mediated by complement bypass systems. Fourth, and probably most important, the type of assay represents the wave of the future in detecting the activity of the complement system in human disease. As these more informative detection systems (like the one used in this report) come into routine clinical use, other examples of these bypass type pathways will likely be uncovered. For human diseases, these more specific and quantitative assay systems will establish which pathway of complement activation is playing a role in disease and elucidate which one to modulate with therapeutic agents.

Finally, a word of caution is in order. These bypass pathways are often not considered by investigators attempting to define the role of the complement system in disease states. For example, C4-deficient animals are widely used to rule out a contribution of the classical pathway and/or lectin pathway in mouse models of human disease. One must be wary of such interpretations in view of bypass cascades that become operative in "deficient" states. Thus the natural maturation of an antibody response to an infectious organism (i.e., to go rapidly into antibody excess) is all that is necessary to trigger these more ancient bypass pathways. Using our current methods, such pathways are not analyzed in clinical medicine or in animal models of human disease.

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The IL-23/IL-17 axis in inflammation
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IL-23 induces the differentiation of naive CD4+ T cells into highly pathogenic helper T cells (Th17/Thb17,γ) that produce IL-17, IL-17F, IL-6, and TNF-α, but not IFN-γ and IL-4. Two studies in this issue of the JCI demonstrate that blocking IL-23 or its downstream factors IL-17 and IL-6, but not the IL-12/IFN-γ pathways, can significantly suppress disease development in animal models of inflammatory bowel disease and MS (see the related articles beginning on pages 1310 and 1317). These studies suggest that the IL-23/IL-17 pathway may be a novel therapeutic target for the treatment of chronic inflammatory diseases.

Th17/Thb17,γ is a new CD4+ helper T cell subset that produces IL-17
Upon antigenic stimulation, naive CD4+ T cells differentiate into 2 subsets, Th1 and Th2 cells, characterized by different cytokine production profiles and effectors.

Nonstandard abbreviations used: CIA, collagen-induced arthritis; IBD, inflammatory bowel disease; IL-1Ra, IL-1 receptor antagonist; R, receptor.
Conflict of interest: The authors have declared that no conflict of interest exists.
The development of autoimmune diseases, such as RA, MS, and IBD, is thought to be mediated by Th1 cells because high levels of IL-12 and IFN-γ are detected in inflammatory sites (9). In addition, treatment with mAbs against IL-12p40 suppresses such disease development in humans and animal disease models (9, 10). However, mice deficient in IL-12p35, IL-12 receptor β2 (IL-12Rβ2), IFN-γ, IFN-γR, or STAT1, which are critical molecules in IL-12/IFN-γ–mediated responses, exhibit an increased severity of diseases such as EAE and CIA (11–13). These observations are inconsistent with the notion that IL-12 is responsible for such disease development. As IL-23, an IL-12 family cytokine consisting of the p19 and p40 subunits, shares the p40 subunit with IL-12 and anti-p40 mAbs inhibit both cytokines, the involvement of IL-23 is suggested. Current evidence suggests that IL-23 is responsible for the differentiation and expansion of Th17/ThIL17 cells from naive CD4+ T cells (2, 4, 14).

In this issue of the *JCI*, Chen et al. report on their use of newly developed anti-IL-23p19 mAbs to demonstrate that IL-23, but not IL-12, is essential for the development of intestinal inflammation (15). They used IL-12p35−/− mice as a model of T cell–mediated IBD (16) and showed that the development of colitis was greatly suppressed by IL-23p19 deficiency but not IL-12p35 deficiency. Exogenous IL-23 administration accelerated the onset of colitis in Rag2−/− mice engrafted with IL-10−/−CD4+ T cells. Notably, IL-17 production was abolished in IL-23p19−/− mice while IFN-γ and IL-4 production were unaffected. IL-17 and IL-6 expression by anti-CD3 mAb–stimulated memory CD4+ T cells were augmented by IL-23, but not by IL-12, indicating that IL-23 can simulate memory CD4+ T cells. This result contrasts with the ability of IL-12 to stimulate naive CD4+ T cells. Moreover, treatment with both anti-IL-6 and anti–IL-17 mAbs significantly ameliorated the severity of the intestinal inflammation induced by IL-23–treated Rag2−/− mice engrafted with IL-10−/−CD4+CD45RBhi T cells. These observations suggest that IL-17 and IL-6 derived from memory T cells are responsible for the development of intestinal inflammation downstream of IL-23.

