

voke a putative GvHD mechanism *per se* but would simply act as the trigger. Indeed the heterogeneous clinical syndromes of GvHD and scleroderma cannot be superimposed, and other autoimmune disease-like syndromes are also observed in GvHD. The same principle could also apply to other ADs seen more commonly in women of child bearing age, such as rheumatoid arthritis and multiple sclerosis. One might predict that we will soon see reports of persisting microchimerism in these and other ADs.

It should also be appreciated that transplantation medicine is currently moving towards the use of microchimerism as a positive tool to induce tolerance by non-myeloablative conditioning and allogeneic hematopoietic stem cell transplantation. This approach attempts to achieve a balance between stable engraftment with reduced recipient immune function, and limited GvHD with reduced donor T-cell numbers⁹. These concepts, successful in certain animal models, are currently being investigated in the treatment of malignant and non-malignant disease, including AD. Their place relative to standard BMT remains to be established¹⁰. It is also hoped and expected that such a stable mixed chimeric state would allow permanent solid organ engraftment without dependency on long-term im-

munosuppressive agents. But clearly, we still fail to understand why some patients with full chimerism do or do not develop GvHD and why other patients with split chimerism may or may not become tolerant to the graft.

The findings of the Nelson and Artlett groups show that the 'absolute' cellular barrier of the placenta is only relative. In the thymus, some thymocytes are capable of escaping thymic control; similarly, the placenta may be leaky enabling a few immunologically reactive fetal cells to cross into the maternal circulation. The capability of such cells to initiate and sustain an autoaggressive process is dependent on class II MHC antigens, cellular, cytokine and growth factor control networks, and other hitherto unknown factors rather than on an 'all or nothing' response. Perhaps, as with low affinity autoreactive T cells that escape thymic deletion, fetal chimeric cells may become activated under certain exogenous conditions, such as concurrent infection, and may then mount an autoaggressive response. This remains speculation.

These reports are a challenge to our understanding of immunological networks. One day we should be able to use microchimerism as a tool for inducing tolerance or for diagnosing it in the quest for selective targeted treatment of AD.

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Two strings to the bow of Ebola virus

Two different forms of Ebola glycoprotein mediate viral entry into endothelial cells and block activation of neutrophils resulting in hemorrhage and an impaired immune response.

MARBURG AND EBOLA virus are filoviruses that cause fulminant hemorrhagic fever in humans and nonhuman primates, killing up to 90 percent of infected individuals. Since the discovery of Marburg virus in 1967 and its better known cousin, Ebola, a few years later, these pathogens have been cause for considerable public and scientific concern. Although it is clear from the recorded history of filovirus outbreaks that they are self-limiting and that no more than one thousand cases have been documented, Ebola's infamy is well-established. With the advent of recombinant DNA technology our knowledge of the filovirus genome and its replication strategies has significantly increased. But these discoveries do not explain why filoviruses are so deadly.

Ebola virus has a single glycoprotein gene that gives rise to different expres-

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sion products (see figure). The transmembrane glycoprotein (GP)—which forms spikes on the viral surface enabling the virus to attach to and invade endothelial cells—is expressed from two reading frames by transcriptional editing^{1,2}. Unedited mRNA derived from the same gene encodes a non-structural smaller glycoprotein (sGP), which is secreted by infected cells. Reporting in *Science*, Yang and co-workers³ now present evidence to suggest that the two different forms, transmembrane GP and sGP, promote disease progression by different mechanisms. They demonstrate that sGP binds to neutrophils and inhibits their activation whereas trans-

membrane GP attaches to endothelial cells, enabling the virus to gain entry. Their study supports the exciting notion that a single viral gene is responsible for paralysis of the host inflammatory response and for damage to the vascular system, which promote, respectively, the rapid progression of infection and the development of hemorrhages characteristic of the disease.

