

# Climate change and waterborne and vector-borne disease

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## 1. SUMMARY

This paper considers the potential impact on human health from waterborne and vector-borne infections. It concentrates on the impact of two possible changes to climate; increased frequency of heavy rainfall events, with associated flooding and increased temperature. Flooding is associated with increased risk of infection in developing nations but not in the West unless water sources are compromised. There have been numerous reported outbreaks that followed flooding that led to contamination of underground sources of drinking water. Heavy rainfall also leads to deterioration in the quality of surface waters that could adversely affect the health of those engaged in recreational water contact. It is also concluded that there may be an increase in the number of cyanobacterial blooms because of a combination of increased nutrient concentrations and water temperature. It is considered unlikely that climate change will lead to an increase in disease linked to mains drinking water, although private supplies would be at risk from increased heavy rainfall events. Although increased temperature could lead to climatic conditions favourable to increases in certain vector-borne diseases such as malaria, the infrastructure in the UK would prevent the indigenous spread of malaria.

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## 2. INTRODUCTION

Despite considerable research activity over a number of years, it must be admitted that the actual impacts of climate change on public health are still far from clear. In part, this uncertainty reflects difficulties in predicting the local effects of global changes in climate. Even if we were certain about future local climates, there is still uncertainty over what impact these climates would have on health. The focus of this paper is on the potential impacts of climate change on both waterborne and vector-borne diseases. In addressing these problems we shall first describe the current burden of disease through these transmission pathways, then go on to discuss what is already known about the impact of climate on their epidemiology and finally we shall attempt to predict what might be the impact of climate change on future epidemiology.

Given the uncertainty in the predictions over how the climate for UK will change, this review can only address the possible effects on health in a general rather than a specific way. In particular, this paper will address the potential health impacts of two likely changes in climate; an increase in the frequency of heavy rainfall events along with associated floods and increase in temperature. Furthermore, the discussion will be restricted to microbial disease. For a wider discussion of the health impacts of climate change on human health, the Department of Health has recently published an expert review for comment (Department of Health 2001).

### 3. WATERBORNE DISEASE

#### 3.1 Current situation

Globally diseases waterborne and sanitation-related infections are one of the major contributors to diseases burden and mortality (Prüss and Havelaar 2001). The main

burden of disease in this category falls on the poorest societies and on children under 5 years. Many different viral, bacterial and parasitic diseases have been associated with waterborne transmission (Hunter 1997). Some diseases have been spread through drinking water and others by recreational water contact (Table 1). The

