Autoimmune and inflammatory uveitis are a group of potentially blinding intraocular inflammatory diseases that arise without a known infectious trigger and are often associated with immunological responses to unique retinal proteins. In the United States, about 10% of the cases of severe visual handicap are attributed to this group of disorders. As I discuss here, experimental models of ocular autoimmunity targeting retinal proteins have brought about a better understanding of the basic immunological mechanisms involved in the pathogenesis of uveitis and are serving as templates for the development of novel therapies.

Introduction

The structure of the eye is highly organized and complex (Figure 1), reflecting the high degree of specialization that is required to support its function. The integrity and transparency of the ocular media (aqueous, lens, and vitreous), which refract, transmit, and sense light, are critical to optimal visual function. Any distortion of the visual axis—from the cornea through the anterior chamber, lens, and vitreous body to the retina (Figure 1)—by inflammatory processes within the eye can adversely affect vision. Uveitis or uveoretinitis is a general term referring to inflammation of the retina and uvea (the pigmented vascular coat of the eyeball, consisting of the choroid, ciliary body, and iris). Uveitis is categorized on an anatomical basis as anterior, intermediate, or posterior, or as panuveitis if it involves both the anterior and posterior parts of the eye. Noninfectious uveitis is believed to be autoimmune or immune-mediated (1). Although the distinction between autoimmune and immune-mediated causality can be blurry, the former is generally believed to be driven by aberrant immune recognition of self, whereas the latter is primarily an innate inflammatory reaction triggered by environmental (microbial) or autologous (tissue damage) “danger” signals. Uveitis, especially if untreated, can result in significant visual deficit and blindness. In the United States alone, uveitic diseases of noninfectious origin have an incidence of 52.4 per 100,000 and a prevalence of 115.3 per 100,000, and are believed to account for about 10% of legal blindness (2).

Some uveitic diseases are confined to the eye, such as sympathetic ophthalmia and birdshot retinochoroidopathy. Others are part of a generalized systemic syndrome in which the eye is one of several organs involved. Examples include anterior uveitis associated with juvenile idiopathic arthritis or ankylosing spondylitis, posterior uveitis in Behçet disease (which also involves skin and mucosal tissues), Vogt-Koyanagi-Harada (VKH) disease (which also targets melanocytes of the skin), and ocular sarcoidosis (part of systemic sarcoidosis, a multisystem inflammatory granulomatous disease that also targets the lymph nodes, lungs, and skin) (1, 3) (Table 1). However, many cases of noninfectious uveitis do not fall under a defined classification and are referred to as “idiopathic.” Patients with noninfectious uveitis frequently exhibit immune responses targeted to ocular antigens such as uveal melanin and proteins involved in its metabolism, retinal arrestin (formerly known as the 48-kDa retinal soluble antigen [S-Ag]), interphotoreceptor retinoid–binding protein (IRBP), and recoverin (4, 5). Although it is unclear whether these immune responses represent the etiological cause or an epiphenomenon of autoimmunization to the products of tissue breakdown (especially in diseases where the initial trigger may be microbial; see Table 1), they are believed to fuel progression of the disease. Diseases in this group tend to have strong MHC associations (6), which is believed to indicate an autoimmune etiology, because MHC molecules select and present antigens for recognition by T cells.

While the transparency of ocular media facilitates observation of the disease process using noninvasive techniques, the delicate nature of ocular tissues makes it difficult to obtain tissue specimens. This limitation has greatly hampered studies of the mechanisms involved in uveitic disease. However, over the past several decades, animal models of autoimmune uveitis directed at eye-specific antigens have been developed and have allowed researchers to approach the study of the basic mechanisms driving ocular autoimmunity at a whole new level (reviewed in refs. 7, 8). In this Review, I attempt to put what we have learned from these models in perspective and discuss what they can teach us about human disease.

The immune system and the eye: a tenuous relationship

The eye has evolved to limit intraocular inflammation so as to protect the delicate visual elements from damage that would be detrimental to visual acuity and therefore survival (9). This ability of the eye to curb and control immune responses is known as ocular immune privilege. This term was initially coined by Peter Medawar in the middle of the 20th century, essentially to describe his observation that tissue grafts from non–genetically identical animals placed in the eye were not rejected (10). It was subsequently found by many groups to be a highly complex phenomenon, resulting from the combined effects of efficient separation of the eye from the immune system by a blood-retinal barrier, local inhibition of immune responses by the unique intraocular microenvironment, and systemic induction of immunosuppressive regulatory T cells by eye-specific mechanisms. Ocular fluids contain the immunoinhibitory cytokine TGF-β and other immunoinhibitory substances and are able to curb the destructive activities of inflammatory cells of both the innate and adaptive immune systems (9). However, while it is generally accepted that immune privilege protects the eye from day-to-day inflammatory insults and contributes to the
extraordinary success of corneal grafts, its role in protection from ocular autoimmunity has been contentious.

