable markers of these dual pathways (55, 56). While this effect is not entirely binary (53), there are some peptides that favor strongly Th1 and Th2 responses (57–59), some of which are MHC-linked (60) and based on strength of peptide/MHC interaction with TCR (51, 61).

Our findings using nephritogenic epitopes in the NC1 domain suggest a different balance among helper T cell phenotypes. We monitored these T cell phenotypes in vivo by noting the presence of IL-12, IL-10, IL-4, or IFN-γ in nephritic kidneys (only IL-12 and IL-4 data were shown), as well as the isotype of anti-GBM antibodies produced by various haplotypes (42). SJL, A.SW, and B10.S mice have a strong cell-mediated effector response in addition to the production of antibody, whereas A/J mice demonstrate only an antibody response that by itself is insufficient to produce glomerulonephritis. In IgG-isotyping experiments with SJL, A.SW, A/J, and (A/J × SJL) F1 mice, all mice had Th2-like responses, whereas only SJL and A.SW mice, strains that develop cellular infiltrates and nephritis, had additional Th1-like responses. These results support the notion that all mice respond to the α3(IV) immunogen with Th2-type response, but only the haplotypes expressing appropriate MHC class II genes launch a Th1 response resulting in disease. This hypothesis is supported further by evidence of IL-12 (and IFN-γ) only in the glomerular lesions of disease-susceptible mice.

Furthermore, when nephritogenic antibodies from SJL and A/J mice were passively transferred into (A/J × SJL) F1 recipients, there was no autoimmune glomerulonephritis. This critical experiment suggests that anti-GBM antibodies, regardless of isotype, are not alone sufficient to produce nephritis. However, in an allotype-compatible and disease-susceptible haplotype, antibodies from nonsusceptible mice can be nephritogenic. Such experiments, performed in (BALB/c × CA21) F1 recipients, clearly speak to the role of nonhumoral mechanisms in the expression of anti-GBM injury. We hypothesize that these nonhumoral mechanisms depend on T cell repertoires directed by MHC class II genes.

The absence of disease in the (A/J × SJL) F1 cross is a curiosity and special interest that has several potential explanations. There is the possibility of a simple reduction in MHC gene dose (62), or of the formation of new hybrid MHC molecules favoring Th2-like effector responses (51, 61). Furthermore, while Th1 responses prevail occasionally in some peptide/MHC interactions (60), others have suggested that IL-4/Th2 effects dominate typically during T cell priming (57, 59, 63). Part of determining this preference is probably shaped by the interaction of peptide with MHC; for example, BALB/c mice immunized with α3(IV) favor a Th1 response, whereas the same mice immunized with leishmania favor a Th2-like effect (64, 65).

The dominant-negative effects of the (A/J × SJL) F1 cross might also be explained by a change in the relative balance or opposing action of anti-α3(IV) NC1 clones (53). In this vein, selected strains like SJL would have strong, bidirectional Th1- and Th2-like responses, B6, BALB/c, and B10 mice a weaker Th1- and similar Th2-like responses, and A/J and AKR mice a
Th2-like response alone. The dominant-negative cross, therefore, could be viewed as favoring heavily a Th2-like effect. Such bias may be self-protective in this autoimmune disease, and fundamental in explaining the low incidence of GPS in humans, where nonfamilial breeding is socially preferred. Further verification of a bidirectional balance in this hypothesis will require special examination of α3(IV) NC1–reactive T cell repertoires. Finally, we were able to downregulate the nephritogenic cell–mediated response in SJL mice by tolerizing them with oral feeding of α3(IV) NC1 target antigen. This modulation of disease has been observed in other systems (35, 43, 44, 47, 66). The primary mechanism by which orally administered antigen induces tolerance is thought to be the attenuation of T cell clones; low doses of orally administered antigen favor active suppression, whereas higher doses favor clonal anergy (43, 47). Our results suggest a preferential effect on the Th1 repertoire responsible for the nephritis, perhaps by reducing the expression of tissue IL-12 or other cytokines in that cascade. Recent findings in other models argue that antigen feeding generates Th2-like responses in gut-associated lymphoid tissue and the emergence of cytokines such as TGF-β and IL-4, which tend to suppress proinflammatory Th1-like responses (35, 43, 44, 47).

Acknowledgments

We wish to thank Michelle C. Werner and Mark Schwartz for their assistance in maintaining mice colonies and collection of tissues and blood from mice.

This work was supported in part by grants DK-46282, DK-07006, DK-30280, DK-02334, and DK-45191 from the National Institutes of Health, the DCI RED fund, and grant B170196 from BioStratrum Inc.; Dr. Kalluri is a founder and consultant in Biostratrum Inc. and owns equity in the company. Dr. Okada was a recipient of a fellowship from Ei Lilly Japan, Ltd.

References


Kalluri et al.

*Murine Anti–Glomerular Basement Membrane Disease* 2275