
*Recent Animal Disease Outbreaks and their Impact on Human Populations*¹

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Animals occupy a special place in human societies. They are utilized for food (*e.g.*, milk and meat), transportation, raw materials (*e.g.*, wool and hides), energy (*e.g.*, manure), recreation, and money (*e.g.*, bartering). Furthermore, animals such as dogs, cats, and horses in some societies often are viewed as “companions.” Their value in long-term care facilities and for the emotional well-being of AIDS patients has been documented (Siegel *et al.*, 1999). In addition to these valuable contributions, there is growing concern about diseases that humans can acquire from animals (*e.g.*, zoonoses). Zoonoses are overrepresented among human diseases that are defined as emerging (Table 1). Taylor *et al.* (2001) documented that 61% of all human pathogens are zoonotic. And of the 175 newly emerging pathogens in humans, 75% are listed as zoonotic (Cleaveland *et al.*, 2001). From 1996 to 2006, eleven of the twelve global emerging diseases originated from animals (Gerberding, 2004).

However, it is also important to remember that some diseases affect animals only, often with economic, environmental and/or societal implications. Recent examples include chronic wasting disease in elk and deer, foot-and-mouth disease, toxoplasmosis in sea otters, and salmonella in song birds. In 1994, canine distemper jumped the “species-barrier” and infected African lions of the Serengeti (Roelke-Parker *et al.*, 1996), killing over a third of the population within 6 months.

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TABLE 1. RECENT ZOOONOTIC AGENTS IDENTIFIED.

Agent	Identified	Common illness in humans	Common illness in animals
<i>Cryptosporidium parvum</i>	1976	Profuse and watery diarrhea	Diarrhea in calves
Ebola virus	1977	Hemorrhagic fever and death (high case-fatality rate)	Hemorrhagic fever and death in primates
Hanta virus	1977	Fever and hypotension	None
<i>Campylobacter</i> spp.	1977	Diarrhea, abdominal pain, and fever	None
<i>Escherichia coli</i> O157	1982	Hemorrhagic enterocolitis	None
<i>Borrelia burgdorferi</i>	1982	Arthritis and skin rash	Arthritis in companion animals
<i>Ehrlichia chaffeensis</i>	1987	Fever, headache and malaise	None
<i>Anaplasma phagocytophilum</i>	1990	Fever, headache and malaise	Fever, lethargy in dogs and horses
<i>Bartonella henselae</i>	1992	Lymphadenitis and fever	Rare illness in cats, fever
Sin nombre virus (Hanta virus)	1993	Pulmonary syndrome, fever, myalgias	None
Hendra virus	1994	Pneumonia/encephalitis	Respiratory disease and death in horses
West Nile virus	1999	Fever, encephalitis	Fever, muscle tremors, encephalitis
Nipah virus	1999	Encephalitis	None
SARS-coV	2003	Pneumonia	None

New diseases emerge for a number of reasons: world trade, animal translocation, ecological disruption, climate change, pathogen adaptation, and agricultural husbandry changes (Smolinski *et al.*, 2003). These factors represent the dynamic relationships among the pathogenic agent, host, and environment (Figure 1). This epidemiologic triangle includes the *intrinsic* characteristics of an individual's susceptibility to disease, including immune status, general health, genetic makeup, lifestyle, age, sex, and socioeconomic status, and *extrinsic* factors, which include the host's biological, social, and physical environment. Coincidentally, the triangle describing this relationship is the same as delta, the symbol for change; change is the one constant in the on-going tension between humans and microbes.

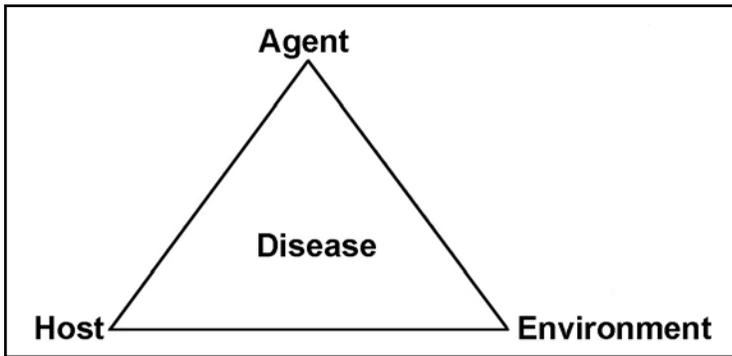


Figure 1. The epidemiologic triangle.

