

## Pathogenesis of autoimmunity in $\alpha\beta$ T cell-deficient lupus-prone mice

S. L. PENG\*§, J. CAPPADONA\*, J. M. MCNIFF‡, M. P. MADAIO¶, M. J. OWEN\*\*, A. C. HAYDAY§† & J. CRAFT\* Sections of \*Rheumatology and †Immunobiology, and ‡Department of Dermatology, Yale University School of Medicine and §Department of Biology, Yale University, New Haven, CT, ¶Renal, Electrolyte, and Hypertension Division, Department of Medicine, University of Pennsylvania School of Medicine, Philadelphia, PA, USA, and \*\*Imperial Cancer Research Fund, London, UK

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### SUMMARY

Murine lupus in MRL mice has been strongly attributed to  $\alpha\beta$  T cell-dependent mechanisms. Non- $\alpha\beta$  T cell-dependent mechanisms, such as  $\gamma\delta$  T cells, have been shown to drive antibody and autoantibody production, but they have not been considered capable of inducing end-organ disease. Here, we have expanded upon the findings of such previous work by examining the mechanism and extent of end-organ disease attainable via  $\gamma\delta$  T cells and/or non- $\alpha\beta$  T cell-dependent mechanisms, assessing two prototypical lupus lesions, renal and skin disease, in TCR  $\alpha$   $-/-$  MRL mice that possessed either functional or defective Fas antigen (Fas + or *lpr*). Observed to 1 year of age, TCR  $\alpha$   $-/-$  MRL mice developed disease characterized by increased mortality, overt renal disease and skin lesions. While delayed in onset and/or reduced in severity compared with TCR  $\alpha$   $+/+$  MRL/*lpr* animals, renal and skin lesions in  $\alpha\beta$  T cell-deficient animals were clearly increased in severity compared with age-matched control non-autoimmune mice. In contrast to TCR  $\alpha$   $+/+$  MRL mice, whose disease reflected pan-isotype immune complex deposition with significant complement fixation, renal disease in TCR  $\alpha$   $-/-$  MRL animals reflected predominantly IgG1 immune complex deposition, with poor complement fixation. Thus, this study demonstrates conclusively that non- $\alpha\beta$  T cell-dependent mechanisms can induce renal and skin injury in murine lupus, but at least in the kidney, only via humoral autoimmunity of a relatively non-pathological isotype which results in the delayed onset of end-organ damage.

**Keywords** autoimmunity T lymphocytes mice autoantibodies nephritis skin

### INTRODUCTION

The MRL model of murine lupus is a particularly useful system to investigate systemic autoimmunity, since its disease closely resembles human systemic lupus erythematosus (SLE), including the development of autoantibodies and renal and skin disease [1–3]. The MRL/Mp-*lpr/lpr* (MRL/*lpr*) mouse, which develops a severely accelerated form of MRL lupus [1] due to a functional defect in the Fas apoptosis antigen [4], provides a convenient congenic strain to examine the role of Fas-mediated immune regulation in lupus. While B cells in MRL/*lpr* mice are intrinsically abnormal [5–8], many studies have used this model to establish the role of T cells in the pathogenesis of murine lupus, focusing upon the role of CD4<sup>+</sup>  $\alpha\beta$  T cells as helpers for autoantibody production [9–14]. Some data suggest that  $\gamma\delta$  T cells may propagate systemic humoral autoimmunity [15–19]; however, none has found that  $\gamma\delta$  T cells serve as significant instigators of end-organ disease.

To evaluate the significance of  $\gamma\delta$  T cell- and/or other non- $\alpha\beta$  T cell-dependent mechanisms in the induction of systemic disease, we assessed renal and skin end-organ disease in MRL mice made deficient in  $\alpha\beta$  T cells via genetic disruption of the T cell receptor (TCR)  $\alpha$  locus [17,20]. TCR  $\alpha$   $-/-$  MRL mice developed increased mortality, renal disease with compromised renal function, and skin disease in association with lupus autoantibodies, although their end-organ disease remained delayed and/or subdued in comparison with wild-type MRL/*lpr* animals. In addition, TCR  $\alpha$   $+/+$  MRL mice developed pan-isotype immune complex deposition associated with complement fixation, while kidneys of TCR  $\alpha$   $-/-$  MRL animals had predominantly IgG1 isotype-restricted immune complex deposition associated with poor complement fixation. Thus, in comparison with previous studies which have shown that non- $\alpha\beta$  T cells, particularly  $\gamma\delta$  T cells, can support autoantibody production [15–19], the current findings demonstrate *in vivo* that non- $\alpha\beta$  T cell-dependent mechanisms are capable of inducing humoral autoimmunity, which, while less aggressive than  $\alpha\beta$  T cell-dependent mechanisms, nevertheless evolves into consequential

Correspondence: Dr J. Craft, PO Box 208031 LCI 609, 333 Cedar Street, New Haven, CT 06520-8031, USA.

autoimmune disease with end-organ dysfunction of the skin and kidneys.