**Table 1** Microbial pathogens linked to drinking water or recreational water contact

| Organism  | Disease                           | Transmission   | Clinical features  |
|---|-----------------------------------|--|--|
| <b>Helminths</b>                                    |                                   |  |  |
| <i>Schistosoma</i> spp.                             | Schistosomiasis                   | Contact with surface water infected with free swimming cercariae     | Urinary and intestinal damage.<br>Bladder cancer   |
| <i>Dracunculus medinensis</i>                       | Dracunculiasis                    | Drinking water   | Painful ulcers on lower limbs and feet   |
| <b>Protozoa</b>                                     |                                   |  |  |
| <i>Giardia duodenalis</i>                           | Giardiasis                        | Faecal oral spread through drinking water or recreational water      | Diarrhoea and abdominal pain, weight loss and failure to thrive                                  |
| <i>Cryptosporidium parvum</i>                       | Cryptosporidiosis                 | Faecal oral spread through drinking water or recreational water      | Diarrhoea often prolonged  |
| <i>Cyclospora cayetanensis</i>                      | Cyclosporiasis                    | Faecal oral spread through drinking water                            | Diarrhoea and abdominal pain, weight loss and failure to thrive                                  |
| <i>Entamoeba histolytica</i>                        | Amebiasis                         | Faecal oral spread through drinking water                            | Diarrhoea, may be severe<br>dysentery  |
| <i>Toxoplasma gondii</i>                            | Toxoplasmosis                     | Drinking water contaminated by feline animals                        | Glandular fever, foetal damage in pregnant women   |
| Free-living amoebae                                 | Amoebic meningoencephalitis       | Aspiration of infected surface water into nose                       | Fatal encephalitis   |
| <b>Algae</b>  |                                   |  |  |
| Cyanobacteria                                       | Various                           | Toxins in drinking water or direct contact with surface water blooms | Dermatitis, hepatitis, respiratory symptoms, potentially fatal                                   |
| <i>Pfiesteria piscicida</i>                         | Estuary-associated syndrome       | Toxins in water  | Respiratory and eye irritation, deficiencies in learning and memory and acute confusional states |
| <b>Bacteria</b>                                     |                                   |  |  |
| <i>Vibrio cholerae</i>                              | Cholera                           | Drinking water   | Watery diarrhoea, may be severe  |
| <i>Salmonella</i> spp.                              | Salmonellosis                     | Occasional outbreaks with drinking water                             | Diarrhoea, colicky abdominal pain and fever  |
| <i>Salmonella typhi</i>                             | Typhoid                           | Drinking water   | Fever, malaise and abdominal pain with high mortality  |
| <i>Shigella</i> spp.                                | Shigellosis (bacillary dysentery) | Both drinking and recreational water                                 | Diarrhoea frequently with blood loss   |
| <i>Campylobacter</i> spp.                           | Campylobacteriosis                | Both drinking and recreational water                                 | Diarrhoea frequently with blood loss   |
| Enterotoxigenic <i>E. coli</i>                      |                                   | Drinking water   | Watery diarrhoea   |
| Enterohaemorrhagic <i>E. coli</i>                   |                                   | Drinking water and recreational water contact                        | Bloody diarrhoea and haemolytic uraemic syndrome in children                                     |
| <i>Yersinia</i> spp.                                | Yersiniosis                       | Drinking water   | Fever, diarrhoea and abdominal pain  |
| <i>Francisella tularensis</i>                       | Tularaemia                        | Drinking water   | Typhoid-like or mucocutaneous with suppurative skin lesions                                      |
| <i>Helicobacter pylori</i>                          |                                   | Drinking water   | Gastritis that can progress to gastric cancer  |
| <i>Mycobacteria</i> spp. not <i>M. tuberculosis</i> | Varies                            | Potable water systems in hospitals, some recreation                  | Varies, includes respiratory disease, wound infections, skin disease                             |

**Table 1** (Contd.)

| Organism                            | Disease                          | Transmission                            | Clinical features      |
|-------------------------------------|----------------------------------|---|------------------------|
| Viruses                             |                                  |   |                        |
| Hepatitis A and Hepatitis E viruses | Viral hepatitis                  | Drinking and recreational water contact | Hepatitis              |
| Various, esp. Norwalk-like viruses  | Viral gastroenteritis            | Drinking and recreational water contact | Vomiting and diarrhoea |
| Enteroviruses                       | Various, including poliomyelitis | Drinking and recreational water contact | Various                |

distribution of many waterborne pathogens varies substantially from one country to another. Some pathogens such as *Vibrio cholerae*, Hepatitis E virus and schistosomiasis are restricted to certain tropical countries, others, such as cryptosporidiosis and campylobacteriosis, are probably widespread. For some pathogens apparent differences in distribution may be due to different availability of diagnostic laboratories or national infectious disease surveillance schemes.

The evidence about the burden of waterborne disease in the UK comes predominantly from recording of outbreaks. In the UK there has been some 65 recorded outbreaks of infection linked to water affecting 4112 people during the years 1991–2000. Of these outbreaks, 25 were associated with public water supplies, 16 with private water supplies, 23 with swimming pools and one with recreational contact with surface waters (Table 2) (Stanwell-Smith *et al.* 2002). By far the commonest reported pathogen was *Cryptosporidium*, although *Campylobacter* was the commonest cause of outbreaks associated with private water supplies.

In the US for the period 1991 to 1998 there were 230 waterborne outbreaks reported affecting an estimated 443 000 people (Craun *et al.* 2002). However, most of the estimated case numbers comes from a single outbreak in Milwaukee and the method of estimating cases numbers has recently been criticised for substantially overestimating the size by at least 10-fold (Hunter and Syed 2001). Of 126 drinking-water outbreaks, 109 were reported in public water systems and 17 in individual water systems. There were also 104 outbreaks were associated with recreational water activities, usually swimming.

