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METHODS

Individual-based simulation model for COVID-19 transmission in Daegu, Korea

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OBJECTIVES: The aims of this study were to obtain insights into the current coronavirus disease 2019 (COVID-19) epidemic in the city of Daegu, which accounted for 6,482 of the 9,241 confirmed cases in Korea as of March 26, 2020, to predict the future spread, and to analyze the impact of school opening.

METHODS: Using an individual-based model, we simulated the spread of COVID-19 in Daegu. An individual can be infected through close contact with infected people in a household, at work/school, and at religious and social gatherings. We created a synthetic population from census sample data. Then, 9,000 people were randomly selected from the entire population of Daegu and set as members of the Shincheonji Church. We did not take into account population movements to and from other regions in Korea.

RESULTS: Using the individual-based model, the cumulative confirmed cases in Daegu through March 26, 2020, were reproduced, and it was confirmed that the hotspot, i.e., the Shincheonji Church had a different probability of infection than nonhotspot, i.e., the Daegu community. For 3 scenarios (I: school closing, II: school opening after April 6, III: school opening after April 6 and the mean period from symptom onset to hospitalization increasing to 4.3 days), we predicted future changes in the pattern of COVID-19 spread in Daegu.

CONCLUSIONS: Compared to scenario I, it was found that in scenario III, the cumulative number of patients would increase by 107 and the date of occurrence of the last patient would be delayed by 92 days.

KEY WORDS: COVID-19, Mathematical model, Model prediction, Infections, Korea

INTRODUCTION

Coronavirus disease 2019 (COVID-19) is an infectious disease caused by the new coronavirus (2019-nCoV) [1]. It began spreading in Wuhan, China, in December 2019, and on January 14,

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2020, the World Health Organization officially stated that there was a potential for inter-personal infection of COVID-19 in a limited range between families, and on January 22, 2020, the officials in Wuhan, China, reported that there was evidence of COV-ID-19 infection among people [2]. In Korea, the first patient was confirmed with COVID-19 on January 20, 2020. Thereafter, 30 confirmed cases appeared by February 16, 2020, and the rate of increase in confirmed cases was limited to one or two cases per day. However, the situation rapidly changed after the #31 confirmed case on February 18, 2020 in Daegu city. As of March 26, 2020, of the 9,241 confirmed cases in Korea, 6,482 confirmed cases occurred in Daegu, accounting for 70.1% of the total cases. Among them, 4,391 (67.7%) confirmed cases in Daegu were related to Shincheonji. Assuming there are 2.5 million citizens in Daegu, and 9,000 members are part of Daegu Shincheonji, the 2019-nCoV infection rate of Daegu citizens was 0.08%, while the infection rate of Daegu Shincheonji members was 48.78%, which is 583 times higher. This shows that the COVID-19 epidemic in

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Daegu was because the Shincheonji Church in Daegu became a hotspot, leading to the spread of COVID-19 in the Daegu community.

A compartmental model [3], which is widely used as a mathematical study model for the spread of infectious diseases, generally divides the entire population into several groups depending on the state of infection, but it is not an appropriate model to reproduce the hotspot and non-hotspot (communities that are not members of Shincheonji) groups that appear in Daegu. In addition, although a two-patch model [4], that divides the entire population into two groups, was proposed to include hotspots and non-hotspots in the compartmental model, there are difficulties in reproducing infections caused by close contact between the two groups.

To overcome the above problems, we simulated the COVID-19 spread in Daegu using an individual-based model suggested by Ferguson et al. [5]. In the individual-based model, each individual can be infected through contact with infected persons in households, workplaces/schools, and communities (religious and social gatherings), and for this, a virtual population of the same size as the population in Daegu was created. The purpose of this study was to use the individual-based model to understand the current COVID-19 epidemic in Daegu, predict the future spread, and analyze how the reopening of schools scheduled for April 6, 2020 will affect the spread of COVID-19.

