

High expression of interleukin-1 β in the corneal epithelium of MRL/lpr mice is under the control of their genetic background

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SUMMARY

MRL/Mp mice bearing the Fas deletion mutant gene, *lpr* (MRL/lpr), spontaneously develop polyarthritis, sialoadenitis and dacryoadenitis, resembling rheumatoid arthritis (RA), and also corneal involvement such as keratopathy and scleritis, which is a major complication in RA patients. In this study, we found that the expression levels of IL-1 β and MMP-1 mRNAs in cornea were high in both MRL/lpr and MRL/Mp-+/+ strains of mice at an age younger than when they develop any inflammatory lesions. This was not true of other inbred strains, even those bearing the *lpr* gene, and also not of (NZB \times NZW) F1 lupus mice. There was no significant difference in the expression of IL-1 α and TGF β in cornea in these strains. Using crosses between MRL/lpr and C3H/HeJ-*lpr/lpr* (C3H/lpr) mice, at least the expression of IL-1 β was found to be under the control of the MRL genetic background, likely with a recessive mode of inheritance. Considering that IL-1 β in cornea was detected particularly in the epithelial layer, the high expression of IL-1 β in cornea is most likely involved in the genetic predisposition for corneal involvement and possibly also for arthritis in an MRL strain of mice.

Keywords rheumatoid arthritis corneal ulcer IL-1 α MMP-1 TGF β genetic

INTRODUCTION

Rheumatoid arthritis (RA) is a systemic autoimmune disease involving polyarthritis, rheumatoid vasculitis, sialoadenitis and dacryoadenitis [1]. Corneal involvement such as scleritis, keratoconjunctivitis sicca and sclerosing keratitis is a major complication in RA patients [2], who frequently have corneal ulceration [3]. The lesion seems to occur rapidly, even in sterile conditions and without trauma [4], and is possibly associated with destruction of collagen matrices of the cornea [2–6].

The major component of corneal stroma is type I collagen [7]. It is well known that MMP-1 and MMP-8 degrade type I collagen, and IL-1 β is a possible inducer of MMP-1 [8–10]. It is likely that IL-1 β is a major factor involved in the destruction of corneal matrices in RA since it is produced significantly in the synovial tissues and other inflammatory lesions in the patients [11,12]. On the other hand, IL-1 β production might occur in the cornea uniquely in RA patients and be followed by corneal ulceration. These possibilities are still controversial.

MRL/Mp-*lpr/lpr* (MRL/lpr) is a unique strain of mice which spontaneously develop polyarthritis, sialoadenitis and dacryoadenitis, and coincidentally corneal lesions such as keratopathy and scleritis [13–15], associated with cytokine abnormalities involving an increased level of IL-1 β in serum [16,17]. In general, MRL/lpr, MRL/+, NZB/WF1, NZB, BXSB and LG/J strains of mice are considered to be multiple murine models of autoimmunity. Among these strains, only those with an MRL background are arthritis-prone and develop corneal lesions, while both MRL/lpr and NZB/WF1 strains are considered useful murine models for Sjogren's syndrome. Thus, this strain of mice may be a model of corneal involvement in rheumatoid arthritis.

In the present study, we found that the production of IL-1 β in cornea is increased in MRL/lpr mice, and we studied its genetic basis. We will present evidence that IL-1 β production in cornea is under the control of the MRL genetic background.

MATERIALS AND METHODS

Animals

MRL/lpr, MRL/Mp-+/+(MRL/+), C3H/HeJ-*lpr/lpr* (C3H/lpr) mice, all of which have an H-2^k haplotype, were obtained from SLC (Sizuoka, Japan). Using MRL/lpr and C3H/lpr mice, we prepared F1 intercross and N2 backcross progenies,

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(MRL/lpr × C3H/lpr) F1 and MRL/lpr × (MRL/lpr × C3H/lpr) F1, respectively. BALB/c (H-2^d), C57BL/6 (H-2^b), DBA/2 (H-2^d), C3H/HeJ (H-2^k), DBA/1 (H-2^q), NZB (H-2^d), and (NZB × NZW) F1 (NZB/W) (H-2^{d/z}) mice were obtained from Japan Charles River (Kanagawa, Japan).

All mice were housed in specific pathogen-free conditions and were used when they were 5–7 weeks old. At this age, MRL/lpr and MRL/+ mice did not have any characteristic histopathological manifestations of inflammatory lesions involving polyarthritis, sialoadenitis, dacryoadenitis and corneal lesions.