This article will discuss some recent outbreaks of disease, lessons learned, and challenges for the future. We will describe:

- the strong connection between animals and humans,
- the challenge of effective risk communication where there is limited knowledge of the risks,
- the dwindling and fragile animal-health and public-health systems,
- the lack of oversight and regulations to prevent disease transmission,
- changes in agricultural practices that result in new or re-emerging diseases, and
- the relationship between culture and disease.

We will discuss the specific examples of foot-and-mouth disease (FMD), chronic wasting disease (CWD), monkeypox, severe acute respiratory syndrome (SARS), and avian influenza.

THE STRONG CONNECTION BETWEEN ANIMALS AND HUMANS

Foot-and-Mouth Disease

Some diseases may not have a direct impact on human health but, nonetheless, exert significant societal pressure by disrupting local economies as well as world trade. This is exemplified by the 2001 outbreak of FMD in the United Kingdom, which spread to other countries in Europe (Figure 2).



Figure 2. The impact of culling sheep during the outbreak of foot and mouth disease in France, 2001.

Foot-and-mouth disease is primarily a disease of cloven-footed domestic and wild animals. It is endemic in Asia, Africa, and parts of South America. However, some areas of the world are free of FMD, including North and Central America, Australia, New Zealand, Japan and most European countries. The causal agent is considered one of the most highly contagious viruses, and its contagiousness has huge implications on trade of livestock and livestock products. The disease may spread by direct or indirect contact with infected animals, aerosol from infected animals or milk trucks, and fomites, as well as through artificial insemination. People who come into contact with infected animals can serve as mechanical vectors, as sufficient FMD virus survives in their upper airways for 24 hours to potentially serve as an ongoing source of infection to livestock (Sellers *et al.*, 1971).

During the FMD outbreak in the United Kingdom in 2001, an estimated 2,000 confirmed cases and an additional 6 million animals were slaughtered to achieve containment (DEFRA, 2005a). The cost of controlling the outbreak and losses due to decreased tourism were estimated at £6.2 billion (DEFRA, 2005a). The postulated source was illegally imported food that eventually ended up as scraps in garbage fed to pigs (DEFRA, 2005b). The psychological and economic impact on the British population—farmers and non-farmers alike—was huge. Increases in suicides among farmers were reported and substantial economic losses were incurred from a trade embargo, travel restrictions, and reduced tourist income (DEFRA, 2005a). This does not take into account the loss of genetic stock and the cost of controlling the outbreak. A psychological assessment of the impact of FMD noted that farmers in the impacted area had significantly higher psychological morbidity scores compared to farmers in non-impacted areas (Peck *et al.*, 2002).

Cryptosporidiosis

Although FMD rarely is detected in humans, human health did not go entirely unaffected by the outbreak. The presence of FMD, an exotic animal disease, correlated with a decreased incidence of an endemic zoonotic disease, cryptosporidiosis, caused

by *Cryptosporidium parvum*. Cryptosporidiosis is the most common parasitic infection among people in the United Kingdom, where an estimated two-thirds of cases are due to *C. parvum*. Two separate reports described a significant drop in *Cryptosporidium* cases during the FMD outbreak. Hunter *et al.* (2003) reported a 69% decline in cases in the northwest of England (Figure 3). Strachan *et al.* (2003) reported a 34% decline in Scotland, with a noticeable difference between FMD-infected areas and FMD-free areas. Reasons for these reductions were restrictions of farm-animal movement, possibly the presence of fewer young animals (the major source of exposure), and fewer animal-to-human interactions that allow transmission.