## MATERIALS AND METHODS

### Mice

TCR  $\alpha$   $-/-$  (TCR $\alpha^-$ ) and TCR  $\alpha$   $+/+$  (TCR $\alpha^+$ ) MRL mice bearing either functional ( $+/+$ ) or defective (*lpr/lpr*) Fas antigen [17], as well as controls including age-matched B10.A mice (Jackson Laboratory, Bar Harbor, ME), were maintained under specific pathogen-free conditions at the Yale University School of Medicine. These animals, of the F4 backcross generation, reflect wild-type MRL lupus-prone animals, since MRL disease-inducing genes are dominant in genetic crosses [21] as confirmed by the penetrance of autoantibodies and end-organ disease in the TCR  $\alpha$   $+/+$  mice of this cohort ([17] and this study). Indeed, in contrast to the present animals, which contain  $\approx 87.5\%$  MRL genes, previous studies have successfully utilized F<sub>2</sub> animals, which contain only  $\approx 50\%$  MRL genes [19,22]. Thus, it is notable that both F<sub>2</sub> and F<sub>4</sub> animals develop renal and skin lesions, as well as anti-snRNP antibodies, since MRL disease itself is stochastic, with incomplete penetrance of autoantibody production and end-organ disease [23–25]. Thus, while these mice may not technically reflect wild-type ('100%') MRL disease, their lupus syndrome closely resembles the severe MRL phenotype. Nevertheless, because of such issues, the various genetic groups in this study have been analysed as populations rather than individual mice: each genetic group as a whole being 87.5% MRL overall, and each population of animals being at similar risk for disease; in this study, we refer to these animals as TCR $\alpha^+$  MRL ( $+/+$  or *lpr*) or TCR $\alpha^-$  MRL ( $+/+$  or *lpr*).

### Antibody detection

Titres of IgM, IgA, and IgG were determined by ELISA on sera using an antibody isotyping kit (Pierce, Rockland, IL). IgE titres were determined by ELISA using separately purchased capture and detection antibodies (PharMingen, San Diego, CA). IgG anti-dsDNA and anti-snRNP, as well as  $\kappa$  anti-IgG, were determined as described elsewhere [17,26–28].

### Renal function tests

Sera were assayed for blood urea nitrogen (BUN) via clinical chemistry at Yale-New Haven Hospital. Proteinuria was assayed by collecting spot urine samples on 3 consecutive days. Total protein content was determined by quantitative Bradford assay [29]; creatinine content was determined by modified Jaffé reaction [30] (Sigma Chemical Co., St Louis, MO). Proteinuria indices were determined as (total urinary protein)/(total urinary creatinine), to normalize for glomerular filtration rate. Statistical significance was evaluated by unpaired, two-tailed Student's *t*-test.

### Histopathology

Renal disease was assessed as described [31,32]. Briefly, for light microscopy, tissue was fixed in 10% buffered formalin and stained with haematoxylin–eosin; glomeruli, tubular interstitium and vessels were examined. For immunofluorescence, specimens were frozen in OCT embedding medium, and sections were stained with fluorescein-conjugated goat anti-mouse IgG, IgG1, IgG2a, IgG2b or IgG3 (Southern Biotechnology Associates, Birmingham, AL),

or fluorescein-conjugated goat anti-mouse C3 (Cappel, West Chester, PA). Immune deposits were sought in glomeruli, tubules, and renal cell nuclei. Individual specimens were read blinded by S.L.P., J.C. and/or M.P.M., and scored on a 0–4+ scale. Cutaneous lesions were processed similarly, and evaluated blinded by J.M.M.

### Correlations of antibody titres with renal disease

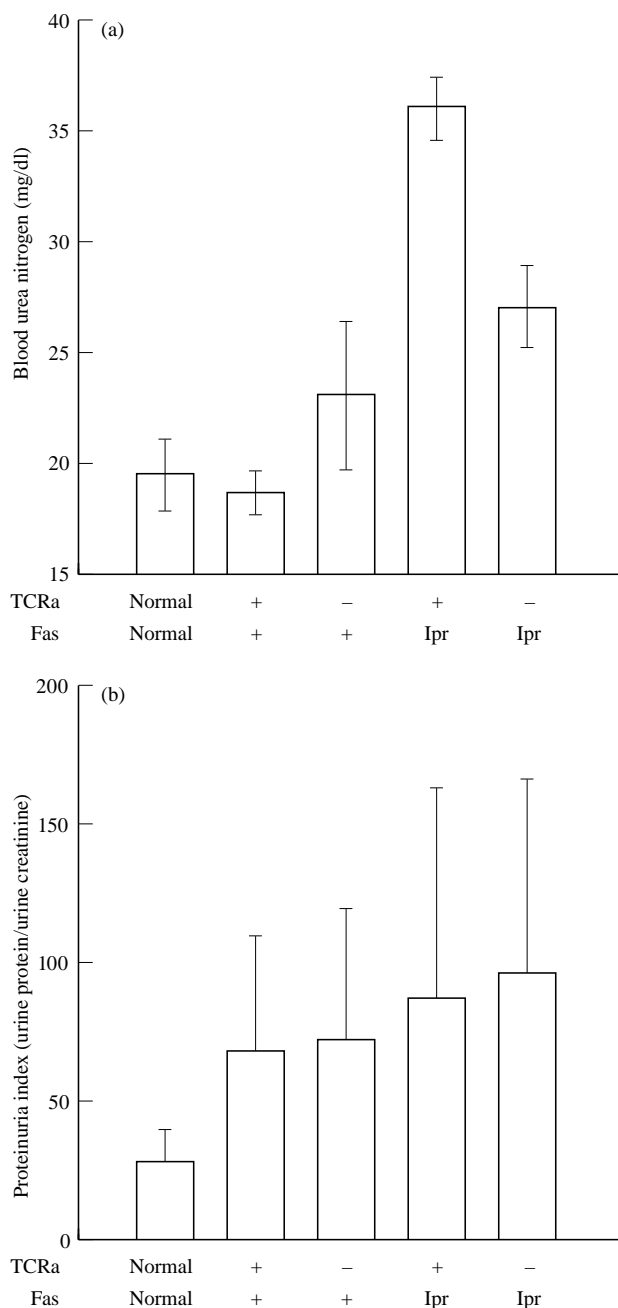
Antibody indices for each mouse (for Fig. 4) were calculated on a 1–10 scale, using one point for each elevated immunoglobulin isotype in  $\mu\text{g/ml}$  (IgM, IgA, IgG and IgE;  $>2$  s.d. above average normal mice), and for each positive autoantibody titre (anti-dsDNA, anti-snRNP, or anti-immunoglobulin;  $>3$  s.d. above average optical density (OD) titres of normal mice). Renal disease indices were calculated as the sum of disease scores (0–4+ scale) as visualized by light microscopy [31,32]. Pearson correlation coefficient was determined by Systat 5.2.1 software (Evanston, IL).