Measurements of cytokines

Total RNA was extracted from the corneas of each strain of mice by a single-step procedure [18] using RNeasyTM (Biotex Laboratory; Houston, TX, USA) and cDNAs were prepared using random-hexamer-primed M-MLV reverse transcriptase (SuperScriptTMII RNase H⁻Reverse Transcriptase, GibcoBRL, Gaithersburg, MD, USA). Specific cDNAs were amplified by polymerase chain reaction (PCR) using primer pairs specific for the mouse nucleotide sequences of IL-1 α , IL-1 β , TGF β , G3PDH (Clontech Laboratory, Palo Alto, CA, USA) and MMP-1 (newly synthesized in our laboratory). A hot start amplification procedure was used whereby the reaction mixture without primers and DNA polymerase was overlaid with mineral oil and heated to 94°C for 4 min. After addition of primers and Taq DNA polymerase, incubation proceeded in a Program Temp Control System (Astec; Fukuoka, Japan) for 30 cycles of denaturation, for 45s at 94°C, annealing for 45s at 60°C and extension for 2 min at 72°C.

After separation by electrophoresis through 2% agarose gels, PCR products were detected by ethidium bromide staining. Digital images of fluorescent bands were quantified using Quantity One image-analysing software (Toyobo, Tokyo, Japan). As shown in Fig. 2, the relative expression units of each sample (from the corneas of 10 mice) were calculated by defining the highest value in each electrophoresis series as '100'. In Fig. 4, the relative expression units of each sample were calculated by defining the value of the sample from MRL/lpr, which was used in the experiment of Fig. 2. The relative expression units were standardized according to the values of G3PDH for each sample.

Immunohistochemical staining of IL-1 β

Mouse corneas were fixed with PLP solution (2% para formaldehyde, 0.1 M lysine, 0.05 M phosphate buffer (PB) pH 7.4, 10 mM NaIO₄). Endogenous peroxidase was quenched by incubating with 40 mM gelatin and 20 ml 30% H₂O₂ in 40 ml 0.1 M PB for 20 min. Non-specific protein binding was blocked by covering the sections overnight with blocking solution which consisted of 5% BSA, 5% normal sheep serum and 0.1% NaN₃ in 0.1 M PB, pH 7.4.

Immunohistochemical staining was performed with 1 mg/ml anti-IL-1 β rabbit polyclonal antibody (Genzyme, Cambridge, MA, USA) and biotin-labelled antirabbit sheep IgG (Vector Laboratories, Burlingame, CA, USA). The sections were washed in 0.1 M PB containing 0.5% BSA, and 0.1% Triton-X 100, and the reacted for 1 h at 4°C with streptavidin-peroxidase solution (Vector Laboratories). Then, the sections were treated at room temperature for 10 min with 0.5% cobalt acetate in 0.1 M Tris buffer (TB) after washing sequentially in 0.1 M PB twice, 0.05 M TB and 0.1 M TB. After washing in 0.1 M TB and 0.1 M PB, antigens were visualized by the DAB reaction.

Transmission electron microscopy

Corneal tissues isolated from mouse eye globes were fixed in an improved fixative containing 2.5% glutaraldehyde, 0.1 M cacodylate buffer (pH 7.3), 50 mM L-lysine, and 1% tannic acid, followed by postfixation with 1% OsO₄. En bloc staining was performed with a 2.5% aqueous solution of uranyl acetate, and embedding was done with a Spurr mixture of vinylcyclohexane dioxide. Thin sections were cut with a glass knife and poststained with uranyl acetate and Sorenson's lead citrate. Micrographs were taken on a JEOL 100CX Electron Microscope (JEOL, Peabody, MA, USA).

RESULTS

High expression of IL-1 β and MMP-1 mRNAs in corneas of MRL mice

IL-1 β and MMP-1 mRNAs were highly expressed in the corneas of both MRL/lpr and MRL/+ mice (Fig. 1). However, the corneas from BALB/c and other strains of mice, even NZB/W, which are known to develop autoimmune diseases involving glomerulonephritis, expressed low levels IL-1 β and MMP-1 mRNAs (Fig. 1). Remarkably IL-1 α and TGF β seemed to be expressed in the corneas of all the mouse strains examined.