It has been convincingly demonstrated that elimination in the thymus of self-reactive T cells, a process known as central tolerance, applies to retinal antigens (11). The mechanisms underlying central tolerance rely on an immature T cell interacting with its cognate tissue antigen through its specific TCR. Tissue antigens, including the retinal antigens IRBP and arrestin, are ectopically expressed in the thymus under the control of the gene-regulatory protein autoimmune regulator (AIRE) (12, 13). Mice made deficient for the \textit{Aire} gene and humans with \textit{AIRE} mutations have absent or reduced expression of tissue antigens in the thymus, fail to eliminate autoreactive T cells, and develop multiorgan autoimmunity (in humans, this condition is known as autoimmune polyendocrinopathy–candidiasis–ectodermal dystrophy [APECED]) (14). Normal thymic expression of IRBP in mice detectably eliminates many T cells expressing TCRs that recognize IRBP, thereby reducing the autoreactive uveitogenic T cell repertoire (11). However, thymic expression of retinal antigens among individual humans and mouse strains is variable (15, 16), and levels of expression insufficently high to induce the elimination of T cells bearing cognate TCRs may permit the escape of retinal antigen–specific T cells into the periphery. Indeed, T cells specific for retinal arrestin can be detected in healthy humans by their proliferation in culture in the presence of retinal arrestin with a frequency of up to 4 per 10 million peripheral blood lymphocytes (17). This is likely to be a gross underestimation of their numbers, as methods that do not rely on proliferation but quantitate the frequency of antigen-reactive T cells directly through the use of MHC-peptide tetramers typically detect 20-fold-higher retinal antigen–specific T cell frequencies (18). Additionally, arrestin is only one of at least 10 retinal antigens known to be uveitogenic in animals (8).

T cells reactive to tissue antigens that escape control in the thymus are normally subject to regulation by peripheral tolerance mechanisms, which induce T cells to become nonresponsive (tolerant) to their specific antigen when they encounter that antigen in healthy tissues, but retinal antigens residing in the eye are relatively inaccessible. Thus, circulating retinal antigen–specific T cells are likely to be “ignorant” of their cognate antigen rather than tolerant and can be activated by a chance encounter with antigen, possibly in the form of a microbial component that structurally mimics their cognate tissue antigen (19). Indeed, it has been demonstrated that forced expression of a retinal antigen in the periphery (as a transgene, by retroviral delivery, or by vaccination with naked DNA) results in tolerance and in resistance to the subsequent induction of autoimmunity (20–23). Therefore, by sequestering retinal antigens within the eye and hindering peripheral tolerance, immune privilege may actually predispose to ocular autoimmunity (24).

\textbf{Animal models of uveitis}

Understanding of uveitis has advanced tremendously as a result of the development of animal models of the disease. Experimental autoimmune uveitis (EAU) is induced by immunization of animals with the retinal antigens known to elicit responses in lymphocytes isolated from patients with uveitis. Elicitation of ocular pathology in animals requires coadministration of bacterial adjuvants, such as pertussis toxin and/or CFA, which contains heat-killed tuberculous bacteria. This triggers bacterial pattern recognition receptors on innate immune cells (monocytes, dendritic cells, neutrophils, NK cells, NKT cells, and γδ T cells) and provides the proinflammatory immunological milieu required to steer the adaptive immune response toward an autopathogenic effector pathway (see below, “Basic mechanisms driving uveitis”). In fact, stimulation of the innate immune response can by itself result in ocular inflammation in animal models (25, 26) and possibly in humans (Table 1).
Because no animal model can reproduce the full complexity of the human disease, it is necessary to develop and use a variety of models to represent the different aspects and diverse clinical/immunological manifestations of uveitis. The major models of uveitis and their salient points are summarized in Table 2. In general, EAU models can be divided into induced (by immunization) and spontaneous (although the mice have been genetically manipulated). The “classical” EAU model, induced by immunization with retinal antigen in CFA, has without a doubt been the most useful one in terms of dissecting basic cellular and molecular mechanisms of uveitis. By combining the use of specific immunological reagents and genetically manipulated strains of mice, this model has provided most of the insights into the pathogenesis of uveitis (reviewed in refs. 1, 5, 19, 24, 27, 28). This EAU model appears to mimic central features of human disease in terms of clinical appearance (Figure 2) and numerous studies discussed below (see “Determinants of susceptibility: who is predisposed?” and “Basic mechanisms driving uveitis”) indicate that it is also a reasonable representation of human uveitis in terms of mechanisms.