MATERIALS AND METHODS

The individual-based model [5] simulates the spread of infectious diseases through close contact between people in households, workplaces/schools, and communities. Through the individual-based model, a virtual population group of the size of

Daegu was created to understand the spread of COVID-19 in Daegu and predict future spread. Each individual in the virtual population group has information regarding the household, work/school, and community. We generated a virtual population of 2,171,000 people from the 2015 census 2% sample data of the MicroData Integrated Service (as of 2015) [6]. Household information and the status of students and workers were set using the age and commuting status of each individual listed in the census data. Assuming that close contact at work or school occurs in the same office or classroom, a student's classroom ID and employee's office ID were virtually generated as follows: the classroom ID was set by randomly selecting students of the same age, city, and district so that an average of 30 students were assigned to the same classroom. Figure 1A shows the histogram according to the number of students in each classroom in Daegu. The office ID was set by randomly selecting the workers in Daegu without the distinction of municipalities and assigning an average of 20 people to the same office. Figure 1B shows the histogram according to the number of workers in each office in Daegu.

Table 1 shows some of the virtual population groups created in this way. Each row represents one individual, and each column represents the attributes of the individual used to simulate the spread of the infectious disease. Individuals with the same household, classroom, and office IDs belonged to the same household, classroom, and office, respectively. If the classroom ID and the office ID are marked as NA, it means that the individual is neither a student nor worker. Items on hotspot indicates whether the individual is a member of Shincheonji or not, and 9,000 people were randomly selected from the population of Daegu and assigned to the hotspot. The infection status in the last column of Table 1 represents the infection status of each individual on a certain date, and the possible infection status were as follows: those susceptible

Figure 1. Histogram of the number of (A) students in the classroom and (B) workers in the office.

right 2. Comparimental studies of our epidemic model for coloriavirus disease 2019 (COVID-19). The infection status is as follows (sus-
ceptible (S), latent (L), infectious (I), hospitalization (H), and recovered or dead and κ , a , η are the latent period, period between symptom onset to confirmation, period from being confirmed to recovery, respectively. **Figure 2.** Compartmental structure of our epidemic model for coronavirus disease 2019 (COVID-19). The infection status is as follows (sus-
All the status of the control of the control of the control of the control of the 320 susceptible (S), latent (L), infectious (I), hospitalization (H), and recovered or dead (R). 320 susceptible (S), latent (L), infectious (I), hospitalization (H), and recovered or dead (R).

individual, and each column represents the attributes of the in-
individual, and each column represents the attributes of the in-
 $\frac{1}{2}$ at a core of 20 per left the latent period of COVID-19, the #31 pat dividual used to simulate the spread of coronavirus disease 2019 $(COVID-19)$ ¹

NA, not available; S, susceptible; I, infectious.

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1 Individuals with the same household, classroom, and office IDs belong to the same nousehold, classroom, and once, respectively. If the classroom
or office ID is NA, it means that she/he is not a student or worker. The for β_{hostpot} and β_{h} are described in the Materia hotspot indicates whether individuals are a member of Shincheonji. A tion. Data on the date of onset of symptoms in confirme to the same household, classroom, and office, respectively. If the classroom The individual-based model and all parameter values except to the same household, classroom, and office, respectively. If the classroom detailed description of the infection status is shown in Figure 2.

fection - latent (L); those that can infect the susceptible -infectious spot). In the individual-based model, changes in infection states that recovered or died from COVID-19 - recovered (R). The individual control of the (I); those isolated after being confirmed with COVID-19 - hospi-
are statistically simulated; that is, the talization (H); and those that recovered or died from COVID-19 ble person becomes a latent person is reali -recovered (R). Figure 2 is a diagram showing the change in each $\frac{1}{2}$ is a diagram showing the change in each infection status. λ is the infection probability of the susceptible and was calculated as follows [5]. talization (H); and those that recovered or died from COVID-19 ble person becomes a latent person is realized as follows. Every

$$
\lambda = \frac{\beta_h/3 \cdot N_h^I}{(N_h)^{0.8}} + \frac{\beta_w/3 \cdot N_w^I}{N_w} + \frac{\beta_s/3 \cdot N_s^I}{N_s} + \frac{\beta_{hotspot}/3 \cdot N_{hotspot}^I}{N_{hotspot}}
$$

