

TABLE 316.8 Summary of 2013 National Vaccine Advisory Committee's Standards for Adult Immunization Practices

AUDIENCE	SUMMARY OF STANDARDS
All providers	Incorporate immunization needs assessment into every clinical encounter. Strongly recommend needed vaccine(s) and either administer vaccine(s) or refer patient to a provider who can immunize. Stay up-to-date on, and educate patients about, vaccine recommendations. Implement systems to incorporate vaccine assessment into routine clinical care. Understand how to access immunization information systems (i.e., immunization registries).
Nonimmunizing providers	Routinely assess the immunization status of patients, recommend needed vaccine(s), and refer patient to an immunizing provider. Establish referral relationships with immunizing providers. Follow up to confirm patient receipt of recommended vaccine(s).
Immunizing providers	Ensure professional competencies in immunizations. Assess immunization status in every patient care and counseling encounter and strongly recommend needed vaccine(s). Ensure that receipt of vaccination is documented in patient medical record and immunization registry.
Professional health care–related organizations, associations, systems	Provide immunization education and training of members, including trainees. Provide resources and assistance to implement protocols and other systems to incorporate vaccine needs assessment and vaccination or referral into routine practice. Encourage members to be up-to-date on their own immunizations. Assist members in staying up-to-date on immunization information and recommendations. Partner with other immunization stakeholders to educate the public. Seek out collaboration opportunities with other immunization stakeholders. Collect and share best practices for immunization. Advocate policies that support adult immunization standards. Insurers, payers, entities that cover adult immunization services should ensure that their network is adequate to provide timely immunization access and augment with additional vaccine providers if necessary.
Public health departments	Determine community needs, vaccination capacity, and barriers to adult immunization. Provide access to all ACIP-recommended vaccinations for insured and uninsured adults and work toward becoming an in-network provider for immunization services for insured adults. Partner with immunization stakeholders and support activities and policies to improve awareness of adult vaccine recommendations, increase vaccination rates, and reduce barriers. Ensure professional competencies in immunizations. Collect, analyze, and disseminate immunization data. Provide outreach and education to providers and the public. Work to decrease disparities in immunization coverage and access. Increase immunization registry access and use by vaccine providers for adult patients. Develop capacity to bill for immunization of injured people. Ensure preparedness for identifying and responding to outbreaks of vaccine-preventable diseases. Promote adherence to applicable laws, regulations, and standards among adult immunization stakeholders.

ACIP, Advisory Committee on Immunization Practices.

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The complete reference list is available online at Expert Consult.

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SHORT VIEW SUMMARY

Definition

- A zoonosis is an infectious disease of humans that originates in animals.

Epidemiology

- Zoonoses account for up to 70% of emerging infectious diseases and include such varied examples as human immunodeficiency virus and acquired immunodeficiency syndrome (HIV/AIDS), Ebola, severe acute respiratory syndrome (SARS), plague, rabies, influenza, Zika, and new variant Creutzfeldt-Jakob disease.

Diagnosis

- The majority of zoonotic diseases are diagnosed through molecular methods. Many appear in unexpected contexts; thus a comprehensive travel and exposure history can be critical.

Treatment

- With the notable exception of HIV/AIDS and influenza, most viral zoonoses have not been a focus for drug development. Accordingly, treatment is primarily supportive. In contrast,

many bacterial zoonoses can be treated with antibiotics.

Prevention

- Vaccines are established for only a minority of zoonotic diseases. Prevention is best achieved by limiting exposure to reservoirs and vectors for transmission of infection.

Zoonoses, derived from the Greek words for animal (*zoo*) and the suffix modification indicating a state or condition (*sis*), are infectious diseases of humans that originate in animals. Infectious diseases that originate in humans and move into other animals are commonly described as reverse zoonoses. The majority of emerging viral diseases, up to 70%, represent zoonoses, with such prominent examples as human immunodeficiency virus and acquired immunodeficiency syndrome (HIV/AIDS), influenza, West Nile virus (WNV) encephalitis, Nipah, Hendra, severe acute respiratory syndrome (SARS), Ebola, Marburg, hantavirus pulmonary syndrome (HPS) (Sin Nombre virus), Zika, and rabies. Common examples of nonviral zoonoses include Lyme disease (*Borrelia burgdorferi* in North America, *Borrelia afzelii* and *Borrelia garinii* in Europe), cat-scratch disease (*Bartonella henselae*), new variant Creutzfeldt-Jakob disease (vCJD), salmonellosis, and toxoplasmosis. Although secondary microbial contamination of agricultural products can cause significant disease, the term *zoonosis* does not apply unless there is direct transmission to humans from an infected animal.¹

The majority of zoonotic disease can be attributed to anthropogenic change. Loss of wildlife habitat to development and consumption of bushmeat, necessitated by poverty or due to cultural preference, increase opportunities for cross-species jumps. Global warming may also increase the geographic range of phlebotomous arthropod vectors, such as mosquitos and ticks that serve as reservoirs and vectors for infectious agents. Given that there are more than 50,000 vertebrate species, for example, if we were to assume an average of 20 endemic viruses per vertebrate species, the potential reservoir of vertebrate viruses could be estimated at 1 million. Although it is unlikely all of them can be transmitted to humans and cause disease, it is sobering to consider the challenge of detecting and responding even to 1% of them—10,000 novel viruses.