Among the other induced EAU models, the “humanized” model of EAU, which is induced in HLA class II transgenic mice, merits particular attention. Arrestin is the ocular antigen to which uveitis patients most frequently respond, but it is not effective at inducing uveitis in wild-type mice. However, mice made transgenic for any one of a number of HLA class II alleles, including HLA-DR3 (DRB1*0301), HLA-DR4 (DRB1*0401), HLA-DQ6 (DQB1*0601), and HLA-DQ8 (DQB1*0302) (29), develop EAU after immunization with retinal arrestin (30). In HLA-DR3 transgenic mice, the region of arrestin responsible for pathology was identified to be the same as that recognized by T cells from patients with uveitis (ref. 31; and M.J. Mattapallil and R.R. Caspi, unpublished observations). Because in mice transgenic for human HLA class II alleles the autoantigen is presented by a human MHC molecule, these animals may help to identify the antigenic regions functionally involved in human disease, which is a prerequisite for defining antigen-specific biomarkers and therapies for human uveitis.

The second category of EAU models comprises those in which disease arises spontaneously. Arguably, in this regard, they are more similar to human disease than the induced EAU models; however, they develop in hosts that have been genetically manipulated and therefore are immunologically very different from normal. Nevertheless, these models can teach us about some of the variables that might bring about susceptibility to disease. Typically, spontaneous uveitis occurs under conditions of enhanced antigen-specific T cell affinity and/or frequency. One example is mice lacking the transcriptional regulator AIRE, in which deficient thymic...
expression of tissue antigens such as IRBP and arrestin results in failure to eliminate retina-antigen–specific T cells. These mice develop antibody and T cell responses directed at IRBP that result in uveitis (13, 32). Another example is double-transgenic mice expressing both a foreign protein as a neo-self antigen in the eye and a TCR specific for that protein on most T cells (Table 2). However, data obtained with neoantigen models should be interpreted with some caution, due to uncertainty about whether the location and level of expression of the neoantigen truly mimic those of a native retinal antigen. Furthermore, TCRs specific for neoantigens

<table>
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<tr>
<th>Table 2</th>
<th>Major animal models of uveitis^</th>
<th>Examples of models</th>
<th>Refs.</th>
<th>Comments</th>
</tr>
</thead>
<tbody>
<tr>
<td>Type of model and method of induction</td>
<td></td>
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<tr>
<td><strong>Monkey models of uveitis</strong></td>
<td>Immunization with retinal arrestin emulsified in CFA or Hunter’s adjuvant</td>
<td>Cynomolgus or rhesus monkeys injected in multiple sites at the nape of the neck</td>
<td>104, 105</td>
<td>Used rarely, primarily for preclinical studies. Clinical and histological appearances indistinguishable from human uveitis.</td>
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<tr>
<td><strong>Rat models of uveitis</strong></td>
<td>Immunization with retinal antigen in CFA or adoptive transfer of T cells from immunized donors to naive recipients</td>
<td>Arrestin-induced model in the Lewis rat. Many other retinal and choroidal (melanin) antigens are also uveitogenic in Lewis rats.</td>
<td>70</td>
<td>Was the major EAU model until the mouse model was reported in 1988. Now used by relatively few groups.</td>
</tr>
<tr>
<td><strong>Induced mouse models of uveitis</strong></td>
<td>“Classical” EAU: immunization of WT mice with ocular antigen in CFA or adoptive transfer of immune cells (or cell lines) from immunized donors to naive recipients</td>
<td>EAU induced in B10.RIII, B10.A, and C57BL/6 mice with IRBP</td>
<td>70</td>
<td>Mycobacteria in CFA provide innate “danger” signals that polarize autoimmune lymphocytes toward a proinflammatory phenotype. PTX is needed as additional adjuvant in less susceptible strains.</td>
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<td></td>
<td>“Humanized” EAU: HLA-DR3 transgenic mice immunized or adoptively transferred as above</td>
<td>EAU in HLA-DR3 transgenic mice immunized with retinal arrestin or its peptide fragments</td>
<td>30</td>
<td>Same as for classical EAU</td>
</tr>
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<td></td>
<td>EAU induced by infusion of antigen-pulsed syngeneic DCs</td>
<td>B10.RIII mice are given splenic DCs elicited with Flt3L, matured in vitro, and pulsed with IRBP p161–180. The model has not been reported in other mouse strains.</td>
<td>83</td>
<td>Requires two injections of DC and PTX. Less severe disease than CFA-EAU and appears mostly dependent on Th1 cells.</td>
</tr>
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<td></td>
<td>EIU</td>
<td>Induced by systemic injection of bacterial LPS in rats and in mice or by local intraocular injection in rabbits</td>
<td>25, 26</td>
<td>Rapid onset and short duration. Purely innate-driven; no autoimmune component. Serves as a model for acute anterior uveitis.</td>
</tr>
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<td></td>
<td>MDP-induced uveitis</td>
<td>Induced by direct injection of MDP, a NOD2 ligand, into mouse eyes</td>
<td>35</td>
<td>Mimics uveitis associated with Blau syndrome. Purely innate; no autoimmune component.</td>
</tr>
<tr>
<td></td>
<td>Neo-self antigen expressed in the mouse eye transgenically under a retinal promoter or by retroviral transduction following intraocular injection. Mice are then immunized with the specific antigen.</td>
<td>Transgenic HEL or p-gal under the control of an eye-specific promoter; or retrovirally introduced influenza HA in the retina.</td>
<td>43, 106, 107</td>
<td>Uveitogenic T cells needed to be activated</td>
</tr>
<tr>
<td><strong>Spontaneous mouse models of uveitis</strong></td>
<td>Deficient central tolerance</td>
<td>Spontaneous uveitis in AIRE-deficient mice directed at IRBP; spontaneous uveitis in nude mice implanted with embryonic rat thymus.</td>
<td>13, 32, 108</td>
<td>Higher frequency and higher affinity of cells bearing TCRs specific for IRBP</td>
</tr>
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<td></td>
<td>Double-transgenic mice expressing a neo-self antigen in the retina and an antigen-specific TCR</td>
<td>Spontaneous EAU-like uveitis in mice expressing ocular HEL and a HEL-specific TCR</td>
<td>42–44</td>
<td>Spontaneous nature of disease in these models may at least in part depend on the frequency and affinity of the TCR</td>
</tr>
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<td></td>
<td>Mice transgenic for a human HLA class I antigen associated with uveitis</td>
<td>HLA-A29 transgenic mice. Birdshot retinocochoroidopathy–like disease develops between 8 and 12 months of age</td>
<td>33, 34</td>
<td>The actual target antigen is unknown</td>
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Fit3L, FMS-like tyrosine kinase 3 ligand; PTX, pertussis toxin. ^With the exception of EIU and MDP-induced uveitis, all models are autoimmune.
typically have a much higher affinity for their cognate antigen than those specific for native antigen that escape the thymic process of central tolerance; they are thus not representative of typical autoreactive T cells. An intriguing model is spontaneous uveitis that develops between 8 and 12 months of age in HLA-A29 transgenic mice (33, 34). HLA-A29 is highly associated with birdshot retinochoroidopathy in humans (Table 1). No information is available as to the antigen(s) targeted in HLA-A29 uveitis in humans, and the etiological mechanisms remain obscure. This model may provide a much-needed tool to approach these questions.

