Immunopathology of highly virulent pathogens: insights from Ebola virus

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Ebola virus is a highly virulent pathogen capable of inducing a frequently lethal hemorrhagic fever syndrome. Accumulating evidence indicates that the virus actively subverts both innate and adaptive immune responses and triggers harmful inflammatory responses as it inflicts direct tissue damage. The host immune system is ultimately overwhelmed by a combination of inflammatory factors and virus-induced cell damage, particularly in the liver and vasculature, often leading to death from septic shock. We summarize the mechanisms of immune dysregulation and virus-mediated cell damage in Ebola virus–infected patients. Future approaches to prevention and treatment of infection will be guided by answers to unresolved questions about interspecies transmission, molecular mechanisms of pathogenesis, and protective adaptive and innate immune responses to Ebola virus.

Ebola virus is an enveloped negative-strand RNA virus of the Filoviridae family, a group of viruses capable of inducing a severe hemorrhagic fever syndrome in humans and nonhuman primates. The virus was first recognized in 1976 during an outbreak in the Ebola River valley in Zaire (now the Democratic Republic of the Congo), Africa. A second outbreak caused by a distinct but related virus occurred in Sudan later the same year1,2. Since its discovery in central Africa, several outbreaks have recurred over the last 30 years, including a current confirmed outbreak (11 September 2007) in the Democratic Republic of the Congo (http://www.who.int/csr/don/2007_09_11/en/index.html). Although the reservoir of virus in nature and the range of intermediate hosts is not fully understood, recent studies have found that fruit bats may support replication of Ebola virus, indicating that these animals may be involved in the life cycle of the virus. However, the natural host of Ebola virus in the absence of active outbreaks, together with the important question of how it is transmitted among various species, represents a continuing subject of investigation.

Human infections usually occur after direct contact with virus in dead or infected people or wildlife, with subsequent person-to-person transmission. Filoviruses enter the body through mucosal surfaces or skin abrasions or through the use of contaminated needles3 (Fig. 1a). The onset of Ebola virus–induced disease is sudden, with a 4 to 10 day incubation period. Patients initially show nonspecific flu-like symptoms such as fever, chills, malaise, muscle aches and headache. Abdominal pain, nausea and vomiting may follow, and a cough, sore throat or diarrhea may also be present. A rash often appears around day five and is a characteristic feature of filovirus infection. Systemic, gastrointestinal, respiratory, vascular and neurologic manifestations result from extensive viral replication and, necrosis is seen in many organs, including the liver, spleen, kidneys and gonads. The terminal stage of the disease is characterized by coagulation disorders such as disseminated intravascular coagulation, fluid distribution problems, hypotension and hemorrhage due to liver inflammation and compromise, tissue disruption and a breakdown in endothelial barrier function that leads to increased vascular permeability. In fatal cases, death occurs typically between 7 and 16 days after infection, the result of multiple organ failure and the onset of a syndrome that resembles severe septic shock4. There are currently no antiviral drugs to treat infection and the mortality rates for the more virulent Zaire and Sudan species of the virus range from 40–90%5.

Host immune response to fatal Ebola infection

The uncontrolled viral replication of Ebola virus is central to its pathogenesis, both because of its cytopathic effects and because it induces prominent dysregulation of the host immune response. Virally induced immune system impairment occurs through a variety of mechanisms. Studies in nonhuman primates as well as guinea pigs raise the possibility that monocytes, macrophages and dendritic cells are early and preferred sites of viral replication6,7, though it remains possible that virus is present on these cells through binding to lectin receptors rather than active replication in vivo. It has been suggested that these cells act as vehicles for the transport of virus through the lymphatics8. Further viral replication ensues, followed by systemic spread to other organs and tissues (Fig. 1b). Although virus is observed in the reticuloendothelial system, little inflammation is seen within the lymphatics or in infected tissues during the course of the infection.

Infection of monocytes and macrophages leads to the release of proinflammatory cytokines and chemokines, including tumor necrosis factor, interleukin-1β, macrophage inflammatory protein-1α and reactive oxygen and nitrogen species9,10,11,12. The expression of these mediators is likely to attract more monocytes and macrophages to the sites of infection and may also attract neutrophils. Although recent data suggests that they are not productively infected, human neutrophils

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Infection, spread and target cell destruction by Ebola virus. (a) Ebola virus (yellow) infects subjects through contact with body fluid or secretions from an infected patient and is distributed through the circulation. Entry can occur through abrasions in the skin during patient care, burial rituals and possibly contact with infected bushmeat, or across mucosal surfaces. Accidental needle stick is the primary route of occupational exposure. (b) Early targets of replication are reticuloendothelial cells, with high replication in several cell types within the lungs, liver and spleen. (c) Dendritic cells, macrophages and endothelium appear to be susceptible to cytopathic effects of Ebola virus gene products in vitro and possibly in vivo through disruption of cellular signaling pathways affected by virus binding, phagocytic uptake or both. Indirect damage may also be inflicted by circulating factors such as tumor necrosis factor and nitric oxide.

