Abstract

The contributions of cytokines to the development and progression of disease in a mouse model of retrovirus-induced immunodeficiency (MAIDS) are controversial. Some studies have indicated an etiologic role for type 2 cytokines, while others have emphasized the importance of type 1 cytokines. We have used mice deficient in expression of IL-4, IL-10, IL-4, and IL-10, IFN-γ, or ICSBP - a transcriptional protein involved in IFN signaling - to examine their contributions to this disorder. Our results demonstrate that expression of type 2 cytokines is an epiphenomenon of infection and that IFN-γ is a driving force in disease progression. In addition, exogenously administered IL-12 prevents many manifestations of disease while blocking retrovirus expression. Interruption of the IFN signaling pathways in ICSBP-/- mice blocks induction of MAIDS. Predictably, ICSBP-deficient mice exhibit impaired responses to challenge with several other viruses. This immunodeficiency is associated with impaired production of IFN-γ and IL-12. Unexpectedly, however, the ICSBP-/- mice also develop a syndrome with many similarities to chronic myelogenous leukemia in humans. The chronic phase of this disease is followed by a fatal blast crisis characterized by clonal expansions of undifferentiated cells. ICSBP is thus an important determinant of hematopoietic growth and differentiation as well as a prominent signaling molecule for IFNs.

Mice infected with the LP-BM5 mixture of murine leukemia viruses (MuLV) develop a syndrome, termed mouse AIDS (MAIDS), characterized by progressive lymphoproliferation and increasing severe immunodeficiency (1,2). A replication-defective virus (BM5def) in this mixture encodes a unique Gag protein that is the critical requirement for disease induction (3,4). Mice injected with BM5def alone develop disease without the requirement for virus spread (5), but most studies of MAIDS have been performed with mice inoculated with ecotropic and MCF helper viruses that mediate cell-to-cell spread of BM5def.

Susceptibility to induction of MAIDS fol-
lowing infection with BM5def is strain dependent, with some mice, such as C57BL/6 (B6) and C57L, developing signs of disease within 2 to 3 weeks of infection, while others, such as A/J, 129 or RIIS, show no signs of disease through a year or more after infection (6-8). These strain differences are genetically determined, and mapping studies have localized strong determinants of resistance to both class I and class II genes of the MHC as well as undefined genes not linked to the MHC (6,8,9). CD8+ T cells contribute to disease resistance (10), but other cells also play an important role, since mice of resistant genotypes deficient in CD8+ cells (β2m knockouts) or perforin (perforin knockouts) develop disease with kinetics that is markedly delayed in comparison with that of fully susceptible mice (Tang Y, Hügin AW, Giese NA, Gabriele L, Kägi D and Morse III HC, unpublished data).

In susceptible mice, the critical viral determinants of disease induction include the MA(p15) and p12 regions of BM5def Gag (11), which differ greatly from the same regions in nonpathogenic ecotropic MuLV. Mature B cells are the primary targets of infection by BM5def, and infection of B cells is required for efficient infection of T cells and macrophages (12,13). Expression of BM5def in B cells and macrophages is not sufficient to induce disease, since mice carrying the nude mutation on a susceptible background express high levels of defective virus but do not develop MAIDS (14,15). However, infected nudes given purified CD4+ T cells do develop disease, indicating a requirement for T/B cell interactions (15). These interactions appear to be antigen-specific, since disease does not develop in mice deficient in expression of MHC class II molecules (15) or CD4 (Morse III HC, unpublished observations). Costimulation would appear to be an additional requirement for development of full-blown disease, as progression of MAIDS is delayed in mice treated with blocking antibodies to CD11a and CD54 (16) or deficient in expression of CD80 (B7-1; Kim Y, Giese NA, Sharpe AH and Morse III HC, unpublished observations) or CD28 (Giese NA, Morse III HC and Abe R, unpublished observations). The CD40/CD40L interactions would also appear to be crucial, since disease is completely blocked in mice treated with antibodies to CD40L (17). In spite of these indications that MAIDS is likely to be antigen driven, the nature of the necessary determinant has not been defined.

In recent years, considerable attention has been focused on the possibility that cytokines may be another critical determinant of T/B cell interactions in MAIDS. Studies of parasitic infections advanced the paradigm that early, polar deviation of responses toward type 1 vs type 2 cytokines determined protective and nonprotective immune responses, respectively. These studies also showed that mice of susceptible strains could be rendered resistant by inhibiting development of the type 2 response or stimulating expression of type 1 cytokines.

