## Calibrated response to emerging infections

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The current flu pandemic raises a public health policy question that could have been asked after the emergence of severe acute respiratory syndrome (SARS): what is the proper response to clinically mild or epidemiologically limited (small number) outbreaks caused by new viruses? Over the past four years, pandemic preparations have focused on responding to worst case scenarios. As a result, officials responded to the H1N1 outbreak as an unfolding disaster. Measures were taken that in hindsight may be seen as alarmist, overly restrictive, or even unjustified. Assumptions about the nature of emerging infections along with advanced laboratory surveillance have changed the way we understand epidemics and we need a new framework for thinking about epidemic disease.

**Predictions that missed the mark**

Before the arrival of novel A/H1N1 virus, pandemics were said to occur when a new subtype of influenza virus to which humans have no immunity enters the population, begins spreading widely, and causes severe illness. Reference was often made to the catastrophic pandemic of 1918 and the ongoing threat of highly pathogenic avian influenza H5N1 that has killed over half of the 456 people with recorded infection since 1997. Without proper preparation, “The loss of human life even in a mild pandemic will be devastating, and the cost of a world economy in shambles for several years can only be imagined,” one highly cited article concluded in 2005. The large sums of public money spent on pandemic preparedness (over $7bn (£4bn; €5bn in the US) underlined the seriousness of the threat, and often repeated phrases such as “not a question of IF a pandemic will happen, but WHEN” characterised the next flu pandemic as a high probability, high consequence event.

But the 2009 pandemic, taken as a whole, bears little resemblance to the forecasted pandemic. Pandemic A/H1N1 virus is not a new subtype but the same subtype as seasonal A/H1N1 that has been circulating since 1977. Furthermore, a substantial portion of the population may have immunity. The US Centers for Disease Control and Prevention (CDC) found that 33% of those aged over 60 had cross reactive antibody to novel A/H1N1, which may explain why cases have been rare in elderly people.

There is also far less certainty today regarding the severity of the threat of pandemic flu. Experts are unsure that the 2009 pandemic—which the World Health Organization previously characterises as moderate—will be any worse than seasonal flu. Since the emergence of novel A/H1N1, descriptions of pandemic flu (both its causes and its effect) have changed to such a degree that the difference between seasonal flu and pandemic flu is now unclear (table). WHO, for example, for years defined pandemics as outbreaks causing “enormous numbers of deaths and illness,” but in early May, removed this phrase from the definition.11

On 29 April 2009, one week after news of the outbreak first surfaced, WHO declared a phase 5 pandemic alert (the highest threat level short of global pandemic), urging all countries to “immediately activate their pandemic preparedness plans.”16 Epidemiological information at this time was mixed, suggesting a severe disease in Mexico but mild everywhere else. Actions were thus taken in an environment of high public attention and low scientific certainty.17 Some countries erected port of entry quarantines. Others advised against non-essential travel to affected areas. Some closed schools and businesses. Many held daily press briefings. The wisdom of many of these actions, particularly in response to what has largely been a clinically mild illness, will undoubtedly be debated in the future. What these actions more clearly show, however, is that the public health response to, as well as impact and social experience of a pandemic, is heavily influenced by longstanding planning assumptions about the nature of pandemics as disaster scenarios.

**Laboratory surveillance drives concern**

One assumption concerned the importance of laboratory surveillance data to help identify and characterise cases, especially during the initial phases of a pandemic.19 This intensive use of the laboratory to understand the epidemiology of an epic epidemic disease is a product of our time.
During the 1918 pandemic, although investigators looked hard for the cause, no distinction was made between pandemic influenza and seasonal influenza as is done today. Influenza was simply influenza—or what today would probably be called influenza-like illness—and diagnosed on clinical grounds.

Much has changed since then. When researchers at the CDC reported the first two cases of A/H1N1 swine flu on 21 April 2009 it was not the clinical illness that worried them—both patients had recovered uneventfully by the time of the report—but the fact that human to human transmission was suspected in two laboratory confirmed cases of novel influenza virus infection.20 On 26 April, with 20 cases and no deaths in the US, the Department of Health and Human Services declared a nationwide public health emergency.21 The subsequent increase in laboratory testing was unprecedented (fig 1).

The sudden emphasis on laboratory testing for H1N1 in the first weeks of the outbreak, particularly in the US, produced what I call concern bias, in which concern and anxiety may drive events more than the disease itself. Concern bias confounds the interpretation of data in important ways. The rapid increase in virological testing amplified the perceived prevalence of A/H1N1 and simultaneously minimised the role other agents may have played in causing the same symptoms. After the declaration of a public health emergency, the percentage of respiratory specimens testing positive for influenza viruses increased for eight consecutive weeks to a peak of 40% (fig 1). This increase, however, may only in part reflect a true increase in prevalence of influenza. It may also be due to behavioural changes in the way respiratory specimens were taken, tested, and reported.22