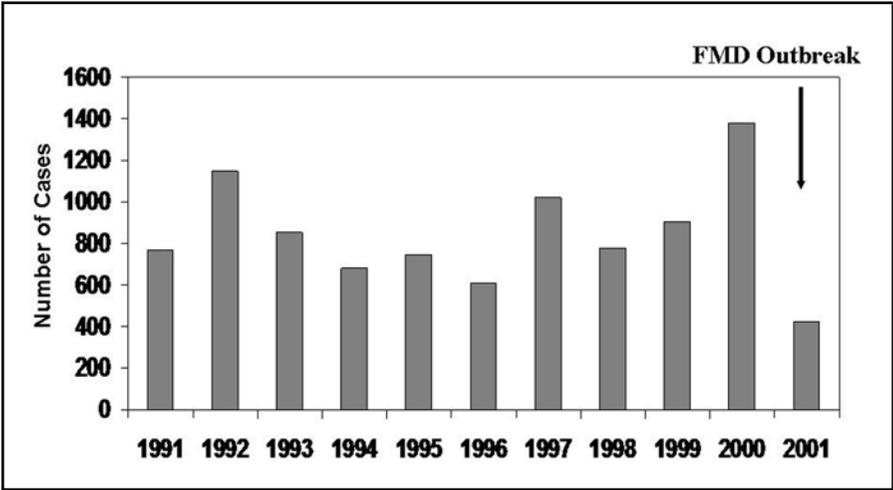


Figure 3. Reported cases of cryptosporidiosis in northwest England, 1991–2001 (Hunter *et al.*, 2003).

This outbreak of FMD highlights the strong and varied interrelationships between animals and humans. Although it is a disease primarily of animals with limited direct transmission to humans, it can have a significant public-health impact in terms of psychological effects, and its presence can send shockwaves through local economies. In addition, FMD is one of the primary agents of concern for agroterrorism, not only because of the economic and trade ramifications it can inflict on the livestock industry, but also because of the severe societal impact it may have. We must never underestimate the societal impact of diseases even when they directly impact the health only of animals.

A CHALLENGE IN EFFECTIVE RISK COMMUNICATION

A second animal disease capturing the headlines is CWD, a disease of the nervous system found in Cervidae: white-tailed deer, mule deer, black-tailed deer, and elk. CWD belongs to the family of diseases known as transmissible spongiform encephalopathies or prion diseases, and is a slowly progressive, invariably fatal neurologic disease in cervids. First

recognized as a new disease among captive mule deer in a Colorado wildlife unit, it was later found to be endemic in both mule deer and elk in Colorado and Wyoming (Williams and Miller, 2003). The origin of the disease is unknown, but some have speculated that CWD 1 (Williams and Miller, 2003):

- is an adapted strain of the scrapie agent found in sheep,
- arose as a spontaneous evolutionary event, or
- originated from a yet unidentified prion reservoir.

CWD has been found in various areas throughout North America, both in captive and in free-ranging cervids (Figure 4). The perceived spread from the initial endemic areas is likely attributable to the movement of deer and elk in commerce, local expansions of farmed herds, and increased surveillance efforts (Williams and Miller, 2003).

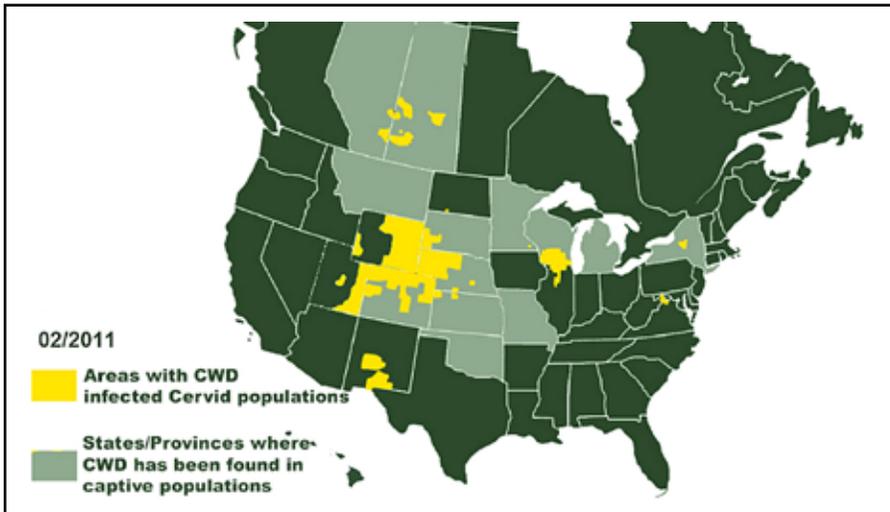


Figure 4. Chronic wasting disease in captive and free-ranging cervids (courtesy of the Chronic Wasting Disease Alliance, <http://www.cwd-info.org/index.php/fuseaction/about.map>).