Figure 1

Like monocytes and macrophages, immature dendritic cells (DCs) are ‘targets’ of Ebola virus, either by means of attachment of viral particles through interactions with DC-expressed C-type lectin DC-SIGN or by means of infection through interaction with other DC-expressed cell-surface receptors (Fig. 1c). Dendritic cells are among the most effective antigen-presenting cells of the immune system, and they secrete critical interleukins and cytokines that provide a critical link between innate and adaptive immune responses to many pathogens; DCs infected with Ebola virus are severely compromised in these critical functions. Human myeloid DCs infected with live virus in vitro, for example, fail to secrete the normal profile of proinflammatory cytokines and costimulatory molecules. These cells do not become mature or activated and are unable to upregulate major histocompatibility complex (MHC) molecules and thus to stimulate T cells16,17. By contrast, treatment with noninfectious Ebola virus–like particles (VLPs) activates DCs and stimulates a robust inflammatory response14, an effect dependent on the mucin-like domain of the envelope glycoprotein15. Inhibition of DC function with live or inactivated virus, but not with VLPs, indicates that the suppression of DC function and maturation is likely to be due to the presence of viral proteins or genomic material not present in the VLPs. Further studies are needed to clarify the detrimental effects of Ebola virus infection on other subpopulations of DCs, in particular on plasmacytoid DCs, which are important in antiviral interferon responses. The consequences of nonfunctional DCs include a diminished ability to stimulate humoral or cell-mediated immune responses, which may contribute to the lack of control of viral replication.

A principal determinant of the inhibitory effect on innate immune function is the resistance of Ebola virus to the antiviral effects of interferon, which is likely to be due to interruption of critical interferon response pathways by the virus itself20–22; interferon production is blocked in macrophages, peripheral blood mononuclear cells and DCs by Ebola virus infection in vitro and in vivo16,23. In addition, the expression of interferon-stimulated genes important in the type-I interferon response is decreased in Ebola virus–infected cells20,22,24. The interferon response has also been shown to be very important for disease outcome in mice. Immunocompetent mice are resistant to Ebola virus infection, but mice that lack the interferon-β receptor or the protein signal transducer and activator-1 (STAT1) or that are treated with antibody to interferon become susceptible to disease25, highlighting the critical role of interferon in protecting uninfected cells. Several mechanisms of Ebola virus–mediated resistance to the interferon response have been identified. Like some other viruses, Ebola encodes specific viral proteins that antagonize the interferon response. Two virally encoded proteins, VP24 and VP35, have been shown to interfere with the induction of the interferon response26–28. Nuclear accumulation of STAT1 is interrupted by VP24, which leads to a block in type-I interferon signaling and renders infected cells insensitive to this antiviral response27. Ebola VP35 blocks the activity of the interferon regulatory factor 3 (IRF-3), thus decreasing interferon responses26,28. More recently, VP35 was shown to counteract the activity of the double-stranded RNA–dependent protein kinase (PKR)29.
response are necessary to protect against fatal infection. If such a host immune response is not generated, the virus evades immune control, and the infection progresses to end-stage disease.

Table 1 Correlative differences in patients who survive versus patients who succumb to Ebola virus infection

| Nonlethal infection | Lethal infection | Refs. |
|---------------------|------------------|-------|
| Prominent CD8⁺ T cell activation | No CD8⁺ T cell activation | 30,33 |
| Above-normal numbers of T cells | Below-normal numbers of T cells | 33 |
| 10⁷ viral genome copies/ml serum | 10¹⁰ viral genome copies/ml serum | 33,47 |
| Detectable antibodies in blood at onset of symptoms | No detectable antibodies in blood at onset of symptoms | 33,65 |
| Low NO | High NO | 33,65 |

In combination, these studies suggest that virus-induced inhibition of the interferon pathway not only decreases interferon-stimulated gene transcription to prevent an antiviral response state, but also contributes to lower numbers of mature and activated myeloid DCs, which in turn hinders the activation of the adaptive immune response.

