

Review

Bench-to-bedside review: Rare and common viral infections in the intensive care unit – linking pathophysiology to clinical presentation

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Critical Care 2008, **12**:219 (doi:10.1186/cc6917)**Abstract**

Viral infections are common causes of respiratory tract disease in the outpatient setting but much less common in the intensive care unit. However, a finite number of viral agents cause respiratory tract disease in the intensive care unit. Some viruses, such as influenza, respiratory syncytial virus (RSV), cytomegalovirus (CMV), and varicella-zoster virus (VZV), are relatively common. Others, such as adenovirus, severe acute respiratory syndrome (SARS)-coronavirus, Hantavirus, and the viral hemorrhagic fevers (VHFs), are rare but have an immense public health impact. Recognizing these viral etiologies becomes paramount in treatment, infection control, and public health measures. Therefore, a basic understanding of the pathogenesis of viral entry, replication, and host response is important for clinical diagnosis and initiating therapeutic options. This review discusses the basic pathophysiology leading to clinical presentations in a few common and rare, but important, viruses found in the intensive care unit: influenza, RSV, SARS, VZV, adenovirus, CMV, VHF, and Hantavirus.

Introduction

Viral infections are common causes for upper and lower respiratory tract infections and a frequent reason for outpatient office visits. Comparatively, viral respiratory infections are less common in the intensive care unit (ICU) setting but still play an important clinical role. Most viral respiratory infections in the ICU are community-associated cases with severe lower respiratory disease that can progress into respiratory failure and acute respiratory distress syndrome (ARDS) [1]. The remainder are infections seen in immunocompromised patients, such as transplantation [2,3]. In some instances (severe acute respiratory syndrome [SARS], influenza, and adenovirus), viral respiratory infections present with fulminant respiratory failure and ARDS, heralding a larger community outbreak [4]. In these situations, the newly recog-

nized illness in an ICU patient might be the first presentation of a larger public health emergency.

The clinical presentation, treatment, outcome, and personal and institutional infection control differ greatly among the most common viral infections in the ICU. These differences are largely based on the viral structure, mode of transmission and cell entry, and host immunology and thus provide the foundation for the clinical presentation, virulence, and medical therapeutics of these viral infections. Therefore, a basic knowledge of the more common ICU viral respiratory pathogens will provide a framework for the clinical and research approaches for these infections. This review will focus on the basic epidemiology, virology, and host immune response for a few common or high-impact viral respiratory pathogens in the ICU: influenza, respiratory syncytial virus (RSV), SARS, varicella-zoster virus (VZV), adenovirus, cytomegalovirus (CMV), and viral hemorrhagic fever (VHF) (Table 1). With this basic foundation, clinical care, public health, and medical therapeutics for these viruses will be enhanced from the laboratory to the bedside.

Influenza

Influenza causes a clinically recognizable, systemic illness characterized by abrupt-onset fever, headache, myalgia, and malaise (the classic influenza-like illness) [5]. Influenza is subdivided into three distinct types: A, B, and C [5,6]. Influenza A infects a variety of species, including birds, swine, horses, marine mammals, and humans [5,6]. Influenza B infects only humans and predominates in children, and both influenza A and B cause yearly outbreaks. Respiratory symptoms are usually self-limited. However, a small number of

ARDS = acute respiratory distress syndrome; CMV = cytomegalovirus; HA = hemagglutinin; HFRS = hemorrhagic fever with renal syndrome; HPS = Hantavirus cardiopulmonary syndrome; ICU = intensive care unit; IL = interleukin; MBL = mannose-binding lectin; MHC = major histocompatibility complex; MIP-1 = macrophage inflammatory protein-1; NA = neuraminidase; RSV = respiratory syncytial virus; SARS = severe acute respiratory syndrome; SARS-CoV = severe acute respiratory syndrome-coronavirus; TNF = tumor necrosis factor; VHF = viral hemorrhagic fever; VZV = varicella-zoster virus.