Also in this issue of the *JCI*, Chen et al. report on their use of newly developed anti-IL-23p19 mAbs to demonstrate the involvement of IL-23 in EAE (17). The authors previously demonstrated that IL-23p19−/− mice are resistant to EAE and CIA; production of IL-17, but not IFN-γ, is almost completely abolished in these mutant mice (3, 18, 19). In contrast, IL-23p35−/− mice exhibited decreased IFN-γ production and increased IL-17 production, suggesting that IFN-γ may suppress IL-17 production. Furthermore, IL-23–induced but not IL-12–induced proteolipid protein peptide–specific T cells are highly encephalogenic (3). Consistent with these reports, the development of EAE was efficiently suppressed by treatment with anti–IL-23p19 or anti–IL-12/IL-23p40 mAbs by inhibiting infiltration of IL-17−, IFN-γ−, and TNF-α–producing CD4+ T cells in the CNS. Disease severity correlated well with serum IL-17 levels; treatment with anti–IL-17 mAbs ameliorated the clinical disease score. Meanwhile, treatment with anti–IFN-γ mAbs exacerbated disease, consistent with previous observations that IFN-γ+ and IFN-γR− mice are highly susceptible to EAE (12, 13). Thus, it is clearly shown that the IL-23/IL-17 pathway, rather than the IL-12/IFN-γ pathway, is critical for the development of autoimmune diseases.
IL-23 causes inflammation through IL-17–dependent and IL-17–independent pathways

Anti–IL-17 treatment did not prevent the onset and relapse of EAE with the same efficiency as anti–IL-23p19 or anti-IL-12/IL-23p40 treatment, suggesting that the effects of IL-23 cannot be explained by the action of IL-17 alone. We also observed that EAE progression is only partially suppressed even in IL-17−/− mice, suggesting the involvement of additional factors (Komiyama et al., unpublished observations). The transcription factor T-bet is responsible for enhancing IFN-γ production and increasing IL-12Rβ2 expression (20). In contrast to IFN-γ−/− mice, T-bet−/− mice are highly resistant to EAE development (11). As Th17/ThIL-17 cells were present in T-bet−/− mice, Chen et al. suggested that Th17/ThIL-17 cells are not sufficient to induce disease and that additional T-bet–dependent factors and/or cell populations play significant roles in disease pathogenesis (17). With respect to this, T-bet expression on dendritic cells, but not on T cells, is required for IL-1 and chemokine production, contributing to the development of collagen antibody–induced arthritis (21).

In the current studies, Chen et al. (17) suggested that IL-23 may directly activate a subset of macrophages and dendritic cells expressing IL-23R, resulting in the production of inflammatory mediators, such as TNF-α and IL-1 (18, 22). However, as progression of diseases such as EAE and CIA is only minimally affected in TNF-α−/− mice and anti–TNF-α treatment of wild-type mice results in more severe MS (23), TNF-α is not critical for the development of this disease. In contrast, IL-6−/− and IL-1−/− mice are significantly resistant, and IL-1 receptor antagonist–deficient (IL-1Ra−/−) mice are more susceptible to EAE (24, 25). Thus, IL-6 and IL-1 likely play important roles in the development of this disease. These observations suggest that IL-23 can induce chronic inflammation through 2 independent pathways: (a) activation of Th17/ThIL-17 cells; and (b) induction of IL-1 and IL-6 production via myeloid cell activation (Figure 2).

Additional IL-17 family molecules may also play an important role in the development of inflammatory diseases. The IL-17 family currently consists of 6 family members; some of these family members, such as IL-17F, share significant amino acid homology with IL-17 (also known as IL-17A), are induced by IL-23, bind the same receptor as IL-17, and are produced by Th17/ThIL-17 cells. Thus, it is possible that these additional IL-17 members may be involved in inflammatory responses. It is important to elucidate the functions of these IL-17 family molecules in normal physiology and in disease pathogenesis.