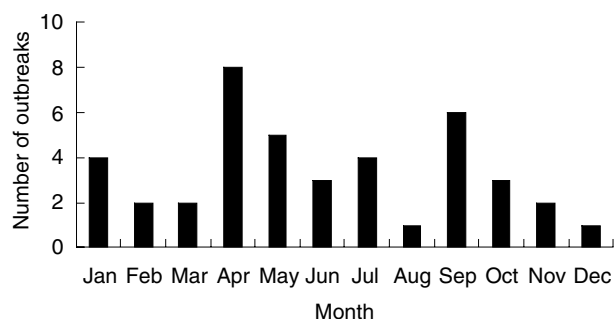
### 3.2 Impact of climate on epidemiology of waterborne disease

The current evidence of the impact of climate on the epidemiology of waterborne disease is considered under three headings; the impact of heavy rainfall events, the impact of flooding and the impact of increased temperature.

**3.2.1 Heavy rainfall events.** In Fig. 1 the month of start of outbreaks associated with water supplies in England and Wales for the last decade is shown. Although, there are probably too few outbreaks to be certain, there does appear to be an increase in Spring (April and May) and again in September. These months do not seem to be those that are typically associated with high rainfall. However, many investigations of drinking water-associated disease have noted that particularly heavy rainfall preceded the start of the outbreak (Smith *et al.* 1989; Joseph *et al.* 1991; Bridgman *et al.* 1995; Atherton *et al.* 1995; Willocks *et al.* 1998; Miettinen *et al.* 2001). Perhaps the most dramatic waterborne outbreak following heavy rainfall and associated

**Table 2** Outbreaks of waterborne disease in England and Wales from 1991 to 2000

| Cause   | Number of outbreaks | Number of cases |
|---|---------------------|-----------------|
| Public water supplies                                 |                     |                 |
| <i>Cryptosporidium</i>                                | 23                  | 2837            |
| <i>Campylobacter</i>                                  | 1                   | 281             |
| Gastroenteritis                                       | 1                   | 229             |
| Total   | 25                  | 3347            |
| Private water supplies                                |                     |                 |
| <i>Campylobacter</i>                                  | 8                   | 178             |
| Mixed <i>Campylobacter</i> and <i>Cryptosporidium</i> | 1                   | 43              |
| <i>Cryptosporidium</i>                                | 3                   | 74              |
| Gastroenteritis                                       | 2                   | 81              |
| <i>Giardia</i>  | 1                   | 31              |
| <i>E. coli</i> O157                                   | 1                   | 14              |
| Total   | 16                  | 421             |
| Recreational contact with surface water               |                     |                 |
| Viral gastroenteritis                                 | 1                   | 7               |
| Swimming pool contact                                 |                     |                 |
| <i>Cryptosporidium</i>                                | 23                  | 337             |
| Overall total   | 65                  | 4112            |



**Fig. 1** Seasonal distributions of outbreaks associated with both private and public drinking water supplies from 1991 to 2000

flooding was the Walkerton outbreak of *Escherichia coli* O157:H7 in Canada which affected over 1000 people of whom 65 were admitted to hospital and six died (Anon 2000).

A recent American investigated the relationship between outbreaks waterborne disease and heavy rainfall events in a more systematic fashion (Curriero *et al.* 2001). They examined records of 548 outbreaks in the US that had been reported from 1948 through to 1994 and linked these outbreaks to rainfall data. Some 51% of all outbreaks were preceded by an extreme rainfall event.

It is not too difficult to understand the reason why heavy rainfall events can help generate impact that heavy rainfall events are associated with outbreaks of waterborne disease. Heavy rainfall may leads to changes in the direction of flow of water systems and flow through channels that would not normally occur. For example, in one outbreak of cryptosporidiosis associated with a borehole extracted groundwater, it was thought that the heavy rainfall led to water running across the surface of fields where cattle were grazing (Bridgman *et al.* 1995). The water (and cattle faeces) then pooled around the head of the borehole and leaked into the water supply. For surface water sources, heavy rainfall can lead to overflow of storm drains that may be combined with the sewage system. This can then allow substantial amounts of faecally polluted water into rivers. Surface water turbidity can also increase dramatically during heavy rainfall events and this can cause additional stress on inadequate water treatment systems.

Further evidence of the impact of heavy rainfall on the epidemiology of enteric pathogen comes from studies of the presence of various organisms in water. For example, there is a correlation between rainfall and the likelihood of detecting *Giardia* or *Cryptosporidium* oocysts in river water (Atherbolt *et al.* 1998) and pathogenic enteric viruses in water (Miossec *et al.* 2000). Heavy rainfall leads to storm water runoff into surface water sources, which has high counts of indicator bacteria as well as potential pathogens (Doran and Linn 1979; O'Shea and Field 1992). Heavy

rainfall is also associated with high counts of indicator bacteria in river waters (Tunncliffe and Brickler 1984) and marine waters (Crowther *et al.* 2001). Given the well described relationships between counts of indicator organisms in surface water and the subsequent risk of predominantly gastrointestinal illness (Kay *et al.* 1994) it should be expected that people swimming in untreated surface waters after heavy rain would be at increased risk of illness.