Table 1

Clinical and immunologic characteristics of major viruses found in the intensive care unit

	Influenza	RSV	SARS-CoV	VZV	Adenovirus	CMV	VHF
Virus family	Orthomyxoviridae	Paramyxoviridae	Coronaviridae	Herpesviridae	Adenoviridae	Herpesviridae	Filoviridae
Epidemiologic link	Seasonal epidemic	Seasonal epidemic, immunocompromised and transplant	Laboratory exposure on known infected individual	Contact with infected individual	Military camps, mental health facilities	Transplantation, immunosuppressive medications	Endemic area or contact with infected individual
Pulmonary clinical findings	Primary alveolar pneumonia	Upper respiratory tract infection, bronchiolitis, pneumonia	Rapid progressive pneumonia, ARDS	Primary alveolar pneumonia	Alveolar pneumonia with bronchiolitis	Interstitial pneumonitis, bronchiolitis	Alveolar edema
Lipid envelope	Yes	Yes	Yes	Yes	No	Yes	Yes
Major receptor for cell entry	Sialic acid	RSV glycoprotein G	CD209L ACE 2	Glycoprotein C and D	Coxsackie-adenovirus receptor	Unknown, involves integrins	Folate receptor alpha
Primary cell of infection	Type 1 respiratory epithelium	Type 1 respiratory epithelium	Type 1 respiratory epithelium	Macrophage and dendritic cells	Type 1 respiratory epithelium	Multiple	Macrophages and dendritic cells
Viremia	No	No	Yes	Yes	Yes	Yes	Yes
Primary host immunity	Humoral	Humoral	Unknown	Cellular	Cellular	Cellular	Humoral

ARDS, acute respiratory distress syndrome; CMV, cytomegalovirus; RSV, respiratory syncytial virus; SARS-CoV, severe acute respiratory syndrome-coronavirus; VHF, viral hemorrhagic fever; VZV, varicella-zoster virus.

individuals can develop primary pneumonia, which can progress to ARDS [5]. The respiratory symptoms will persist or progress, and in a minority of cases ARDS can develop [5,7-9]. The combination of pneumonia and ARDS usually occurs in at-risk individuals, like individuals with chronic lung diseases, but has been described in healthy individuals as well.

The structure of influenza's viral envelope is important in viral infection and thus host cell immunity [10,11]. The envelope contains surface glycoproteins essential for virus entry into the host cell. The trimeric hemagglutinin (HA) structure undergoes limited proteolysis by host cellular proteases such as furin. HA then binds to specific sialosaccharides found on the surface of respiratory epithelial cells to initiate cell entry [12]. The neuraminidase (NA) is an enzyme that catalyzes the removal of terminal sialic acids from glycoproteins [12]. This helps degrade respiratory tract mucus and release viral progeny after cell infection and thus is necessary for subsequent viral entry to viral escape from the host cell [12]. Influenza A is divided into subtypes based on H and N antigenicity [11]. All H subtypes have been found in multiple avian species and other animals. H1, H2, and H3 predominate in human disease seasonally, and more recently, avian subtypes such as H5 and H7 have increased in humans over the past decade [13-15].

Infection occurs when viruses containing aerosols are deposited into the upper respiratory tract epithelium [5]. In experimental volunteers, inoculation with small-particle aerosols more closely mimics natural disease than large drops into the nose, illustrating the easy transmission with coughing or sneezing [16,17]. The virus can attach (HA) and penetrate the columnar epithelial cells. Predominantly human subtypes (H1, H2, and H3) bind to alpha-2,6-galactose sialic acid found in ciliated human respiratory tract epithelium [18]. On the other hand, avian influenza subtypes (for example, H5N1) bind preferentially to alpha-2,3-galactose sialic acid, which is found in the gastrointestinal tract of water fowl, epithelial cells on human conjunctivae, and on human type 2 pneumocytes [18-20]. This preferential binding for specific sialic acid receptors illustrates the differences in clinical presentation seen with avian influenza infections in humans: conjunctivitis, diarrhea, and fulminant alveolar pneumonia [20]. Additionally, it underlies the difficulty with human-to-human transmission of avian strains as preferential binding to type 2 pneumocytes requires smaller particle aerosolization and deep inhalation into the alveoli rather than larger droplets seen with seasonal influenza transmission [20].