The Th17/ThIL-17 differentiation mechanism is not yet known

Naïve CD4+ T cells isolated from mice deficient in STAT1, STAT4, or T-bet...
retained the ability to differentiate into Th17/ThIL-17 cells in vitro following TCR stimulation in the presence of IL-23 (2). The generation of Th17/ThIL-17 cells following immunization with antigen stimulation was also normal in mice deficient for STAT4, STAT6, or T-bet (4). Thus, these Th1- and Th2-specific transcriptional factors are not involved in the differentiation of Th17/ThIL-17 cells, indicating that the Th17/ThIL-17 lineage is independent from these classical Th cell lineages. In this issue, Chen et al. (17) demonstrate that T-bet−/− lymph node cells produced IL-17 upon stimulation with anti-CD3 mAbs, consistent with previous reports (2, 4). Exogenous IL-23, however, did not further enhance IL-17 production in T-bet−/− lymph node cells, suggesting that T-bet may influence IL-23 responsiveness during early Th17/ThIL-17 development (17). Thus, the transcriptional factors involved in Th17/ThIL-17 cell development still need to be elucidated (Figure 1). With regard to this, STAT3 was recently implicated in the IL-23R signaling pathway (22).

As IL-12Rβ1, the common subunit of IL-12R and IL-23R, is constitutively expressed in naive CD4+ T cells, IL-12Rβ2 and IL-23R expression are critical for the responsiveness to IL-12 and IL-23 and development of Th1 and Th17/ThIL-17 cell lineages. Only memory and/or activated T cells express IL-23R; naive Th17/ThIL-17 progenitor cells are devoid of this molecule (22). However, when naive CD4+ T cells were stimulated with IL-23 in the presence of anti–IL-4 and anti–IFN-γ mAbs, a large IL-17-producing population was observed, indicating that IFN-γ and IL-4 inhibit the differentiation of Th17/ThIL-17 cells from naive CD4+ T cells. IFN-γ and STAT1 signaling inhibit the differentiation by downregulating the expression of IL-23R (2). Although IL-4 also inhibits Th17 cell expansion, the mechanism governing this suppression is not known (2, 4). Thus, the identification of the signals that induce IL-23R expression on naive CD4+ T cells is crucial in elucidating the mechanisms of Th17/Th17 cell lineage differentiation.

Recently, Veldhoen et al. reported that Treg-derived TGF-β induces the differentiation of Th17/ThIL-17 cells from naive CD4+ T cells in the presence of IL-6 in vitro (Figure 1) (26). TGF-β-mediated Th17/ThIL-17 cell differentiation is promoted by dendritic cell–derived IL-1β and TNF-α. They showed that IL-23 is not essential for the development of Th17/ThIL-17 cells, but required for their survival and expansion through the positive feedback loop that upregulates IL-6, IL-1β, and TNF-α. Thus, current evidence provides us with 2 Th17/ThIL-17 cell differentiation pathways; one is IL-23 dependent and the other is IL-23 independent. Further studies are definitely required to address the precise roles of IL-23 and other factors in the development of Th17/ThIL-17 cells in vivo.

Concluding remarks
While the importance of IL-12 in host defense against bacteria is widely accepted, the role of IL-23 in host defense is not well understood (Figure 2). Recent studies have revealed that IL-12 and IL-23 have different roles in host defense. Mice deficient in IFN-γ, IFN-γR, or STAT1 are highly susceptible to many different pathogens, including Leishmania major, Listeria monocytogenes, and Mycobacterium tuberculosis (27). IL-23 and IL-17 are also important in host defenses against infection. It should be noted that IL-12/IFN-γ are primarily involved in host defenses against intracellular pathogens while IL-23/IL-17 are important for defenses against extracellular pathogens, including Klebsiella pneumoniae (28). This is because IFN-γ stimulates the immune system to kill intracellular bacteria and infected host cells while IL-17 recruits and activates neutrophils. The detailed host defense mechanisms involving IL-23 and IL-17, however, still remain to be elucidated.