Initial studies of supernatants of concanavalin A-stimulated spleen cells from MAIDS-susceptible B6 mice showed that at 1 week after infection there was a burst of cytokine expression with a profile that could be interpreted as mixed type 1/type 2 or Th0 (18). A similar pattern of cytokines was expressed transiently by unstimulated spleen cells from mice infected 2 weeks or more showed that expression of IL-2 and IFN-γ decreased progressively, while expression of IL-4 and IL-10 increased in a reciprocal fashion (18). The observation that serum IgE levels increased in concert with induced IL-4 expression was taken to be an indication of IL-4 expression in vivo and thus a validation of the in vitro assays. These observations had some similarities to those in studies of peripheral blood lymphocytes of HIV+ subjects, suggesting that there might be common
mechanisms contributing to retrovirus-induced immunodeficiency in mice and humans (19). The model proposed was that type 2 cytokines were responsible for downregulation of host cell-mediated immunity governed by type 1 cytokines.

This model was tested in MAIDS-susceptible mice in three ways: first, by determining whether enforced expression of type 1 cytokines would be protective against disease, and second, by judging whether mice impaired in their ability to produce type 2 cytokines would be resistant to MAIDS. A third approach was to study MAIDS-resistant mice for their pattern of cytokine expression with the expectation that they would exhibit a type 1 profile.

The first approach involved treatment of B6 mice with IL-12 to stimulate expression of IFN-γ and to bias T helper development toward Th1. Treatment was initiated at the time of infection or at 2 to 4 weeks after virus inoculation (20). Treatment beginning at the time of infection markedly inhibited the development of lymphoproliferation, as well as signs of B cell activation, and protected the ability of stimulated spleen cells to produce IFN-γ and IL-2. IL-12 – used as a therapeutic agent – also had strikingly beneficial effects in limiting lymphoproliferation and restoring immune function. The beneficial effects of IL-12 were dependent on the expression of IFN-γ, as infected IFN-γ knockout mice were not protected from disease when treated with IL-12 (20). These results were in keeping with the model that type 1 cytokines would mediate a resistant phenotype in mice genetically susceptible to MAIDS.

The first reported studies of mice deficient in expression of IL-4 appeared to lend further strong support to this model (21). Progeny of F₂ mice from crosses between strain 129 IL-4-deficient mice and B6 showed that almost all IL-4+/+ mice developed disease within 12 weeks of infection, whereas IL-4-/- mice generally developed disease after more than 20 weeks. Interpretation of this study is complicated by the facts that strain 129 mice are completely resistant to MAIDS and that the +/+ and -/- mice used in the study were progeny of crosses between selected +/+ and -/- animals instead of being strictly F₂ mice (Kanagawa O, personal communication). This raises the possibility that genes determining the resistance to MAIDS of strain 129 might inadvertently have become fixed in the -/- mice.

At the time of this report, we had backcrossed an independently generated IL-4 defect from strain 129 onto a B6 background. Analyses of these -/- congenics showed them to be indistinguishable from B6+/+ mice for their sensitivity to MAIDS, leading us to conclude that IL-4 is not required for induction or progression of disease (22). This conclusion has since been supported by studies of yet a third IL-4 knockout strain generated, this time, using B6 ES cells. Mice of this strain, which carries no 129 genes, showed the same sensitivity to induction of MAIDS as the congenics tested previously (Morawetz RA, Noben-Trauth N and Morse III HC, unpublished data). In addition, detailed studies of the response to infection of mice from a series of genetic crosses between B6 and 129 mice, some including mice with mutant IL-4 genes and some not, have revealed no association between IL-4 deficiency and resistance to MAIDS (Morawetz RA, Coffman RL and Morse III HC, unpublished data); however, mice homozygous for the Fv-1a allele of strain 129 were much more resistant to disease than mice carrying the B6 allele, Fv-1b. This would be expected from the restriction to spread of B-tropic helper viruses in the LP-BM5 mixture imposed by the allele of Fv-1 (Morawetz RA, Coffman RL and Morse III HC, unpublished data). These findings strongly suggest that MAIDS-resistant mice in the Kanagawa study were homozygous Fv-1a, in addition to being IL-4b<sup>c</sup>, while sensitive mice were most likely Fv-1b and IL-4b<sup>c</sup>.

During the course of these studies, it was found that both IL-4b<sup>c</sup> and IL-4b<sup>c</sup> mice with
MAIDS had comparably high levels of serum IgE (23), even though IL-4 has long been considered to be an absolute requirement for class switching to IgE. The mechanisms involved in IgE expression are fairly well understood, consisting of induction of germline transcripts (ge) followed by switch recombination and the generation of productive transcripts (pe), which are translated to yield IgE protein. Induction of ge following IL-4 engagement of its receptor is thought to occur by activation of JAK1 and JAK3, leading to phosphorylation and activation of STAT6, which enters the nucleus and binds to γ-activated sequences (GAS) in the IgE heavy chain promoter. STAT6 would appear crucial to IgE switching, since STAT6 knockout mice have no serum IgE (e.g., Ref. 24).