## RESULTS

### TCR $\alpha^-$ MRL mice develop renal disease

End-organ disease was first assessed in 1-year-old mice by BUN and proteinuria (Fig. 1). In comparison with normal and TCR $\alpha^+$  MRL  $+/+$  animals, both TCR $\alpha^-$  MRL/*lpr* and TCR $\alpha^-$  MRL  $+/+$  mice contained elevated BUN, although levels in only the former group reached a statistically significant difference ( $P < 0.05$ ). At the same time, neither TCR $\alpha^-$  MRL group developed as high BUN as TCR $\alpha^+$  MRL/*lpr* mice ( $P < 0.05$ ). In addition, all groups of lupus-prone mice, TCR $\alpha^-$  MRL  $+/+$  and TCR $\alpha^-$  MRL/*lpr*, as well as TCR $\alpha^+$  MRL  $+/+$  and TCR $\alpha^+$  MRL/*lpr*, developed elevated proteinuria indices in comparison with normal, age-matched B10.A mice ( $P < 0.05$ ). In the TCR $\alpha^+$  MRL/*lpr* group, end-stage renal disease probably caused decreased protein excretion (data not shown), even though those animals remaining alive at this age probably represented a biased group with milder disease.

In accordance with the renal function studies, both MRL  $+/+$  and MRL/*lpr* mice lacking  $\alpha\beta$  T cells developed glomerular, interstitial, and sometimes perivascular lesions (Table 1 and Fig. 2). While these were limited compared with their  $\alpha\beta$  T cell-intact MRL/*lpr* counterparts, they were still significant in comparison with age-matched normal mice. They also developed substantial renal immune deposits, sometimes comparable to the severe glomerular, tubular, and/or renal nuclear deposition characteristic of  $\alpha\beta$  T cell-intact disease (Table 2 and Fig. 3 and data not shown). Isotyping of the immune deposits in TCR $\alpha^+$  MRL/*lpr* animals consistently revealed pan-isotype accumulation by 12 weeks old, associated with significant complement (C3) deposition (Table 2 and Fig. 3). In contrast, deposits in TCR $\alpha^-$  MRL/*lpr* animals consisted of predominantly IgG1 antibodies, which required  $\approx 6$  months or more to reach levels which were consistently comparable to TCR $\alpha^+$  MRL/*lpr* mice. TCR $\alpha^-$  MRL/*lpr* animals furthermore had a relative paucity of IgG2a, IgG2b, IgG3 and C3 deposition, although these molecules were occasionally, but not predictably, detected. Nevertheless, histological abnormalities showed a correlation of light microscopic renal disease with serum antibody and autoantibody levels in both  $\alpha\beta$  T cell-intact and -deficient animals (Fig. 4), suggesting that immunoglobulin deposition was responsible for renal disease in all groups of mice.



**Fig. 1.** Renal function tests in 1-year-old TCRα+ MRL and TCRα- MRL mice. Mouse sera were measured for blood urea nitrogen levels. Urine samples were measured for total protein content and creatinine, and proteinuria index was calculated as protein/creatinine to normalize for glomerular filtration rate. Standard deviations are shown for five to seven mice in each group; normals are age-matched B10.A mice.

*TCRα- MRL mice develop cutaneous disease*

Skin disease developed in parallel with renal disease (Figs. 5 and 6). Whereas nearly all TCRα+ MRL/lpr mice characteristically developed severe hair loss, scab formation, and purpuric lesions of the dorsal neck and ears by 24 weeks old, TCRα- MRL mice of both Fas phenotypes appeared grossly normal. At this age, however, nearly all of TCRα- MRL/lpr mice developed subclinical lesions as determined by histology, which developed into grossly apparent cutaneous lesions and microscopic patterns

resembling T cell-intact disease by 7–8 months old. Approximately 50% of TCRα- MRL +/+ mice also developed clinically apparent skin lesions by 8–9 months old, while none of the TCRα+ MRL +/+ animals developed skin lesions even beyond 1 year of age (Figs. 5 and 6 and data not shown). Thus, as determined by renal and skin lesions, TCRα- MRL mice develop end-organ disease, albeit with less intensity compared with age-matched TCRα+ MRL/lpr animals.

*Mortality of TCRα- MRL mice*

In conjunction with end-organ disease, TCRα- MRL mice of both Fas phenotypes demonstrated increased mortality in comparison with TCRα+ MRL +/+ mice (Fig. 7;  $P < 0.05$  for MRL +/+,  $P < 0.01$  for MRL/lpr), although they demonstrated decreased mortality in comparison with TCRα+ MRL/lpr animals ( $P < 0.05$ ). The mortality rates correlated with renal and skin disease, since TCRα+ MRL/lpr mice, which had the highest mortality, developed the highest BUN and severity of skin disease, while TCRα- MRL/lpr mice, of second highest mortality, had the second most severe azotaemia and skin disease, and so forth (Figs. 1 and 7). These data establish a correlation between compromised renal function, skin disease and mortality in these animals, verifying the presence of significant, albeit limited, end-organ disease despite the absence of  $\alpha\beta$  T cells. Accordingly, more substantial renal and skin disease was incidentally noted in TCRα- MRL mice that died or were moribund (data not shown).