The most widely used model of autoinflammatory uveitis driven largely by innate immune mechanisms is endotoxin-induced uveitis (EIU), which can be elicited in rats and in mice by systemic injection of bacterial LPS. The response is acute anterior uveitis of short duration (25, 26) (Table 2). While EIU has been useful for dissecting mechanisms of innate inflammation and examining therapeutic modalities, it is not quite clear which human disease the model represents. In that regard, the recently developed uveitis model induced by intraocular injection of muramyl dipeptide (MDP), a ligand of nucleotide-binding oligomerization domain 2 (NOD2; also known as CARD15), is arguably a better representation of autoinflammatory uveitic diseases such as Blau syndrome, which is associated with mutations in NOD2 (35).

**Determinants of susceptibility: who is predisposed?**

Uveitis in humans has thus far been associated primarily with HLA genes. Strong HLA class I and class II associations have been reported for some uveitic diseases (Table 3). Particularly striking are the associations of sympathetic uveitis and VKH disease with HLA-DR4 (although VKH is also strongly associated with HLA-DQ4) and of birdshot retinochoroidopathy with HLA-A29 (relative risk of 49–224, depending on the study). Because HLA molecules are involved in antigen presentation, HLA associations are thought to reflect recognition of particular antigen(s) and epitopes. Immune responses in sympathetic uveitis and VKH disease target melanin-related antigens (4, 36). In birdshot retinochoroidopathy, Behçet disease, and intermediate uveitis, immune responses to retinal arrestin have been reported (5, 37).

In mice, a “susceptible” H-2 MHC haplotype is a prerequisite for EAU to develop, reminiscent of the strong HLA associations in humans. Examples of EAU-susceptible haplotypes are H-2b (which is found in C57BL/10 and C57BL/6 mice), H-2k (which is found in B10.BR mice), and H-2r (which is found in B10.RIII mice). The hierarchy of susceptibility to EAU after immunization with the retinal protein IRBP is B10.RIII > B10.BR > C57BL (38). Clearly, however, MHC accounts for only a part of the genetic influences predisposing to uveitis. The availability of H-2 congenic mouse strains, which share the same genetic background but have a different H-2 haplotype or share the same H-2 haplotype but differ in genetic background, has allowed researchers to demonstrate that background genes have a major effect on penetrance and severity of disease (38, 39). The effects of these genes are complex and involve all aspects of the immune response, including hormonal responses to stress; levels of ectopic tissue antigen expression in the thymus (which translate to efficiency of central tolerance); and innate responses to environmental stimuli acting as adjuvants, which in turn affect the outcome of adaptive immunity. Integration of these diverse genetic effects with those provided by the environment shapes disease pathogenesis.