IL-17 is produced not only by Th17/ThIL-17 cells, but also by activated CD8+ T cells, TCRγδ+ T cells, and neutrophils (5, 29). We have observed that CD4+ CD8− TCRγδ+ T cells also produce IL-17 in IL-1Rα−/− mice, which spontaneously develop autoimmune arthritis (Komiyama et al., unpublished observations). Development of arthritis in these mice can be completely suppressed by IL-17 deficiency (6). Thus, IL-17 production by cells distinct from Th17/ThIL-17 cells may also be involved in inflammatory responses and host defense mechanisms. It remains unclear, however, which of these producer cells are involved in the different allergic and infectious diseases and how the differentiation pathways of these cell lineages are controlled.

Taken together, accumulating evidence suggests that 3 independent pathways are involved in inflammatory responses: IL-12/IFN-γ, IL-4/IL-5/IL-13, and IL-23/IL-17. These pathways are largely exclusive, although the effect of Th17/ThIL-17 cells on Th1 and Th2 cells is not well understood. Identification of the major immune pathways responsible for the development of each disease is important for its treatment because suppression of 1 pathway may accelerate the others. Therapeutic targeting of the newly discovered IL-23/IL-17 immune axis may prove effective for the treatment of autoimmune and allergic inflammatory responses.

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In blood vessels, endothelia are submitted to constant shear effects and are, under normal conditions, capable of responding to any variation in hemodynamic forces. Caveolae — 50- to 100-nm plasma membrane invaginations present at the surface of terminally differentiated cells and particularly enriched in ECs — are composed of a high sphingolipid and cholesterol content and the protein caveolin-1 (Cav-1). Previous studies have suggested that caveolae and endothelial Cav-1 may regulate the vascular response to altered shear stress. In this issue of the JCI, Yu et al. have examined the role of Cav-1/caveolae in the regulation of flow-induced alterations (i.e., mechanotransduction) in vessels from wild-type mice, Cav-1–deficient mice, and Cav-1–deficient mice re-expressing Cav-1 only in ECs. Their data suggest that caveolae/Cav-1 may act as sensors of altered shear stress and that they also organize the signaling response in stimulated ECs (see the related article beginning on page 1284).

Shear stress: an important regulator of endothelial cell function

In the vasculature, blood vessels must respond rapidly to any external stimuli and especially to any physical change related to modifications in shear stress, which is a function of the blood viscosity and the velocity gradient at the arterial wall. In this context, blood vessels need to adapt and adjust their luminal diameters and their physical properties. ECs are the primary targets of these changes, as they are the first cell type exposed to these forces. One of the earliest findings that suggested an important role for ECs in this process was the observation of the ability of ECs to reorient and change shape during exposure to shear stress conditions. When submitted to steady laminar shear stress, ECs reorient in the direction of the flow and become remarkably elongated. These observations were made both in vitro (1) and in vivo (2, 3). These findings have suggested that ECs can respond and adapt to changes in blood flow. In fact, ECs act as sensors to transduce hydrodynamic forces. Not only does the morphology of ECs change, but other important signaling pathways have been shown to be regulated in response to altered shear stress. Several studies have shown that under laminar shear stress, the rate of EC proliferation is reduced compared with static conditions (4, 5). Oscillating and/or disturbed conditions have major effects on the pathology of the vasculature. In this regard, atherosclerotic lesions have been shown to develop primarily at sites of disturbed or altered blood flow, i.e., at bifurcations, branch ostia, and curved regions (6). Under oscillating and/or disturbed conditions, EC proliferation is increased compared with cells submitted to laminar shear stress and may allow ECs to repair injuries (7, 8). Other important regulatory pathways have now been shown to be activated under shear stress conditions (9–11). These pathways include those involved in changes in endothelial apoptotic, migratory, and permeability prop-

Nonstandard abbreviations used: Cav-1, caveolin-1.

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Role of caveolin-1 in the regulation of the vascular shear stress response

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