**DISCUSSION**

This study demonstrates that MRL mice without  $\alpha\beta$  T cells

**Table 1.** Renal pathology in 1-year-old TCRα+ MRL and TCRα- MRL mice, and in normal B10.A controls

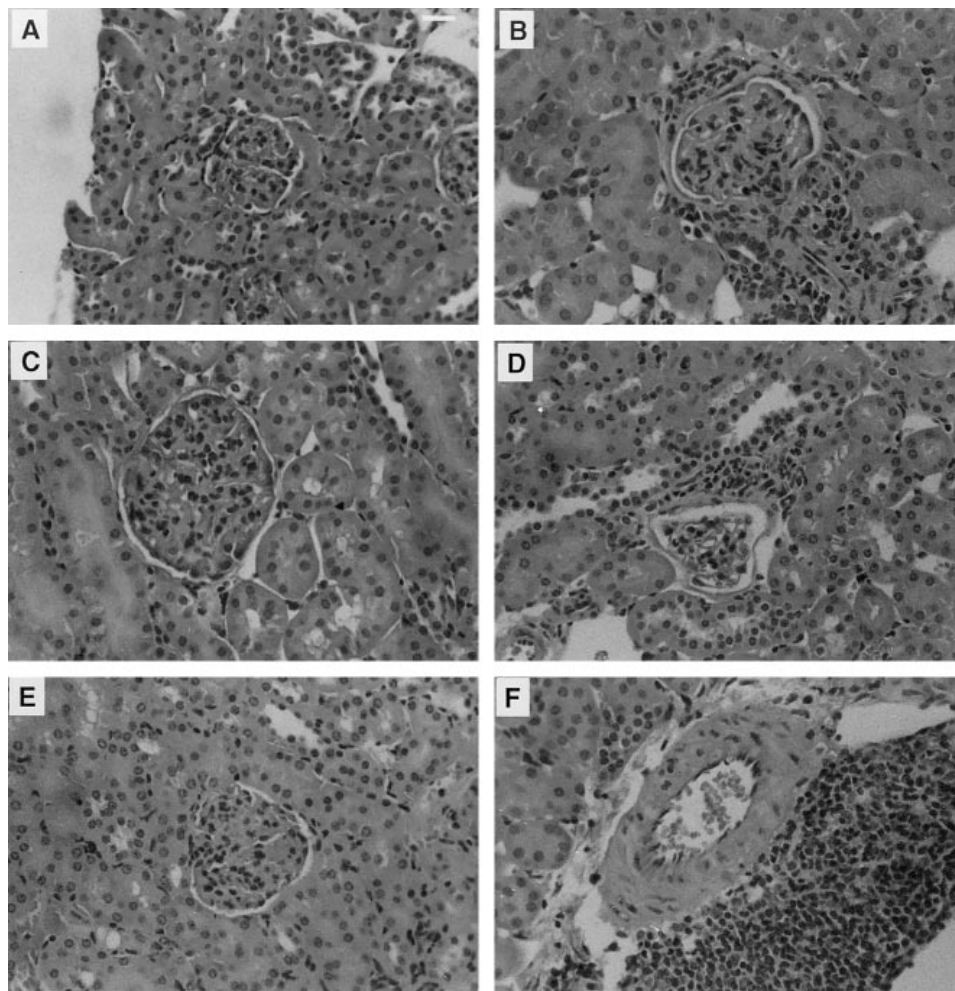
Strain/genotype	Mouse no.	Light microscopy		
		Glomerular	Interstitial	Perivascular
B10.A	1	+/-	+	-
B10.A	2	+/-	+	-
B10.A	3	+/-	+	-
B10.A	4	+/-	+	-
TCRα+ MRL +/+	1	+++/+	+	-
TCRα+ MRL +/+	2	+++ +/+	+++ +/+	+++ +/+
TCRα+ MRL +/+	3	+++ +/+	+	++
TCRα+ MRL +/+	4	+	-	+
TCRα- MRL +/+	1	+	+	-
TCRα- MRL +/+	2	+	+	-
TCRα- MRL +/+	3	+	+	-
TCRα- MRL +/+	4	-	-	-
TCRα- MRL +/+	5	-	+	-
TCRα+ MRL/lpr	1	+++ +/+ +/+ +/+	+++ +/+ +/+	+++ +/+
TCRα+ MRL/lpr	2	+++ +/+ +/+ +/+	+++	+++ +/+
TCRα- MRL/lpr	1	+++ +/+ +/+	+	-
TCRα- MRL/lpr	2	+/-	+/-	-
TCRα- MRL/lpr	3	+/-	-	-
TCRα- MRL/lpr	4	+	+++ +/+ +/+	-
TCRα- MRL/lpr	5	+	++	+++ +/+ +/+

Lesions were scored on a 0–4+ scale in blinded fashion.

develop overt renal and skin disease, dispelling models that consider only these T cells capable of inducing end-organ disease in murine lupus, and suggesting that  $\gamma\delta$  T cell-dependent helper functions are capable of inducing lupus nephritis and cutaneous lesions. While a contribution to antibody production and disease by  $\text{TCR}\alpha^{-}\beta^{+}$  cells cannot be entirely ruled out [33,34], previous studies have established that  $\gamma\delta$  T cells are the primary source of T-dependent help in  $\alpha\beta$  T cell-deficient mice [17–19]. In contrast to prior studies on non- $\alpha\beta$  T cells in autoimmunity [15–19], this study provides a clear *in vivo* demonstration of the pathological consequences of such non-classical mechanisms.