Surprisingly, patients who succumb to Ebola virus infection show little evidence of an activated adaptive immune response. Adaptive immunity is severely compromised not only because of a lack of functional DCs and other important antigen-presenting cells, but also because lymphocytes undergo massive apoptosis in infected humans and nonhuman primates²⁵,³⁰,³¹. Although lymphocytes are not targets of the virus, substantial numbers—with the exception of B cells—undergo apoptosis during the illness³²; as a result, numbers of CD4⁺ and CD8⁺ T cells are substantially reduced in fatal human and nonhuman primate infections before death³⁰,³¹,³³. Lymphocyte apoptosis is also a common manifestation of other viral hemorrhagic fevers and is frequently observed during septic shock³⁴.

Studies with lymphocytes in vitro indicate that several molecules involved in triggering apoptosis are present on these cell populations, including TRAIL and Fas-FasL.¹⁵ However, the mechanisms responsible for this ‘bystander’ apoptosis are still under investigation. It may be that inflammatory mediators and other factors, such as the pro-apoptotic soluble factor nitric oxide (NO) secreted by infected macrophages, are capable of inducing the observed lymphocyte apoptosis.

Alternatively, impaired DC function and an overall immunosuppressive state may contribute to the phenomenon³¹. Yet another possibility is that cell death is actively triggered by direct interactions between lymphocytes and Ebola virus or soluble gene products. The importance of early responses involving cells of the innate immune system and/or a rapid adaptive antibody response is highlighted by a recent study showing protection of nonhuman primates with postexposure vaccine administration³⁵.

Although filoviruses are among the most virulent and fatal pathogens known, some patients infected with Ebola virus recover from the infection. Identifying the differences in the immune response between fatal and nonfatal cases is important for future development of effective therapies and vaccines. Specific differences in clinical presentation and immune responses have been noted in those who succumb contrasted with those who recover from Ebola virus infection (Table 1). This comparison makes it clear that the development of an antigen-specific cell-mediated immune response correlates with clearance of the virus. Studies demonstrating antigen-specific cellular immune responses in vaccinated nonhuman primates that survived infectious Ebola virus challenge⁶–⁸ support this finding. Additionally, induction of a humoral and CD8⁺ T cell response was found to be required for protection in mice challenged with lethal Ebola virus infection⁹,¹⁰. However, the protective role of immunoglobulins remains uncertain, as a recent report indicates that the passive transfer of the neutralizing human monoclonal antibody KZ52 is unable to control infection in a macaque model¹⁰. Based on these considerations, it is becoming increasingly evident that an early and robust, but transient, innate immune response and the subsequent activation of adaptive immune response are pivotal to the systemic dissemination of the virus during human infection⁸,¹¹. Infected monocytes and macrophages travel from the site of infection to lymph nodes, where more monocytes and macrophages are recruited and then become targets of infection. Infection of these cells leads to further amplification and dissemination of the virus through the lymphatic system¹². In addition, infection and necrosis of hepatocytes causes impairment of liver function. Liver enzymes are elevated in most filovirus infections⁴⁸–⁵⁰, and decreased liver function could account for the decreased synthesis of coagulation factors and development of coagulation disorders prominent during fatal infection. Finally, the development of shock at later stages of the disease is multifactorial and, along with hemorrhage, may be due in part to the infection and resulting necrosis of cells of the adrenal cortex⁵⁰, as these cells are important in the regulation of blood pressure.

Pathogenesis of infection

The pathological changes seen in patients dying from Ebola virus infection include coagulation abnormalities, vascular permeability, hemorrhage and organ necrosis and failure. The current hypothesis is that the fundamental mechanism of Ebola virus pathogenesis is vascular injury and damage secondary to coagulation abnormalities and increased vascular permeability, due to the release of inflammatory cytokines and chemokines by infected and activated monocytes and macrophages, and to direct endothelial cell damage from viral replication late in infection⁴¹,⁴². It is evident that in addition to the ‘cytokine storm’, the virus itself can also induce immunosuppression and damage host cells directly⁴³,⁴⁴. Thus, the deleterious manifestations of the infection stem in part from the factors secreted from dysfunctional immune cells and in part from virally induced damage of host tissues and organs.