Figure 2 shows the results of semiquantitative analyses of these results. The expression levels of IL-1 β and MMP-1 mRNAs were almost the same in MRL/lpr and MRL/+ mice, the latter having the same genetic background as MRL/lpr mice except for the *lpr* mutation. This indicates that the *lpr* mutation is not associated with the levels of expression of these cytokines. These values were considerably lower in other strains. However, IL-1 α and TGF β were not particularly highly expressed in both MRL strains compared with other strains examined.

High expression of IL-1 β and MMP-1 mRNAs in the corneas of the MRL strains seemed not to be related to the H-2^k haplotype itself since their expression levels in corneas of C3H/HeJ mice, whose H-2 haplotype is k, the same as that of MRL strains, were much lower and almost the same as those of BALB/c mice (data not shown).

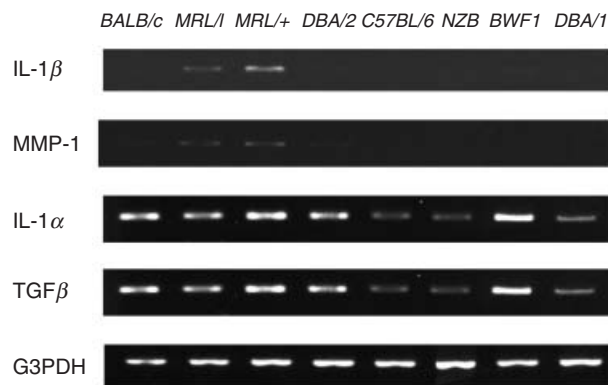


Fig. 1. RT-PCR analyses of IL-1 β and MMP-1 expression in the corneas of various strains of mice. The PCR products from 0.5 mg of total RNA extracted from corneas pooled from 10 eyes of each strain were electrophoresed through 2% agarose gels and detected by ethidium bromide staining. Amplified PCR fragment sizes with specific primers for IL-1 β , TGF β , and G3PDH were confirmed with 563, 525, and 983 bp fragments, respectively, of ϕ X174/Hae III.

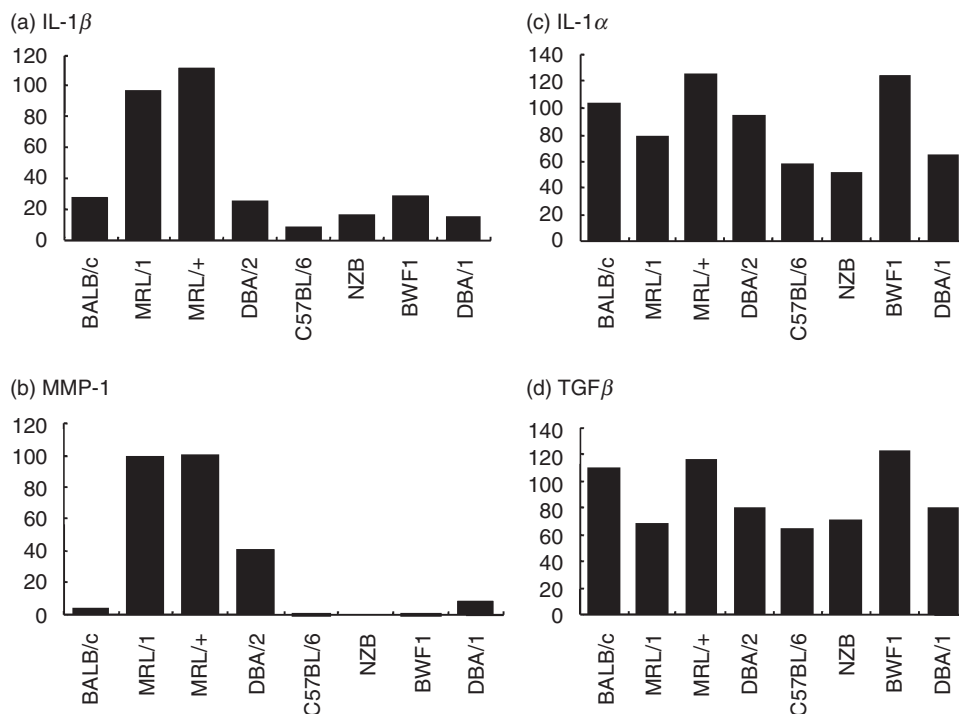


Fig. 2. Semi-quantitative analyses under 'Night Hawk' of PCR products following the procedures described in Fig. 1.