ONE HEALTH INITIATIVE

The connection between human and animal microbiology was originally recognized by Virchow and Osler in the 1800s. The relationship has been emphasized through the One Health Initiative, which promotes

coequal collaborations between practitioners and researchers in human and comparative medicine, and surveillance programs involving wildlife, domestic animals, and humans.^{2,3} The catalytic event for the Initiative was the 1999 WNV outbreak in New York City, where two independent lines of research investigation converged. One that focused on high mortality in native corvids and exotic birds in the Bronx Zoo ultimately led to the culture and identification of the virus by the US Department of Agriculture (USDA). The other focused on human encephalitis cases that were initially attributed to the St. Louis encephalitis virus based on serology. WNV was subsequently implicated when brain material of victims was sequenced by Lipkin and colleagues at the University of California.⁴ The cause of the human outbreak would have been ascertained earlier if the veterinary and medical practitioners had collaborated in sharing data and samples. A series of national and international meetings focused on emerging infectious diseases and concerns that the use of antibiotics as growth promoters in agriculture was promoting the development of antimicrobial resistance culminated in the One Health Resolution, signed by the presidents of the American Medical Association and American Veterinary Medical Association, with endorsements from the Centers for Disease Control and Prevention, the USDA, and the European Union.⁵⁻⁷

MECHANISMS OF TRANSMISSION

Mechanisms of transmission of zoonotic agents vary widely. Many are linked to food collection, processing, or consumption. The most dramatic examples are diseases associated with the slaughter and consumption of wild animals (bushmeat), such as Ebola, Marburg, SARS, monkeypox, and tularemia; however, less exotic meats are associated with transmission of *Salmonella*, highly pathogenic *Escherichia coli*, and prion diseases (new vCJD). Phlebotomous arthropods such as mosquitos, ticks, and flies may also serve as vectors for transmission of bacteria and viruses from birds and mammals to humans. Prominent examples include WNV, St. Louis encephalitis virus, tick-borne encephalitis virus (TBEV), Zika virus, Powassan virus, and *B. burgdorferi* (Lyme disease). Direct transmission can occur through exposure to infected urine or feces, as in

leptospirosis, toxoplasmosis, lymphocytic choriomeningitis, or HPS; through wound inoculation as in *Bartonella* infection (cat-scratch fever); or in rabies, the oldest recorded example of zoonotic disease and the only common one transmitted directly from infected animals to human by bite.

FACTORS IN THE EMERGENCE OF ZOOONOTIC DISEASES

Travel and trade are increasingly global, carrying pathogens in addition to passengers and products to new locations. John F. Kennedy International Airport, for example, one of two international airports in the greater New York metropolitan area, receives nonstop flights from >100 international destinations and annually serves >12 million international passengers.⁸ Similar traffic data apply in other major metropolitan areas. Given that an infected individual, mosquito, or other cargo can cross the world in less than 24 hours, clinicians and public health practitioners must be prepared to encounter known and novel agents in virtually any context.

The advent of global agribusiness and urbanization are also important factors in zoonotic diseases. The international food trade has burgeoned by more than 200% since 1975.⁹ A hundred years ago, most fresh food was produced and consumed within a radius of a few kilometers. It is now not unusual for individuals to consume plants and animals harvested thousands of kilometers away.¹¹ Contamination of meat with prions, influenza viruses, and Rift Valley fever virus has been documented in international trade of livestock.¹² The centralization of food production and processing, particularly of ground meat or raw fruits and vegetables, has resulted in outbreaks of infectious diseases that may be distributed over large geographic areas.

Illegal trafficking in wildlife, as pets or food, is difficult to monitor. However, annual sales estimates in the United States alone exceed \$10 billion for pets and \$15 billion for bushmeat.¹³ Analysis of bushmeat from bats, rodents, and primates confiscated in major ports has revealed evidence of foamy viruses, herpesviruses, and pathogenic bacteria.¹⁴ Imported pets have been linked to outbreaks of human infection with poxviruses, *Salmonella*, and other pathogens.¹⁴

Attention has also increasingly focused on the role of land-use dynamics in infectious disease emergence.¹⁵ Deforestation, the expansion of agriculture, and the extractive industries, particularly in tropical regions with high wildlife biodiversity, have led directly or indirectly to the emergence of HIV/AIDS, Nipah virus, and filoviruses.^{16,17} The growth of suburbs in areas once only sparsely populated, particularly in the Northeastern United States, has been associated with an increase in the incidence of Lyme disease.