Host immunity occurs via a number of mechanisms. Upon receptor binding, a large cytokine response occurs, with interleukin (IL)-2, IL-6, and interferon gamma predominately

[21]. This leads to extensive local inflammation with neutrophils and macrophages infiltrating the subepithelium of the respiratory tract. In cases of severe avian subtypes, a hemophagocytic syndrome and severe diffuse alveolar damage occur, causing the clinical findings of severe pneumonia and respiratory failure [21]. Within the alveolar macrophages and pneumocytes, major histocompatibility complex (MHC) I up-regulation leads to antigen presentation of the HA and other subcapsular proteins [22,23]. This eventually leads to natural killer cell destruction of infected cells and the development of neutralizing antibodies (largely against HA) by day 14 of infection [22].

Treatment of active influenza involves antiviral agents and supportive care. The most effective therapy is prevention via vaccination and infection control [4,5,13]. Two types of antiviral medications have been used: (a) M2 inhibitors (amantadine and rimantadine) inhibit the M2 ion channel needed for viral replication [24]. These are not active against influenza B and C and resistance is common in seasonal influenza. Thus, they should be used only in cases of known susceptibility. (b) The NA inhibitors, oseltamivir and zanamivir, have less resistance and prevent cleavage of sialic acid, which is necessary for a new virus to exit from the host cell [24]. Studies with the NA inhibitors show a reduction in symptom time and viral shedding, with peak effect when started within 48 hours of symptom onset [4,5,13]. However, treatment with NA inhibitors after 48 hours may provide some additional benefit but has not been fully studied [13]. Resistance is low within the community, but NA inhibitor resistance has already been described in clinical isolates from human cases of avian influenza.

Respiratory syncytial virus

Respiratory syncytial virus (RSV) is the most common cause of lower respiratory tract infections in children under 1 year of age, and healthy adults are infected repeatedly throughout their lives [25,26]. Adults typically have upper respiratory tract symptoms; however, some adults will develop lower respiratory tract infections, including bronchiolitis, pneumonia, and (rarely) ARDS [25-28]. The elderly and immunocompromised, particularly bone marrow transplant patients, are at highest risk of lower respiratory tract infection and respiratory failure [28]. In these cases, upper airway infection usually precedes lower tract infection by 1 to 3 days. Infection follows a pattern similar to influenza, with epidemics occurring in the winter months [25].

Inoculation occurs at the nasal or ocular mucosa via direct contact with secretions or infected fomites [29,30]. RSV has a lipoprotein envelope with surface glycoproteins that are important in host infection [31,32]. These glycoproteins act as cell fusion proteins, ultimately forming multinucleated giant cells ('syncytia'), assisting in cell-to-cell spread [31,32]. The virus replicates locally and then spreads to the epithelium of the bronchioles. From the bronchioles, the virus can then

extend to the type 1 and 2 alveolar pneumocytes [31,33]. Infection leads to cellular (neutrophils, monocytes, and T cells) infiltration of the epithelium and supplying vasculature, with subsequent necrosis and proliferation [31,33]. This will cause the airway obstruction, air trapping, and increased airway resistance that are characteristic of RSV infection [25,31,33]. RSV infection is more specifically associated with IL-6 and macrophage inflammatory protein-1 (MIP-1) release [34-36]. Elevated levels of IL-6 and MIP-1 in the bronchioles have correlated with more severe disease [37].

Both droplet and contact transmissions are the main methods of spread, and thus hand washing, droplet isolation, and the use of personal protective equipment are all important in reducing viral spread [29,30]. Specific genotypes will predominate during a seasonal outbreak, and since the genotypes change annually, adult reinfections occur [32]. Treatment usually is focused on controlling bronchospasm and preventing spread to other patients and health care workers [25,28]. Bronchodilators and corticosteroids are used for bronchospasm, and aerosolized ribivirin has been used in severe and high-risk cases such as bone marrow transplants [25,28]. However, a recent study evaluating bronchiolitis in infants, in which over 50% of cases were caused by RSV, showed that corticosteroids had no effect on outcome [38].