The development of both renal and cutaneous lesions in  $\text{TCR}\alpha^{-}$  MRL mice contrasts with the conclusions of previous studies, which have strongly implicated a requirement for helper  $\alpha\beta$  T cells in the generation of pathogenic autoantibodies which may mediate end-organ disease [2,9–14,35,36]. Indeed,  $\text{TCR}\alpha^{-}$  MRL mice contained delayed and/or subdued renal and cutaneous lesions, which may indicate an inability of non- $\alpha\beta$  T cell-dependent mechanisms to promote pathogenic, affinity-matured auto-

antibodies [17]. Likewise,  $\alpha\beta$  T cells may be required for the development of complement-fixing pathogenic antibodies, particularly IgG of the IgG2a, IgG2b or IgG3 subclasses [37]. Thus, the T-dependent help in  $\text{TCR}\alpha^{-}$  MRL mice is notable for the presence of IgG antibodies, autoantibodies and end-organ immune deposits predominantly of the IgG1 subclass ([17] and Table 2 and Fig. 3), which fixes complement poorly ([37] and Table 2 and Fig. 3), suggesting that  $\text{TCR}\alpha^{-}$  MRL pathogenesis interestingly reflects a primarily complement-independent mechanism for glomerular disease. Accordingly, the apparently decreased renal interstitial infiltration in  $\text{TCR}\alpha^{-}$  MRL animals may reflect their relatively diminished glomerulonephritis, which may provide cytokines or other inflammatory mediators that recruit inflammatory cells to the interstitium. Regardless of the specific mechanism of renal injury, these results demonstrate that the  $\text{TCR}\alpha^{-}$  MRL phenotype reflects a delayed and subdued autoimmune disease syndrome in comparison with  $\text{TCR}\alpha^{+}$  MRL/*lpr* animals, but nevertheless reflects a consequential pathological response. As an intriguing possibility,  $\text{TCR}\alpha^{-}$  MRL disease may in fact represent a protective form of T