Global warming is already extending the geographic range of mosquitos and ticks that harbor and transmit plasmodia and arboviruses, resulting in outbreaks of malaria, dengue, and yellow fever in new locations.¹⁸ Recent examples in North America include the appearance of dengue fever in Florida in 2010^{19,20} and a surge in cases of West Nile encephalitis in Texas in 2012.²¹ Mass migration due to war, natural disaster, poverty, and desertification can lead to increases in population density not only of humans but also of disease vectors, such as rodents and ectoparasites that carry pathogenic viruses and bacteria. In concert, these factors, malnutrition, lack of access to or refusal of vaccines, and exposure to contaminated food and water have enabled the emergence and transmission of infectious diseases.^{22,23}

PREDICTION AND EARLY DETECTION OF EMERGING AND REEMERGING ZOOONOTIC DISEASES

Rapid recognition and response are critical to reducing the mortality, morbidity, and economic and social costs of zoonotic outbreaks. An important factor in early outbreak detection has been the advent of tools for internet-based infectious disease surveillance. The first of these was ProMED-mail (Programme for Monitoring Emerging Infectious Diseases), created in 1994, which provides continuous, free email updates on new or evolving outbreaks and epidemics²⁴ in people, domestic animals, and wildlife. Reports submitted by readers are curated by a panel of experts who post submissions with commentary in several languages to a listserv of subscribers in close to 200 different countries.

Another program, GPHIN (Global Public Health Intelligence Network),²⁵ a fee-based service, scans news services worldwide in several languages for information concerning outbreaks.

HealthMap²⁶ integrates reports from news media, ProMED-mail, and official documents into a user-friendly map that displays real-time updates of disease emergence. HealthMap also allows for public submission observations via its website or cellular phone applications.

An ideal surveillance system for zoonotic disease is one that allows identification of potential health threats before they move into the human population. By considering factors implicated as drivers in the emergence of zoonotic diseases, such as human demographics, agricultural production, land-use change, travel and trade patterns, climate, and wildlife distribution, risk algorithms can be developed and used to focus surveillance on sites, populations, professions, species of domestic animals, and wildlife in which there is an increased probability of known or novel high-threat pathogen emergence.²⁷ The two largest research programs with this focus are PREDICT, sponsored by the US Agency for International Development, and PREEMPT, sponsored the Defense Advanced Research Projects Agency.^{27a,27b} PREDICT supports zoonotic surveillance and laboratory capacity in the developing world.²⁸ PREEMPT will develop models of viral emergence and strategies for eliminating potential human pathogens in wildlife before they jump to humans.²⁹ The global acknowledgment of the importance of rigorous, coordinated infectious disease surveillance, early detection, and transparency in data sharing was underscored in the International Health Regulations of 2005, signed by all member states of the United Nations.³⁰

ZOOONOTIC DISEASES

A chapter concerned with infectious agents that share the capacity to jump host species from domestic animals or wildlife into humans could be organized according to agent, mechanism of transmission, or clinical presentation. This chapter includes all three approaches. The following text provides an overview of a representative set of zoonotic diseases associated with bats, rodents, and other wildlife and domesticated animals. It is by no means complete, and others might highlight different choices. Nonetheless, it will provide a framework for thinking about the range of zoonotic diseases and the factors that contribute to their emergence and control. Table 317.1 indicates routes of transmission and associated syndromes.

INFECTIONS ASSOCIATED WITH WILDLIFE

Bats

Bats, order Chiroptera, comprising more than 1200 species, represent approximately 20% of mammalian species diversity and are found on every continent except Antarctica. They are divided into two suborders: the Megachiroptera (fruit bats and flying foxes) and the Microchiroptera (insectivorous and vampire bats). Bats are unique among mammals in their ability to fly, tendency to aggregate at high density, and capacity to harbor a wide range of viruses without apparent disease. The physiology of bats is poorly understood, and their tolerance for persistent infection with viruses that cause fatal disease in other mammals is an enigma. However, this is likely to change with recent investments in bat genomic sequencing and systems biology.

A wide range of highly pathogenic RNA viruses have been found to be carried by bats: henipaviruses associated with equine and human infections in Australia (Hendra virus); Nipah virus, responsible for human and pig infections in Malaysia and human infections in Bangladesh; filoviruses associated with human and great ape infections in central Africa (Ebola virus and Marburg virus); rabies virus; and the coronaviruses implicated in SARS and Middle East respiratory syndrome (MERS), which caused a recent outbreak of fatal respiratory and renal disease in the Middle East. The route of transmission can vary, even for the same virus. In Malaysia, farm workers became infected with Nipah virus through exposure to pigs that had consumed contaminated droppings of frugivorous bats; in Bangladesh, where pigs are not farmed, human Nipah virus infection has been associated with consumption of palm sap beverages contaminated by bats. In North America, silver-haired bats transmit rabies virus via a bite wound. Humans are typically infected with filoviruses (e.g., Ebola) through exposure to infected

