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Control dynamics of severe acute respiratory syndrome transmission

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Abstract Severe acute respiratory syndrome (SARS) is a serious disease with many puzzling features. We present a simple, dynamic model to assess the epidemic potential of SARS and the effectiveness of control measures. With this model, we analysed the SARS epidemic data in Beijing. The data fitting gives the basic case reproduction number of 2.16 leading to the outbreak, and the variation of the effective reproduction number reflecting the control effect. Noticeably, our study shows that the response time and the strength of control measures have significant effects on the scale of the outbreak and the lasting time of the epidemic.

Keywords: SARS, basic reproduction number, effective reproduction number, scale of epidemic, lasting time of epidemic.

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The epidemic of severe acute respiratory syndrome (SARS) has spread rapidly throughout the world ever since it was reported in Guangdong Province of China last November. As of May 26, 2003, 8202 cases have been reported to WHO from 28 countries with 725 deaths recorded^[1]. The Asia-Pacific region, especially the mainland of China, has been badly affected by SARS. "If China is not able to deal with SARS, then it will be very problematic to deal with globally"^[2]. Since the end of April, Chinese authorities have introduced very strict measures to contain the epidemic. As a result, the number of daily reported new cases has declined drastically since May 22, 2003. However, quantitative understanding of the transmission of the epidemic and effectiveness of the control measures is still limited.

In fact, SARS is a new disease with many puzzling features. For this kind of sudden disease with a lot undetermined details, we need a quantitative model to rapidly assess the scale of an outbreak, evaluate the effectiveness of control measures and estimate the trend of epidemic. In this report, we present a simple, dynamic model to assess the probability of SARS transmission and the effectiveness of control measures. Although the model is very simple, it possesses some basic, common characteristics of nonlinear dynamics. Hence, with very few fitted parameters, it can successfully delineate the evolution of nonlinear system with unknown details. With this model, we analyzed the SARS epidemic data in Beijing. We also investigated the impacts of response time and strength of control measures on the scale of the outbreak and the lasting time of the epidemic. The results show that the model can properly characterize the development of epidemic, and evaluate the effectiveness of control measures.

1 Mathematical model

There are many models available for infectious disease transmission^[3]. Choosing or building a right model for a disease is a trade-off between accuracy, generality and simplicity. In this section, we propose a mathematical model to evaluate the SARS transmission and impacts of the control strategies on the epidemic. Before going ahead to the model, it is helpful to look at the main features of SARS transmission. Since Beijing, the capital of China, is the city hardest-hit by SARS, we would focus our study on the epidemic in Beijing. The following are some facts about SARS spreading there:

(1) Beijing has a population of thirteen millions. But the cumulative number of SARS cases is about 2.5 thousands and the cumulative number of cured cases is over 1.2 thousands till the end of May^[1]. Both the infectious and the recovered are a small fraction in the total population.

(2) SARS coronavirus is a novel virus^[4] and no vaccine is available now. Hence, except for those recovered from SARS, almost all individuals may be susceptible to the disease.

(3) Infected individual seems not to be infectious during his or her incubation period. The incubation period is about $2-10 d^{[5]}$.

(4) Outburst individuals are capable of transmitting disease. The infectious period depends on the time of treatment^[5].

Based on these, we assume that in the control of SARS transmission (CT) model, the number of daily new SARS cases *i* is proportional to the existing infectious cases, $i \propto K \cdot [I(t) - I(t - \tau)]$, where *I* is the cumulative number of SARS cases and $i = \frac{dI}{dt}$, *t* is time, *K* is the average rate at which the secondary cases are generated by a primary case in a susceptible population and τ is the average infectious period.

As for the controlling measures to fight SARS, the main experience in Beijing is early case detection, swift contact tracing, prompt isolation and quarantine, timely treatment, etc.^[6]. Then, it is reasonable to introduce a control reduction factor $0 \le \kappa < 1$, to reduce the infect rate *K* to $K \cdot \kappa$. Generally speaking, the reduction factor κ depends on two aspects of the public, institutional and governmental efforts. Some measures are enhanced with the increasing cumulative number *I*, such as the improvement of public health measures. Others will be let up with de-

ceasing daily new cases i when the epidemic appears to be under control, like restriction on travel and school closing. Hence, without losing generality, we introduce the following mode of controlling measures,

$$\kappa = \exp\left[-\left(\frac{I}{I_{\rm c}}\right)^m \left(\frac{i\tau}{I_{\rm c}}\right)^q\right],$$

where I_c is the cumulative number of cases when controlling measures take effects, *m* and *q* (*m*, *q* >0) are two indexes describing the strength of the controlling measures. The parameter *m* relating to the cumulative number *I* represents persistent control measures, whilst the parameter *q* relating to the new cases *i* represents changeable control, namely tight or loose quarantine depending on the number of current new cases.