Severe acute respiratory distress syndrome

SARS is caused by a novel coronavirus (SARS-CoV) that was first detected in 2003 [39,40]. The initial outbreak rapidly spread into a global epidemic, with cases reported from 29 countries. The fatality rate was 11%, with most deaths in patients older than 65 and no deaths in children [39-41]. Since the initial epidemic in 2003, no new cases have been reported. SARS appears to clinically present as a two-stage illness. The initial prodrome, characterized by fever with or without rigors, malaise, headache, and myalgias, occurs an average of 7 days after contact with infected individuals [40-42]. Some patients also have mild respiratory symptoms or nausea and diarrhea. The respiratory phase appears to develop approximately 8 days after the onset of fever [40-42]. Forty-five percent of patients will develop hypoxemia and approximately 20% of these patients will progress to acute lung injury and require mechanical ventilation [40-42]. SARS-CoV appears to have originated from the horseshoe bat. The horseshoe bat appeared to be a natural reservoir for the virus and the civet cat acted as an intermediate host, allowing transmission to humans [43,44].

Like RSV and influenza, SARS-CoV has a lipoprotein envelope, but unlike the RSV and influenza, the virus is assembled and obtains its envelope from the endoplasmic reticulum [45]. SARS-CoV, like other coronaviruses, starts with infection of the upper respiratory tract mucosa [40]. SARS-CoV binds to CD209L (L-SIGN) and ACE-2, two functional receptors on the respiratory tract epithelium [46,47]. After binding, local inflammation and edema increase.

ACE-2 has a key protective role in acute lung injury by reducing alveolar fluid, and thus the binding of SARS-CoV to ACE-2 may contribute to the dysregulation of fluid balance in the alveolar space [48]. Additionally, low mannose-binding lectin (MBL) levels are thought to play a role in SARS pathogenesis [49]. In many respiratory infections, MBL prevents receptor attachment, activates complement, and enhances phagocytosis. In SARS-CoV infections, low or deficient levels of MBL have been noted, particularly associated with an MBL haplotype [49]. The binding of SARS-CoV to ACE-2, along with lower levels of MBL, leads to higher viral levels, increased alveolar edema, and the severe acute respiratory failure associated with SARS-CoV.

Viral spread is by droplet transmission, although many cases suggest that airborne and contact routes also occur [39]. Spread to health care workers who wore appropriate personal protective equipment suggests an airborne mode, and additional spread by aerosol-generating procedures, such as resuscitation (cardiopulmonary resuscitation), medication nebulization, and noninvasive ventilation, further supports this mode [39,50-52]. The treatment for SARS is largely supportive with low tidal volume mechanical ventilation [40,53]. Numerous treatment strategies, including corticosteroids, ribavirin, immunoglobulin, and interferon, have been investigated in SARS: none has been demonstrated to provide clinical evidence of benefit.

Varicella-zoster virus

VZV infection routinely occurs during childhood, presenting with low-grade fever, malaise, pharyngitis, and a vesicular rash [54,55]. Primary disease occurs throughout the year and usually is self-limited in immunocompetent host. VZV pneumonia is rare in children. However, it is the most frequent complication in adults (20%) and accounts for the majority of hospitalizations from VZV [56,57]. Varicella pneumonia develops insidiously, usually a few days after the onset of rash, and can progress to respiratory failure and ARDS [56,57]. Risk factors for VZV pneumonia and ARDS include pregnancy, smoking, and immunosuppression (malignancy, corticosteroids, HIV, and solid-organ transplant), but young healthy adults rarely develop ARDS [54,58]. Mortality for VZV pneumonia is 10% to 30%, with a mortality of 50% when respiratory failure ensues [54,58]. Additional complications include encephalitis, hepatitis, and secondary skin and soft tissue infections.

VZV is a herpes virus, a common group of DNA viruses that have a lipid-containing envelope with surface glycoproteins [59]. Infection starts in the upper respiratory tract mucosa as the surface glycoproteins allow for fusion of the lipid envelope with the respiratory cell membrane [60,61]. Upon cell entry, viral replication and assembly occur after integration of the viral genes into the cellular DNA [60,61]. Naked capsids then acquire their envelope at the nuclear membrane and are released into the perinuclear space where

large vacuoles are formed, leading to the clinical vesicles [60,61]. Local replication and spread lead to seeding of the reticuloendothelial system and ultimately viremia, which leads to diffuse and scattered skin lesions associated with primary varicella [62,63]. Viral shedding can last from onset of fever until all lesions have crusted and pneumonia has improved. Both humoral immunity and cell-mediated immunity are involved in protection [62,64]. Antibodies are directed at the surface glycoprotein and lead to viral neutralization. Cellular immunity drives local inflammation, leading to cell repair and vacuole removal. The virus becomes latent within the dorsal root ganglia [59,63]. During latency, the viral DNA is located in the cytoplasm rather than integrated into nuclear DNA.