Since cervids were found to have CWD, hunters, farmers and venison consumers have become concerned about the risk of zoonotic transmission, largely because of the connection between bovine spongiform encephalopathy (BSE) and variant Creutzfeldt-Jakob disease (vCJD). Creutzfeldt-Jakob disease occurs around the world at a rate of 1–2 per million humans. The majority of cases has occurred among British citizens and persons who have resided in the United Kingdom. All vCJD cases to date have lived in countries with BSE.

If CWD is a zoonotic disease, what would it look like in humans? Would people living in endemic areas be at greater risk? To date, investigators have not seen higher numbers of human spongiform encephalopathies in CWD-endemic areas. However, prion diseases

are rare, have long incubation periods, and can be difficult to detect. Because of recent concerns about prion diseases, epidemiologists are investigating neurologic diseases focusing on young people with unusual clinical presentations or neuropathology. Several documented clusters of cases have been investigated, often in response to concern from family members believing that deer-meat consumption was linked to illness (Belay *et al.*, 2004). These cluster investigations are a challenging exercise in risk communication about human and animal health. One investigation involved three elderly men, all of whom had a history of eating venison, who died of degenerative neurologic illnesses (CDC, 2003a). However, further diagnostic work-up revealed that only one actually had evidence of a prion disease. Currently, it is the consensus of the World Health Organization and the Centers for Disease Control and Prevention that there is no scientific evidence that CWD causes human illness (Belay *et al.*, 2004; WHO, 2005).

As with FMD, CWD has a psychological impact on humans although it does not directly harm human health. No definite link has been found between CWD and human brain disease, yet the detection of CWD in free-ranging deer in Wisconsin and Illinois in 2002 had a substantial impact on the human psyche. Nine months after CWD was discovered in Wisconsin, there was an 11% drop in deer-license sales (Heberlein, 2004). Also, similar to FMD, the discovery of CWD hurt local economies. Businesses that served Wisconsin hunters saw sharp declines in sales, as did feed dealers and local butcher shops. The decrease in license sales resulted in reduced revenues for the State of Wisconsin, and state expenditures increased \$14.7 million to control CWD; overall, the estimated economic impact in 2002 was between \$53 and \$79 million (Bishop, 2004). This situation illustrates the emotional and economic impacts of infectious diseases and the challenge of effectively communicating evolving risk with reference to emerging animal diseases.

DWINDLING AND FRAGILE ANIMAL- AND PUBLIC-HEALTH INFRASTRUCTURE

National economies are vulnerable to outbreaks of animal disease, both intentionally malicious and accidental. Recent terrorist attacks have exposed the vulnerability of our transportation, food, and medical infrastructure. Several episodes have been documented in which food was intentionally contaminated for terrorist purposes (Manning *et al.*, 2005). However, in recent years, the most dramatic impact on national economies has not come from terrorism, but from the accidental introduction of foreign animal diseases. The threat is very real when we consider the volume of travelers and traffic that enter the United States each year, both legally and illegally. There is no feasible way for each vehicle and piece of luggage to be thoroughly checked for microscopic travelers. In addition, millions of animals and animal products are imported. They can serve as silent disease carriers or can harbor insects and ticks that serve as disease vectors. Clearly, we need to give greater attention to training of, and cooperation among, veterinarians, livestock producers, extension personnel, and healthcare professionals. Specifically, since some of these diseases can be zoonotic, veterinarians and people who work to protect human health need to combine forces to quickly diagnose and control their spread, especially in rural communities.

