Severe acute respiratory syndrome (SARS): breathtaking progress

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Reports of a new severe respiratory disease, now defined as severe acute respiratory syndrome (SARS), began to emerge from Guangdong, in southern China, in late 2002. It burst upon international attention through an explosive outbreak in Hong Kong of what was soon to be defined as severe acute respiratory syndrome (SARS) in March 2003 (Lee et al., 2003). The appearance of cases throughout South-East Asia and in Toronto, the spread of SARS being accelerated by international air travel. A global emergency was declared by the World Health Organization, bringing together an international team of epidemiologists, public health physicians and microbiologists to study and contain the disease. This response has enabled the nature of the infectious agent to be identified, its mode of transmission to be established and diagnostic tests to be created rapidly (Drosten et al., 2003; Ksiazek et al., 2003). Furious efforts are being made to determine the optimal treatment regimen and to develop therapeutic agents and vaccines. Nonetheless, it is a paradox that, despite these technological achievements, we remain as vulnerable to this new agent as our ancestors were to previous plagues.

Abbreviations: HCW, health-care worker; SARS, severe acute respiratory syndrome.

Introduction

Reports of a new severe respiratory disease began to emerge from Guangdong, in southern China, in late 2002. It burst upon international attention through an explosive outbreak in Hong Kong of what was soon to be defined as severe acute respiratory syndrome (SARS) in March 2003 (Lee et al., 2003). The appearance of cases throughout South-East Asia and in Toronto was accelerated by international air travel (Poutanen et al., 2003). The World Health Organization declared a global emergency, bringing together an international team of epidemiologists, public health physicians and microbiologists to study and contain the disease. The coordinated response to SARS by the medical and scientific community has enabled the nature of the infectious agent to be identified, the mode of transmission established and diagnostic tests to be created rapidly (Drosten et al., 2003; Ksiazek et al., 2003). Furious efforts are being made to determine the optimal treatment regimen and to develop therapeutic agents and vaccines. Nonetheless, it is a paradox that, despite these technological achievements, we remain as vulnerable to this new agent as our ancestors were to previous plagues.

Early phase

SARS initially appeared as a small cluster of cases of viral pneumonia in rural Guangdong province in the People’s Republic of China, which, although out of the ordinary, was not considered exceptional. As far as can be ascertained, the first recognized case of SARS occurred in Fushan City and was reported on 16 October 2002. Three members of the index case’s family were also affected, but no medical staff. Small clusters of cases were noted over the next 3 months. The event that was to lead to the current global alert was the admission of a patient to a tertiary referral hospital in Guangzhou with presumed viral pneumonia who had been referred from Zongshan city outside Guangzhou. He was admitted to the 2nd Affiliated Hospital of Sun Yet-San Medical University for 2 days, then transferred to another hospital. Following his admission at the 2nd Affiliated Hospital, 28 medical and nursing staff fell ill with SARS, as well as the ambulance driver who transferred him. SARS then spread rapidly to five different hospitals and this has just been documented (Zhao et al., 2003). The early and high attack rate amongst hospital staff was to become a constant feature in all further outbreaks in different countries. The spread of SARS outside Guangdong province occurred when a nephrologist from Guangzhou travelled to Hong Kong on 21 February 2003. He had a 5-day history of respiratory symptoms and was admitted to hospital on the second day of his stay in Hong Kong. Unfortunately, while staying at a hotel in Hong Kong, he is presumed to have infected a couple from Toronto, Canada, a businessman who travelled to Hanoi, Vietnam, three people who travelled to Singapore and a number of Hong Kong residents. All of these people went on to initiate large outbreaks (Poutanen et al., 2003; Lee et al., 2003).