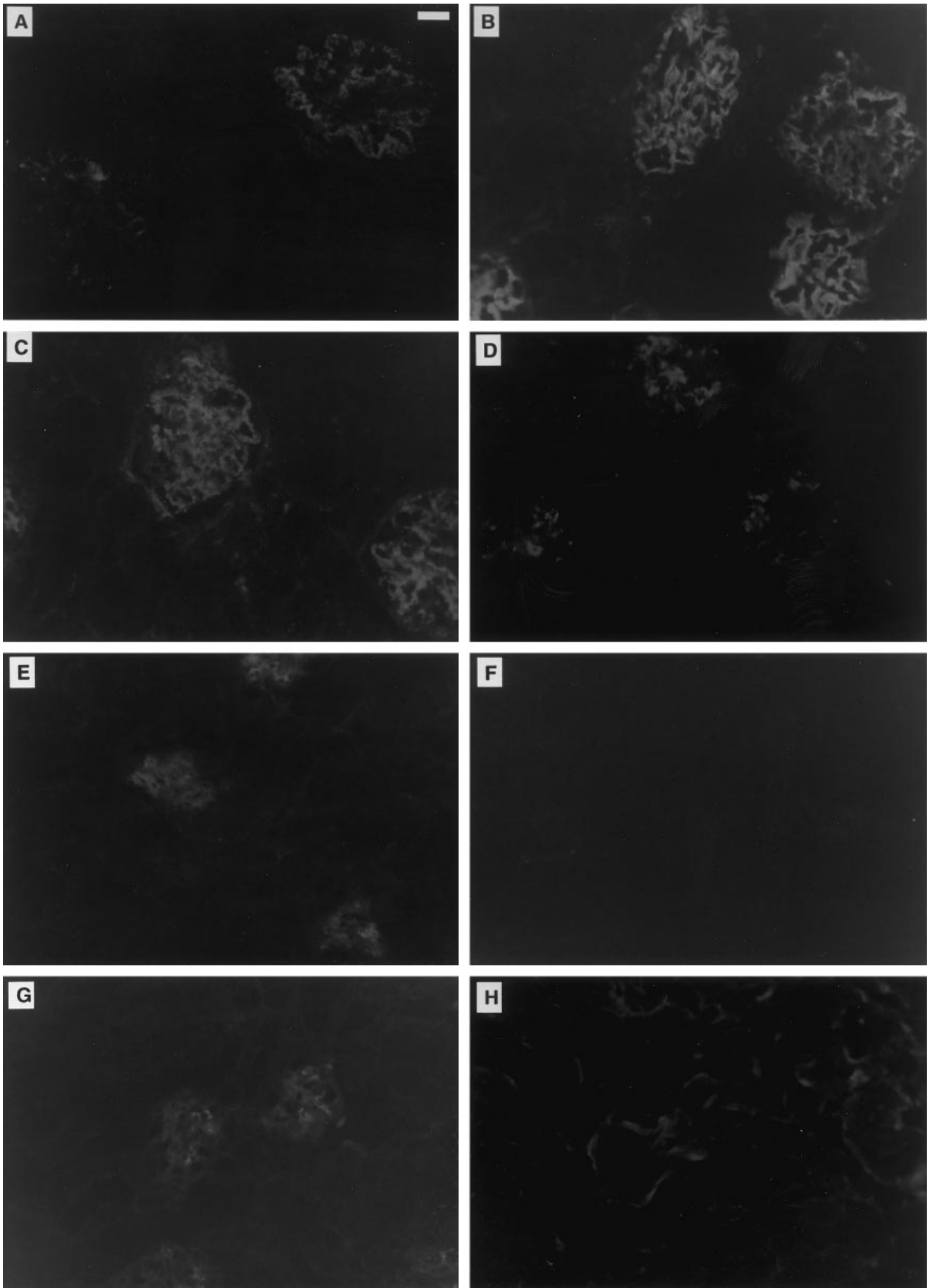


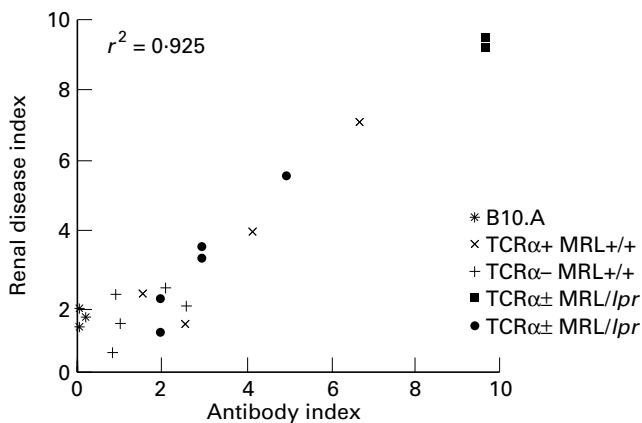
**Fig. 2.** Renal disease in 1-year-old  $\text{TCR}\alpha^{+}$  MRL and  $\text{TCR}\alpha^{-}$  MRL mice. Shown are light microscopic specimens from 48-week-old animals, including: (A) normal histology of a B10.A mouse; (B) focal glomerulonephritis and periglomerular infiltrate in a  $\text{TCR}\alpha^{+}$  MRL/*lpr* mouse; (C) glomerular hypercellularity in a  $\text{TCR}\alpha^{+}$  MRL +/+ mouse; (D) periglomerular infiltrate in a  $\text{TCR}\alpha^{-}$  MRL +/+ mouse; (E) focal glomerular hypercellularity in a  $\text{TCR}\alpha^{-}$  MRL/*lpr* mouse; and (F) perivascularitis in a  $\text{TCR}\alpha^{-}$  MRL/*lpr* mouse. Scale bar, 25  $\mu\text{m}$ ; all panels are of the same magnification.

**Table 2.** Isotyping of immune deposits in TCR $\alpha$ - MRL/lpr mice

Age (weeks)	Total IgG			IgG1			IgG2a			IgG2b			IgG3			C3		
	G	T	A	G	T	A	G	T	A	G	T	A	G	T	A	G	T	A
42	+++	+	+/-	+/+	+/+	-	-	-	-	-	-	-	+/+	-	-	+/+	++	-
24	++	+	-	+/+	+/+	+/+	-	-	-	-	-	-	+/+	-	-	-	+/+	-
12	+/+	-	-	+	-	-	-	-	-	-	-	-	-	-	-	-	+	-
WT	+++	++	++	++	-	++	++	-	++	+	+	++	++	-	++	+++	-	-

\* Ratings indicate relative levels of detectable staining in: G, glomerular; T, tubular; and A, antinuclear patterns, with values representative of five mice in each group. Immune deposits were rated on a 0-4+ scale, with at least two mice in each group positive for a certain intensity of deposition to be scored; those deposits scored as -, +, ++, or +++ are representative of at least four of the five mice in each group; those scored as +/-, +/+, +/+++ are representative of two mice with one scale of deposits and three mice with another. WT, 12-week-old TCR $\alpha$ + MRL/lpr.





**Fig. 4.** Antibody and autoantibody correlations with renal disease in TCR $\alpha$ + and TCR $\alpha$ - MRL mice, and in age-matched B10.A mice. Elevated serum immunoglobulins and/or autoantibodies show a correlation with renal disease severity (Pearson correlation coefficient,  $r$ , is shown). Renal disease indices were calculated as the sum of light microscopic renal disease scores; antibody indices were calculated on a 1–10 scale, using one point for each elevated immunoglobulin level or positive autoantibody titre (see Materials and Methods).

cell help, in which  $\gamma\delta$  T cells provide an IL-4-rich, Th2-like environment which protects against end-organ disease, skewing the immune response away from a disease-inducing, interferon-gamma (IFN- $\gamma$ )-rich, Th1-like environment characteristic of murine lupus (which may induce complement-fixing IgG antibodies) [8]. Such a view correlates with the substantial development of autoantibodies and accumulation of renal immune deposits despite the delayed and limited progression of renal disease in TCR $\alpha$ - MRL mice, all in association with poor end-organ C3 accumulation, as well as with the results of a recent study describing the protective role of B cell-produced IL-4 in murine lupus [38].

This model also may account for the time course of cutaneous disease, which appears delayed but not necessarily subdued in TCR $\alpha$ - MRL animals. IgG1 autoantibodies have been implicated in the pathogenesis of cutaneous lesions, since they are found at disproportionately high frequencies in MRL/lpr skin [2]. At the same time, however, lesional skin does not contain IgG or complement deposits, probably because the histological architecture has been destroyed by the autoimmune processes ([2] and data not shown). It was therefore difficult to assess the degree of cutaneous immune complex deposition in TCR $\alpha$ - MRL mice, in order to draw conclusions regarding any relationship between

autoantibody production, immune complex deposition, and/or cutaneous lesion development. The autoimmune skin disease in these mice therefore probably reflects the effects of autoantibodies of affinity and/or isotypes that are preserved despite the absence of  $\alpha\beta$  T cells; alternatively, these lesions may reflect primarily cellular autoimmune processes not previously described as important in these mice [36]. Regardless of the particular mechanisms of skin disease, these data emphasize the potential for non- $\alpha\beta$  T cell-dependent mechanisms to enforce overt cutaneous lesions which may be virtually indistinguishable from wild-type disease when fully matured.