**Basic mechanisms driving uveitis**

**Critical checkpoints in the disease process: lessons from animal models.** Animal models of uveitis have helped to identify critical checkpoints in the pathogenesis of the disease (Figure 3). Activated T cells specific for retinal antigens mediate EAU in animals. These cells can transfer disease from immunized donors to genetically compatible naive recipients and orchestrate the entire process of uveitis. It is believed that T cells are also central to the pathogenesis of human uveitis. Strong support for this notion comes from the clinical success of approaches that directly target T cells, includ-
ing macrolide antibiotics such as cyclosporine, FK-506, and myco-
phenolic acid, which inhibit signaling through the IL-2 receptor
pathway that is needed to sustain T cell activation and function,
and antibodies specific for the IL-2 receptor (40, 41). The role of
antibodies to ocular antigens in disease pathogenesis is less clear.
In animals, antibodies in the form of hyperimmune serum are, by
themselves, unable to transfer disease to naive recipients, prob-
ably because they are too large in molecular size to penetrate the
blood-retinal barrier. However, if uveitogenic T cells disrupt the
blood-retinal barrier, serum antibodies exacerbate EAU (30). Here,
I attempt to systematically present what we have learned about
the stages of the disease process using animal models, following
the order presented in Figure 3.

As mentioned above, deficiencies in central and peripheral tol-
erance lead to the presence in healthy individuals of nontolerant
retinal antigen–specific T cells that can be activated to acquire
effector function. The number of these T cells and the affinity
of their TCR for cognate antigen, both of which are determined by
the effectiveness of autoreactive T cell elimination in the thymus,
and the effector T cells that they control share the same antigenic
expression in the thymus. This suggests that some low level of pre-
vention lead to the presence in healthy individuals of nontolerant

Table 3
HLA associations in uveitis

<table>
<thead>
<tr>
<th>Disease; race (refs.)</th>
<th>HLA genes</th>
<th>Relative risk</th>
</tr>
</thead>
<tbody>
<tr>
<td>Birdshot retinochoroidopathy; European descent (37, 109, 110)</td>
<td>A29</td>
<td>49.9–224.35</td>
</tr>
<tr>
<td>Behçet disease; European descent, Asian (111–113)</td>
<td>B51</td>
<td>4.0–18.2</td>
</tr>
<tr>
<td>Anterior uveitis in ankylosing spondylitis; European descent (114–116)</td>
<td>B27</td>
<td>3.8–20.0*</td>
</tr>
<tr>
<td>VKH disease; Asian (117, 118)</td>
<td>DR8</td>
<td>34.3</td>
</tr>
<tr>
<td>Sympathetic ophthalmia; various (119, 120)</td>
<td>DRB1*0405</td>
<td>46.7–74.5</td>
</tr>
<tr>
<td>Sympathetic ophthalmia; various (119, 120)</td>
<td>DQA1*0302</td>
<td>12.0</td>
</tr>
<tr>
<td>Sympathetic ophthalmia; various (119, 120)</td>
<td>DQB1*0401</td>
<td>18.9–41.3</td>
</tr>
<tr>
<td>Sympathetic ophthalmia; various (119, 120)</td>
<td>DPB1*0501</td>
<td>3.8</td>
</tr>
<tr>
<td>Pars planitis associated with MS; European descent (121)</td>
<td>DRB1*0404</td>
<td>5.6</td>
</tr>
<tr>
<td>Pars planitis associated with MS; European descent (121)</td>
<td>DRB1*0405</td>
<td>13.7</td>
</tr>
<tr>
<td>Pars planitis associated with MS; European descent (121)</td>
<td>DQA1*03</td>
<td>3.9–12.8</td>
</tr>
<tr>
<td>Pars planitis associated with MS; European descent (121)</td>
<td>DR2</td>
<td>3.2*</td>
</tr>
<tr>
<td>Pars planitis associated with MS; European descent (121)</td>
<td>DR51</td>
<td>4.8*</td>
</tr>
<tr>
<td>Pars planitis associated with MS; European descent (121)</td>
<td>DR17</td>
<td>14.3*</td>
</tr>
<tr>
<td>Pars planitis associated with MS; European descent (121)</td>
<td>DR3</td>
<td>Not reported</td>
</tr>
</tbody>
</table>

*Relative risk calculated from odds ratio as reported in the reference.
may be immunologically similar in structure to their cognate retinal antigen (antigenic mimicry) (48–50). Microbial components also interact with innate pattern recognition receptors on APCs, generating “danger” signals that are necessary to elicit inflammatory immune reactions. Animal models of uveitis based on immunization of laboratory animals with retinal antigens in CFA, which contains heat-killed mycobacteria, seek to duplicate this principle by introducing the uveitogenic antigen in the context of microbial “danger” signals.

