



Risk factors associated with COVID-19 infection: a retrospective cohort study based on contacts tracing

Tao Liu , Wenjia Liang , Haojie Zhong , Jianfeng He , Zihui Chen , Guanhao He , Tie Song , Shaowei Chen , Ping Wang , Jialing Li , Yunhua Lan , Mingji Cheng , Jinxu Huang , Jiwei Niu , Liang Xia , Jianpeng Xiao , Jianxiong Hu , Lifeng Lin , Qiong Huang , Zuhua Rong , Aiping Deng , Weilin Zeng , Jiansen Li , Xing Li , Xiaohua Tan , Min Kang , Lingchuan Guo , Zhihua Zhu , Dexin Gong , Guimin Chen , Moran Dong & Wenjun Ma

To cite this article: Tao Liu , Wenjia Liang , Haojie Zhong , Jianfeng He , Zihui Chen , Guanhao He , Tie Song , Shaowei Chen , Ping Wang , Jialing Li , Yunhua Lan , Mingji Cheng , Jinxu Huang , Jiwei Niu , Liang Xia , Jianpeng Xiao , Jianxiong Hu , Lifeng Lin , Qiong Huang , Zuhua Rong , Aiping Deng , Weilin Zeng , Jiansen Li , Xing Li , Xiaohua Tan , Min Kang , Lingchuan Guo , Zhihua Zhu , Dexin Gong , Guimin Chen , Moran Dong & Wenjun Ma (2020): Risk factors associated with COVID-19 infection: a retrospective cohort study based on contacts tracing, *Emerging Microbes & Infections*, DOI: [10.1080/22221751.2020.1787799](https://doi.org/10.1080/22221751.2020.1787799)

To link to this article: <https://doi.org/10.1080/22221751.2020.1787799>



© 2020 The Author(s). Published by Informa UK Limited, trading as Taylor & Francis Group, on behalf of Shanghai Shangyixun Cultural Communication Co., Ltd



[View supplementary material](#)



Accepted author version posted online: 01 Jul 2020.



[Submit your article to this journal](#)



Article views: 188

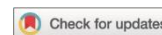


[View related articles](#)



View Crossmark data [↗](#)

Publisher: Taylor & Francis & The Author(s). Published by Informa UK Limited, trading as Taylor & Francis Group, on behalf of Shanghai Shangyixun Cultural Communication Co., Ltd
Journal: *Emerging Microbes & Infections*
DOI: 10.1080/22221751.2020.1787799



Risk factors associated with COVID-19 infection: a retrospective cohort study based on contacts tracing

Tao Liu^{1*}, Wenjia Liang^{2*}, Haojie Zhong^{2*}, Jianfeng He^{2*}, Zihui Chen^{1*}, Guanhao He^{1*}, Tie Song², Shaowei Chen¹, Ping Wang¹, Jialing Li², Yunhua Lan², Mingji Cheng², Jinxu Huang², Jiwei Niu², Liang Xia², Jianpeng Xiao¹, Jianxiong Hu¹, Lifeng Lin², Qiong Huang², Zuhua Rong¹, Aiping Deng², Weilin Zeng¹, Jiansen Li², Xing Li¹, Xiaohua Tan², Min Kang², Lingchuan Guo¹, Zhihua Zhu¹, Dexin Gong¹, Guimin Chen¹, Moran Dong¹, Wenjun Ma^{1†}

1. Guangdong Provincial Institute of Public Health, Guangdong Provincial Center for Disease Control and Prevention, Guangzhou, China.
2. Guangdong Provincial Center for Disease Control and Prevention, Guangzhou, China.

*These authors contributed equally to this work.

†The author was corresponding author.

Correspondence to:

Dr. Wenjun Ma, Guangdong Provincial Institute of Public Health, Guangdong Provincial Center for Disease Control and Prevention, Guangzhou 511430, China

mawj@gdiph.org.cn

Running title: Risk factors of COVID-19 infection.

Total word count: 4941

Abstract

Objectives To estimate the attack rates, and identify the risk factors of COVID-19 infection.

Methods Based on a retrospective cohort study, we investigated 11,580 contacts of COVID-19 cases in Guangdong Province from January 10 to March 15, 2020. All contacts were tested by RT-PCR to detect their infection of SARS-COV-2. Attack rates by characteristics were calculated, and logistic regression was used to estimate the risk factors of infection for COVID-19.

Results A total of 515 of 11,580 contacts were identified to be infected with SARS-COV-2.