There is also some evidence that heavy rainfall events may be followed by coliform re-growth in water distribution systems, presumably because of increased nutrients in water (LeChevallier *et al.* 1991). However, the health significance of this observation is unclear as non-faecal coliforms in drinking water do not appear to be associated with disease in the community (Edberg *et al.* 1986; Zmirou *et al.* 1987; Hellard *et al.* 2001).

A further issue related to heavy rainfall events is the additional nutrient input into water bodies that accompany heavy rainfall. When other conditions are appropriate, this can lead to the rapid growth and blooming of various planktonic species. This is considered further below.

**3.2.2 Flooding.** Flooding may follow heavy rainfall, although floods can also follow from tidal surges and rapid snow melt. Flooding of low-lying lands caused by rise in sea level secondary to melting of the ice caps is also likely to happen. Of all the types of natural disaster world-wide, flooding is the commonest and can cause enormous damage and financial loss (Malilay 1997; Greenough *et al.* 2001). No one in the UK will be unaware of the floods of recent years and the suggestion that flood events are likely to increase in future.

The health effects of flooding can be divided into those associated with the acute event and those arising after the flood has resolved. The main acute threat to health is drowning, although in the UK deaths from drowning as a result floods are few. On the other hand some 15 000 people are killed each year by floods in Bangladesh (Malilay 1997).

Although most concerns about subsequent health risk tends to focus on infectious disease, the evidence is relatively scant, at least for developed countries. The impact that heavy rainfall on the genesis of outbreaks of disease associated with drinking water systems has already been discussed. In general these outbreaks have been associated with flows of contaminated water into the groundwater or the interference with the effectiveness of water treatment. An additional outbreak was one of *Acanthamoeba* keratitis in Iowa which followed flooding that inundated a water treatment works (Meier *et al.* 1998).

For developing nations there is evidence of outbreaks following floods. Outbreaks of leptospirosis in Rio de Janeiro (Barcellos and Sabroza 2001) and in the Philippines (Easton 1999) have followed floods. Hepatitis E, malaria and

diarrhoeal disease have followed floods in Khartoum (Homeida *et al.* 1988; Novelli *et al.* 1988; Shears 1988; McCarthy 1994). Both acute diarrhoea and acute respiratory disease increased in Nicaragua following Hurricane Mitch and the associated flooding (Campanella 1999). By contrast epidemics do not tend to follow floods in developed countries (Bissell 1983; Aavitsland *et al.* 1996; Malilay 1997; Greenough *et al.* 2001).

It would seem to be the case that when infrastructure is adequate, epidemics of infection do not follow flood events unless the flooding has directly compromised the security of the water supply.

**3.2.3 Temperature effects.** Probably the most obvious link between waterborne disease and increased temperature relates to the blooms of various planktonic species that are directly or indirectly hazardous to human health. Several planktonic species have been implicated in causing disease in humans:

- Cyanobacteria (Blue-green algae) have been implicated in causing various clinical syndromes such as dermatitis, respiratory problems and hepatitis (Hunter 1998). Illness follows consumption of water containing toxins or contact with water or blooms during bathing.
- Dinoflagellates and diatoms are implicated in a variety of neurotoxic, diarrhetic and amnesic shell fish poisonings (Hungerford 2001). These are not directly waterborne diseases, although the shellfish accumulate toxins whilst filter feeding.
- *Pfiesteria piscicida* is a dinoflagellate which is responsible for a recently recognised syndrome known as 'estuary-associated syndrome'. This causes acute respiratory and eye irritation and has been associated with deficiencies in learning and memory and acute confusional states in people who have had contact with contaminated water or aerosols (Morris 1999).

A major factor in the appearance of blooms is the increase in nutrients in water bodies especially in inland freshwater lakes and estuaries (Reynolds 1984). However, most blooms occur during the summer months and relatively high water temperatures are necessary for algal growth and bloom formation (Maier and Dandy 1997; Jacoby *et al.* 2000; Saker and Griffiths 2001). There is a strong impression that algal blooms are becoming more common in Europe (Skulberg *et al.* 1984; Fastner *et al.* 1999).

