The Intrathymic Pathogenesis of Myasthenia Gravis

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The thymus is considered to play an important role in the pathogenesis of Myasthenia gravis, an autoimmune disease characterized by antibody-mediated skeletal muscle weakness. However, its role is yet to be defined. The studies described herein summarize our efforts to determine how intrathymic expression of the neuromuscular type of acetylcholine (ACh) receptors is involved in the immunopathogenesis of this autoimmune disease. We review the work characterizing the expression of neuromuscular ACh receptors in the thymus and advance a new hypothesis that examines the intrathymic expression of this autoantigen in disease pathogenesis.

Keywords: Myasthenia gravis; Thymus; Acetylcholine receptor; Intrathymic expression

INTRODUCTION

Myasthenia gravis (MG) is a disease characterized by weakness of striated muscles. The weakness is due to impaired neuromuscular transmission resulting from a reduction of the number of receptors for the neurotransmitter, acetylcholine (ACh) at the postsynaptic myoneural junction. This reduction is caused by the action of anti-acetylcholine receptor (anti-AChR) antibodies, reviewed in Levinson et al. (1987). MG is a prototypic autoimmune disease; the immune effector mechanisms and autoantigenic target have been delineated. However, the events leading to the abrogation of self-tolerance to the neuromuscular type of AChR (nAChR) remain a mystery. The thymus gland has long been considered to hold the key to solving this mystery, although the nature of its involvement remains to be elucidated.

Interest in a pathogenic role for the thymus in MG has been fueled by pathologic, clinical and immunologic lines of evidence as reviewed by Levinson and Wheatley (1995). Briefly, thymus glands of 60–70% of MG patients demonstrate the histological pattern of germinal center hyperplasia whereas another 10% display cortical epithelial cell thymomas. Thymectomy, particularly in young patients (<40 years of age) with thymic hyperplasia, is followed by clinical improvement and remains a first-line therapeutic intervention. nAChR-specific B and T cells have been recovered from MG thymus specimens but not control thymus tissue. This indicates that the autoimmune effector cells populate the diseased thymus.

A major feature of the thymus that very likely represents an important pathogenic link to MG is the expression of nAChRs on cells in this organ as reviewed in Levinson and Wheatley (1995). The issue of thymic expression of AChR has attracted considerable interest, in part, because of the pivotal role that self-antigen expression in the thymus plays in tailoring the T cell repertoire. Moreover, the expression of nAChRs on thymic cells represents the first description of the “promiscuous” intrathymic expression of an organ-specific self-antigen that is the target of an autoimmune attack in the periphery. The early reports prompted some investigators to propose that the thymus might actually serve as a site of sensitization for this autoantigen (Wekerle et al., 1978). Taken at face value, this idea might be viewed as being incongruent with the cardinal immunologic construct, noted above, namely that thymic self-proteins, particularly those expressed on epithelial cells orchestrate the induction of self-tolerance (Klein and Kyewski, 2000).

The nAChRs are expressed in two major forms as reviewed in Levinson (2001a). The so-called mature or junctional form is expressed on innervated muscles and the immature or fetal form is expressed on non-innervated tissue. At the mature (innervated) myoneural junction, nAChRs are comprised of four subunits labeled α, β, δ and ε. Two alpha subunits and one each of the other subunits are assembled, like the whalebone in a corset, to form an asymmetric hourglass channel spanning the membrane. Two alternatively spliced alpha subunit
isoforms have been characterized, P3A− and P3A+. The larger P3A+ isoform, which includes an additional sequence of 25 amino acids between exons 3 and 4, is found only in humans and other primates. In fetal muscle, as in adult denervated muscle or nonjunctional membrane, a γ subunit replaces the ε subunit found in mature, innervated muscle endplates.

In the analysis of nAChRs in the thymus, many investigators have focused on the nAChR alpha subunit (nAChRα) since it is the source of the important pathogenic T and B cell epitopes for the pathogenic autoimmune response in MG (Oshima et al., 1990; Zhang et al., 1990; Conti-Fine et al., 1998; Fuji and Lindstrom, 1988). Expression of this subunit was originally reported on a variety of thymic cells including epithelial cells (Engel et al., 1977), thyocytes (Fuchs et al., 1990), and myoid cells (Kao and Drachman, 1977; Schluep et al., 1987). Myoid cells, which share phenotypic properties with skeletal muscle cells, were originally viewed as the principal AChR-expressing cells in thymus (Schluep et al., 1987). They are found in the medulla of both normal and MG thymus.