TABLE 317.1 Examples of Zoonotic Diseases: Agents and Vectors

VECTOR		AGENT FAMILY	AGENT	DISEASE
Arthropod	Mosquito	Bunyaviridae	La Crosse encephalitis virus, California encephalitis virus	Encephalitis
		Flaviviridae	Rift Valley fever virus	Hemorrhagic fever
		Flaviviridae	Japanese encephalitis virus, St. Louis encephalitis virus, West Nile virus	Encephalitis
	Tick	Togaviridae	Dengue virus, yellow fever virus, Zika virus	Hemorrhagic fever
		Togaviridae	Eastern equine encephalitis virus, western equine encephalitis virus, Venezuelan equine encephalitis virus, chikungunya virus, o'nyong-nyong fever virus	Encephalitis
		Bunyaviridae	Crimean-Congo hemorrhagic fever virus	Hemorrhagic fever
	Flea	Flaviviridae	Tick-borne encephalitis virus, Powassan encephalitis virus	Encephalitis
		Spirochaetaceae	Omsk hemorrhagic fever virus, Kyasanur Forest virus, Langat virus	Hemorrhagic fever
		Spirochaetaceae	Borrelia	Lyme disease
		Enterobacteriaceae	<i>Yersinia pestis</i>	Hemorrhagic fever (plague)
Mammal	Rodent	Arenaviridae	Lassa fever virus, Guanarito virus, Junin virus, Machupo virus, Sabia virus, Lujo virus	Hemorrhagic fever
		Bunyaviridae	Lymphocytic choriomeningitis virus	Meningitis, encephalitis
		Bunyaviridae	Dobrava-Belgrade virus, Hantaan virus, Puumala virus, Seoul virus	Hemorrhagic fever
	Rabbit	Spirochaetaceae	Sin Nombre virus	Pulmonary
		Spirochaetaceae	<i>Leptospira</i>	Leptospirosis
		Francisellaceae	<i>Francisella tularensis</i>	Tularemia
	Bat	Filoviridae	Ebola virus, Bundibugyo virus, Sudan virus, Tai Forest virus, Marburg virus	Hemorrhagic fever
		Rhabdoviridae	Rabies virus, Chandipura virus	Encephalitis
		Paramyxoviridae	Hendra virus, Nipah virus	Pulmonary, encephalitis
	Cattle	Coronaviridae	Severe acute respiratory syndrome (SARS) coronavirus, Middle Eastern coronavirus	Pulmonary
		Prions	Bovine spongiform encephalopathy prion	Neurodegeneration
	Cat	Bartonellaceae	<i>Bartonella</i>	Hemorrhagic fever
		Sarcocystidae	<i>Toxoplasma</i>	Toxoplasmosis
Avian	Bird	Orthomyxoviridae	Influenza virus	Pulmonary

primate meat; however, there are instances in which infection may have occurred through consumption of infected bats or exposure to bat excreta in close quarters (e.g., caves and abandoned mines).

The SARS coronavirus is presumed to have entered the human population through intermediate species (civets) in live-animal markets. The major reservoir for the MERS coronavirus is the dromedary camel.^{31,32} Although there are examples of human-to-human transmission of filoviruses and henipaviruses through contact with blood or other bodily fluids, the only bat-associated pathogen known to have established efficient transmission within the human population is the SARS coronavirus. Nipah virus, Hendra virus, SARS coronavirus, and MERS coronavirus cause respiratory disease and spread via aerosol. All are associated with high morbidity and mortality. However, the primary site of pathology in SARS and MERS is the lung, whereas Nipah and Hendra viruses cause encephalitis. Ebola virus and Marburg virus cause disseminated intravascular coagulation, resulting in hemorrhagic shock and multiorgan failure. Ebola virus, Marburg virus, and the arenaviruses and hantaviruses, described in the next section, are known collectively as hemorrhagic fever viruses in accord with the clinical manifestations of disease.

A diagnosis can be made based on clinical presentation in contexts in which others with similar presentation (including animals) have had a laboratory diagnosis. Although serology can be definitive, a laboratory diagnosis is typically made by means of an agent-specific polymerase chain reaction (PCR) assay. There is no established effective therapy for the diseases caused by SARS, MERS, Nipah, Hendra, Ebola, or Marburg viruses. A recombinant vesiculostomatitis virus vaccine expressing the surface glycoprotein of the Zaire ebolavirus was 100% effective in preventing disease in an open label trial in Guinea.³³ Monoclonal antibodies may reduce morbidity and mortality in disease caused by Nipah and Hendra viruses; therefore it is likely that a vaccine that elicits responses to similar antigenic targets would be effective. Rabies is typically fatal once the virus has entered the central nervous system; however, postexposure prophylaxis is effective, particularly when the exposed individual receives not only the vaccine, but also rabies immune globulin injected in proximity to the puncture wound.^{34,35}