The West Nile virus is another dramatic example of the animal- and public-health challenges of understanding an emerging disease with only limited personnel dedicated to understanding insect vectors and viral spread through wild-bird hosts. Originally a disease of Africa and Europe, it was first observed in New York in 1999 (Lanciotti *et al.*, 1999). Initially misdiagnosed as St. Louis encephalitis, this disease, new to the Western hemisphere, was astutely diagnosed with the combined efforts of a veterinary pathologist, a physician, and epidemiologists. The virus now has been documented in all states of the continental United States. Migratory birds and competent mosquito vectors were instrumental in the rapid westward spread. The ensuing epizootic has had a dramatic effect on horse, bird, and human populations. In 2002, over 15,000 horses were reported ill, and 30% died as a result of the infection (CDC, 2002a). The impact on raptors and corvids (blue jays and American crows) has also been well documented (Wunschmann *et al.*, 2004). However, the broader impact within ecosystems, especially on wildlife, is unknown. From 1999 through 2004, over 16,700 human cases and 666 deaths were reported in the United States (Hayes and Gubler, 2005). This disease highlights some of the new challenges for human clinicians of unusual disease presentations (*e.g.*, acute flaccid paralysis syndrome) and new routes of transmission (*e.g.*, blood transfusion and organ transplantation). The appearance of West Nile virus required the training and funding of public-health officials in mosquito trapping, vector control, and close collaboration with academic institutions for disease surveillance and public education.

LACK OF OVERSIGHT AND REGULATIONS

Monkeypox was first documented in 1958 in a colony of primates (hence the term). The first human cases were identified in 1970 in Zaire by local health officials on the lookout for the re-emergence of the smallpox virus. This rare disease was documented among people who lived where hunting was an integral aspect of their lifestyle. The natural disease hosts are likely several species of squirrel.

In 2003, an outbreak in the United States associated with legally imported African “pocket pets” led to seventy-two suspected human cases in six states (CDC, 2003b). Eighteen persons were hospitalized, some because of the potential for human-to-human spread. Interestingly, a number of the cases were veterinarians and veterinary technicians exposed while treating ill pets, highlighting potential occupational risk. The majority of patients had direct or close contact with prairie dogs that were infected by close contact with imported animals from Ghana, shipped to a distributor in Texas. The shipment included six genera of African rodents, including rope squirrels (*Funisciurus* sp.), tree squirrels (*Heliosciurus* sp.), Gambian giant rats (*Cricetomys* sp.), brushtail porcupines (*Atherurus* sp.), dormice (*Graphiurus* sp.), and striped mice (*Hybomys* sp.). There was a real concern of spillover of the virus from these imported animals to susceptible wildlife populations in the United States.

Even though this outbreak was not directly related to agriculture, it exemplifies the problem of both legal and illegal animal movements. The US Fish and Wildlife Service estimates that the global trade in endangered wildlife is \$4.2 billion annually, second only to illegal drugs. Other examples of emerging diseases linked to live-animal trade, include

the spread of rabies from trapping raccoons in Florida for game farms in West Virginia (CDC, 1981), the collection of prairie dogs for pet markets that were subsequently diagnosed with tularemia (CDC, 2002b), and the shipping of elk infected with CWD to Korea (Sohn *et al.*, 2002). All of these examples clearly demonstrate potential consequences when humans move animals from one area to another and the need for regulations and federal policies that control the transfer/exchange of exotic animals. Currently, there are regulations for rodents from Africa and poultry from Southeast Asia, but numerous animals still pass through US ports unregulated (DHHS, 2003). Currently, no regulations control the interstate movement of exotic animals or wildlife within the United States.

CHANGES IN AGRICULTURAL PRACTICES AND FOOD PROCESSING

The emergence of BSE demonstrated the role of animal-feed commodities such as meat and bone meal (MBM) in the spread of disease. Meat and bone meal is an important recycled byproduct used as an inexpensive protein source. Since the 1950s, this protein source has increasingly been added to the diets of high-producing or rapidly growing animals, for example, beef and dairy cattle. While the BSE outbreak has largely been confined to Great Britain, the movement of affected animals and/or contaminated MBM spread the disease throughout Europe and beyond including sporadic cases in Japan and North America. As a result, “firewalls” were devised to decrease the amplification and spread of the disease when a clear understanding of the risks was identified.