Laboratories were overwhelmed with a large volume of respiratory specimens, often from patients who under ordinary circumstances would not have had a specimen taken (an extension of the “worried well” effect).23 The early screening out of samples unlikely to be A/H1N1 positive (for example, specimens positive for influenza B by rapid diagnostic testing at the bedside) is one way to reduce the workload at more sophisticated laboratories focused on confirmatory testing. However, this is likely to lead to overstatement of the proportion of influenza-like illness caused by A/H1N1 reported by laboratories. By contrast, in Sweden, of 79 travellers meeting the suspected novel H1N1 case definition (flu-like symptoms and recent travel to the US or Mexico) between 24 April and 10 June, only four had A/H1N1 infection. Non-influenza viruses were diagnosed in 40 samples, and 32 had unknown cause.24

As cause can affect treatment decisions, timely laboratory surveillance is essential. The apparent discordance between Swedish and US laboratory data suggests that without ongoing randomised sampling, it will be difficult to understand the effect of any single aetiological agent that causes clinically non-specific symptoms, such as influenza-like illness.

The high concern also makes it difficult to determine whether this epidemic revealed itself or whether its presence came to light only because of heightened awareness triggered by official announcements. During 19-25 April—nearly three weeks after the first two US cases and when the virus was presumably spreading—respiratory specimen testing was tapering off at laboratories around the country (fig 1). But in the week after the emergency declaration on 26 April, reports increased nearly sevenfold.

Large and geographically dispersed surveillance systems allow us to see more than ever. Whether they do us more good than harm, however, depends on the future course of the event.

**Calibrating the response to the threat**

If the 2009 influenza pandemic turns severe, far exceeding the impact of seasonal influenza, early and enhanced surveillance may prove to have bought critical time to prepare a vaccine that could reduce morbidity and mortality. The negative effect on the pork and travel industries, the discrimination some felt for the “crime” of catching a new disease, the mandatory isolation of uninfected people, and the substantial public money invested into pandemic preparations will probably be said to, on balance, have been fair better than being caught unprepared for a severe pandemic.

But if this pandemic does not increase in severity, it may signal the need to reassess both the risk assessment and risk management strategies towards emerging infectious diseases. The SARS outbreak showed that large numbers of infected people are not necessary to generate concern and fear over disease. The SARS virus is known to have affected only 8096 people globally, but the fear of infection, involuntary quarantine, travel restrictions and subsequent political antagonisms, and at least $18bn in losses were felt by far more. It was not the virus but the response to it that caused these social and economic harms.

Future responses to infectious disease may benefit from a risk assessment that broadly conceives of four types of threat based on the disease’s distribution and clinical severity (fig 2). Infectious diseases, whether caused by new or old pathogens, may infect few people or they may infect many (distribution). Furthermore, these pathogens can produce a clinical illness of variable severity, from mostly mild (or even asymptomatic) illness at one extreme to mostly severe illness at the other. Distribution
and severity are independent variables and together produce a matrix of four possible impacts. A single, one size fits all public health strategy cannot respond to the vastly different challenges these four clinical-epidemiological combinations present.

The commonality between the SARS epidemic and the present flu pandemic (at least so far) is that both were responded to with a public health strategy that may be more suitable to an epidemic of severe disease infecting many people (type 1). But SARS (which killed around 10% of infected people) was a type 2 epidemic (infecting few, mostly severe disease), and the H1N1 pandemic may prove to be type 3 (affecting many, mostly mild). Recent historical evidence suggests that most new viruses do not have constituted type 1 threats. While the 1918 pandemic surely qualifies as type 1, the 1957 and 1968 pandemics do not. Most people did not even notice the 1968 pandemic, and the recorded mortality in both pandemics was similar to that in contemporary non-pandemic influenza seasons. Despite this, pandemic preparedness strategies have largely considered only type 1 (catastrophic) epidemics. Public health responses not calibrated to the potential impact of new influenza A (H1N1) virus after vaccination with seasonal influenza vaccine. MMWR Mortal Wkly Rep 2009;58:521-4.


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See RESEARCH, pp 618, 619

ANSWERS TO ENDGAMES, p 639. For long answers use advanced search at bmj.com and enter question details

STATISTICAL QUESTION

Intention to treat analyses

C

Kasyer-Fleisher rings, visible as a greenish ring at the outer corneal surface (arrow)

PICTURE QUIZ

Eye sign in an 18 year old man with psychosis

1. The eye sign shown is Kayser-Fleisher rings—greenish discoloration at the outer corneal circumference. This abnormality was named after ophthalmologists Bernhard Kayser and Bruno Fleischer, who described the sign independently in the early 1900s. The rings were later recognised to be copper deposits and diagnostic of Wilson’s disease.

2. The combination of psychosis, extrapyramidal features (dyskinesia), and Kayser-Fleisher rings is classic for Wilson’s disease. A positive family history, low serum ceruloplasmin, high 24 hour urinary copper excretion, high liver copper, and the results of brain magnetic resonance imaging will support the diagnosis.

3. Untreated Wilson’s disease is fatal. Early diagnosis and lifelong copper chelation, with close clinical monitoring, are essential. The chance of neurological recovery is high.