VZV is highly contagious and transmission is via respiratory droplets and direct contact with lesions [56,62]. The envelope is sensitive to detergent and air drying, which account for the lability of VZV on fomites. In adults who progress to pneumonia or ARDS, treatment with acyclovir and corticosteroids has been shown to lessen hospital and ICU stays [62,65,66]. In immunocompromised persons not previously exposed to VZV, varicella-zoster immune globulin has been shown to be useful for both prevention of disease and symptomatic improvement [62,65,66].

Adenovirus

Adenovirus is the one of the most common causes of upper respiratory tract infections in adults and children [67,68]. Clinical disease usually is a self-limited upper respiratory tract infection associated with conjunctivitis; however, severe lower respiratory disease can occur in both high-risk and healthy individuals [67,69-71]. The combination of pneumonia and ARDS develops in a minority of individuals and usually is associated with conjunctivitis and other extrapulmonary manifestations, such as gastrointestinal disease, hepatitis, meningitis, and hemorrhagic cystitis [68]. The extrapulmonary complications, along with ARDS, are more frequent in transplant recipients. Pneumonia and ARDS appear to be more common with subtype E type 4 and subgroups B type 7, but serogroup 35 also has been documented in mental health facilities [69-71]. Recent increases in respiratory diseases in adults have been noted over the past year with serotype 14 [72].

Over 51 human adenovirus subtypes exist and clinical syndromes vary among subtypes [53]. However, certain subtypes appear to have an increased likelihood of lower respiratory tract involvement and this appears to be related to the viral capsid proteins [73]. Unlike influenza, RSV, and SARS, adenovirus is a DNA virus covered by a protein capsid without a lipid envelope. Rodlike structures called fibers are one of three capsid protein types (hexons, pentons, and fibers) and these fibers are the attachment apparatus for viral adsorption to the cell [73]. Attachment occurs at the coxsackieadenovirus receptor, the same receptor as the coxsackie B virus. The hexon capsid protein appears to have

some antigenic sites that are common to all human adenoviruses and contains other sites that show type specificity [73]. Fiber antigen seems to be primarily type-specific with some group specificity, whereas the penton base antigen is common to the adenovirus family. Upon infection, respiratory epithelial cells express these capsid proteins on their surface, leading to direct CD8⁺ cytotoxic T-cell MHC class 1 killing of these cells [74]. Thus, epithelial destruction associated with submucosal edema drives the clinical findings of lower respiratory disease [67]. Additionally, neutralizing antibody is directed at the hexon type-specific antigen and provides some future protection against serotypes [74].

Adenovirus is relatively stable on environmental surfaces for long periods of time, and thus viral spread is largely associated with infected fomites [53,67]. Spread also occurs via droplet transmission. Treatment is largely supportive. For severe cases, especially in immunosuppressed patients, antiviral therapy has been attempted but no clinical studies exist [69-72]. In severe cases, especially in immunocompromised patients, ribavirin and cidofovir antiviral therapy has been attempted, but no controlled clinical trials exist.

Cytomegalovirus

CMV is a common viral infection that causes both primary and latent infections. Seroprevalence rates range from 60% to 70% in US adult populations [75,76]. CMV causes a wide spectrum of illness, ranging from an asymptomatic infection to a mononucleosis syndrome, organ-specific complications, and fulminant multisystem disease [77-79]. Immunocompetent patients are more likely to present with minimal to no symptoms, whereas immunocompromised patients are more likely to develop organ-specific complications and fulminant disease [77-79]. The most significant and severe disease syndromes are found in lung, liver, kidney, and heart transplant recipients [80]. Significant morbidity and mortality usually are confined to immunocompromised persons; however, previously healthy individuals can present with organ-specific complications or even present with fulminant disease [78,80].