The observation that IL-4 is not required for IgE expression in mice with MAIDS suggests that the MAIDS-defective virus may substitute for some protein in the IL-4 signaling pathway, serving to activate JAK/STAT signaling or perhaps acting as a surrogate transcription factor. As an initial approach to understanding IL-4-independent IgE induction in mice with MAIDS, we have studied B cell lymphomas expressing BM5def Gag (Morawetz RA, Rothman P and Morse III HC, unpublished data). Some of these lymphomas were obtained from mice with MAIDS (25), while others were non-MAIDS lymphomas infected with BM5def. Remarkably, several MAIDS lymphomas expressed ge and pe transcripts. In addition, CH12.LX cells infected with BM5def expressed substantially increased levels of ge transcripts, again suggesting that the Gag protein is involved in signaling to the nucleus. None of the lines constitutively expressed activated STAT6, however, even though all were inducible for STAT6 when treated with IL-4. This strongly suggests that the Gag protein does not affect activation of JAK1 and JAK3, which would lead to activation of STAT6. The possibility that the virus might affect expression of IL-13 - which can signal IgE switch recombination in human cells - seems unlikely; B cells from normal or IL-4−/− mice could not be induced to switch to IgE following stimulation with high doses of IL-13 in vitro (Coffman RL, personal communication). While the mechanisms governing IL-4-independent induction of IgE thus remain poorly understood, the initial interpretation of the basis for IgE expression in MAIDS was clearly wrong. Studies are in progress to further define this phenomenon.

The demonstration that IL-4 is not required for normal development of MAIDS left open the possibility that other type 2 cytokines could be required for induction and progression of this disease. Strong evidence against this possibility was generated in studies of IL-10 knockouts and IL-4/IL-10 double knockouts on a B6 background. Both strains of mutant mice developed lymphoproliferation and immunodeficiency with kinetics nearly identical to those of wild-type mice (Morawetz RA, Rajewsky K, Kuhn R, Muller W and Morse III HC, unpublished observations). In addition, MAIDS-susceptible BALB.B mice carrying an IL-6 knockout were normally susceptible to induction of MAIDS (McCarty TC and Morse III HC, unpublished observations). These findings demonstrated that expression of type 2 cytokines in MAIDS was an epiphenomenon of infection rather than a driving force for development of disease.

During the course of these studies, the question of cytokine expression during the induction and progression of MAIDS was readdressed using semiquantitative RT-PCR to evaluate cytokine transcripts. These studies showed that in spleens of infected B6 mice, the levels of transcripts for IL-2, IL-4, IL-10, IL-12, IFN-γ, and TNF-α were all significantly increased at 1 week after infection and that transcripts for all cytokines, except IL-2, continued to increase 4 months after infection (26). PCR analyses of mRNA in spleens of the MAIDS-resistant strain BALB/c showed a pattern of cytokine expression simi-
lar to that of B6 at 1 week postinfection, but after that, the levels of transcripts for all cytokines returned rapidly to preinfection levels. Since it was known that the type 2 cytokines contributed little, if anything, to the development of MAIDS, we examined the possible contributions of IFN-γ expression to disease sensitivity or resistance in studies of infected B6 and BALB/c IFN-γ knockouts, respectively (26). The B6 IFN-γ−/− mice developed MAIDS but with a delayed time course for progression of lymphoproliferation, indicating that IFN-γ was needed for characteristic disease progression. This conclusion is strongly supported by the observation that B6 mice treated with neutralizing mAb to IFN-γ were protected against lymphoproliferation and immunodeficiency (27). IFN-γ was found to play no role in resistance to MAIDS, however, since the BALB/c IFN-γ−/− mice showed no signs of disease at any time after infection.

Taken together, the studies of B6 mice treated with IL-12 and B6 IFN-γ knockouts showed that high-level expression of this cytokine could protect against disease, while low-level expression contributed to disease progression. The potential contributions of type I IFNs (IFN-α/β) to MAIDS sensitivity are controversial. Early studies of serum IFN showed that mice infected for several weeks were unable to produce IFN-α/β after in vivo challenge with Newcastle disease virus (28). Subsequent RT-PCR analyses of IFN-α/β transcripts in spleens of infected mice showed a slow, low-level accumulation of transcripts over time in B6 mice and transient, modest levels of expression in BALB/c and 129 mice. MAIDS-resistant mice treated with neutralizing antibodies to IFN-α/β were unchanged in their response to infection (Giese NA, Morawetz RA, Gresser I and Gabriele L, unpublished observations). This led us to conclude that expression of type I IFNs is induced by infection but is not of importance to resistance to disease.