Obviously, $K \cdot \tau$ is the basic reproduction number R_0 , the most important quantity in the disease transmission. I_c or $I_c/(K \cdot \tau)$ is an indicator of the timeliness of the controlling measures. More importantly,

$$R_{\rm t} = K \cdot \kappa = K\tau \cdot \exp\left[-\left(\frac{I}{I_{\rm c}}\right)^m \left(\frac{\dot{I}\tau}{I_{\rm c}}\right)^q\right]$$

is the effective reproduction number. Generally, R_t is lower than R_0 because of the effect of control measures in cutting off transmission. For an epidemic to be controlled, R_t has to be kept below 1, the self-sustaining threshold.

2 Results and discussions

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We obtained these parameters in the CT model by fitting it to the reported daily new cases in Beijing^[1], only data from Apr. 21 to May 24, 2003 are adopted in the model fit. Fig. 1(a) and (b) show the model estimates for the number of daily new cases and accumulated cases in Beijing, respectively. The comparison between the estimates and the reported data demonstrates that the model can delineate the development of the epidemic properly. In addition, the model clearly shows the decreasing daily new cases since mid-May, which reflects the effectiveness of the anti-SARS measures.

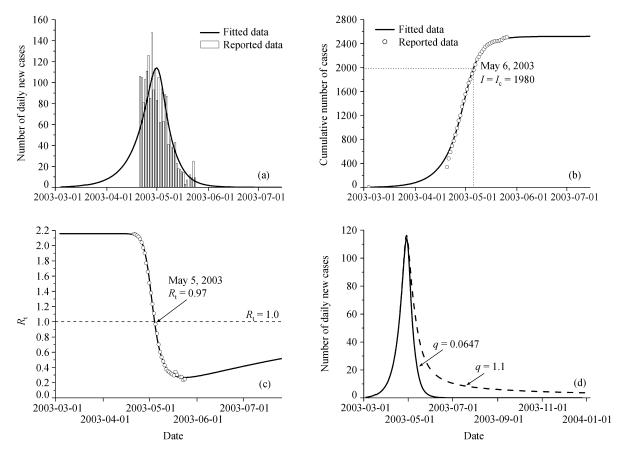


Fig. 1. Quantitative evaluation of SARS epidemic in Beijing. Fitting parameters: K=0.135/day, $\tau=16$ d, $I_c=1980$, m=4.14, and q=0.065). (a) Comparison between the reported and estimated number of daily new cases (*i*) in Beijing; (b) comparison between the reported and estimated cumulative number of cases (*I*) in Beijing; (c) variation of the effective reproduction number R_t in the epidemic. The continuous curve is the R_t calculated from the estimated *i* and *I*, while the scattered points are calculated from the reported *i* and *I*. (d) variation of the epidemic potential with different strength of anti-SARS measures. It is noticeable that the two curves are of the same *m* and I_c but different *q*. The *q* (1.1) causing epidemic curve with long tail is more than 17 times higher than the *q* (0.065) fitted from the data of Beijing.

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Our model fit gives the parameters K=0.135/d, $\tau=16$ d, $I_c=1980$, m=4.14 and q=0.065. Then we derived the basic reproduction number as $R_0=K\tau=2.16$, which is in the same order as Lipsitch^[7] and Riley^[8] reported for Canada and Hong Kong. Therefore, the SARS coronavirus can spread swiftly in susceptible population, if timely diagnosis and isolation measures are not taken by authorities. The value of parameter I_c of about 1980 suggests that the anti-SARS measures took distinct effect on May 6, 2003, 64 days after the first SARS case reported in Beijing (Mar. 4).