In the past several years, there has been renewed interest in thymic cells as a source of nAChR expression. Several investigators, including ourselves, have taken a molecular approach in characterizing thymic AChRs and identifying cell populations expressing them. Using reverse-transcription-PCR (RT-PCR) technology, we reported that mRNA for the AChRα was expressed in normal mouse (Wheatley et al., 1992), normal human and MG thymus (Wheatley et al., 1993; Zheng et al., 1999). We also reported that AChRα mRNA was expressed on transformed murine thymic cortical and medullary epithelial cell lines and thymic dendritic cell lines (Wheatley et al., 1992). We found that mRNAs encoding both major isoforms of the human AChRα, i.e. P3A+ and P3A−, were expressed in normal and MG thymus and normal human thymic epithelial cells (Wheatley et al., 1993; Zheng et al., 1999). Sequencing of P3A− and P3A+ cDNA clones recovered from control and MG thymus indicated that they share the same nucleotide sequence as their respective counterparts at the myoneural junction. Thus, unless there are posttranslational changes, the structure of the AChRα proteins expressed in the thymus and the periphery are likely to be identical. These results provided a structural basis for proposing that an immune response directed at thymic nAChRα may be responsible for initiating or perpetuating disease. Berrihof-Akinin subsequently reported that AChRα protein as well as mRNA was expressed on human thymic epithelial cells (Wakkach et al., 1978).

However, there is still a controversy over the expression of the other subunits on thymic cells and whether they are expressed as components of intact receptors. Some of the reported discrepancies may reflect differences in the ages of the thymus donors and differences in the design of the RT-PCRs. At this time, it appears as if ε and β mRNAs are expressed in most normal and MG thymus specimens with variable expression of δ and γ subunits (Navaneetham et al., 2001; Bruno et al., 2004). Expression of the AChR subunits appears to be concentrated in the thymic medullary compartment.

To gain a better understanding of how intrathymic expression of nAChRα might be linked to the development of disease, we addressed additional features of nAChRα expression in the thymus. We observed that the smaller P3A− isoform is present in approximately a five-fold excess in both MG and control thymic tissue and a 2.5-fold excess in a non-transformed human thymic epithelial cell (TEC) line relative to the larger P3A+ isoform (Fig. 1) (Zheng et al., 1999). The greater expression of the P3A− isoform in thymus does not parallel its expression in healthy and MG muscle tissue where both isoforms show equivalent expression (Beeson et al., 1990). These observations suggest that the expression of mRNAs encoding the P3A− and P3A+ isoforms is regulated differently in human thymus and muscle. Since the same disproportionate expression of P3A− was observed in control and MG thymus, it appears that the differential pattern of expression observed in thymus relative to muscle is not a manifestation of thymic pathology in MG. Rather, this pattern may reflect control processes that are unique to these two distinct tissue compartments. Presently, it is not known whether the disproportionate expression of P3A− vs. P3A+ isoforms has pathogenic significance.

We also observed that expression of both P3A+ and P3A− mRNAs are increased in MG thymus compared to control thymus (Fig. 2) (Zheng et al., 1999). This finding parallels that reported for skeletal muscle where AChR mRNA expression was found to be greater in MG muscle than in control muscle (Guyon et al., 1993). The finding of increased AChRα mRNA expression in MG thymus may represent an attempt to compensate for the destructive action of locally secreted anti-AChR antibodies. It is also possible that the increased AChR mRNA expression may reflect the antecedent action of other local environmental factors, e.g. cytokines.

![FIGURE 1](https://www.nature.com/nature/journal/v372/n6507/fig_tab/372739a01.jpg)

**FIGURE 1** Relative expression of AChRα P3A− and P3A+ isoforms in thymus and TEC. Compilation of data from 14 MG thymuses, 7 control thymuses, and 4 separate TEC experiments. The signals for the P3A− and P3A+ bands on Southern blots were quantitated on a phosphorimager. The P3A−/P3A+ ratios are shown. Expression of P3A− exceeded that of P3A+ by a factor of 5.5 ± 0.9 (mean ± SEM) in Control thymus, 4.7 ± 0.05 in MG thymus, and 2.8 ± 0.2 in TEC. (Copyright: Clinical Immunology, 1:1999).
REGULATION OF INTRATHYMIC nAChRα EXPRESSION