The SARS coronavirus

It has now been firmly established that the cause of SARS is a
coronavirus (Drosten et al., 2003; Ksiazek et al., 2003), and Koch’s postulates were fulfilled fully when experimental infection was achieved in macaques (Fouchier et al., 2003). The sequence contains nine novel viral proteins for which sequence analysis has predicted functions. Coronaviruses typically have narrow host ranges and are significant veterinary pathogens, causing severe bronchitis, peritonitis and gastroenteritis in different animal species. Human coronaviruses fall into groups 1 and 2 and are responsible for about one-third of upper-respiratory-tract infections (Rota et al. 2003). At least two examples of this virus have been fully sequenced, and this work demonstrates that the SARS agent is not related to any of the three previously described coronavirus groups (Rota et al., 2003; Marra et al., 2003).

It should not be a surprise that the SARS coronavirus is different from previous members of this genus, as the behaviour of the pathogen differs markedly from human coronaviruses described previously. The virus may have acquired a conserved motif, s2m, from avian infectious bronchitis virus but, apart from this, there is no evidence of any exchange of genetic material with non-Coronaviridae (Marra et al., 2003; Rota et al., 2003). This is consistent with the hypothesis that a previously unknown animal coronavirus has recently mutated, developing the ability to productively infect humans. There are some indications that the civet cat, eaten as a delicacy in southern China, was the source animal.

Not only did this virus differ genetically from other coronaviruses isolated in humans, but it behaves differently from other human coronaviruses and most other respiratory pathogens. The mortality rate appears to be very high (Tsang et al., 2003; Lee et al., 2003). Elderly patients with pre-existing respiratory complaints are especially vulnerable. This must be compared with acute community-acquired pneumonia, which has an approximate case fatality rate of 5% for those patients admitted to hospital, but this represents the tip of a substantial iceberg of clinical disease in the community with negligible mortality. It is not yet known whether some patients may be infected with the SARS agent and develop few or no symptoms. Only when these data are available will it be possible to calculate an accurate case mortality rate. However, it is likely that the mortality rate of this infection is over 10% (Booth et al., 2003). In view of its ability to spread in the hospital environment to health-care workers (HCW), it is likely that SARS will alter medical practice in many ways.

Virus transmission

Some patients, like the index patient at the hotel in Hong Kong, appear to shed large amounts of virus, resulting in a very large number of secondary cases. These patients have been described in other viral infections such as Ebola haemorrhagic fever (Khan et al., 1999) and are termed ‘super-spreaders’. This epidemiological model assumes that there is no asymptomatic carrier state and, although there is no evidence of such a state, it has been noted that, in some outbreaks, there are mild cases of infection (Poutanen et al., 2003). This difference has been ascribed to age, genetic predisposition, smoking, previous immunity and co-infection with another pathogen. Should it be found that asymptomatic carriage occurs, control of the SARS coronavirus is going to be extremely difficult.

The route of transmission seems to be via aerosolized droplets. The portal of entry is thought to be inhalation or contact with mucous membranes and/or conjunctiva. Virus is also present in faeces, and about 25% of patients complain of diarrhoea (Zhou et al., 2003; Lee et al., 2003; Poutanen et al., 2003). The high attack rate in medical staff is probably a function of the stage at which patients present and the production of virus-containing aerosol when medical procedures (such as suction and intubation) are undertaken without adequate protection (Donnelly et al., 2003).

Transmission to health-care workers (HCW)