The most evidence of the effect of temperature on risk from waterborne disease is in relation to cholera (Colwell 1996). There is now good evidence that *V. cholerae* survives in marine waters in a viable but non-cultural form that seems to be associated with algae and plankton (Islam *et al.* 1990; Colwell 1996). When temperatures rise, plankton bloom and, in appropriate areas, such blooms are followed

by increases in reported cases of cholera. For example, increases in sea-surface temperature as a result of El-Niño-events have been shown to predate increases in cholera incidence in both Asia and South America (Lobitz *et al.* 2000; Pascual *et al.* 2000; Speelman *et al.* 2000).

There has been some suggestion that higher water temperatures will lead to more prolonged survival of pathogens in the environment (Chief Medical Officer 2001). However, the evidence does not support this concern. Indeed, survival of enteric pathogens in water is usually reduced at higher temperatures. This has been shown to be the case for *Campylobacter* (Buswell *et al.* 1998; Thomas *et al.* 1999; Obiri-Danso *et al.* 2001), enterohemorrhagic *E. coli* O157:H7 (Wang and Doyle 1998; Rice and Johnson 2000), *V. cholerae* (but see above) (Okitsu *et al.* 1994) and enteroviruses (Hurst *et al.* 1989).

### 3.3 The future impact of global warming on waterborne disease in the UK

Making predictions of future events is always a dangerous pastime. Nevertheless, one can make at least some intelligent guesses about what the future may hold for waterborne disease.

It seems unlikely that global warming will have a major impact on the risk of disease associated with mains water supplies in the UK. The standards of water treatment and distribution within the UK is of a very high standard compared with many nations even elsewhere in Europe. Although outbreaks of waterborne disease do still occur, they are nearly all due to *Cryptosporidium* and are associated with supplies that are still inadequately treated. Most of these high-risk supplies will have been closed or had new filtration plants installed by about 2005. Provided that the UK does not suffer serious economic collapse it is difficult to see how global warming could have a major impact on risk of waterborne disease associated with mains drinking water.

On the other hand, there are still many private supplies in use in the UK and these are frequently poorly treated. The microbiological quality of such supplies is generally very poor and can deteriorate markedly after heavy rainfall (Shepherd and Wyn-Jones 1997; Fewtrell *et al.* 1998; Rutter *et al.* 2000). If heavy rainfall events are to become more common, then the safety of private supplies (already poor) will deteriorate even further.

Heavy rainfall events are more likely to lead to marked decline in microbiological quality of inland and marine recreational waters as a result of heavy runoff. Furthermore, increased temperatures may add to the continued rise in problems due to algal blooms in Europe. Both these mechanisms would increase the probability of disease from recreational use of waters in and around the UK. However,

the epidemiological evidence of infectious disease associated with recreational water contact is that it is generally very mild and not likely to cause significant disease burden in the population.

## 4. VECTOR-BORNE DISEASE

### 4.1 Current situation

Vector-borne diseases are those diseases that are spread by insect vectors. The insects act as an essential stage in the

transmission of the infection from one person to another or from animal hosts to humans. There are a large number of viral, rickettsial, bacterial and parasitic diseases that can be transmitted by insect vectors (Cook 1996) (Table 3). Various vector-borne are amongst some of the major microbial causes of morbidity and mortality in the World today. Malaria alone causes some one million deaths and 273 million cases worldwide each year (Rogers and Randolph 2000). However, with the exception of Lyme disease, vector-borne diseases do not currently cause problems for public health in the UK.