As an additional consideration, some components of the  $\alpha\beta$  T cell-independent autoimmunity seen here may result from the absence of regulatory  $\alpha\beta$  T cells [16,17,39]. Accordingly, TCR $\alpha$ - MRL +/+ mice demonstrated increased mortality in association with increased autoantibody production, renal dysfunction and skin disease in comparison with TCR $\alpha$ + MRL mice, but TCR $\alpha$ - MRL/lpr mice demonstrated decreased mortality and autoantibody production in comparison with  $\alpha\beta$  T cell-intact counterparts. Such findings suggest a model for  $\alpha\beta$  T cells in MRL disease in which  $\alpha\beta$  T cells both help and regulate autoreactive B cells, the latter via a Fas-dependent mechanism [40–42]. As such, in MRL +/+ mice,  $\alpha\beta$  T cell deficiency may allow the proliferation of  $\gamma\delta$  T cell-driven B cell autoimmunity, which progresses unchecked in the absence of  $\alpha\beta$  T-dependent, Fas-dependent regulation. In contrast, MRL/lpr  $\alpha\beta$  T cells, unable to regulate other cells via Fas, may only be capable of providing help to B cells; consequently,  $\alpha\beta$  T cell deficiency in the absence of Fas significantly reduces autoantibody production and end-organ disease, decreasing mortality. The revealed  $\gamma\delta$  T cell-dependent autoimmunity would then induce limited disease processes.

$\alpha\beta$  T cell-independent autoimmunity therefore plays a significant role in murine lupus by maintaining the capability to drive systemic autoimmunity. These findings introduce a complexity in the  $\alpha\beta$  T cell-based models for lupus pathogenesis [11,13,14,36], calling for a consideration of both  $\alpha\beta$  and  $\gamma\delta$  T cells in the T cell-directed therapy of human SLE [43].

#### ACKNOWLEDGMENTS

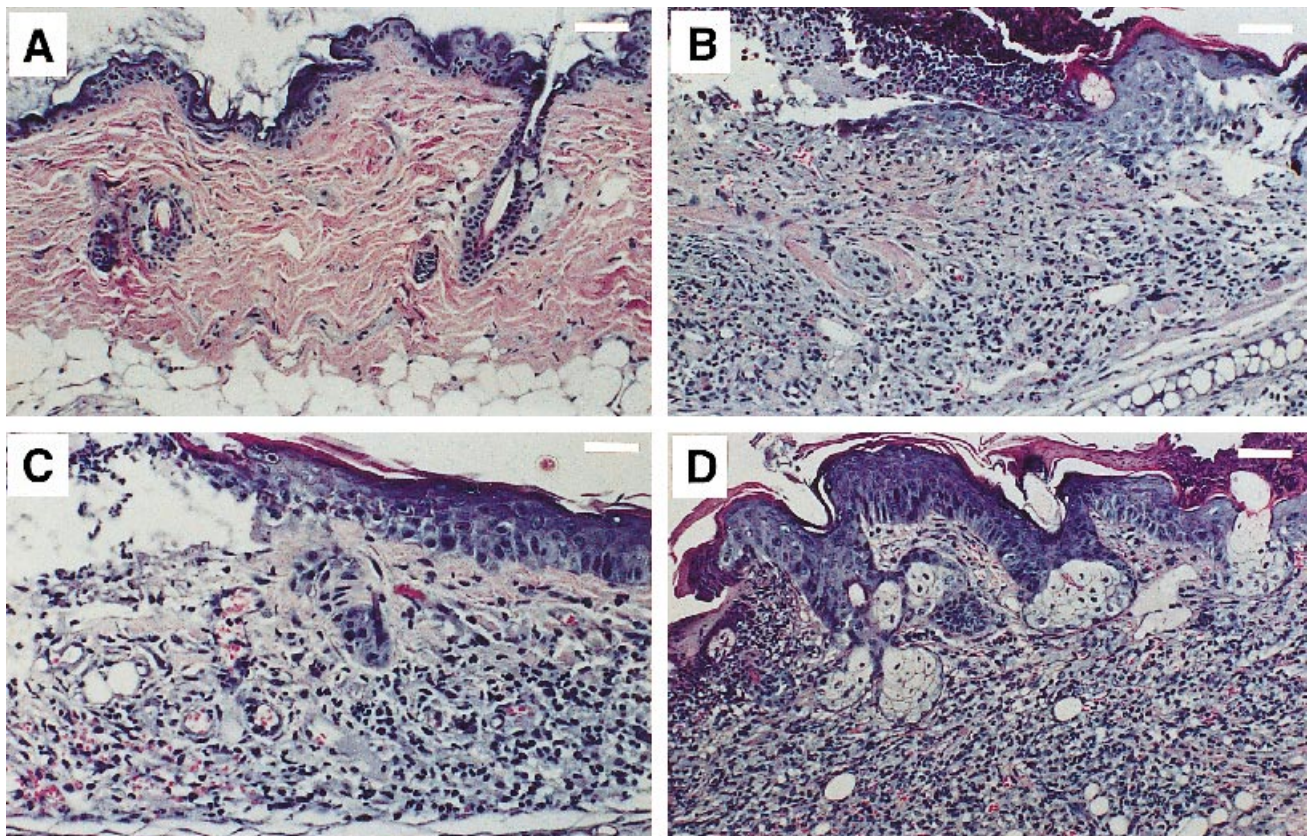
This study was supported in part by grants from the NIH (AR40072 and AR44076 to J.C., and AI27855 to A.C.H.) and from the Arthritis and Lupus Foundations, their Connecticut chapters, and donations to Yale Rheumatology in the memories of Irene Feltman, Albert L. Harlow, and Chantal Marquis (to J.C.). S.L.P. was supported by the Medical Scientist Training Program, Yale University School of Medicine. This study is based partially upon a dissertation submitted to fulfil in part the requirements for the degree of Doctor of Philosophy in Yale University.