IL-1 and IL-6 production by epithelial cells is increased in hyperplastic thymic tissue obtained from MG patients compared to thymus from control subjects (Cohen-Kaminsky et al., 1978; Emilie et al., 1991). Since cytokines produced in vitro by thymic epithelial cell (TEC) lines demonstrate autocrine function (Galy and Spits, 1991), it seemed plausible that these cytokines or perhaps others produced by cells in the thymus might regulate TEC expression of AChR. We examined this possibility by incubating a human non-transformed TEC line with either IL-1β, IL-4, IL-6 and interferon-γ (IFN-γ). We found that neither IL-1, IL-4, nor IL-6 altered the expression of AChRα mRNA by this cell line (Zheng et al., 1999). By contrast, IFN-γ increased expression of the P3A− and P3A+ isoforms by factors of 2.7 and 2.8, respectively. It is known that IFN-γ up-regulates the expression of MHC class II antigens on thymic epithelial cells (Berrí-Aknin et al., 1985; Galy and Spits, 1991). This dual effect of IFN-γ on AChRα and MHC antigens raises the possibility that this cytokine, and perhaps others, may alter expression of thymic AChRα in vivo in a manner that leads to the development or perpetuation of MG. Before addressing this idea, it would be helpful to briefly review the role of self-antigen expression in thymus plays in the development of T cell tolerance and consider how the thymus could serve as a site of immune activation.

INTRATHYMIC EXPRESSION OF SELF ANTIGEN AND THE DEVELOPMENT OF T CELL TOLERANCE

The thymus plays a fundamental role in the generation of the peripheral T cell repertoire as reviewed in Klein and Kyewski (2000), Sprent et al. (1988), Kisielow and Boehmer (1990), Alam et al. (1996) and Anderson et al. (1996). It is generally believed that thymocytes with low affinity receptors for self are positively selected for export to the peripheral lymphoid tissues where they comprise the T cell repertoire that recognizes exogenous antigens (Kisielow and Boehmer, 1990; Anderson et al., 1996). By contrast, T cell tolerance to self is effected largely by the process of central deletion/inactivation. Developing thymocytes with high affinity receptors for self-peptide are silenced by apoptosis or anergy. There is widespread agreement that presentation of self-peptides by cortical epithelial cells is necessary for positive selection (Kisielow and Boehmer, 1990; Alam et al., 1996; Anderson et al., 1996). Thymic medullary epithelial cells and to a lesser extent, bone marrow-derived macrophages and dendritic are considered to be the major APCs involved in negative selection (Blackman et al., 1990; Bonomo and Matzinger, 1993; Hugo et al., 1994; Hoffmann et al., 1995; Klein and Kyewski, 2000). However, central deletion is not complete even though a broad array of self-peptides is “promiscuously” expressed on medullary thymic epithelial cells (Klein and Kyewski, 2000). Self-reactive T cells escape from the thymus in small numbers, perhaps due to the fact that limiting levels of self-antigen limit the efficiency of tolerance induction (Adelstein et al., 1991; Iwabuchi et al., 1992; Oehen et al., 1994). However, such self-reactive T cells are silenced by their anergic or ignorant status, i.e. they never encounter self-antigen in the periphery in a manner that leads to immune activation, or they are suppressed by regulatory T cells (Shevach, 2000).
THE THYMUS AND T CELL TRAFFICKING

Based on the classic studies of Gowans, traffic of lymphocytes is generally considered to be unidirectional, i.e. out of the thymus into the blood and peripheral lymphoid organs (Gowans and Knight, 1964). However, small numbers of peripheral immunocompetent T cells migrate to the thymus, entering via the medulla (Naparstek et al., 1982, 1993; Michie et al., 1988; Hirokawa et al., 1989; Agus et al., 1991; Gossmann et al., 1991; Westermann et al., 1991). Most of the thymic immigrants are T cells activated in the peripheral immune system although even resting T cells may gain access to the thymus (Hirokawa et al., 1989; Agus et al., 1991; Gossmann et al., 1991; Westermann et al., 1991). It is not known if the rate or number of thymic immigrants is increased by an inflammatory reaction in the thymus. Furthermore, it is not known if self-reactive T cell immigrants are activated when they encounter their specific antigens in the thymus. Thymus T cell immigrants specific for the lymphocytic choriomeningitis virus (LCMV) clear infectious foci from the thymus (Gossmann et al., 1991). This observation indicates that peripheral T cells can be activated when they engage specific foreign antigens in the thymus in an appropriate context. When self-reactive T cells encounter their antigens in other tissue compartments in the presence of requisite co-stimulatory signals, they can be activated to express their differentiation program, reviewed in Mondino et al. (1996). One mechanism that leads to a milieu that promotes the abrogation of tolerance peripherally is infection. Local infection can lead to the upregulation of MHC antigens and co-stimulatory molecules on cells that express low levels of self-antigens and thereby lead to activation of autoreactive T cells (Mondino et al., 1996).