**Table 3** Some examples of vector-borne diseases of clinical importance

| Disease                          | Pathogen                      | Vector          | Geographical distribution  | Clinical effects                                |
|----------------------------------|-------------------------------|-----------------|--|---|
| <b>Viruses</b>                   |                               |                 |  |   |
| Dengue                           | Flavivirus                    | Mosquito        | Africa, Caribbean, Pacific, Far East                               | Haemorrhagic fever                              |
| Japanese Encephalitis            | Flavivirus                    | Mosquito        | Japan, Far East  | Encephalitis                                    |
| West Nile                        | Flavivirus                    | Mosquito        | Africa, India, Europe and North America                            | Encephalitis                                    |
| Murray River encephalitis        | Flavivirus                    | Mosquito        | Australia, New Guinea  | Encephalitis                                    |
| St Louis encephalitis            | Flavivirus                    | Mosquito        | America  | Encephalitis                                    |
| Yellow fever                     | Flavivirus                    | Mosquito        | Africa, South and Central America                                  | Hepatitis & Haemorrhagic fever                  |
| Eastern Equine Encephalitis      | Alphavirus                    | Mosquito        | North America  | Encephalitis                                    |
| Crimean-Congo haemorrhagic fever | Nairovirus                    | Ixodic tick     | Europe, Africa, Middle East, Central Asia                          | Haemorrhagic fever                              |
| Tick-borne encephalitis          | Flavivirus                    | Ixodic tick     | Former USSR, Europe  | Encephalitis                                    |
| Rift Valley Fever                | Phlebovirus                   | Mosquito        | Africa   | Haemorrhagic fever                              |
| <b>Rickettsia</b>                |                               |                 |  |   |
| Murine typhus                    | <i>Rickettsia typhi</i>       | Flea            | Tropical countries   | Typhus  |
| Rocky Mountain Spotted fever     | <i>R. rickettsii</i>          | Ticks           | USA  | Spotted fever                                   |
| Boutonneuse fever                | <i>R. conorii</i>             | Tick            | Africa, Mediterranean, Middle East                                 | Spotted fever                                   |
| <b>Bacteria</b>                  |                               |                 |  |   |
| Bartonellosis                    | <i>B. bacilliformis</i>       | Sandflies       | Western Slopes of the Andes  | Oroya fever                                     |
| Plague                           | <i>Yersinia pestis</i>        | Flea            | Africa, Asia, South America, USA                                   | Bubonic, pneumonic or septicaemic plague        |
| Lyme disease                     | <i>Borrelia burgdorferi</i>   | Ticks           | Europe, North America  | Arthritis                                       |
| Relapsing fever                  | <i>B. recurrentis</i>         | Lice            | Ethiopia, Burundi, Peru, Bolivia, North Africa, India, Asia, China | Severe relapsing fever with high mortality      |
| Relapsing fever                  | e.g. <i>B. duttoni</i>        | Ticks           | East, Central & South Africa                                       | Less severe relapsing fever                     |
| <b>Protozoal parasites</b>       |                               |                 |  |   |
| Malaria                          | <i>Plasmodium</i> spp.        | Mosquitos       | Widespread in tropics  | Febrile illness with high mortality             |
| African trypanosomiasis          | <i>Trypanosoma brucei</i>     | Tsetse flies    | Africa   | Sleeping sickness                               |
| American trypanosomiasis         | <i>T. cruzi</i>               | Triatomine bugs | Central and South America  | Chagas disease                                  |
| Leishmaniasis                    | <i>Leishmania</i> spp.        | Sandflies       | Africa, Central & South America                                    | Cutaneous and mucosal lesions, visceral disease |
| <b>Helminths</b>                 |                               |                 |  |   |
| Lymphatic filariasis             | eg. <i>Wucheria bancrofti</i> | Mosquito        | Tropics  | Elephantiasis                                   |
| Onchocerciasis                   | <i>Onchocerca volvulus</i>    | Blackflies      | Africa, Central & South America                                    | Dermatitis, Blindness                           |
| Loaiasis                         | <i>Loa loa</i>                | Tabanid flies   | West & Central Africa  | Calabar swellings                               |

## 4.2 Impact of climate on epidemiology of vector-borne disease

The geographical distribution of vector-borne diseases is influenced by the geographical distribution of both vertebrate host (where one exists) and the distribution of the vector.

**4.2.1 Temperature effects.** Temperature can affect both the distribution of the vector and the effectiveness of pathogen transmission through the vector. Gubler *et al.* (2001) list a range of possible mechanisms whereby changes in temperature impact on the risk of transmission of vector-borne disease:

- Increase or decrease in survival of vector
- Changes in rate of vector population growth
- Changes in feeding behaviour
- Changes in susceptibility of vector to pathogens
- Changes in incubation period of pathogen
- Changes in seasonality of vector activity
- Changes in seasonality of pathogen transmission