 β_h (β_w , β_s , $\beta_{hotspot}$) is the probability of encountering an infected lation results using different random P_h , P_w , P_s , $P_{hostpol}$, is the probability of encountering an infected and results using different random seeds rather than a sm
individual in the household (work, school, hotspot) and getting simulation. We performed intervalue in the notational (work, sensor, holdpot) and getting
infected. We set this as $\beta_h = \beta_w \beta_s = 2\beta_h$, same as that set by Ferguson dom seeds and confirmed whether the median of this result reet al. [5]. N_h (N_w , N_s , $N_{hotspot}$) represents the total number of produces the statistics of cumulative confirm rouseholds (work, school, hotspot), and N_h (N_w , N_s , $N_{hotspot}$) \rightarrow 2.0, 2020. This $p_{hostoot}$ and p_h , the parameter with a higher determined cases in represents the total number of those infected in households (work, school, hotspot). $1/\kappa = 5.2$ [7], $1/\alpha = 4.3$ [8], and $1/\eta = 14$ are the av-
Daegu was β_{hostpot} . When β_{hostpot} was determing the confirmation (days), average period between symptom onset to communicate dases were realized, and to confirmation (days), and average period from being confirmed β_{μ} , a parameter value for reproducing the no with COVID-19 to recovery (days), respectively. erage latent period (days), average period between symptom onset cumulative confirmed cases were realized, and while adjusting for erage latent period (days), average period between symptom onset \mathcal{R} and *β_h*, *b* 11_{*Wh*}, *l* 3_{*b*}, *l* hotspot) represents the total number of *produces* the statistics of cumulative commence cases as of material households (work, school, hotspot), and N_h^I (N_w^I , N_s to confirmation (days), and average period from being confirmed β_h , a parameter value for reproducing the non-hotspot cumulative in a continuous confirmation (days), and average period from being confirmed β_h , a par

In February 1, 2020, 10 infected individuals in the hotspot were $\beta_h = 0.33$. This shows to set as the initial confirmed patients. The #31 patient was confirmed with COVID-19 on February 18, 2020, but it was assumed

Table 1. Each row of the synthetic population data represents one that the symptoms started on February 7, 2020, and considering $(COVID-19)$ ¹
2020. The individual-based model was simulated on a daily basis, 2 1 44 NA NA True 1 COVID-19 infections from other regions in Korea and from individual, and each column represents the attributes of the in-
the latent period of COVID-19, the $\#31$ patient was assumed to d to simulate the spread of coronavirus disease 2019 be infected from the initial patients of the hotspot on February, 1 15 individual, and each column represents the attributes of the individual used to simulate the individual u that the symptoms started on February 7, 2020, and considering and it was assumed that the population inflow into Daegu using public transportation was minimal after the COVID-19 epidemic $\overline{1}$ 1 48 NA 3 False 5 in Daegu. That is, it was assumed that there was no influx of new abroad.

87 group has information regarding the household, work/school, and community. We generated a 88 virtual population of 2,171,000 people from the 2015 Census 2% sample data of the Microdata

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RESULTS

to COVID-19 - susceptible (S); those in the latent stage after in-
March 26, 2020 (4,391 cases in the hotspot, 2,091 in the non-hot- $\frac{1}{100}$ are not can infect the latent statent state after the set $\frac{1}{100}$ patients in Daegu until households (work, school, hotspot), and $N_h^I(N_w^I, N_s^I, N_{hotspot}^I)$ 26, 2020. Among $\beta_{hotspot}$ and β_h , the parameter with a higher deter-Follows: the possible infection status were assumed to COVID-19 $P_{n\text{obsport}}$ and P_{n} are assessed in the individuals are a member of Shincheonji. A tion. Data on the date of onset of symptoms in confirmed patients detailed description of the infection status is shown in Figure 2. ${}$ with COVID-19 are not currently available. We set β_{hostpot} and β_{h} -19 - susceptible (S); those in the latent stage after in-
March 26, 2020 (4,391 cases in the hotspot, 2,091 in the non-hotent (L); those that can infect the susceptible -infectious spot). In the individual-based model, changes in infection status $B \cdot N_h^I$, $\beta_W/3 \cdot N_W^I$, $\beta_S/3 \cdot N_s^I$, $\beta_{hotspot}/3 \cdot N_{hotspot}^I$ ble person becomes a latent patient. Since the random number �������� changes every time it is generated, the simulation results can be β_{hostspot}) is the probability of encountering an infected lation results using different random seeds rather than a single in the household (work, school, hotspot) and getting simulation. We performed 100 simulations using different ran- W_h $(N_w, N_s, N_{hotsnot})$ represents the total number of households the statistics of cumulative confirmed cases as of March 121 **(work)** $1/\kappa = 5.2$ [7], $1/\alpha = 4.3$ [8], and $1/\eta = 14$ are the av-
Daegu was $β_{hotspot}$. When $β_{hotspot}$ was determined, the results of the 123 the average period from being communed *P_p*, a parameter value for reproducing the fior flotspot cumulative CD-19 to recovery (days), respectively. cases was found. The parameter results were *β_{hotspot}*= 3.06 and for β_{hostoot} and β_h are described in the Materials and Methods secare statistically simulated; that is, the changes in which a susceptiday, a uniform random number between 0 and 1 is generated for all susceptible people, and if this value is less than lambda, which is the probability of becoming latent after infection, that susceptidifferent even with the same parameters and initial patient settings. Therefore, it is necessary to check the distribution of simudom seeds and confirmed whether the median of this result reminant power for reproducing the cumulative confirmed cases in β_h = 0.33. This shows that the probability of infection between the hotspot and household and workplace differed by more than nine times. Figure 3 shows the cumulative confirmed cases and simu-