IL-1 β expression localized in the corneal epithelial layer

We employed immunohistochemistry to examine the localization of IL-1 β expression in cornea. As shown in Fig. 3, IL-1 β was expressed in the corneal epithelial layer of MRL/lpr mice although there was no inflammatory cell infiltration into the cornea or degenerative changes of the epithelial cells. Transmission electron microscopy indicated that there were no inflammatory cells in any layer of the cornea of MRL/lpr mice (Fig. 4). Another interesting transmission electron microscopy finding was the decrease in corneal epithelial thickness along with disarrangement of basal epithelial cells in the MRL strains. The corneal epithelium of BALB/c mice consisted of 6–7 cell layers, while that of MRL/lpr mice consisted of only 4–5 cell layers possibly due to the decreased wing cell layers. In addition to the decreased cell density in the basal epithelial layer, the basal cells were spherical in shape (BALB/c mice have a single layer of columnar basal cells). There were no differences in the structure or dimensions of corneal stroma or endothelium between MRL/lpr and BALB/c mice.

Genetic control of IL-1 β expression in cornea

The results described above lead us to speculate that the expression of IL-1 β in cornea may be under the control of the MRL genetic background. To explore this, we examined IL-1 β expression levels in corneas from the F1 and N2 progenies prepared from MRL/lpr and C3H/lpr strains. As shown in Fig. 5, IL-1 β expression levels in the corneas of all (MRL/lpr \times C3H/lpr) F1 mice were significantly lower (less than 80 units) than in MRL/lpr mice (more than 90 units) and almost the same level as in C3H/lpr mice. In MRL/lpr \times (MRL/lpr \times C3H/lpr) F1 mice, there were regular variations in the expression levels as a few mice showed high expression at the same level as some MRL/lpr mice and the majority expressed less than 10 units, the same as the F1 progeny. These findings indicate that IL-1 β expression in cornea is under

the control of the MRL genetic background and that is likely has a recessive mode of inheritance.

DISCUSSION

In the present study, we found that IL-1 β is highly expressed in the corneas of MRL strains of mice under the control of the MRL genetic background. Such IL-1 β expression was not observed in any other strains of mice including NZB/W mice, which spontaneously develop a variety of autoimmune disorders, but not arthritis [19]. With the same PCR amplification conditions, we previously observed a transient elevation of IL-1 β expression after wounding the cornea of BALB/c mice with photoblinding, indicating the validity of the comparison of IL-1 β expression in mouse strains [20].

In the past decade, extensive studies of the genetic basis of arthritis in an MRL strain of mice [21,22] have indicated that arthritis might be under the control of multiple genes in an additive fashion. Those studies did not identify the polymorphism at the IL-1 promoter loci and did not show an association of IL-1 expression with any polymorphism. Since the overproduction of IL-1 β in the synovial cells of joints has been shown to play an essential role in the development of arthritis [11,23], some gene loci controlling the development of arthritis may also be involved in the high expression of IL-1 β in the corneas of these particular strains. Our results from crosses between MRL/lpr and C3H/lpr mice showed that only less than 15% (4/28) of MRL/lpr \times (MRL/lpr \times C3H/lpr) F1 mice had high expression of IL-1 β in their corneas, suggesting that two or more gene loci might affect IL-1 β production in the corneas of MRL mice. Although we performed sequence analysis of the region spanning –500 bp upstream of the transcription start site, which includes IL-1 β promoter [42], there was no polymorphism between MRL and C3H strains. An association