**Fig. 3.** Isotyping of immune deposits in TCR $\alpha$ - MRL/lpr mice. Shown are representative kidney specimens from (A–D) TCR $\alpha$ + MRL/lpr and (E–H) TCR $\alpha$ - MRL/lpr mice. All TCR $\alpha$ + MRL/lpr animals consistently had (A) punctate mesangial IgG1 deposition, in contrast to the punctate and diffuse mesangial deposition of (B) IgG2a, (C) IgG2b and IgG3 (not shown) isotypes. Occasionally, tubular and/or antinuclear staining were evident as well in these animals (not shown). TCR $\alpha$ + MRL/lpr mice also consistently developed substantial (D) punctate mesangial C3 deposits. TCR $\alpha$ - MRL/lpr mice, on the other hand, developed predominantly (E) mesangial IgG1 deposition, with a paucity of other IgG isotypes, resulting in negative (F) IgG2a as well as IgG2b or IgG3 (not shown) deposition. Occasionally, however, TCR $\alpha$ - MRL/lpr animals developed substantial immune deposits of IgG2a (not shown), (G) IgG2b or IgG3 (not shown) isotypes. (H) C3 deposition was generally absent in the mesangium of these animals, correlating with their relatively mild glomerular disease, and contrasting with the substantial C3 deposition and severe glomerular disease apparent in their TCR $\alpha$ + MRL/lpr counterparts. Interestingly, the glomerular and tubular membranes of TCR $\alpha$ - MRL/lpr animals had C3 staining; the significance of this finding remains unclear. Shown here are samples from a 12-week-old TCR $\alpha$ + MRL/lpr and a 42-week-old TCR $\alpha$ - MRL/lpr mouse, representative of four to six animals of each kind. Control 12-week-old TCR $\alpha$ + MRL +/+ animals stained negative for all IgG isotypes and C3, similar to (F) (not shown). Scale bar, 25  $\mu$ m.





**Fig. 5.** Cutaneous lesions in  $\text{TCR}\alpha^+$  MRL and  $\text{TCR}\alpha^-$  MRL mice. (A)  $\text{TCR}\alpha^+$  MRL/*lpr* mice generally developed severe hair loss, scab formation and purpuric lesions on the dorsal neck and ears by 24 weeks old. (B) In contrast,  $\text{TCR}\alpha^-$  MRL mice of either Fas genotype at the same age lacked gross skin disease, but soon developed visible lesions, as shown by (C) this 28-week-old  $\text{TCR}\alpha^-$  MRL/*lpr* mouse.  $\text{TCR}\alpha^-$  MRL/*lpr* mice generally developed gross lesions by 7–8 months of age, whereas  $\text{TCR}\alpha^-$  MRL  $+/+$  animals generally developed lesions at 8–9 months of age (not shown). Within 2–3 months of onset, such  $\text{TCR}\alpha^-$  MRL  $+/+$  lesions progressed to phenotypes indistinguishable from  $\text{TCR}\alpha^+$  MRL/*lpr* disease, as in (A).



**Fig. 6.** Dermatohistopathology in  $\text{TCR}\alpha^+$  MRL and  $\text{TCR}\alpha^-$  MRL mice. (A) Some young  $\text{TCR}\alpha^-$  MRL/*lpr* animals had normal skin pathology, as seen in this neck specimen from a 24-week-old animal (scale bar, 200  $\mu\text{m}$ ). (B) In contrast, 24-week-old  $\text{TCR}\alpha^+$  MRL/*lpr* animals consistently developed ulceration, hyperkeratosis, acanthosis, and mononuclear cell infiltration into the dermis and epidermis by 24 weeks old (here, a neck specimen; scale bar, 200  $\mu\text{m}$ ). (C) Many 24-week-old  $\text{TCR}\alpha^-$  MRL mice, however, developed subclinical ulceration, fibrosis, and mild dermal infiltration (here, a  $\text{TCR}\alpha^-$  MRL/*lpr* neck; 50–70% of  $\text{TCR}\alpha^-$  MRL/*lpr* versus 20–30% of  $\text{TCR}\alpha^-$  MRL  $+/+$  animals developed such lesions; scale bar, 125  $\mu\text{m}$ ), which soon developed into (D) florid lesions, such as in this 28-week-old  $\text{TCR}\alpha^-$  MRL  $+/+$  neck, which contained striking infiltrates, as well as hyperkeratosis, acanthosis, and dermal fibrosis (scale bar, 200  $\mu\text{m}$ ).



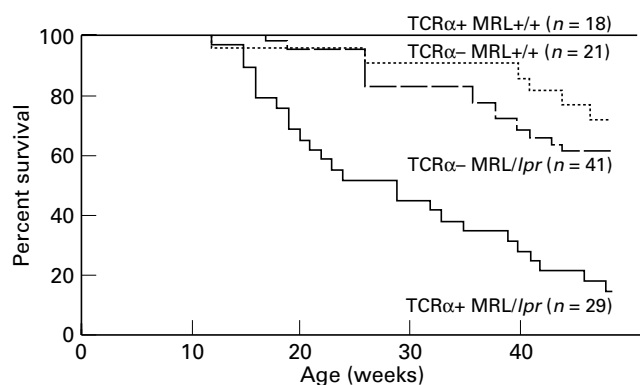


Fig. 7. Life-table analysis of lupus-prone TCR $\alpha$ + MRL and TCR $\alpha$ - MRL mice.

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