Ebola virus shows tropism in vitro for cells of the innate immune system as well as endothelial cells, dendritic cells and several types of epithelial cells. Replication occurs at an unusually high rate in infected cells. The ability of virus to replicate in different cell types is less well characterized in vivo. In addition, viremia in infected patients is generally difficult to quantify⁶; however, a viral load exceeding 10⁹ plaque-forming units per milliliter of serum (PFU/ml) was noted in at least one outbreak of the Zaire species of Ebola⁴⁵. Viremia in infected nonhuman primates can reach up to 10⁷ PFU/ml⁴⁶. Humans with fatal infections have up to 10¹⁰ copies of viral RNA per milliliter, whereas much fewer (10⁷ copies/ml) are found in the serum of those who survive Ebola virus infection (see Table 1)⁴⁷. High rates of viral replication lead to lysis and necrosis in cells of many organs, including the liver, spleen, kidneys and gonads. Much of the observed necrosis is virally induced, as there is little infiltration within infected tissues, and extraordinary numbers of viral particles are present in the necrotic debris. In addition, microscopic examination of infected human tissues shows a correlation between tissue damage and the presence of viral antigens, nucleic acid and sites of viral replication⁴,⁴³,⁴⁴. This observation indicates that direct viral damage of tissues and organs may lead to organ failure and shock.

The infection of certain cell types plays an important role in Ebola virus pathogenesis. Infection of innate immune cells is thought to be pivotal to the systemic dissemination of the virus during human infection⁸,¹¹. Infected monocytes and macrophages travel from the site of infection to lymph nodes, where more monocytes and macrophages are recruited and then become targets of infection. Infection of these cells leads to further amplification and dissemination of the virus through the lymphatic system¹². In addition, infection and necrosis of hepatocytes causes impairment of liver function. Liver enzymes are elevated in most filovirus infections⁴⁸–⁵⁰, and decreased liver function could account for the decreased synthesis of coagulation factors and development of coagulation disorders prominent during fatal infection. Finally, the development of shock at later stages of the disease is multifactorial and, along with hemorrhage, may be due in part to the infection and resulting necrosis of cells of the adrenal cortex⁵⁰, as these cells are important in the regulation of blood pressure.
Vascular impairment and Ebola glycoproteins

Endothelial impairment is a prominent feature of Ebola hemorrhagic fever. A loss of vascular integrity is often observed in humans and nonhuman primates during late stages of the disease and is associated with bleeding and the imbalance of fluid between tissue spaces. The complete mechanisms leading to endothelium permeability have not been elucidated. Several studies have shown that the virally induced release of inflammatory mediators increases vascular permeability in vitro\textsuperscript{1,15,1}. However, endothelial cells are targets of infection during later stages of the disease and direct virus-induced cytotoxicity of endothelial cells cannot be ruled out as a contributory mechanism for increased hemorrhagic manifestations. Indeed, the viral envelope glycoprotein GP has been implicated as one of the major determinants of vascular cell injury. GP is one of the most studied Ebola virus proteins because of its importance in viral entry and potential as a target for vaccine development. As mentioned, it is also under intense research because of its possible role in pathogenesis. The glycoprotein is responsible for targeting the virus to cells that are relevant to pathogenesis. GP is likely to have a role in immune suppression through its effects on downregulation of cell surface proteins essential for lymphocyte adhesion and antigen presentation\textsuperscript{52,53}. Although some have suggested that soluble GP may compete for neutralizing antibodies that might otherwise target viruses or infected cells\textsuperscript{54,55}, a protective role for such antibodies has not been demonstrated, and the biochemistry and antibody reactivity of soluble GP differ from those of the membrane-bound trimer spike\textsuperscript{56,57}. Soluble GP does inhibit neutrophil activation\textsuperscript{57}, providing a further mechanism by which viral immunity may affect the innate inflammatory response. Ebola virus entry depends also on endosomal cathepsins, enzymes critical for antigen presentation\textsuperscript{58,59}, and release of cathepsins may contribute to virus-induced cell damage\textsuperscript{60}.

Several groups have shown that GP has a direct cytotoxic effect. Yang and colleagues found that of the seven viral gene products, GP was responsible for cell rounding and detachment in endothelial cells both in vitro and ex vivo and that this can lead to a substantial increase in vascular permeability\textsuperscript{61}. Expression of GP from all four species of Ebola virus induces variable degrees of cytotoxicity in cell lines and primary cells in vitro that are characterized by cell rounding and detachment, followed by cell death\textsuperscript{61}. These effects are mediated by a heavily glycosylated mucin-like domain of the glycoprotein. Although there is some debate over the role of GP cytotoxicity during live viral infection\textsuperscript{62}, differences in GP-induced cytotoxicity are correlated with the mortality rates of the different viral species\textsuperscript{52,61}, suggesting that this gene product is important in the pathogenesis of the disease. Expression of membrane-bound GP seems to be precisely controlled during viral replication by a mechanism involving transcriptional editing by the viral polymerase\textsuperscript{63}. This indicates that the glycoprotein may be a key viral determinant of pathogenicity during infection.