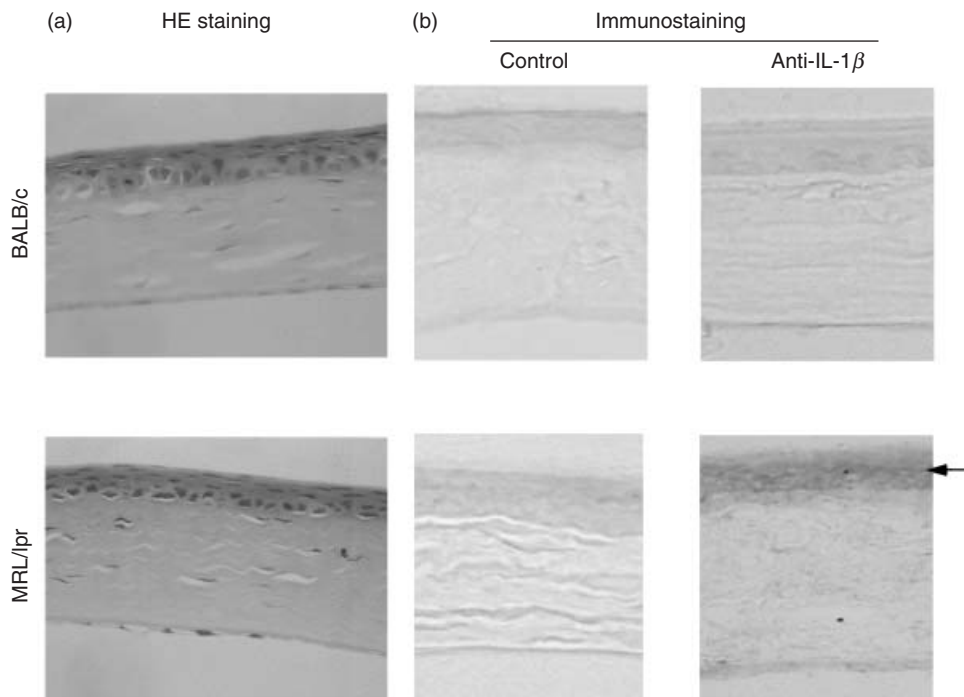


Fig. 3. Histological and immunohistochemical examinations of corneas from MRL/lpr and BALB/c mice. (a) Representative histological pictures of corneas showing no significant differences between the two strains. (H&E staining, $\times 400$) (b) Representative pictures of IL-1 β expression in corneas: left; without specific antibodies, right; stained with anti-IL-1 β antibodies. IL-1 β is specifically localized in the corneal epithelial layer and was observed only in an MRL/lpr strain. ($\times 400$)

study of IL-1 β expression with polymorphic microsatellite markers is needed to further investigate the genetic mechanisms.

Although the mechanism responsible for the high expression of IL-1 β remains unclear, our finding may partly explain the predisposition to corneal pathology common to MRL/lpr mice and RA patients. It is well known that IL-1 β stimulates the production of MMP-1 by chondrocytes [9,10,24,25] or keratocytes [26], which is known to be involved in the degradation of type I collagen, the major structural protein of the corneal stroma [8]. In this study, the expression of MMP-1 was also found to be high in the corneas of MRL strains, while TGF β was not expressed as much as IL-1 β . TGF β is known to suppress collagenase production [27] as well as to promote collagen production by fibroblasts [28,29] and TIMP-1 production by keratinocytes [26] or chondrocytes [30]. Thus, the homeostasis of collagen turnover in the corneal stroma may become imbalanced in MRL strains of mice, leading to an elevation of collagenase activity in the corneal stroma [31,32] or the induction of IL-1-associated keratocyte apoptosis as shown by the study of Wilson *et al.* [33].

It was clearly demonstrated by immunohistochemistry that IL-1 β was intensely expressed in the corneal epithelial layer of an MRL strain of mice. This is consistent with the report by Stephen *et al.* [34] who showed IL-1 expression in cultured human corneal epithelial cells, but not in cultured stromal cells. It is unlikely that IL-1 β was derived from infiltrating inflammatory cells in the cornea or dendritic cells at the limbus because MRL/lpr and MRL/+ mice did not show any inflammatory lesions at those sites. It is still unclear as to what kinds of factors actually promote IL-1 β production in corneal epithelial cells. Extrinsic factors, such as proinflammatory cytokines [17], which might be released from the

surrounding tissue, or mouse tear proteins [35], may exhibit MRL allelic polymorphism and/or be regulated by the MRL genetic background. Lemay *et al.* [36] reported that IL-1 expression depended on the onset of disease and the *lpr* mutation in the kidneys, liver, lymph node and spleen of MRL strain mice. They also reported that lymphocytes highly expressed IL-1 in the MRL/lpr strain mice [37]. To the best of our knowledge, this is the first report investigating the expression of IL-1 β in an organ that is isolated from blood cells and blood vessels.