CMV is a member of the herpes virus family and, like other members of this family, is known for causing latent infections [75]. Like other herpes viruses, CMV is an enveloped virus with multiple surface glycoproteins. These glycoproteins are important for viral entry into host cells and are targets for host cell humoral and cell-mediated immunity [75,81]. The cellular protein that serves as the specific receptor for CMV entry has not been identified, but CMV infects cells by a process of endocytosis [37]. Once entry has occurred, CMV alters host immunity through the activation of multiple genes. One important CMV protein prevents cellular HLA-1 molecules from reaching the cell surface, preventing recognition and destruction by CD8⁺ T lymphocytes [82]. Thus, the CMV genome can remain in infected cells and avoid immune

destruction, which accounts for its latency in clinical disease. Eventually, a cellular immune response, driven by high levels of anti-CMV CD4⁺ and CD8⁺ T cells, leads to control of the disease [37,82,83]. Antibodies against CMV do not provide significant immunity [83].

Avoiding immune detection gives CMV the ability to remain latent after infection, which contributes a great deal to serious CMV disease. Evidence for persistent CMV genomes and antigens exists in many tissues after initial infection, and CMV has been found in circulating mononuclear cells and in polymorphonuclear neutrophils [84]. The virus can be cultured from most bodily fluids, including blood, urine, stool, tears, semen, and breast milk, and from mucosal surfaces, including the throat and cervix [85-88]. Detection of cells that contain CMV intranuclear inclusions in renal epithelial tissue and in pulmonary secretions provides evidence that CMV may persist in these tissues as well. CMV antigens have also been detected in vascular endothelial cells; this site has been suggested as a cause of vascular inflammation and development of atherosclerosis [89]. When immune suppression occurs in patients by means of HIV infection or through immunosuppressive therapy, such as antilymphocyte antibody infusion, CMV can reactivate, producing end-organ disease [80,83]. Specifically from a pulmonary standpoint, CMV is common after lung transplantation, causing an acute pneumonitis or contributing to a chronic bronchiolitis [90]. In HIV patients, CMV pneumonitis is rare but postmortem studies suggest that pulmonary disease from CMV occurs at higher rates than previously recognized [90].

CMV is transmitted via many routes. Transmission has been observed among family members (thought to be secondary to close contact and viral shed from the upper respiratory tract), among children and employees at daycare centers, from sexual contact, blood and tissue exposure (seroconversion after transfusion of blood products or organ transplantation), and perinatally (during birth or from breast milk) [85-88]. There are several antiviral agents available for systemic treatment of CMV. These agents include ganciclovir, valganciclovir, foscarnet, and cidofovir [9,37,91].

Viral hemorrhagic fevers

The VHF's include a wide number of geographically distributed viruses found worldwide, including Ebola and Marburg viruses, Rift Valley fever, Crimean Congo hemorrhagic fever, Lassa fever, yellow fever, and dengue fever. Ebola and Marburg viruses are in the family filoviridae [92-95]. Although the underlying pathophysiology differs slightly between the VHF's, Marburg and Ebola viruses serve as a classic template [92-95].

Marburg virus has a single species whereas Ebola has four different species that vary in virulence in humans [92-95]. The clinical manifestations of both Marburg and Ebola viruses are similar in presentation, with a higher mortality with Ebola Zaire

(75% to 90%) than Marburg (25% to 40%) virus being the only major difference between them. The initial incubation period after exposure to the virus is 5 to 7 days, with clinical disease beginning with the onset of fever, chills, malaise, severe headache, nausea, vomiting, diarrhea, and abdominal pain [92-94,96]. With this initial infection, macrophages and dendritic cells initially are the site of viral replication, followed by spread to the reticuloendothelial system heralding the initial onset of symptoms [97]. As macrophages and other infected tissues undergo necrosis, an overwhelming cytokine response occurs, leading to abrupt prostration, stupor, and hypotension [92,93,96,98]. Particularly, tumor necrosis factor (TNF), IL-1, IL-6, macrophage chemotactic protein, and nitric oxide levels are markedly increased [98]. VHF-infected macrophages, along with noninfected macrophages stimulated by cytokines, release cell surface tissue factor, which subsequently triggers the extrinsic coagulation pathway [97,98]. The clinical and laboratory findings of impaired coagulation with increased conjunctival and soft tissue bleeding shortly follow [95,98]. In some cases, more massive hemorrhage can occur in the gastrointestinal and urinary tracts, and in rare instances, alveolar hemorrhage can occur [95,96,98,99]. The onset of maculopapular rash on the arms and trunk also appears to be classic and may be a very distinctive sign. Along with the bleeding and hypotension, multiorgan failure occurs, eventually leading to death [95,96,98,99]. The overwhelming viremia resulting in macrophage and dendritic cell apoptosis leads to impaired humoral immunity, which in turn leads to increase viral production [98]. This ultimately results in the rapid overwhelming shock seen with VHFs.

