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Coaxing the liver into preventing autoimmune disease in the brain

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The liver has several unique immunological properties that affect T cell activation and immune regulation. Recent studies have uncovered opportunities for the treatment of genetic disease by directing expression of the functional therapeutic protein to hepatocytes. In a new study in this issue of the *JCI*, Lüth and colleagues demonstrate that hepatic expression of a brain protein is protective against neuroinflammatory disease in a mouse model of human MS (see the related article beginning on page 3403). Suppression of autoimmunity was dependent on transgene expression in the liver and was mediated by induction of antigen-specific CD4⁺CD25⁺Foxp3⁺ Tregs. These findings suggest that the introduction of antigens to the liver may have potential as a preventative or therapeutic intervention for autoimmune disease.

Immune responses are substantially influenced by the context of the tissue in which they occur. For example, brain and liver have both been characterized as immune-privileged sites. The liver is constantly exposed to food antigens and therefore must control inflammatory responses to exogenous antigens. Furthermore, injection of antigen into the portal vein can induce tolerance, and spontaneous acceptance of liver allograft accompanied by donor-specific T cell tolerance has been documented previously (1). The immune system in the brain parenchyma lacks dendritic cells and therefore the ability to prime T cells. However, T cell activation upon immunization at an ectopic site can cause lymphocyte infiltration and inflam-

matory responses targeting antigens in the brain (2). For example, immunization against myelin basic protein (MBP) — a protein important in the process of nerve myelination — causes EAE in certain strains of mice, and these animals have been used as a model for human MS.

On the other hand, it is also known that antigen administration to specific organs can induce suppressive immune responses that ultimately promote tolerance. Oral and nasal tolerance are extensively studied scenarios in which antigen introduced to mucosal surfaces at optimal doses causes activation of Tregs such as Tr1 or Th3 cells (3–5). These subsets of CD4⁺ T cells can suppress immune responses by secretion of the cytokines IL-10 and TGF- β . Similarly, antigen administered to the anterior chamber of the eye, another immune-privileged site, is transported to the spleen via a macrophage-like F4/80⁺ cell (6). Antigen presentation by this cell type in the lymphoid tissue causes activation of CD4⁺ and CD8⁺ Tregs and NKT cells, resulting in immune

suppression. These examples illustrate how the immune system has evolved to generate cellular mediators of tolerance, which are exposed to antigens in the context of one organ and are capable of subsequently transferring tolerance to a different organ.

Gene transfer to the liver induces immune tolerance

The question of how to establish immune tolerance is critical for gene- and cell-based therapies for genetic and autoimmune diseases. Interestingly, a number of reports in recent years have shown that hepatic gene transfer can induce tolerance to a variety of therapeutic proteins (7–13). While there has been evidence of deletion and anergy of transgene product-specific T cells following hepatocyte-derived gene expression, a finding key to our understanding of the mechanism of tolerance induction was the observation of CD4⁺ Treg activation (14–16). Using a T cell receptor transgenic model, the induction of antigen-specific CD4⁺CD25⁺Foxp3⁺ T cells by liver-directed gene transfer with an adeno-associated viral vector has been previously demonstrated (14). These induced T cells appeared phenotypically comparable to naturally occurring Tregs, which are required to prevent autoimmune disease. In other experiments, Tregs induced by hepatic adeno-associated viral gene transfer were capable of suppressing antibody formation and CD8⁺ T cell responses against a coagulation Factor IX gene product and were found to be required for tolerance induction (14, 17).

Nonstandard abbreviations used: MBP, myelin basic protein.

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Figure 1

Prevention of EAE in mice by hepatic expression of MBP. **(A)** Immunization with MBP-specific peptide normally causes an inflammatory T cell response in the brain in the EAE mouse model of human MS. **(B)** In the study by Lüth et al. in this issue of the *JCI* (20), the authors show that hepatic expression of MBP (normally expressed in the central nervous system), accomplished constitutively in liver-specific MBP transgenic mice (not shown) or transiently after gene transfer to liver cells in vivo, results in induction of MBP-specific CD4⁺CD25⁺Foxp3⁺ Tregs. These MBP-specific CD4⁺CD25⁺Foxp3⁺ Tregs suppressed effector CD4⁺CD25⁻ T cells and converted conventional CD4⁺CD25⁻ T cells into CD4⁺CD25⁺Foxp3⁺ Tregs, thereby preventing neuroinflammatory disease. Introduction of antigens to the liver may have potential as a preventative or therapeutic intervention for autoimmune disease.

Moreover, tolerance established by antigen expression in the liver was upheld when the antigen was subsequently expressed in other organs, such as skeletal muscle, or was delivered i.v. (18, 19). Therefore, hepatic gene expression can suppress antibody and cellular immune responses to a specific protein in extrahepatic sites. These findings have important practical implications for the treatment of genetic disease. For example, tolerized animals can now receive supplementary therapies such as enzyme replacement therapy.

Tregs induced by hepatic antigen expression prevent neuroinflammatory disease

In this issue of the *JCI*, Lüth et al. report exciting data that illustrate the far-reaching implications of this hepatic tolerance model (20). The authors demonstrate that expression of MBP in the livers of mice via transgenic and gene transfer approaches prevents the neuroinflammation characteristic of EAE, providing what is believed to be the first evidence that the tolerogenic potential of the liver can abrogate development of autoimmune disease even if the target antigen for the inflammatory T cell

response is located in a distant organ, in this instance the brain (Figure 1).