Rodents

Rodents, like bats, are globally distributed. They are the largest order of mammals, comprising more than 2200 species, and are implicated in transmission of both viruses and bacteria. The rodent-associated arenaviruses (named for their sandy appearance with electron microscopy; *arena* means "sand" in Latin) causing human disease include the Lassa fever virus, endemic in Western Africa; the lymphocytic choriomeningitis

virus, found in both the Eastern and Western Hemispheres; and several South American viruses including Guanarito, Junin, Machupo, and Sabia. Another African arenavirus, Lujo virus, is presumed, but not proven, to have a rodent reservoir. Lassa fever is clinically similar to Ebola and Marburg; however, Lassa fever virus infection may be asymptomatic in up to 80% of cases, whereas asymptomatic filovirus infection is uncommon. The South American arenaviruses can cause hemorrhagic fever, but can also be associated with asymptomatic infection. Lymphocytic choriomeningitis virus has been implicated in aseptic meningitis and birth defects, particularly in the central nervous system. Although there is no specific drug for arenavirus infection, the nucleoside analogue ribavirin may have efficacy. Passive immunotherapy with hyperimmune plasma has been useful in Argentinean hemorrhagic fever (Junin virus infection).³⁶

Leptospira, a genus named for its morphology (*leptos* is "thin" in Greek; *spiral* is "coil" in Latin), consists of spirochetes that persistently infect rodents, concentrating in the kidneys. Infectious organisms are excreted in urine, collect in standing water, and enter humans and other animals either through breaks in the skin or via the alimentary canal. Infection is correlated with the potential for exposure; hence, leptospirosis is common year round in the tropics and during warm wet months in temperate climate zones. Food markets, particularly those where rodents congregate and where there is opportunity for urine from infected animals to contaminate standing water, are high-risk environments for contraction of disease. The onset of disease is heralded by nonspecific symptoms such as fever, malaise, and headache. It may progress to meningitis, vasculitis, renal and liver failure, and disseminated intravascular coagulation. Antibiotics (penicillin, amoxicillin, and doxycycline) are effective in prophylaxis and as a therapeutic intervention. There are vaccines for some serotypes.

The Bunyaviridae family of RNA viruses includes several important human pathogens—hantaviruses,airoviruses, and phleboviruses. Hantaviruses are transmitted through contact with rodent excreta; the other clinically significant bunyaviruses are transmitted through blood exchange by arthropod vectors, including mosquitoes, ticks, and flies. Hantaviruses (named for the Hantan river in South Korea, where the first hantaviruses were identified) may cause hemorrhagic fever with renal failure (hemorrhagic fever with renal syndrome [HFRS])³⁷ or an acute respiratory distress syndrome (HPS).³⁸ HFRS is more common in Africa, Asia, and Europe. HPS has been reported only in North America. There is no hantavirus-specific drug or vaccine. Recovery from HFRS is not uncommon with supportive care; however, HPS is typically fatal. Crimean-Congo hemorrhagic fever virus, aairovirus,

is distributed throughout Asia and Africa.³⁹ A wide range of small mammals may serve as reservoirs; domestic sheep, goats, cattle, and horses are typically intermediate hosts. The agent is most commonly transmitted by ticks; however, outbreaks have been linked to preparation or consumption of infected meat.⁴⁰ Manifestations of disease range from mild influenza-like illness to hemorrhagic fever.

Tick-borne encephalitis is a meningoencephalitis due to infection with the flavivirus TBEV.⁴¹ TBEV is found in Europe and Asia. It is transmitted to humans from a wide range of infected hosts, including sheep, deer, and rodents. Infection is rarely fatal; however, neurologic sequelae are not uncommon. There are effective vaccines but no known antivirals.

The most common tick-borne illness in the United States is Lyme disease.⁴² In North America, the causative agent is the spirochete *B. burgdorferi*, carried by the tick *Ixodes scapularis*. In Europe and Asia, the corresponding agents are *B. afzelii* and *B. garinii*, carried by *Ixodes ricinus* and *Ixodes persulcatus*, respectively. Early signs and symptoms of disease may include fever, lassitude, and headache. Some individuals have a classic "bull's-eye" skin rash known as erythema chronicum migrans, characterized by redness in the center and the periphery with central clearing. Later stages of the disease may include facial weakness (Bell palsy), peripheral neuropathy, and encephalomyelitis. Rodents are the reservoir for the spirochete; however, deer are also important to its life cycle as alternate reservoir hosts. Thus, human transmission is most common in suburban and rural areas, particularly where the density of vegetation is sufficiently thick to harbor large concentrations of rodents. Diagnosis is straightforward in the presence of the characteristic rash when there is a history of travel to an endemic area and a tick bite. However, many patients will not recall a tick bite, and up to 20% may not have the rash. Direct detection of the causative organism by means of culture or PCR analysis of the rash is definitive. Serology, the mainstay of laboratory diagnosis, is frequently inconclusive or misleading.⁴³