As mentioned above, the effective reproduction number R_t can reflect the interaction between the transmission and control vividly. To trace the tendency of the effective reproduction number can give some practical clues to control the epidemic. The variation of the effective reproduction number R_t in the epidemic is shown in Fig. 1(c). At the beginning of the curve (before Apr. 24), $R_{\rm t} = R_0 \sim 2.16$. This indicates that there were not effective control measures and the disease spread rapidly. The value of R_t drops drastically since Apr. 30, after the anti-SARS measures started to take effects. On May 5, R_t dropped down to less than 1, the self-sustained threshold of the epidemic. This means that the disease has been subdued since then. Therefore, if the present control measures are still kept in force, the disease will be put under control in mid-June, as shown in Fig. 1(a). Till then, the cumulative number of SARS cases will reach about 2530. However, if the effective control measures are let up, the epidemic curve would present a tail lasting a long time even bounce. For an illustration of the danger, Fig. 1(d) provides a fictitious epidemic with the parameter q 17 times higher than its real value.

 Table 1
 Model fitting parameters for Beijing and Hong Kong

	<i>K</i> /d	t∕d	$I_{\rm c}$	m	q
Beijing	0.135	16	1980	4.138	0.0647
Hong Kong	0.155	18	1036	1.30	0.0394

In addition, we applied this CT model to the epidemic in other areas and obtained similar and reasonable results. Table 1 compares the fitting parameters in the model for Beijing and Hong Kong. It is clear that the basic reproduction number in the two areas ranges from 2 to 3, but the response delay (characterized by I_c) and the two indexes *m* and *q* are different, reflecting the difference in control measures in the two areas.

Now, we intend to understand the mechanism governing the transmission and control of SARS. We characterize an epidemic curve with the total cumulative number of patients I^* and the epidemic lasting time *T*. It is found that the parameters I_c , *m*, and *q* are the corresponding indicators of the effectiveness of control measures.

Fig. 2(a) illustrates the effects of I_c on the total number of SARS cases I^* and the lasting time of the epidemic

T. Obviously, I^* is directly proportional to I_c , which is reasonable because the CT model can be rescaled with I_c . In addition, the epidemic lasting time increases with I_c . The curves in Fig. 2(a) suggest that the timeliness of the control measures, i.e. smaller I_c , leads to a shorter, smaller-scaled, and less intense epidemic.

The impacts of parameters m and q on the epidemic are shown in Fig. 2(b) and (c), respectively. Their effects are not so straightforward as that of I_c . However, numerical simulations show that I^* and T increase with m decreasing (Fig. 2(b)). That is, the more intensive anti-SARS measures result in a shorter and smaller-scaled epidemic. On the contrary, Fig. 2(c) shows that I^* and Tincrease with increasing q. Higher values of parameter qmay even lead to oscillations. Hence, the laxer the control

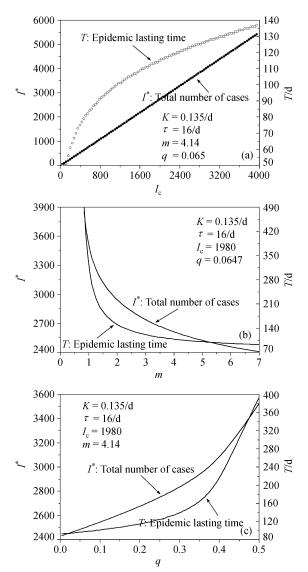


Fig. 2. Impacts of parameters I_c , m, and q on the scale and lasting time of the epidemic. (a) Effects of I_c on I^* and T; (b) effects of m on I^* and T; (c) effects of q on I^* and T.

measures, the longer time the epidemic will last.

3 Summary

Actually, to control the SARS transmission is an interplay between the mankind and the coronavirus. The basic and effective reproduction numbers, depending on the nature of SARS and the infrastructure of public health, play the key role in an outbreak. Unless the vaccine specific to SARS becomes available then greatly reduces the reproduction number, strict quarantine remains the most effective control measure. The timely response can halt the outbreak significantly. The two behavior modes in control, characterized by parameters m and q respectively, govern the late phase of the epidemic. These may need more attention in the late-term measures to eradicate the disease.

Additionally, the CT model, a simple nonlinear dynamic model with a few parameters only, proves to be a reasonable representation of the interaction between transmission and control in an epidemic. As long as the parameters are determined by data fitting, the model can help rapidly assess the scale of an outbreak, evaluate the effectiveness of control measures and estimate the trend of the epidemic. This approach may also be helpful to deal with sudden events with a lot of undetermined details.

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