Following exposure to a uveitogenic stimulus, circulating retinal antigen–specific cells become activated and acquire effector function. Experiments with adoptive transfer of retinal antigen–specific T cells versus nonspecific T cells, activated ex vivo and fluorescently labeled, have led to the conclusion that a very small number of infused cells infiltrate the eye within hours after transfer, reaching the eye entirely by chance (51, 52). They recognize their specific antigen there and cause changes within the eye that bring about development of disease several days later.

The APCs within the retina have still not been positively identified. Although there are resident dendritic cells in the retina that could serve as APCs, their expression of MHC class II molecules is low, at least until inflammation is underway (53). CD45 cells derived from the circulation are sufficient to provide APC function, even without eye-resident APCs (54). It can be calculated by infusing labeled retina-antigen–specific T cells that fewer than ten such T cells must reach the eye to trigger the early changes leading to EAU (24). In the meantime, the bulk of the transferred cells take up residence in the spleen, where they proliferate and undergo a maturation process that includes upregulation of the chemokine receptor CXCR3 (55). On day four, the retinal antigen–specific T cells leave the spleen and migrate to the eye, where they reactive and drive the inflammatory process started by the initial infiltrating cells, by secreting cytokines and chemokines, activating the retinal vasculature, and promoting massive recruitment of inflammatory leukocytes from the circulation. It is these recruited leukocytes, which include neutrophils, monocytes, and “nonspecific” polyclonal T cells, that serve as the final mediators of tissue damage (Figure 3). Each of the recruited cell types apparently provides an essential function, as depletion of any one of them reduces the severity of EAU (53, 56, 57).

It should be emphasized that innate immune cells such as γδ T cells, NK T cells, and monocytes/macrophages are important not only as final effectors of inflammatory tissue damage, but also as active participants in shaping the effector T cell response that leads to uveitis. Thus, NK and NKT cells may produce innate IFN-γ that is protective in EAU (58, 59). Monocytic cells, in addition to functioning as inflammatory cells, can also differentiate to a regulatory phenotype, known as myeloid-derived suppressor cells, which inhibit autoregressive T cells (60). Conversely, γδ T cells appear to promote differentiation of autopathogenic effector T cells in EAU by producing cytokines and possibly by acting as APCs (61). The nature of these autopathogenic effector T cells has recently been the subject of much interest and debate and merits its own section. It is discussed in more detail in the section below dealing with pathogenic effector T cells (Pathogenic effector T cells: who is directing the action?).

Figure 3
Critical checkpoints in uveitis, as defined from studies with animal models. Retinal antigen–specific T cells that have not been eliminated in the thymus encounter an activating stimulus in the context of costimulatory “danger” signals, escape from the control of nTregs, and differentiate into pathogenic effector T cells. These undergo clonal expansion, migrate to the eye, break down the blood-retinal barrier, and recruit inflammatory leukocytes from the circulation. The resulting inflammation results in damage to the tissue and release of ocular antigens, which triggers eye-specific regulatory mechanisms that terminate the disease and limit pathology. mTECs, medullary thymic epithelial cells. Modified with permission from Immunological Reviews (24).
While natural regulatory mechanisms set the threshold of susceptibility to disease (46), its termination likely depends on induced regulatory mechanisms. It is well established that T cells exposed to their cognate antigen in the presence of the appropriate environmental cues, such as TGF-β, can acquire a regulatory function that permits them to suppress the effector function of other T cells (62). The eye, as part of its immune-privileged status, is able to promote in vivo induction of systemic regulatory responses such as anterior chamber–associated immune deviation (ACAID) and post-recovery tolerance (reviewed in ref. 63). ACAID is a complex regulatory phenomenon whereby an antigen injected into the anterior chamber of the eye is transported via the bloodstream by eye-derived APCs. There, it induces the generation of antigen-specific CD4+ and CD8+ regulatory T cells through a complex mechanism involving B cells and NKT cells (64). These CD4+ and CD8+ regulatory T cells inhibit the acquisition and expression, respectively, of effector function by antigen-specific T cells. Although its relevance to autoimmunity has been contested (partly due to the fact that most studies used ovalbumin as antigen and delayed-type hypersensitivity as the only readout), ACAID or ACAID-like mechanisms might limit immune reactions to retinal antigens following ocular trauma. In addition, in animals that recovered from EAU, post-recovery CD4+ regulatory T cells have been described, and these cells appear to be distinct from ACAID-induced regulatory cells (65). Their induction is dependent on the presence of the eye (they are not induced in enucleated animals immunized as for EAU) and on expression of melanocortin-5 receptor by T cells (66). It has been proposed that the dependence on the eye reflects involvement of α-melanocyte-stimulating hormone (α-MSH) present in ocular fluids, but direct evidence to support this is lacking. Importantly, some types of syndromes that include uveitis (such as Behçet and VKH) are reported to be associated with reduced regulatory T cell number and/or function, although the origin of the affected regulatory T cells (natural, induced, or both) is not clear (67, 68).