The decreased thickness of the corneal epithelial layer in the MRL background mice, characterized by the loss of wing cell layers and the disarrangement of basal epithelial cells, is another interesting observation. These changes in the corneal epithelium were not related to the *lpr* mutation, because similar pathological changes were also seen in MRL/+ mice. The decreased number of wing cell layers probably reflects disturbed cell supply or accelerated desquamation of the corneal epithelium. Several investigators reported that IL-1 β induces apoptotic cell death of fibroblasts [38], anterior pituitary cells [39], islets of Langerhans [40], macrophages [41] and keratocytes [33] through nitric oxide production. Thus, one explanation of our finding is that overproduction of IL-1 β from corneal epithelial cells may lead to abnormalities in corneal epithelial structure.

Based on the results of this study, we speculate that the excessive production of IL-1 β by the cornea may be associated with genetic susceptibility and corneal involvement in RA through the acceleration of MMP-1 production. However, there is no report concerning penetrating keratoplasty due to the corneal perforation in this mouse model, although other ocular manifestations in RA patients, such as keratopathy, scleritis and dry eye have been

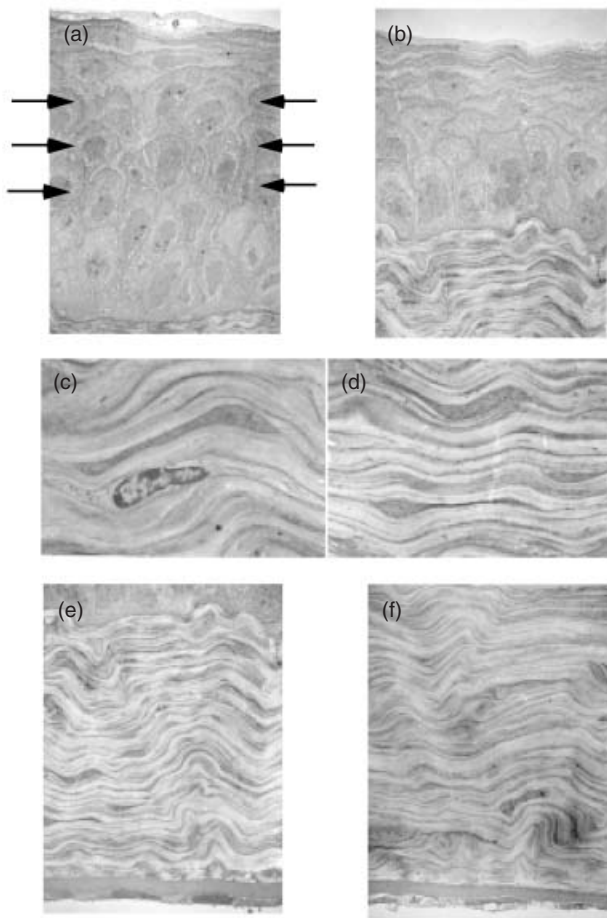


Fig. 4. Transmission electron microscopy of the corneas of BALB/c and MRL/lpr mice (Original magnification $\times 2700$). No inflammatory cells were seen in any layers of the corneas of both strains. The wing cells in (b) the corneal epithelium of MRL/lpr mice were decreased compared with (a) those of BALB/c (arrows). No differences were seen in the structure or dimensions of the corneal stroma or endothelial cells between MRL/lpr (c,e) and BALB/c (d,f) mice.

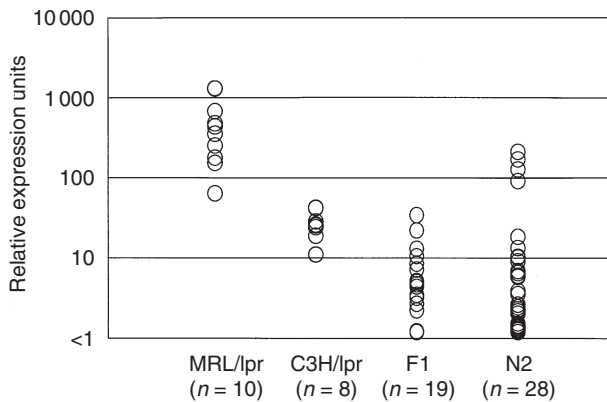


Fig. 5. Relative expression of IL-1 β in corneas of MRL/lpr and C3H/lpr mice and their intercrosses.

investigated. Our preliminary study disclosed that corneal epithelial healing was significantly delayed in MRL/lpr mice compared with BALB/c mice following excimer laser photoablation. This animal model could be a valuable tool for elucidating the

significance of high IL-1 β expression in the corneal pathology of RA patients and its genetic basis.

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