The investigators analysed binding of vector-expressed sGP to normal and transformed human cell lines by immunofluorescence and found that sGP did not interact with endothelial cells but did bind neutrophils through CD16b, the neutrophil-specific form of Fcγ receptor III. They provide good evidence that sGP inhibits activation of neutrophils (although the block could be overcome using very potent activators such as phorbol ester). The interac-

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The transcriptional, translational and proteolytic products of the glycoprotein gene of Ebola virus. Transmembrane glycoprotein (GP) is encoded by two overlapping frames and is produced by transcriptional editing. It consists of subunits GP₁ and GP₂ (linked by S-S disulfide bonds) and mediates binding of Ebola virus to endothelial cells (and presumably to other susceptible cells). Significant amounts of GP₁ are released from infected cells but its binding specificity is not yet known. The secreted small glycoprotein, sGP, is expressed from unedited transcripts of the glycoprotein gene. It binds to neutrophils and appears to inhibit their activation. The first 295 amino acids of sGP and transmembrane GP are identical whereas the carboxy terminal 69 amino acids of sGP are different.

tion of sGP with neutrophils was particularly surprising because it had previously been thought that sGP served as a decoy, binding to protective antibodies and interfering with the specific immune response to transmembrane GP on the viral surface.

To determine the binding specificity of transmembrane GP, Yang and colleagues incorporated the protein into an avian retrovirus (the Moloney leukemia virus) and examined which cells were invaded by the pseudotyped retroviral particles. They showed that transmembrane GP expressed on the retroviral surface enabled the Moloney viral particles to enter endothelial cells. This confirms previous observations demonstrating that transmembrane GP mediates virus entry into cells⁴ and that endothelial cells support filovirus replication⁵.

As clearly indicated by the Yang study, neutrophils and endothelial cells are important targets of the Ebola virus. Of course, there are other cell types that also play an important role in its pathogenesis—cultured human macrophages can be readily infected⁶ and morphological studies in monkeys and guinea pigs reveal early and extensive virus replication in cells of the mononuclear phagocytic system^{7,8}. These findings led to the

concept that the mononuclear phagocytic system is the primary target of Ebola virus. This view is further supported by the observation that virus-induced release of cytokines, such as TNF- α , by mononuclear cells appears to be the major cause of increased endothelial leakiness⁹. Filovirus infection results in depletion of lymphocytes and necrosis of antigen-presenting cells further supporting the notion that filoviral disease has an immunopathological component^{7,9}. It also appears that, as the virus spreads through the organism, the spectrum of target cells expands to include endothelial cells, fibroblasts, hepatocytes, and other cell types.

There is evidence that transmembrane GP plays a crucial role in determining spread of infection. GP is

cleaved at the post-translational level into the disulfide-linked fragments GP₁ and GP₂ (see figure)¹⁰. The same type of processing occurs with many other viral glycoproteins and is usually a precondition for virus infectivity. Interestingly, the transmembrane GP of the Reston strain of Ebola, which at least in man apparently does not cause disease, is less efficiently cleaved (because it has a different cleavage site) than the transmembrane GP of highly pathogenic strains. Such differences in glycoprotein susceptibility to cleavage are responsible for variations in the pathogenicity of avian influenza viruses and paramyxoviruses. Thus, efficient cleavage of transmembrane GP may be an important determinant of the pantropism of filoviruses.

Another secreted soluble glycoprotein, cleavage fragment GP₁, is also shed in large amounts from cells infected with Ebola virus. Notably, the highly pathogenic Marburg virus—which causes comparable disease in man and monkeys—releases GP₁ in similar amounts to Ebola but does not produce sGP because its glycoprotein gene is organized differently. Furthermore, there are highly pathogenic Ebola virus variants⁷ that only secrete minute amounts of sGP (ref.10). However, the apathogenic Reston sub-

type produces large quantities of sGP (ref.2) but less GP₁ than other pathogenic strains¹⁰. So, the significance of sGP and GP₁ in the pathogenesis of filovirus infections remains to be clarified.

The work of Yang and colleagues raises several important points. Because transmembrane GP has at least some specificity for endothelial cells, it has potential as an agent for targeting retroviral vectors (or other viral vectors) carrying therapeutic genes to the vascular endothelium. This could translate into treatments for cancer or cardiovascular disease. Moreover, blocking the binding of sGP to neutrophils and of transmembrane GP to endothelial cells may constitute new approaches to treating Ebola virus infection. Developing drugs that interfere with proteolytic cleavage of transmembrane GP may be another therapeutic avenue. However, the molecular mechanisms of filovirus pathogenesis must be fully elucidated before therapeutic strategies for interfering with the different steps of infection can be successfully designed and implemented.

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