Mechanistically, the authors showed again that peripheral CD4⁺CD25⁺Foxp3⁺ Tregs mediated tolerance, while the results of experiments in T cell receptor transgenic animals showed no evidence for clonal deletion of MBP-reactive effector T cells, despite some level of expression in the thymus of liver MBP-transgenic mice (20). Importantly, expression of MBP in the skin did not protect against EAE, which indicates that expression of this protein in the hepatic environment is critical. In an elegant set of experiments based on adoptive transfer of Tregs and effector T cells labeled with fluorescent dye, the authors showed that Tregs induced by hepatic expression can turn effector T cells into Tregs by inducing expression of the transcription factor Foxp3. Effector T cells lacking TGF- β receptor II were resistant to this infectious tolerance mechanism, indicating dependence on TGF- β signaling. These findings are reminiscent of TGF- β -dependent suppression of CD8⁺ T cells by CD4⁺CD25⁺Foxp3⁺ Tregs previously reported by Mempel, von Andrian, and colleagues (21) and further substan-

tiate the importance of this cytokine for in vivo suppression.

The current study by Lüth et al. provides insight into how tolerance is manifested following initial induction of Tregs. Consistent with previously published data (12), hepatic-induced tolerance was found to be robust and not broken by immunization with antigen in complete Freund's adjuvant. In the EAE model, tolerant mice lacked the striking inflammation as well as T cell and macrophage infiltrate seen in the brains of control animals after immunization with MBP (20). The present study opens the door to the treatment of autoimmune disease by antigen administration to the liver. More work will be required to test whether ongoing disease can be reversed, as opposed to prevented.

The study also raises several interesting questions about the mechanism of Treg generation, including whether antigen presentation in the liver or in a lymphoid tissue induces CD4⁺CD25⁺Foxp3⁺ Tregs and which antigen-presenting cells mediate Treg induction. It should be noted that the liver represents a unique microenvironment with several specialized antigen-presenting cells, such as Kupffer cells (resident macrophages), liver sinusoidal endothelial cells, and hepatic dendritic cells, which have all been implicated in T cell activation and tolerance induction (1, 22, 23). Furthermore, suppressive cytokines such as IL-10 and TGF- β are expressed in the liver and have important roles in regulating antigen uptake and presentation as well as T cell activation. In the present study, suppression of MBP-specific effector T cells by Tregs likely occurred at the level of the draining lymph nodes of the site of immunization and perhaps in the spleen. It would be exciting if this model could be further developed to study potential interactions between ectopically induced Tregs and the immune system of the brain.

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Toward a broadly protective influenza vaccine

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The current inactivated influenza virus vaccines induce antibodies that protect against closely related virus strains. They do not, however, protect against antibody-escape variants of seasonal influenza A viruses or new pandemic influenza A viruses emerging from non-human reservoirs. Might boosting influenza A virus-specific CD8⁺ T cell memory diminish the danger posed by these variant viruses? Pre-existing CD8⁺ T cell-mediated immunity directed at peptides from conserved internal proteins of the influenza A virus does not prevent infection, but it can promote early virus clearance and decrease morbidity in mice. In this issue of the *JCI*, Lee et al. show that people who have not been exposed to avian influenza A (H5N1) viruses have cross-reactive CD8⁺ T cell memory to a wide range of H5N1 peptides (see the related article beginning on page 3478). These peptides could be used to add a CD8⁺ T cell component to current antibody-focused vaccine strategies with a view to reducing the impact of infection with novel influenza A viruses.

The recent spread of the extremely virulent avian influenza A subtype H5N1 viruses, herein referred to as H5N1, through Asia and to North Africa and Europe has raised serious concerns about the possibility of a novel human influenza pandemic (1, 2). Though the severe disease that can develop

in humans exposed to H5N1-infected birds is rare and sustained human-to-human transmission of the virus has not yet been observed, the three influenza pandemics of the 20th century were all caused by influenza A viruses that originated from birds (3). Variant influenza A (H1N1), A (H3N2), and B viruses also cause regular seasonal epidemics that are associated with substantial morbidity and economic loss. It is bad enough that some 250,000–500,000 (particularly elderly) people die annually from influenza, but what if we should face an event like the 1918–1919 influenza pandemic? That pan-

demic killed in excess of 40 million people worldwide — before the era of rapid air travel and at a time when the global population was less than a third of that today.

Limitations of current influenza vaccines

Inactivated influenza vaccines elicit neutralizing antibody responses that provide reasonable protection against the homologous H1N1, H3N2, and B viruses (4). However, antibody-mediated selection drives changes (known as *antigenic drift*) in the viral HA (H) and neuraminidase (NA; N) surface glycoproteins, which in turn dictate the frequent production of a new vaccine, sometimes as often as annually, as has been the case in each of the last five years. The WHO recommends candidate vaccine virus strains that have been identified among the collections of emerging field isolates supplied by a global network of 124 WHO National Influenza Centers and other diagnostic laboratories and characterized by the four WHO Collaborating Centers for Influenza (in London, Atlanta, Melbourne, and Tokyo). The WHO's recommendations also inform the composition of the live vaccines produced from “cold-adapted” viruses that

Nonstandard abbreviations used: M1, matrix protein 1; NA, neuraminidase; NP, nucleoprotein.

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