In addition to changes in feed ingredients such as those that led to the spread of BSE, other agricultural and food-production factors that might appear to be innocuous can also provide a mechanism for disease transmission. For example some have speculated that the move from pasture feeding in the mid-20th century to intensive grain feeding has altered the gastrointestinal tracts of cattle in a way that favors the growth of *Escherichia coli* O157:H7 (Russell *et al.*, 2000). A second example is *Listeria monocytogenes*, a bacterium recognized as an animal pathogen more than 100 years ago, but seen as a significant cause of human illness only since the 1980s. The emergence of *L. monocytogenes* as a food-borne pathogen is due to pathogen survival at refrigeration temperatures, the increasing number of immunocompromised individuals in the population, the centralization and consolidation of food production, and changes in consumer food habits (*e.g.* consumption of ready-to-eat foods) (Swaminathan, 2001). This disease reflects the impact of changing food-processing techniques, with which post-contamination of cooked foods can be a source of infection. These factors demonstrate the complex and evolving nature of pathogens and the need for animal- and public-health surveillance systems to quickly identify and characterize new and emerging pathogens.

CULTURAL PRACTICES AND DISEASE EMERGENCE

In many communities, there exist cultural or societal practices that can inadvertently encourage disease transmission by artificially causing animals to congregate. Recently, *Mycobacterium bovis* was identified among deer in northern Michigan, and its presence was attributed to the congregation of the deer due to “baiting” or feeding by deerhunters (Miller *et al.*, 2003). As a result, Michigan passed legislation prohibiting the feeding of

deer in an attempt to limit the transmission of *M. bovis*. A similar phenomenon is occurring with birds: when songbirds congregate at feeders, their increased proximity can lead to the spread of salmonellosis and their subsequent illness and death.

Examples of global problems of disease transmission abound. In November 2002, the detection of an atypical pneumonia quickly challenged the world public-health system. SARS caused illness in over 8,000 persons around the world with 774 documented deaths. The identification of this rapidly spreading disease had a dramatic impact on healthcare workers and patients' willingness to utilize medical services (Emanuel, 2003; Chang *et al.*, 2004; Maunder, 2004). Half of the first sixty cases identified were healthcare workers, but, despite the risk, they continued to care for patients. The impact was felt globally with cancelled air flights and record low hotel occupancy rates; for example, in Hong Kong hotels, they dropped to 17% compared to 83% a year earlier (Emanuel, 2003). The economic cost to Toronto, Canada, was estimated at nearly \$1 billion in 2003 (Blendon *et al.*, 2004).

SARS is a corona virus that likely emerged from a wild-animal source (Lau *et al.* 2005). This is supported by the detection of initial cases among restaurant workers handling exotic animals in Guangdong Province (Zhong *et al.*, 2003). SARS-CoV has also been isolated from masked palm civets and other wild animals in a live-animal market (Guan *et al.*, 2003; Lau *et al.*, 2005). Seroepidemiology of animal traders and handlers further supports this; 13% of animal traders had IgG antibody to SARS-CoV, as compared to 1 to 3% from community control groups (CDC, 2003c).

Researchers speculate that SARS-CoV likely originated from animals with which humans have infrequent contact, such as exotic species. The zoonotic link has been attributed to the phylogenetic relationship between corona viruses and those isolated from wild animals such as the palm civet and the raccoon dog. Contact likely occurred among southern Chinese who periodically consume wild-game meat for medicinal purposes. Zhong *et al.* (2003) have suggested that viruses that are transmitted between species tend to undergo more rapid genetic change as they adapt to new hosts. It is likely that novel viruses such as Ebola, HIV, and SARS-CoV will continue to appear with increased human interaction with wild animals. The lucrative wild-animal markets in Southeast Asia, a smorgasbord of wild and domestic animals, are often unregulated (Karesh *et al.*, 2005).

Avian influenza is another example that illustrates the relationship of cultural and social practices and the appearance of animal disease. Southeast Asia is considered the epicenter of recent influenza outbreaks. This is linked to agricultural practices in a highly populated area. Rice fields often have standing water that attracts waterfowl. These waterfowl are natural reservoirs, potentially spreading the disease to other domestic animals (*e.g.* chickens, ducks, and pigs) raised outdoors. In 2005 it was estimated that there were 1.3 billion humans, 508 million pigs and 13 billion chickens in China (Osterholm, 2005). The identification of novel avian influenza strains over the past 15 years documents the continual re-assortment of influenza viruses among birds, pigs and humans. Fortunately, sustained human-to-human transmission has not been documented (Ungchusak *et al.*, 2005). But with aquatic wild birds as the natural reservoir, it will be nearly impossible to eradicate this disease. The H5N1 strain responsible for the 1997 Hong Kong outbreak