One of the biggest concerns around the spread of SARS has been the high rate of infections amongst HCW and in healthcare settings. In the Greater Toronto area, 111 of the 144 cases admitted to hospital had exposure to SARS in a hospital, and 73 (51%) were HCWs (Booth et al., 2003). In Hong Kong, among 138 cases of secondary and tertiary spread, 85 (62%) involved HCWs (Lee et al., 2003). Most of this spread was prior to the recognition of the aerosol, environmental and fomite spread of the virus. This has led to specific recommendations of nursing of suspected cases in negative-pressure isolation or a single side room. Protocols have been set up for patients meeting the clinical case definitions to be recognized early and isolated appropriately. HCWs in contact with patients have adopted specific infection-control measures, including the use of face masks with filter efficiencies of 95% or above, use of gloves and eye and head protection (Seto et al., 2003) and careful attention to hand-washing. Not since the advent of HIV have such stringent infection-control procedures been adopted so universally by HCWs. However, despite these measures, there have been reports of transmission to HCWs in the intensive-care setting (Hong Kong Department of Health, 2003). This is thought to be as a result of aerosol-generating procedures like non-invasive ventilation, endotracheal intubation, open tracheal suction, use of nebulizers and bronchoscopy. The recognition that there is a significant increase in viraemia in the second week of the illness (Peiris et al., 2003) and that patients requiring intensive care may therefore have high levels of circulating virus support this increased risk. Moreover, it is now apparent that the virus can survive for over 24 h outside the body on environmental surfaces (WHO, 2003a). This has led to many treatment centres minimizing or avoiding certain high-risk procedures and adopting protocols that include the use of hood respirators with a filtered air-supply for unavoidable procedures. Such equipment is expensive and requires extensive training to gown and de-gown safely and, unless teams are comfortable with and fully versed in their use, this may lead to increased risk of contamination. A recent report of the investigation of one
Clinical features

Infection

Incubation period ranges from 2 to 10 days and early symptoms are fever, myalgia and headache. Fever appears to be almost universal and is often the presenting feature, although some patients present with mild respiratory symptoms (Booth et al. 2003). The respiratory phase starts after 3–7 days, with a dry cough and dyspnoea. In some patients, hypoxia develops and progressive pulmonary infiltrates appear on the chest X-ray. In China, this has been called the ‘blossoming flower’ lesion. The infiltrates are typically peripheral and a study using thin-section CT has shown this, together with a characteristic ‘crazy-paving’ appearance due to thickening of interlobular septa (Wong et al., 2003). Some patients develop respiratory failure and require ventilation. This presents considerable cross-infection control problems (see above). There is some evidence that non-invasive methods of ventilation may be associated with a better outcome, but there are concerns that this approach to ventilation may be associated with the generation of aerosols and an increased risk of viral transmission (Zhao et al., 2003).

In some patients, the illness is biphasic in that, 4–7 days after defervescence of pyrexia, new infiltrates appear on X-ray and respiratory failure worsens; these patients often do badly. Consistent laboratory findings are a thrombocytopenia and leucopenia, which particularly affects lymphocytes. Consistent with virus replication in organs other than lung is the elevation of creatine kinase, lactate dehydrogenase and liver transaminases (Tang et al., 2003; Booth et al., 2003). In a multivariate analysis, the independent predictors of an adverse outcome were advanced age (odds ratio per decade 1·80; 95 % confidence interval, 1·16–2·81; P = 0·003) and an absolute neutrophil count that exceeded the upper limit of the normal range on presentation (odds ratio, 1·60; 95 % confidence interval, 1·03–2·50; P = 0·04) (Lee et al., 2003).

Diagnosis

The laboratory tests that can be applied for routine diagnosis are PCR and immunofluorescent ELISA detection of serum antibodies using SARS coronavirus from Vero cell culture. Early efforts at the University of Hong Kong using random RT-PCR provided 646 bp of genomic sequence, from which a diagnostic PCR was developed (Peiris et al., 2003). Parallel efforts in Hamburg yielded three fragments that, when sequenced, did not overlap with a 400 bp fragment reported the day before (24 March 2003) by the CDC (Drosten et al., 2003). Nested primers were designed by the Hamburg group within the orflab region, which encodes the viral replicase 1B (Marra et al., 2003). The Hamburg group found five of the patients with probable SARS studied to be positive with all of their primers, both nested and non-nested. Only three of 13 suspected patients were positive. The group also developed a real-time quantitative PCR that showed that the concentration of viral RNA was highest in sputum and low in nose and throat swabs, suggesting that sputum is the best diagnostic sample. The index patient had very high viral loads \[8 \times 10^6 \text{ virus copies (ml sputum)}^{-1}\]. This fits with the highly infectious nature of patients at presentation. Low levels of virus were found on day 9 in serum, suggesting a long viraemic phase and replication of the virus outside the respiratory tract (exemplified by deranged liver function tests). Viral RNA was also found in the faeces of patients late in convalescence, a feature of other coronaviruses (Cho et al., 2003). Extended testing of patients and contacts for antibody to SARS coronavirus will be important in the future to delineate spread and to identify asymptomatic carriage. The CDC group also developed RT-PCR and an ELISA as well as IFA for antibody detection (Ksiazek et al., 2003). Through the efforts of the WHO, a standardized PCR protocol is now available in participating national reference laboratories.