Figure 3. (A) Cumulative confirmed coronavirus disease 2019 (COVID-19) cases in the city of Daegu and (B) cumulative number of hospitalizations in the simulation. Here, we show the median of 100 simulation results for different random seeds.

lation results of the individual-based model by March 26, 2020. The reo Since a list of members of Shincheonji Church and the screening and quarantining of them for COVID-19 infection began at the March 26, 2020. The assumption of scenario III that end of February, in the above simulation, $\beta_{\text{hotspot}} = 0$ as of February 29, 2020. In addition, 1/α was set to 2.7 instead of 4.3 after February 29, 2020 to reflect the effect of the shortened $1/\alpha$, the average period from symptom onset to confirmation, after massive screening tests for the Shincheonji Church members. last patient $\frac{1}{2}$

> The statistics of the confirmed cases as of March 26, 2020 were reproduced, and to predict the spread of COVID-19 in Daegu thereafter, we considered the three following scenarios.

- Scenario I: Maintaining vacations in elementary/middle/high schools
- Scenario II: Reopening of elementary/middle/high schools on April 6, 2020
- Scenario III: Reopening of elementary/middle/high schools on April 6, 2020 & after April 6, 2020, 1/α, the average period from symptom onset to confirmation, increases again to 4.3 days

26, 2020. The reopening of elementary/middle/high schools on April 6, 2020, mentioned in scenarios II and III, is the current plan as of March 26, 2020. The assumption of scenario III that the average period from symptom onset to confirmation would increase again to 4.3 days considers the students' relatively passive expression of symptoms after the reopening of schools. The individual-based model simulation results for the above three scenarios are shown in Table 2, Figures 4 and 5. In scenario I, the number of cumulative confirmed cases in Daegu was 6,677 (4,394 in hotspot, 2,322 in non-hotspot), and the last newly confirmed cases occurred on April 26, 2020. In scenario II, the number of cumulative confirmed cases in Daegu was 6,716 (4,394 in the hotspot, 2,322 in the non-hotspot) and compared with scenario I (based on the median), 39 Daegu citizens were additionally infected (non-hotspot, not a member of Shincheonji). The last newly confirmed cases occurred on May 3, 2020, 7 days later than in scenario I. In scenario III, the number of cumulative confirmed cases were 6,784 (4,394 in hotspot, 2,390 in the non-hotspot), and 107 addi-

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Figure 4. Cumulative number of hospitalization cases in the simulation. Here, we show the median and 5th to 95th percentile range for (A) scenario I, (B) scenario II, and (C) scenario III.

Figure 5. Cumulative and daily hospitalization cases in the simulation (A, B) scenario I, (C, D) scenario II, and (E, F) scenario III. Here, we show the median of 100 simulation results for different random seeds.

tional citizens of Daegu that were not members of Shincheonji were infected. The last newly confirmed cases occurred on July 27, 2020, 92 days later than in scenario I. Figure 4 shows the cumulative daily confirmed cases for each scenario and the interval except for the median value and the top and bottom 5% of 100 simulations using different random seeds. Figure 5 shows the median values for cumulative confirmed cases in the hotspot and non-hotspot.

