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Analysis

Calibrated response to emerging infections

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WHO has revised its definition of pandemic flu in response to current experience with A/H1N1. **Peter Doshi** argues that our plans for pandemics need to take into account more than the worst case scenarios

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.^{1 2} 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 2005.³ 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”⁴ 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/H1N1,⁵ 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 presently characterises as moderate⁶—will be any worse than seasonal flu.^{7 8 9} 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 ↓).¹⁰ WHO, for example, for years defined pandemics as outbreaks causing “enormous numbers of deaths and illness,”¹⁰ but in early May, removed this phrase from the definition.¹¹

Aspect	Before A/H1N1	Since A/H1N1
One line summary	WHO 2003-9: “An influenza pandemic occurs when a new influenza virus appears against which the human population has no immunity, resulting in epidemics worldwide with enormous numbers of deaths and illness” ¹⁰	WHO: “An influenza pandemic may occur when a new influenza virus appears against which the human population has no immunity” ¹⁰
Virus and immunity	WHO 2005: “Most people will have no immunity to the pandemic virus” ¹	WHO: “The vulnerability of a population to a pandemic virus is related in part to the level of pre-existing immunity to the virus” ¹²
	US CDC 1997: “When antigenic shift occurs, the population does not have antibody protection against the virus” ¹³	US CDC: “Cross-reactive antibody [to A/H1N1] was detected in 6%-9% of those aged 18-64 years and in 33% of those aged >60 years” ⁵

Impact (health, social, economic)	WHO 2005: “Large numbers of deaths will occur . . . WHO has used a relatively conservative estimate—from 2 million to 7.4 million deaths . . . Economic and social disruption will be great” ¹	WHO: “H5N1 has conditioned the public to equate an influenza pandemic with very severe disease and high mortality. Such a disease pattern is by no means inevitable during a pandemic. On the contrary, it is exceptional” ¹⁴
	CDC 1997: “The hallmark of pandemic influenza is excess mortality” ¹³	CDC: “There are some pandemics that look very much like a bad flu season” ⁸
	Canada 2006: “An influenza pandemic results if many people around the world become ill and die from such a [new form of influenza] virus” ¹⁵	Canada: “An influenza pandemic does not necessarily cause more severe illness than seasonal influenza” ⁹

Changing views of pandemic flu, before and after emergence of influenza A/H1N1 virus

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 18} 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 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↓).

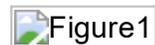


Fig 1 CDC data on numbers of respiratory specimens testing positive for influenza virus in US, January-June 2009 (www.cdc.gov/flu/weekly/weeklyarchives2008-2009/data/whoAllregt31.htm)

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 on.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↑). In the week after the emergency declaration on 26 April, however, 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 far 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^U). 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.



Fig 2 Proposed classification of impact of new infectious diseases

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 have not 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,²⁵ and the recorded mortality in both pandemics was similar to that in contemporary non-pandemic influenza seasons.²⁶ Despite this, pandemic preparedness strategies have largely considered only type 1 (catastrophic) epidemics. Public health responses not calibrated to the threat may be perceived as alarmist, eroding the public trust and resulting in people ignoring important warnings when serious epidemics do occur.

Advanced laboratory capabilities allow us to track epidemics at an unprecedented level of detail. Such information must not be allowed to obscure a broader perspective that places importance on the severity of the clinical illness most people experience and knowledge of how many people are being infected. The success of public health strategies today depends as much on technical expertise as it does on media relations and communications. Strategies that anticipate only type 1 epidemics carry the risk of doing more harm than they prevent when epidemiologically limited or clinically mild epidemics or pandemics occur.

Notes

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Footnotes

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