Francisella tularensis, the causative agent of tularemia (also known as rabbit fever), infects a wide range of vertebrate and invertebrates. The most important reservoirs are rodents, deer, and rabbits. Humans become infected through contact with infected animals or hematophagous vectors, including mosquitoes, ticks, or flies. Thus, hunters, hikers, and farmers are at highest risk in endemic areas. Outbreaks of severe respiratory disease have been described in landscapers. The manifestations of infection vary with the route of exposure. Cutaneous inoculation typically results in a vesicular rash, which may or may not ulcerate, and regional lymphadenopathy. Ingestion results in oropharyngeal lesions. Inhalation can lead to fatal pneumonia. Ocular inoculation results in conjunctivitis. Without antibiotic treatment, some strains disseminate systemically and are associated with up to 30% mortality. The diagnosis is typically made based on clinical presentation, with confirmation by means of PCR or culture. The agent is sensitive to ciprofloxacin, aminoglycosides, and chloramphenicol. Vaccines are in development due to concerns that *F. tularensis* may be weaponized.⁴⁴

Yersinia pestis, the causative agent of plague, is found in rodents throughout the world with the exception of Oceania. Infection is transmitted to humans primarily by fleas. However, transmission can also occur through ingestion and inhalation. Periodic pandemics, including the Black Death, have cost hundreds of millions of lives. Although outbreaks continue in the developing world, the majority of current reports in the United States note infections in single individuals. The manifestations of disease can vary from an influenza-like mild fever and lymphadenopathy (buboes) to pneumonia, shock, disseminated intravascular coagulation, and tissue necrosis. The diagnosis is typically made based on clinical presentation, with confirmation by means of PCR or culture. Like *F. tularensis*, *Y. pestis* is sensitive to ciprofloxacin, aminoglycosides, and chloramphenicol. It has the dubious distinction of being one of the first microbes to be used as a biologic weapon, in the 14th century, and remains a biosecurity concern.⁴⁵

DISEASES ASSOCIATED WITH AGRICULTURAL AND OTHER DOMESTIC ANIMALS

Zoonotic agents associated with livestock, poultry, and companion animals include bacteria, fungi, parasites, and prions. Industrialization

of food production has increased the probability of meat cross-contamination, in addition to animal-to-animal infection on feedlots. Outbreaks of disease due to *E. coli*, including the potentially fatal hemolytic-uremic syndrome, are commonly linked to contaminated ground beef. Infected poultry and eggs are frequently associated with outbreaks of *Salmonella* infection. The use of antibiotics as growth promoters in agriculture also encourages the emergence of antibiotic-resistant bacteria that contribute to human morbidity and mortality associated with infectious disease.⁷

vCJD is a progressive, fatal brain disorder caused by exposure to misfolded proteins (prions) in cattle. vCJD and an analogous disease known as bovine spongiform encephalopathy (BSE) catalyze conformational changes in host proteins. BSE was first identified in the United Kingdom in 1986 and peaked in incidence in 1993. The original infection is presumed to have been introduced when cattle were fed with meat (offal) from infected sheep or cows. An estimated 400,000 cows with BSE are estimated to have entered the food chain before the outbreak was terminated by culling cattle with behavior indicative of BSE, reducing the age of slaughter (so that titers would be lower in the event that an infected animal was inadvertently used for human consumption), and eliminating the practice of feeding offal to ruminants in 1997. At least 227 people exposed to BSE contracted vCJD. Susceptibility to disease is determined at least in part by the prion protein (PrP) genotype. The peak incidence of vCJD was observed in 2000 and tapered thereafter. Some investigators hold that a large wave of new cases is unlikely in the absence of recurrence of BSE. However, others note that the incubation period of kuru, a prion disorder associated with ritual cannibalism, due to essentially the same agent, may be as long as 27 years. The diagnosis of vCJD is made through brain or tonsillar biopsy based on histologic findings and/or the presence of protease-resistant PrP. There is no effective treatment or vaccine for prion diseases.

Rift Valley fever virus, a bunyavirus transmitted by mosquitoes, chiefly affects livestock but can cause human disease (Rift Valley fever), ranging from a mild febrile illness to meningitis or hemorrhage and multiorgan failure. Originally reported in Kenya, the virus has expanded in distribution across sub-Saharan Africa and the Arabian Peninsula.