T cells with regulatory phenotype are found in the eyes of rats and mice with EAU and may act locally to control inflammation (69). Although regulatory T cells may enter the eye from the circulation, in vitro data suggest that the ocular microenvironment could be conducive to local induction of regulatory T cells. Ocular fluids contain high levels of TGF-β2 and appear to promote regulatory T cell induction in culture in a TGF-β–dependent manner (63). Furthermore, ocular pigmented epithelia cocultured with “conventional” T cells has been reported to promote conversion of those T cells to regulatory T cells (63). It remains to be demonstrated, however, that such a conversion occurs in vivo, within the living eye.

Pathogenic effector T cells: who is directing the action? Considerable attention has been directed toward determining the lineage and cytokine profile of uveitogenic effector T cells. This issue has direct clinical significance, as it identifies the T cell subset(s) and/or lineage(s) that might be a therapeutic target. CD4+ but not CD8+ T cells are necessary for EAU development. Depletion of CD4+ T cells prevents EAU, and highly purified CD4+ T cells and T cell lines specific for retinal antigens transfer EAU into naive recipients (70). Although retinal antigen–specific CD8+ T cells cultured in the laboratory can induce retinal pathology in rodents (71–73), they are not necessary to elicit EAU. Mice and rats depleted of CD8+ T cells develop full-blown EAU upon immunization with a uveitogenic inducing antigen (74). Furthermore, antigen-specific CD8+ T cells can also act as regulatory cells to control EAU (75–77).

A body of evidence has accumulated over the years implicating CD4+ Th1 effector cells, which predominantly produce the cytokine IFN-γ, in experimental and clinical uveitis (3, 27, 78). However, it has recently become clear that CD4+ effector cells that predominantly produce the cytokine IL-17 (so called Th17 cells) may have an important role in ocular pathology (79, 80). Indeed, neutralization of IL-17 in mice prevents and reverses EAU, in contrast to neutralization of IFN-γ, which exacerbates it (80, 81). This is a still unresolved paradox, which has been seen also in experimental arthritis and experimental allergic encephalomyelitis, and has lent apparent credence to the notion that IL-17–producing Th17 cells, and not IFN-γ–producing Th1 cells, are the true mediators of tissue-specific autoimmunity (78). However, accumulating evidence indicates that Th1 and Th17 cells can both have a pathogenic role (reviewed in ref. 82). In EAU, one or the other lineage can predominate as a pathogenic effector, depending on the model studied. Thus, while EAU induced by injection of IRBP and CFA is IL-17 dependent, and is prevented using neutralizing IL-17–specific antibodies, EAU induced with antigen-pulsed mature dendritic cells seems to require IFN-γ–producing effector T cells, as Ifng-knockout mice fail to develop EAU when injected with uveitogenic dendritic cells that induce EAU in wild-type mice (80, 83). Notably, polarized Th1 and polarized Th17 cells are both able to transfer EAU into naive recipients, independent of the ability of the recipient to produce the reciprocal effector cytokine, suggesting that Th1 and Th17 each represent a fully competent uveitogenic effector phenotype. It thus appears that either a Th1 or a Th1 effector response can drive ocular autoimmunity.

Extrapolating from the models of EAU induced by IRBP in CFA versus antigen-pulsed dendritic cells, the dominant effector Th phenotype appears to be largely determined by the milieu present during initial recognition of the autoantigen by the immune system. Presentation of antigen in the context of mycobacteria (which are present in CFA) by the diverse APC population present in the lymph node appears to promote a dominant Th17 response, whereas antigen presented by dendritic cells matured in vitro promotes a Th1 response. The initial eliciting event(s) connected to human uveitis is largely unknown, but if the two EAU models discussed above are applicable to humans, the events surrounding initial antigen exposure may be critical to the nature of the subsequent disease.

An important lesson to be learned from these findings is that different autoimmune response profiles can be associated with destructive pathology in the eye. Human uveitis is clinically heterogeneous, even though patients may respond to the same retinal antigen(s). Ocular inflammation in humans has been associated in some cases with a Th1-like cytokine profile, and more recently the presence of Th17 responses has been reported (79, 84–87). However, it is still unclear whether such responses in these patients are causally related to uveitis and represent a valid therapeutic target, especially in the case of patients whose uveitis is part of a systemic syndrome, as they could simply be a consequence of an overactive systemic immune response. Ongoing clinical trials targeting IL-17 in uveitis (http://clinicaltrials.gov/ct2/show/NCT00685399) may soon provide answers to these questions.