Transmission appears to occur through contact with nonhuman primates and infected individuals [95]. No specific therapy is available and patient management includes supportive care [92,93,95,98]. In a few cases in the Zaire outbreak of Ebola in 1995, whole blood with IgG antibodies against Ebola may have improved outcome, although subsequent analysis suggests that these patients were likely to survive even without this treatment [100].

Hantavirus

Hantavirus is one of four major genera within the family bunyaviridae, a family of more than 200 animal viruses spread via arthropod-vertebrate cycles [101-103]. Hantavirus causes two severe acute febrile illnesses: hemorrhagic fever with renal syndrome (HFRS) (found in the Old World) and Hantavirus cardiopulmonary syndrome (HPS) (found in the New World) [101-103]. HPS was first classified in the Southwestern US. A new species termed Sin Nombre virus was identified after an outbreak in the Four Corners region of the Southwestern US in 1993 [101-103]. In North America, disease largely has been reported in the Southwest and California, with cases reported in Canada, Europe, China, Chile, Argentina, and other parts of South America. Outbreaks are often cyclical and focal and are affected by weather and climatic variables and the effect this has on rodent populations [104].

Symptoms begin with a prodrome of fever, chills, and myalgias; HFRS and HPS also can be accompanied by abdominal pain and gastrointestinal disturbances [101-104]. In HPS, initially, there is an absence of upper respiratory symptoms. At about day 5, modest dry cough and dyspnea will develop. Due to the severe increase in vascular permeability associated with HPS, disease progresses rapidly (within hours) to respiratory failure, shock, ARDS, coagulopathy, and arrhythmias [104,105]. Resolution also can occur rapidly. If hypoxia is managed and shock is not fatal, the vascular leak reverses in a few days and recovery is apparently complete. Notably, thrombocytopenia with an immunoblast-predominant leukocytosis is characteristic of the early cardiopulmonary phase [104,105].

The exact mechanism for ARDS, shock, and coagulopathy is unclear, but it is suspected that the immune response, rather than the virus itself, causes the capillary leak and shock. The intense cellular immune response alters endothelial cell barrier function and is harmful. Hantavirus causes an increased release of TNF and alpha interferon and increased MHC I antigen presentation [106,107]. There is also a more intense CD8⁺ T-cell response in sicker patients [106,107]. It appears to result from a massive acute capillary leak syndrome and shock-inducing mechanisms thought to be due to the release of kinins and cytokines [106,107]. The syndrome's clinical presentation, rapid resolution, and histopathologic findings of interstitial infiltrates of T lymphocytes and alveolar pulmonary edema without marked necrosis support this underlying process. Treatment mainly is supportive, with extracorporeal membrane oxygenation being used in some cases [104,105]. Ribavirin has been effective in HFRS, but not HPS. Mortality remains at roughly 20%.

Conclusion

Viral infections in the ICU are common in the outpatient setting but become less common in the ICU. However, a small number of viral infections can lower respiratory tract disease and subsequent respiratory failure. These viral pathogens vary greatly in clinical disease, from rapid and fulminant respiratory failure and shock (VHF) to chronic latent disease of immunosuppression (CMV). However, most of these viruses commonly have lipid envelopes, except for adenovirus, and all have surface proteins or glycoproteins that allow for attachment, cell entry, and virulence. Host response to these infections varies from primarily cellular to

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humoral. All can cause respiratory disease but a few are of great public health concern, particularly novel strains of influenza, adenovirus, SARS, and VHF. An understanding of the basic viral pathogenesis, along with host response, allows for a foundation in treatment and public health response within the ICU.

Competing interests

The authors declare that they have no competing interests.

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