Outcome

One study has determined a more accurate estimate of the case fatality rate than initial work early in the epidemic. Depending on the statistical method used, mortality in < 60-year-old patients was 3·2 % and, for > 60 years, 43·3 or 55·0 % (Donnelly et al., 2003). This rate was higher than the cumulative rates given previously by the WHO and local authorities. One of the explanations is that the study was able to allow for new patients that had been added to the denominator but that had not yet experienced mortality.

Global control

In an effort to co-ordinate a global public health effort to curb the spread of SARS, early recognition and isolation of cases remain key elements in formulating local, national or international strategies. Epidemiological monitoring of the outbreak in terms of numbers of cases, clustering of cases and documenting exposures has been crucial in identifying modes of transmission and incubation period (Donnelly et al., 2003). The WHO devised case-definitions based on close contact with a suspected case or travel to an area reporting local transmission associated with symptoms of a respiratory tract infection and radiological changes. Now that the first-generation tests for the SARS coronavirus have been developed, these have been incorporated into the case-definitions (CDC, 2003). However, viral cultures and viral detection in patient secretions and excretions by RT-PCR remain relatively insensitive for a reliable early diagnosis. Serological diagnosis is likely to provide definitive diagnosis based on acute and convalescent specimens. Detailed clinical features and prognostic markers for disease progression have been characterized (Booth et al., 2003; Hsu et al., 2003; Lee et al., 2003).
of the first major outbreak in the world in Guangzhou shows some important trends (Zhao et al., 2003). Some 190 patients were randomized to four different treatment regimens. It was found that very-high-dose steroids given early with non-invasive ventilation offered the best outcome; there was no convincing evidence that ribavirin helped. Recently, a very small study has supported these findings (So et al., 2003). The Guangzhou group went on to treat a further 160 patients with the optimal regimen, with only three deaths. At the time this study was undertaken, no diagnostic tests were available, so only clinical case-definitions could be used. This means there may be an excess of suspect cases, leading to a lower mortality.

The clinical features of SARS are now well recognized. Many of the characteristic features were described in early reports (Tsang et al., 2003; Booth et al., 2003). Treatment remains supportive; early hopes that ribavirin would be effective (Koren et al., 2003) have not been borne out by experience (Zhao et al., 2003). The use of corticosteroids, particularly at high doses, is important and two studies, one with 31 patients and the other with 190, support this action (So et al., 2003; Zhao et al., 2003). Much effort will doubtless be expended in developing antiviral agents in the future. The major protease of SARS coronavirus is a potential target and has been expressed in Escherichia coli (Anand et al., 2003). Preliminary characterization shows that it retains many of the structural motifs found in other coronavirus proteases. Molecular modelling suggests that inhibitors such as AG7098 would serve as good lead compounds for development of specific antivirals.

The fate of the epidemic

China was the source of SARS coronavirus and remains the worst affected country, with 5013 of the world’s 7447 cases on 12 May 2003. Almost half (252) of the total deaths (552) have occurred in China. The other major clusters include the eastern coastal region to inland rural areas which may make control and eradication much harder. Reports that SARS has been identified in Hebei, Anhui, Guangxi and Henan (Parry, 2003a), with a new outbreak after the epidemic was declared over, mean that we cannot be complacent about controlling this new threat to public health.

References