A NEW MODEL OF THYMIC INFLAMMATION AND ITS IMPACT ON “RETROGRADE” T CELL MIGRATION

Delineation of the molecular events, particularly in the thymus, that trigger MG has been hampered by the lack of a model system. Thymic pathology is not a feature of experimental models of MG in rodents. Although such models have provided insight into the pathogenesis of MG, they have not served to elucidate the role played by the thymus (Christdassos et al., 2000). To address this issue we have developed a model of inflammation targeted to the thymic medulla, the site of thymic entry by peripheral T cells (Levinson et al., 2001b). We generated molecular variants of the well characterized thymotropic Gross murine leukemia virus (G-MLV), GD17, that had previously been shown to exclusively infect medullary thymic epithelium following their intrathymic injection in naïve mice. The variants were constructed to allow for easy casetting of a broad array of genes of interest. The thymo-tropic MLV vectors were created by ligating a 425 bp fragment containing the U3 region of GD-17 into the LTR backbone of the well defined M-MLV vector LXSH.

The vectors used in our studies are presented in linear form in Fig. 4. The parental LXSH vector includes 5' M-MSV LTR, the psi packaging site and 5' gag region, the hygromycin resistance gene under control of the SV40 promoter, and the 3' LTR of M-MLV. For our experimental protocol, we modified this vector by insertion of the Lac z gene (LBSHG). We utilized LBSHG and LXSHG as our experimental and control vectors, respectively. As was true for GD17, we found that these vectors also target expression of encoded genes to the thymic medullary epithelium.

Balb/c mice were immunized to β-galactosidase (β-gal) and then injected intrathymically (i.t.) with the β-gal encoding vector LBSHG or the control vector LXSHG. Hematoxylin and eosin stained sections of thymus obtained four days after i.t. injection of LBSHG, but not LXSHG, showed obliteration of the cortical/medullary architecture with marked cellular expansion of the medulla. To determine whether this local inflammatory reaction non-specifically augmented the entry of peripheral T cells into the thymus, β-gal immunized mice were injected i.v. with a population of CFSE-labeled CD4+ T cells specific for an unrelated antigen four days after i.t. injection of LBSHG or LXSHG. The CD4+ T cells were derived from a transgenic mouse.

![DIAGRAM]

FIGURE 4 Schematic diagram of MLV-based vectors. The vectors used in these studies are presented in linear form. The parental LXSH vector includes the 5' M-MSV LTR, the psi packaging site and the 5' gag region, the hygromycin resistance gene under the control of the SV40 promoter, and the 3' LTR of M-MLV. Vectors are modified by insertion of either GD-17 U3 or LacZ. (Copyright: Annals of the New York Academy of Sciences, 998:2003).
Peripheral T cells traffic to thymus where they enter the medulla. This facilitates entry of peripheral AChR-reactive CD4+ T cells homing to the thymus is also augmented by a concurrent intrathymic inflammatory response to an unrelated antigen and (2) AChR-reactive T cell immigrants undergo activation following their engagement of autoantigen in this inflammatory milieu, provide help for the production of anti-AChR antibodies by immigrant autoreactive B cells, and thereby promote the development of a myasthenic syndrome.

**CONCLUSION**

There is considerable circumstantial evidence that the thymus plays a pivotal role in the pathogenesis of MG. Nevertheless, the pathogenic link remains to be forged.

We are re-examining the hypothesis that AChR expressed in the thymus drives the pathogenic autoimmune response. We have established a model of intrathymic inflammation that is localized to the thymic medulla and demonstrated that such an inflammatory process promotes the nonspecific entry of peripheral CD4+ T cells into the thymus. We are currently exploiting this model to determine whether (1) AChR-reactive CD4+ T cells entering the thymus is driven by the pathogenic autoimmune response.

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