Therapeutic implications

Ideally, an experimental therapy should be able to prevent disease and reverse a disease process that has already set in. However, therapies that only prevent disease may be of value, since chronic autoimmunity involves constant priming of new lymphocytes.
This process may include immune recognition over time of new antigentic specificities as a result of ongoing exposure to autoantigens released from damaged tissue (88). Historically, therapeutic approaches used successfully to modulate disease in EAU models have often shown efficacy in the clinic; for example, T cell-targeting therapies such as cyclosporine, FK-506, and rapamycin, which are already FDA approved and in clinical use (3, 27, 89). Anti-TNF therapy, which was effective in experimental uveitis (90), has been efficacious in controlling uveitis associated with seronegative spondyloarthopathies and juvenile idiopathic arthritis (91). IFN-α is being used with considerable success to treat uveitis in patients with Behçet disease and has been approved in Europe (although not yet in the United States) for this indication (92). Two other therapeutic approaches that are showing promising results in clinical trials are oral tolerance, which should be explored further, and IL-2 receptor–directed therapy using the humanized CD25-specific antibody daclizumab (27).

Next-generation approaches to therapy might take advantage of tolerogenic administration of retinal antigen to correct defects in peripheral tolerance and to enhance regulatory T cell numbers and/or function. The utility of antigen-specific therapy in humans is suggested by encouraging results in a clinical trial mentioned above using oral tolerance to arrestin (27). Peripheral de novo tolerogenic expression of retinal antigens can be achieved in the adult by naked DNA vaccination with an IRBP expression plasmid or by infusion of autologous B cells engineered by retroviral transduction to express a uveitogenic fragment of IRBP (22, 23). Importantly, this tolerogenic B cell–based therapy has been shown to control EAU in HLA-DR3 transgenic mice (93) and could have significant clinical potential.

The pathogenic effector cytokines produced by uveitogenic T cells and recruited inflammatory leukocytes may serve as targets for inhibition or neutralization. As mentioned above, TNF-α neutralization is already in clinical use (91). The finding that neutralization of IL-17 by monoclonal antibodies in mice aborts EAU, even when instituted after the uveitogenic effector T cells have already been generated (80), suggests potential clinical efficacy. As mentioned above, clinical trials examining IL-17 neutralization therapy in uveitis are ongoing.

Inhibition of migration and recruitment of inflammatory cells into the eye could be achieved by targeting adhesion molecules and/or chemokines and chemokine receptors. Blockade of the integrin very late activation antigen–4 (VLA4) ameliorates EAU in mice (94). As VLA4-specific monoclonal antibody is an FDA-approved treatment for multiple sclerosis and Crohn disease (95), it is conceivable that the approach could be considered as a therapy for uveitis. Similarly, blocking the integrin lymphocyte function–associated antigen 1 (LFA-1) or its ligand intercellular adhesion molecule 1 (ICAM-1) inhibits EAU, as does blockade of CD44 (the receptor for hyaluronan, osteopontin, collagen, and matrix metalloproteinases) (94, 96, 97). Amelioration of EAU has also been achieved by blockade of the chemokine receptors CXCR3 and CXCR5, which are important for migration of uveitogenic and recruited cells into the eye (55, 98).

Future directions

Knowledge of the critical checkpoints in the pathogenesis of uveitis (Figure 3) is providing the clinician with relevant targets for immunotherapeutic intervention. Therapies have been moving away from general immunosuppression toward the use of increasingly specific biologicals, targeting defined aspects of the immune response. Nevertheless, therapies directed at components shared by activated immune cells, such as adhesion molecules or cytokines/chemokines and their receptors, may have the unwanted consequence of inhibiting antimicrobial defense and antitumor surveillance mechanisms (99). Furthermore, they are only symptomatic and do not treat the underlying cause of the breakdown of self tolerance. The “holy grail” for clinicians treating patients with uveitis is a therapy that will uniquely target the retinal antigen–specific T cells that drive the disease. Our knowledge of the retinal proteins that may drive uveitis in humans is increasing; however, to devise effective antigen-specific approaches to diagnosis and treatment, we need to know more about their HLA dependence and epitope specificity, and whether antigen recognition may change during the course of disease. “Humanized” models of uveitis in HLA transgenic mice may be of help in these studies. The examples discussed in this Review underscore the contribution of basic research in animal models to unraveling the mechanisms driving uveal autoimmune and autoimmune inflammatory disease and highlight their importance in the continued development of novel therapies to combat these sight-threatening conditions.

Acknowledgments

I am grateful to Chi-Chao Chan for providing EAU scoring criteria as well as histology pictures of EAU and human uveitis.

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