A detailed discussion of influenza virus is beyond the scope of this chapter. Furthermore, influenza is not typically considered a zoonotic disease because human-to-human transmission is efficient and represents the large majority of cases of influenza A. Nonetheless, both birds and pigs play key roles in the maintenance of existing strains and the emergence of new ones. Only influenza A viruses are associated with pandemic disease, presumably because they have a higher propensity to evolve toward antigenic diversity and thus evade host immune surveillance than influenza B and C viruses. Wild aquatic birds are the natural hosts of the classic seasonal influenza viruses H1N1 and H3N2. Infection of free-range pigs by migrating aquatic birds or in live-animal markets where they may be housed near aquatic birds can lead to genetic reassortment (when more than one type of influenza virus is present) and adaption to humans. H5N1, popularly known as avian influenza, only rarely results in human disease but is frequently fatal when it does. A wide range of diagnostic assays is available for detection of influenza viruses. Vaccines are designed twice each year to reflect strains predicted to predominate during the influenza season in the Southern and Northern Hemispheres. Their efficacy fluctuates with the accuracy of those predictions, but, as a rule, they are less effective in inducing a protective immune response in the elderly. Efforts are underway to create universal influenza vaccines based on conserved regions of the virus hemagglutinin rather than those that evolve to evade the immune system.⁴⁶ The drug oseltamivir is effective for many strains.

Up to 30% of the world's population may have been exposed to the protozoan *Toxoplasma gondii*, including more than 60,000,000 people in the United States. Infection is contracted through ingestion of contaminated food or contact with feces of domestic cats, the natural host. In most instances, infection is asymptomatic or associated with a transient influenza-like illness; however, with immunosuppression in the context of HIV/AIDS, chemotherapy, or pregnancy, the parasite can cause a mass lesion in the brain, leading to focal neurologic deficits and seizures. Infection of the fetus during pregnancy can result in

miscarriage and abnormal brain development. Treatment varies with the immunologic status of the patient.

Eastern equine encephalitis virus (EEEV), western equine encephalitis virus (WEEV), and Venezuelan equine encephalitis virus (VEEV) are mosquito-borne alphaviruses that infect a wide range of vertebrates. Birds are the natural reservoirs. The nomenclature reflects the observation that horses with encephalitis are frequent sentinels for risk of human disease and roughly approximates the geographic distribution of the virus. EEEV is found throughout North, Central, and South America but is concentrated in the United States east of the Continental Divide. WEEV is found in the western United States. VEEV is found primarily in Central and South America. All are associated primarily with an influenza-like illness but may, in a small proportion of cases, progress to encephalitis. WNV is a mosquito-transmitted flavivirus associated with only mild, sporadic disease until the 1990s, when virulent strains appeared first in Europe and then in North America. Many vertebrate species are susceptible to infection; however, the primary reservoir is birds. Although most infections are asymptomatic or associated with mild influenza-like illness (West Nile fever), encephalitis and poliomyelitis can occur in individuals who are immunologically compromised because of advanced age or underlying medical conditions. Only supportive therapy is available for EEEV, WEEV, VEEV, and WNV encephalitis. Vaccines have been developed but are typically not used in humans, except where biomedical researchers concentrate VEEV and are at high risk for exposure.

Zika virus, identified in a rhesus monkey in 1947, is named for its site of isolation within the Zika forest in Uganda. It has been implicated in microcephaly and other neurodevelopmental damage in children

whose mothers were infected during pregnancy and in Guillain-Barré syndrome in infected adults. Zika virus has both a sylvatic and an urban cycle. In the former, the virus is maintained in nonhuman primates by *Aedes* and *Anopheles* mosquitoes. Humans appear to be the only reservoir in the urban cycle. Human-to-human transmission is most commonly mediated by *Aedes* mosquitoes, but humans also can become infected through exposure to infected semen or blood products.⁴⁷ There is no known treatment for Zika; however, several vaccines are in development.

CONCLUSIONS

The majority of emerging and reemerging infectious diseases are zoonoses. Zoonoses are likely to increase in frequency owing to anthropogenic factors that include incursion into wildlife habitats, globalization of travel and trade, and climatic changes that influence the distribution of arboviral vectors, including mosquitoes, phlebotomous flies, and ticks. Antimicrobial resistance is not a zoonosis in the strict sense of the term. Nonetheless, the threat of antimicrobial resistance is clearly exacerbated by the misuse of antibiotics in animal husbandry.

In viral zoonoses, interventions are currently limited to supportive care and in some instances, such as rabies and some hemorrhagic fevers, to immunotherapy with hyperimmune globulin or monoclonal antibodies. In contrast, specific antibiotic therapy has utility in bacterial zoonoses. Investments in biodefense focused on understanding viral biology and the development of antiviral drugs are rapidly leading to specific interventions that will reduce the morbidity and mortality of emerging viral infections. Accordingly, early differential diagnosis will become increasingly important not only for outbreak containment but also for ensuring that patients receive appropriate treatment.

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F Protection of Travelers

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Protection of Travelers

David O. Freedman

SHORT VIEW SUMMARY

- A pretravel office visit with an adult traveler to the developing world should follow a structured approach.