There is also good epidemiological evidence of disease risk in humans linked to climate variability. Most of this evidence comes from studies of malaria. For example, there is evidence that malaria prevalence in both South America and Asia varies in relation to El-Niño-events (Bouma and van der Kaay 1996; Bouma *et al.* 1996, 1997; Bouma and Dye 1997). This relationship between El-Niño and increased malaria risk is partly due to increased temperature and partly due to increased rainfall leading to increased mosquito breeding sites because of surface water collections. Climate has also been shown to have a direct effect on the risk of outbreaks of St Louis encephalitis (SLE). SLE is generally restricted to areas south of the 20°C June isotherm, although it has spread northerly in particularly warm years (Hess *et al.* 1963; Shope 1980). Furthermore outbreaks of SLE tend to follow periods of hot weather where the temperature exceeds 30°C for seven consecutive days (Murray *et al.* 1985; Monath and Tsai 1987). Evidence for other vector-borne pathogens is less clear. Indeed, a study of reported cases of Lyme disease in the UK showed no correlation with mean summer temperatures (Cannell *et al.* 1999).

**4.2.2 Rainfall effects.** Gubler *et al.* (2001) list a range of possible mechanisms whereby rainfall can impact on the risk of transmission of vector-borne disease:

- Increased surface water can provide breeding sites for vectors
- Low rainfall can also increase breeding sites by slowing river flow
- Increased rain can increase vegetation and allow expansion in population of vertebrate host

- Flooding may eliminate habitat for both vectors and vertebrate hosts
- Flooding may force vertebrate hosts into closer contact with humans

The link between El-Niño and malaria has already been discussed. This link is due, in part, to the increased amount of surface water providing breeding sites for mosquitoes (Bouma and van der Kaay 1996; Bouma *et al.* 1996, 1997; Bouma and Dye 1997). Decreased rainfall has been shown to be associated with epidemics of SLE when the vector, *Culex pipiens*, starts to breed in urban drainage systems (Mitchell *et al.* 1980). Heavy rains have also been shown to precipitate epidemics of SLE in Florida as the rainfall event stimulates blood-feeding by *Culex nigripalpus* (Day and Curtis 1989).

## 4.3 The future impact of global warming on vector-borne disease in the UK

Most concern about the potential impact of climate change on vector-borne disease has focused on the potential reintroduction of malaria to the UK. Indeed Malaria was endemic to the UK until the early part of the last century (Dobson 1994). Climate scenarios suggest that within the next few decades, summer temperatures could become conducive to the spread of vivax malaria in some areas of England (Rogers *et al.* 2001). However, climate alone is not a sufficient requirement for the spread of malaria. There has to be a pool of parasitaemic individuals to act as the reservoir of infection. There are many regions that have adequate climate for transmission of malaria but do not experience indigenous malaria, such as the United States and Southern Europe. The absence of malaria in these regions is due to adequate public health and medical infrastructure where mosquito breeding is controlled and infected individuals diagnosed and treated. It is difficult to envisage a situation where the public health infrastructure would breakdown and indigenous malaria return to the UK without a significant economic collapse. One possible exception to this is the potential for small localised outbreaks of 'airport malaria' whereby infected mosquitoes or individuals arrive in the UK from infected areas abroad. Such outbreaks are already well documented (Guillet *et al.* 1998; Van den Ende *et al.* 1998). Whether a warming climate will increase the risk of such events occurring remains to be seen.

With the current knowledge about Lyme Disease, it is not possible to predict what if any effect climate change will have on its epidemiology.

## 5. CONCLUSIONS

With the current available evidence and scenarios for climate change in the UK it would appear that the public health effects of climate change, at least as far as water- and

vector-borne disease is concerned, are likely to be relatively negligible when compared to other public health concerns. However, we cannot afford to be complacent. The recent introduction of West Nile encephalitis into New York and elsewhere in the United States serve as a warning of other, unanticipated, threats of vector-borne disease (Epstein 2001). Such sudden emergence of infectious disease can rarely be predicted in advance. We do not know sufficient about the epidemiology of many infectious diseases to adequately predict the full impact of climate change. Furthermore, we have no idea how many potential pathogens are sharing this planet with us that are awaiting the right conditions for their chance at stardom by becoming an emerging infectious disease.

There is also another very important point to make. This review has concentrated on the potential impact of climate change on the UK. The adverse impact of any change is likely to be far more damaging on less wealthy societies. Adverse health consequences of climate change may be yet another example of the observation that often the adverse consequences of the lifestyles of the wealthy are felt, not by those enjoying the benefits of the those lifestyles but by the poor. In this author's view the West has a moral responsibility towards the poorest nations for the potential adverse impacts that they are likely to experience as a result of our inability to control our own excess energy consumption.

## 6. REFERENCES

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