DISCUSSION

The individual-based model was selected to explain the characteristics demonstrated by the spread of COVID-19 in Daegu, specifically, that the cumulative infection rate of Shincheonji members was about 583 times higher than that of non-Shincheonji members. Compared with the compartmental model which is widely used as a mathematical model for the spread of infectious diseases [3], the individual-based model simulates the transmission of infectious diseases through close contact among people using the socio-demographic information of each individual (household, workplace/school, community such as religious and social gatherings). Therefore, it has the advantage of being able to analyze the effect of quarantine policies, such as closing schools and implementing shifts at work, in preventing infection spread in more detail.

Using the individual-based model, we reproduced the cumulative COVID-19 confirmed cases of Daegu until March 26, 2020. The number of newly confirmed cases per day in Daegu sharply increased from February 21, 2020, after the #31 confirmed case, and the largest number of confirmed cases was reported on February 29, 2020, with 656 cases. Since then, and after March 11, 2020 with 131 confirmed cases, a decreasing trend was maintained with less than 100 confirmed cases per day. However, it cannot be said that the rapid increase in the number of confirmed cases during this period reflects the actual rate of COVID-19 spread in Daegu. Because of the intensive large-scale screening of the members of Shincheonji, it is likely that the rate increased more steeply than the actual rate of spread because more confirmed cases arose in a short period of time compared with the previous order and rate of infection. Since data on the date of symptom onset in patients in Daegu are not available, the results of this study using the cumulative data may be different from the actual COVID-19 transmission patterns in Daegu. If the data on the date of symptom onset are collected in the future, a follow-up study using this information should be conducted. In addition, because we did not assume any other additional group infections, such as in nursing homes, other than Shincheonji, this study may show a different pattern from the actual transmission of COV-ID-19.

The above results assume that newly infected COVID-19 cases in Daegu did not come from abroad or other regions in Korea. For a more accurate prediction and analysis of the effect of quarantine policies, such simulations should be expanded to reflect the whole country and consider the entry of latent patients from other regions and abroad. Studies on this are currently under way.

SUPPLEMENTARY MATERIALS

Korean version is available at http://www.e-epih.org/.

CONFLICT OF INTEREST

The authors have no conflicts of interest to declare for this study.

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AUTHOR CONTRIBUTIONS

All work was done by WSS and RISEWIDs Team.

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REFERENCES

- 1. World Health Organization. Naming the coronavirus disease (COVID-19) and the virus that causes it [cited 2020 Apr 1]. Available from: https://www.who.int/emergencies/diseases/novel-coronavirus-2019/technical-guidance/naming-the-coronavirus-disease-(covid-2019)-and-the-virus-that-causes-it.
- 2. World Health Organization. WHO timeline-COVID-19; 2020 Apr 27 [cited 2020 Jun 20]. Available from: https://www.who.int/ news-room/detail/27-04-2020-who-timeline---covid-19.
- 3. Brauer F, van den Driessche P, Wu J. Mathematical epidemiology. Berlin: Springer; 2008, p. 19-79.
- 4. Dowdy DW, Golub JE, Chaisson RE, Saraceni V. Heterogeneity in tuberculosis transmission and the role of geographic hotspots in propagating epidemics. Proc Natl Acad Sci U S A 2012;109:9557- 9562.
- 5. Ferguson NM, Cummings DA, Cauchemez S, Fraser C, Riley S, Meeyai A, et al. Strategies for containing an emerging influenza pandemic in Southeast Asia. Nature 2005;437:209-214.
- 6. MicroData Integrated Service. Data in service [cited 2020 Jul 20]. Available from: https://mdis.kostat.go.kr/infoData/detailData. do?statsConfmNo= 101001%20(2020) (Korean).
- 7. Li Q, Guan X, Wu P, Wang X, Zhou L, Tong Y, et al. Early transmission dynamics in Wuhan, China, of novel coronavirus-infected pneumonia. N Engl J Med 2020;382:1199-1207.
- 8. Ki M; Task Force for 2019-nCoV. Epidemiologic characteristics of early cases with 2019 novel coronavirus (2019-nCoV) disease in Korea. Epidemiol Health 2020;42:e2020007.