PERFORM RISK ASSESSMENT

- Exact itinerary, including regions within each country to be visited, dates of travel to assess risk of seasonal diseases, age, past vaccination and travel history, immune status, underlying illnesses, current medications, pregnancy status, allergies, purpose of trip, risk exposures (blood, body fluids, adventure or extensive outdoor exposures), urban versus rural travel, type of accommodations, travelers' risk tolerance, and financial limitations that may necessitate prioritization of interventions

ADMINISTER IMMUNIZATIONS

- Routine vaccinations that are not up-to-date, including measles-mumps-rubella (MMR), tetanus-diphtheria–acellular pertussis (Tdap), pneumococcal, varicella zoster virus
- Indicated routine travel vaccines, including hepatitis A, hepatitis B, typhoid, and influenza
- Indicated specialized vaccines, including yellow fever, rabies, polio, meningococcal, cholera,

Japanese encephalitis, and tick-borne encephalitis in certain countries

PROVIDE MALARIA PREVENTION (IF INDICATED)

- Several equally effective drugs of choice may be indicated, including atovaquone-proguanil, mefloquine, and doxycycline. Ascertain which drug is best suited to the individual patient and itinerary.
- Educate on personal protection against arthropods.

TRAVELER'S DIARRHEA

- Recommend food and water precautions, as well as hand hygiene.
- Advise on use of loperamide and oral hydration if needed.
- Prescribe and educate on standby therapy with azithromycin (sometimes a quinolone) only for diarrhea that is disabling and prevents normal activities.

VECTOR-BORNE DISEASES

- Educate on personal protection measures for malaria, dengue, chikungunya, Zika virus infection, leishmaniasis, rickettsial disease, and sleeping sickness

TEACH ESSENTIAL PREVENTIVE BEHAVIORS

- Most travel-related health problems, including vaccine-preventable diseases, can be avoided through simple behaviors initiated by the traveler.
- Educate on appropriate strategies in the following categories (some topics are not applicable to all destinations): bloodborne and sexually transmitted diseases, safety and crime avoidance, transportation-associated and other injury prevention, swimming safety, rabies, skin/wound care, and tuberculosis.

DISCUSS OTHER APPLICABLE HEALTH ISSUES

- Educate on and prescribe for altitude illness, motion sickness, traveler's thrombosis, or jet lag.
- Discuss minimal-risk conditions (e.g., hemorrhagic fevers) that are a frequent cause of patient anxiety.

MEDICAL KIT AND MEDICAL CARE ABROAD

- Contents of a personal health kit
- Finding destination medical facilities
- Evacuation insurance and supplemental health insurance

Pretravel management of an international traveler should be based on risk management principles. Prevention strategies and medical interventions need to be individualized according to both the itinerary and factors that are dependent on the traveler. A structured approach to patient interaction (Table 318.1) is the most efficient way to cover the necessary educational and preventive interventions. Because many of these measures will not be initiated until much later at the traveler's destination, clearly printed instructions in lay language are advisable. The worldwide epidemiology of travel-related diseases is constantly changing. A body of knowledge in travel medicine has been published, and online and print resources (Table 318.2) should be consulted frequently¹⁻³ to keep current.

EPIDEMIOLOGY OF TRAVEL-RELATED ILLNESS

In 2015 international tourist arrivals in all countries exceeded 1.2 billion people. In 2014 the total number of arrivals in countries with emerging markets nearly surpassed the number in developed countries. Depending on the destination, 6% to 87% of travelers become ill across all studies.⁴⁻⁶ Of four studies that provide the best estimate, 43% to 79% of travelers who frequently visited developing nations (e.g., India, Tanzania, and Kenya) became ill; most of these illnesses were mild and self-limited such as

diarrhea, respiratory infections, and skin disorders.⁷ Some travelers return to their own countries with preventable life-threatening infections.⁸ Yet 20% to 80% of travelers do not seek pretravel health consultation.^{9,10} Data about the effect of pretravel advice are limited, although such advice has had a positive effect on the prevention of malaria. Travelers visiting friends and relatives in their country of origin constitute the group with the highest morbidity, especially from malaria and typhoid; this group requires special approaches to illness prevention and education.^{11,12} Rates are significantly higher in summer. Approximately 8% of travelers consult a physician either during or after a trip, but less than 1% require hospitalization. Infectious diseases account for up to 10% of morbidity during travel but only 1% of deaths, with malaria the most common disease. Causes of death vary according to the population studied. At destinations that attract elderly adults, cardiovascular events predominate, whereas in developing countries, motor vehicle accidents and drowning prevail.

IMMUNIZATION

The choice of vaccines for an individual traveler is based on risk of exposure to vaccine-preventable diseases on the chosen itinerary; the severity of disease, if acquired; and any risks presented by the vaccine itself. Travelers differ in their tolerance of risk. Requests for immunization