Thus, virus-induced and host factors combine to result in a destructive pathway in which a fatal response to Ebola virus infection invariably correlates with suppression of both B and T cell–mediated immunity. Patients who fail to recover have virtually no viral antigen–specific antibodies. Low amounts of specific immunoglobulin Ms are present in only 30% of fatally infected patients, and no specific immunoglobulin Gs are detected\textsuperscript{60,64,65}. There seems to be limited initiation of cytotoxic T cell or CD4\textsuperscript{+} helper T cell responses, most likely owing to their depletion in fatal cases. Lymphocyte depletion is likely to exacerbate the uncontrolled viral replication within macrophages and other inflammatory cells. Therefore, fatal Ebola virus infection is characterized by broad immunosuppression typified by the development of an aberrant nonspecific and deleterious innate immune response and little or no stimulation of an antigen-specific adaptive response. This lack of response leads to an overwhelming viral burden and resultant immune- and virus-mediated pathology.

Relevance to other highly lethal pathogens and future research

Valuable insights into critical features of the host immune system can be gained from an examination of the immune response to highly virulent pathogens such as Ebola virus. One trend that seems to be emerging is that lethal, acute pathogens tend to kill rapidly, before the development of an adaptive immune response, whereas chronic pathogens can survive and replicate despite an adaptive immune response. In this regard, there are interesting parallels between Ebola virus infection and the highly pathogenic 1918 influenza virus (see the accompanying review by Ahmed and colleagues\textsuperscript{66}). For example, Kobasa and co-workers found that a reconstituted 1918 strain of influenza shows high levels of viral replication, which correlates with macroscopic lesions in lung tissue of infected cynomolgus macaques\textsuperscript{67}. Infection in this animal model culminated in acute respiratory distress and an overwhelmingly fatal outcome. Interestingly, infected animals were able to mount an immune response that was in many ways similar to the responses observed during Ebola infection in nonhuman primates. The immune response to 1918 influenza was characterized by an aberrant interferon response and the expression of abnormally high levels of cytokines and chemokines. The authors concluded that the high lethality of the 1918 influenza strain can be attributed in part to the generation of an atypical and harmful innate immune response that is insufficient for protection.

A comparison of the immune responses to Ebola and 1918 influenza viruses indicates that the high lethality of these viruses may stem from a combination of the deleterious effects of high viral titers and direct viral damage and a nonspecific and abnormally sustained innate immune response. A similar picture of overwhelming viremia, lack of control by the innate immune response and failure to develop adaptive immunity has been observed with other highly lethal viruses as well, including the severe acute respiratory syndrome (SARS) coronavirus, Marburg virus, Lassa fever virus and others. In each case, it would seem that the virus causes lethal infection through its overwhelming replication, though the specific receptors, cell and organ tropisms, mechanisms of evading inflammation and immunity and natural reservoir may differ.

Many questions remain unresolved regarding the mechanisms and full extent of virus-induced immune dysregulation. For example, the mechanism of lymphocyte apoptosis is unknown. Ebola virus does not target these cells directly, yet their numbers are rapidly exhausted once viral titers are measurable in the host. Do these cells enter an anergic terminal differentiation due to local cytokine imbalances, or is there aberrant target destruction by other immune cells? It is also unknown whether Ebola virus shows tropism for a particular DC that may enhance evasion of the antiviral response. The mechanism by which DC antigen presentation is impaired is unknown. The role of cathepsin in immunopathogenesis is also not completely understood; because cathepsins also contribute to antigen processing, it is possible they affect the adaptive immune response as well. Similar questions remain regarding the details of viral replication in vivo. Although Ebola virus can be found by immunostaining of a variety of cell types, including macrophages, DCs and endothelial cells, the virus binds to ubiquitous lectin receptors on many of these cells; thus, it is unclear whether the presence of the virus in a given cell represents active replication or merely binding to the cell surface. Finally, the role of cytokine storm versus direct viral cytotoxicity to endothelial cells remains the subject of much speculation but, unfortunately, very little data.
Ultimately, many of those questions, including the roles of specific parts of the immune system in protection, may be resolved with studies that use antibody depletion in vivo against cytokines, cytokine receptors and lymphocyte subsets in nonhuman primate models of infection. Until these important issues are addressed, the current hypotheses explain Ebola virus–induced pathophysiology in broad terms: a combination of factors, including uncontrolled and nonspecific inflammatory responses, virally induced immunosuppression and direct viral destruction of several cell types, collectively contribute to the collapse of the vascular system, multiorgan failure and shock–like syndrome of lethal Ebola virus infection.

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