of influenza in domestic poultry resulted in the culling of 1.5 million birds and the identification of eighteen human cases with six deaths (Bridges *et al.*, 2002). Similarly in the Netherlands, 28 million birds were culled with eighty-nine reported human cases and one death (Fouchier *et al.*, 2004). The 2003–2005 H5N1 outbreaks in Asia affected eleven countries, with 109 reported human cases and fifty-five deaths (CIDRAP, 2005). Like SARS, the economic impact in Southeast Asia was substantial. The South Korean Ministry of Health and Welfare estimated that the cost of avian influenza to Asian countries at about \$130 billion. Unlike SARS, influenza is a potentially greater problem with a common wildlife reservoir (*e.g.* aquatic birds). This is complicated by minimal public-health and medical infrastructure and large numbers of other potential reservoirs, such as pigs and domestic poultry commingling with humans in village settings. Avian influenza demonstrates the immediate need for international cooperation and interdisciplinary interventions for disease detection, control, and prevention. It also illustrates the need to engage local farmers in the development of sustainable strategies to identify suspect cases and prevent the commingling of domestic and wild-bird populations.

SUMMARY

We face some critical needs as we combat emerging diseases. We must understand the global consequences of moving animals and animal products around the world and assess the impact of an increasing human population on the environment. This combination sets the stage for potential mixing of microorganisms around the globe in contact with susceptible populations. The influenza epidemic of 1918–1919 killed 50 to 100 million people worldwide, but since the 1960s, many of us have had the luxury of forgetting about the enormous death toll brought by outbreaks of infectious diseases (Osterholm, 2005). Even today, however, we cannot disregard the possible catastrophic effects of currently emerging diseases.

To control emerging diseases requires early detection and intervention. The phenomenal speed in the diagnosis and identification of the SARS-CoV demonstrates how technologies have improved our response and mitigation efforts. These rapid diagnostic tests need to be incorporated in the field to shorten detection and response times. This is especially true for exotic animal diseases that can harm our domestic livestock. These tests could also be used to quickly identify exposed individuals for early treatment or isolation. Another important learning point from both SARS and avian influenza is that agricultural workers may often be the first to acquire these new or re-emerging diseases. Therefore, it is imperative to have adequate healthcare for workers. With healthcare, timely information needs to be collected by public-health personnel to also assess the population health of agricultural workers.

Our public-health and veterinary infrastructure needs to be improved. We must build the expertise, resources, and tools necessary for developing the capacity to respond to threats posed by vector-borne and zoonotic diseases (Smolinski *et al.*, 2003). Our universities need to train more medical entomologists, vector ecologists, mammologists and ornithologists who have a thorough understanding of the interactions among human, animal, and ecosystem health. There is a need to develop interdisciplinary infectious-

disease centers for training, research, diagnostic systems and data sharing. Furthermore, public-health authorities should look beyond traditional disciplines and training when hiring new epidemiologists and microbiologists. These and other recommendations have been clearly outlined (Smolinski *et al.*, 2003; NRC, 2005).

In the 19th century, Rudolf Virchow stated that animal and human health are inextricably intertwined. Our common environment is where this weaving of lives takes place, hence, we must guard the health of our ecosystems. Recent examples include decreasing wetlands and the subsequent congregation of waterfowl in smaller areas, resulting in outbreaks such as avian influenza and Newcastle disease. Deforestation and the greater interaction of wildlife with domestic animals and humans are likely factors for the emergence of novel viruses such as hendra, lyssavirus, and Nipah (Parashar *et al.*, 2000). Conversely, reforestation and suburbanization are likely contributing factors for the emergence of Lyme disease in the northeastern portion of the United States (LoGiudice *et al.*, 2003). Dramatic weather events have also been linked to disease emergence. This was documented with the outbreaks of Rift Valley fever among ruminants and people in East Africa and the Arabian peninsula (CDC 1998, 2000), following periods of above-normal precipitation and subsequent increases in mosquitoes. We can be sure that diseases will continue to emerge, and the complex relationship between animals, plants, and humans will require the interaction and cooperation of a broader range of scientists and medical professionals. The time to train them is now.

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