More recently, others have indicated that an IFN-α/β response is generated within hours of infection of MAIDS-resistant but not MAIDS-sensitive mice (28). They also showed that sensitive mice stimulated to express high levels of IFN-α/β at the time of infection exhibited reduced lymphoproliferation and less pronounced immune defects when compared with untreated infected mice (28). The interpretation was that type I IFNs are important to disease resistance. This view was inconsistent with our findings but, together with the documented importance of type I IFN in MAIDS (29), prompted us to investigate pathways of type I and type II IFN signaling in infected mice.

Both types of IFN signal to the nucleus by activation of different JAK/STAT pathways (reviewed in Ref. 30). Engagement of type I IFN receptors induces activation of JAK1 and Tyk2, which phosphorylate STAT1a, STAT1b, and STAT2. The STATs form heterodimers that associate with p48 (ISGF3γ) to form a complex, ISGF3, that has DNA-binding specificity for sequences designated IFN-stimulated response elements (ISRE). Binding of the complex to ISRE results in transcriptional induction of specific genes that confer cellular responses to type II IFNs. Binding of IFN-γ to its receptor stimulates activation of JAK1 and JAK3, which activate STAT1. Homodimers of STAT1 are transported to the nucleus, where they bind to DNA sequences, designated IFN-γ activation sites (GAS), in the promoters of IFN type II-responsive genes.

These two signaling systems were once thought to be independent, but it is now recognized that there is considerable crosstalk between them. For example, IFN type II response factors (IRF) besides ISGF3γ have specificity for ISRE. One IRF, designated interferon consensus sequence binding protein (ICSBP), has GAS sequences in its promoter and its expression is regulated by IFN-γ (31).

Using electrophoretic mobility shift assays (EMSA) of different IRF family members in nuclear extracts prepared from spleens of mice with MAIDS, it was found that
bands associated with binding of IRF-2 or of complexes formed between IRF-2 and ICSBP were lost within a few weeks after infection (Giese NA, Ozato K, Nelson N and Morse III HC, unpublished data). The loss of these binding specificities was unique to infection with BM5def since it was not observed in mice infected with nonpathogenic ecotropic virus. T cells were critical for this effect as it was not seen in nude mice infected with the LP-BM5 virus mixture. The effect extended to some transcription factors not involved in IFN signaling; YY1 binding was reduced in EMSA. In contrast, other transcription factors such as NFKB were normal (31). Of considerable interest was the finding that, while STAT1 activity was upregulated in MAIDS-resistant mice infected with the virus mixture as well as in sensitive mice infected with ecotropic virus only, it was reduced in MAIDS-sensitive mice.

During the course of these studies, our laboratories were involved in analyses of mice carrying a mutant ICSBP gene generated by gene knockout technology. These studies showed the mice to have two distinct phenotypes, enhanced susceptibility to several virus infections associated with reduced expression of IFN-γ by non-T cells, and an unusual dysregulation of hematopoiesis (32). The latter syndrome resembled chronic myelogenous leukemia in humans in several respects. The parallels included a prolonged phase of neutrophilia followed by a malignant blast cell crisis that was clonal and transferable to normal mice (32).

By EMSA, IRF family binding activities to ISRE by nuclear extracts from spleens of ICSBP−/− mutant mice resembled, in some ways, those seen in mice with MAIDS. ICSBP activity was absent as expected but IRF-2 binding was undetectable, while NFKB binding to a control probe was normal (32). To determine whether signaling pathways regulated by ICSBP were important for the development of MAIDS, ICSBP−/− mice were infected with LP-BM5 viruses. Infection induced striking splenomegaly that, by histopathology, resembled exacerbated chronic myelogenous leukemia with no signs of MAIDS (Giese NA, Masumi A, Ozato K and Morse III HC, unpublished observations). The means by which retroviruses convert a chronic myeloid disease into an acute one are not known but are under study.

Based on this study, MAIDS can be seen as a disease of disordered extracellular and intracellular signaling. Abnormalities induced in B cells by infection with BM5def are not expressed to any great extent without signals provided by T cells. At least some of these abnormalities can be attributed to altered expression of IRF family members and changes in JAK/STAT expression. These observations indicate that development of MAIDS is dependent on modulation of cellular functions regulated by genes with ISRE or GAS sequences in their promoters.

References