Compared to young adults aged 20-29 years, the infected risk was higher in children (RR: 2.59, 95%CI: 1.79-3.76), and old people aged 60-69 years (RR: 5.29, 95%CI: 3.76-7.46). Females also had higher infected risk (RR: 1.66, 95%CI: 1.39-2.00). People having close relationship with index cases encountered higher risk to be infected (RR for spouse: 20.68, 95%CI: 14.28-29.95; RR for non-spouse family members: 9.55, 95%CI: 6.73-13.55; RR for close relatives: 5.90, 95%CI: 4.06-8.59; RR for other relatives: 3.37, 95%CI: 2.15-5.28). Moreover, contacts exposed to index case in symptomatic period (RR: 2.15, 95%CI: 1.67-2.79), with critically severe symptoms (RR: 1.61, 95%CI: 1.00-2.57), with symptoms of dizzy (RR: 1.58, 95%CI: 1.08-2.30), myalgia (RR: 1.49, 95%CI: 1.15-1.94), and chill (RR: 1.42, 95%CI: 1.05-1.92) had higher infected risks.

Conclusion Children, old people, females and family members are susceptible to be infected with COVID-19, while index cases in incubation period had lower contagiousness. Our findings will be helpful for developing targeted prevention and control strategies to combat the worldwide pandemic.

Key words: COVID-19; attack rate; risk factors; close contact; China

Introduction

Since the Coronavirus Disease 2019 (COVID-19) outbreak on 31 December, 2019 [1], it has hit more than 200 countries, areas or territories with 8,525,042 cases and 456,973 deaths as of June 20, 2020 [2]. World Health Organization (WHO) has declared COVID-19 as a pandemic on March 11, 2020 [3]. Owing to the effective measure taken in China, the chain of transmission has been broken and the epidemic has been under control.

Contact tracing is a major public health response to imports of rare or emerging infectious diseases.

The main objectives of contact tracing are to identify potentially infected individuals before onset of severe symptoms, and to prevent onward transmission from the secondary cases. Contact tracing has decisively contributed to the control of many infectious diseases worldwide including severe acute respiratory syndrome (SARS), Ebola virus disease, and Middle East respiratory syndrome (MERS) [4-7]. Report of the WHO-China Joint Mission on COVID-19 pointed out that China has

a policy of meticulous case and contact identification for COVID-19 [8]. Previous studies using mathematical modeling also theoretically demonstrated that contact tracing and quarantine play important roles in controlling the spreading of COVID-19 [9,10]. In addition to this, contact tracing also provides a unique opportunity to investigate the epidemiological features of COVID-19.

Previous researches have analyzed the data of COVID-19 patients and found some risk factors of mortality, such as older age, pre-existing cardiovascular or cerebrovascular diseases, low levels of CD3⁺CD8⁺ T-cells, high levels of cardiac troponin I, higher Sequential Organ Failure Assessment score and d-dimer [11,12]. Unfortunately, limited study has paid attention to the risk factors related to COVID-19 infection. Recent studies conducted among 1,286 close contacts (98 of them were infected by SARS-CoV-2) in Shenzhen and among 2098 close contacts in Guangzhou (134 of them were infected by SARS-CoV-2) explored the risk factors for COVID-19 infection, like older age, travelling to Hubei, etc [13,14]. Another recent study among 2761 close contact of 100 selected index cases in Taiwan identified exposure to index case with severe symptoms as a risk factor [15]. However, their limited sample size, especially the limited cases, may restrict their ability to perform detailed analysis, and reduce the power to detect significant risk factors. Additionally, findings within a single city or selected sample may restrict its ability of generalization.

In the current study, we employed a large dataset including 11,686 close contacts of COVID-19 cases (449 of them were infected) in Guangdong Province, China to estimate the attack rates, and identify risk factors for infection of COVID-19. Under the context of worldwide pandemic, understanding this issue can identify high-risk groups and provide evidence to develop targeted prevention.

Methods

Setting and definitions

Guangdong, a province with large population size located in Southern China, is a place early affected by COVID-19. The first confirmed case was reported on January 15, 2020, and a total of 1,361 confirmed cases were reported by March 15, 2020. Since the very early stage of COVID-19 outbreak, an intensified surveillance was implemented across Guangdong Province to detect suspected and confirmed COVID-19 cases, and their close contacts following standardized protocols released by the National Health Commission of China. Suspected and confirmed COVID-19 cases were defined based on the Diagnosis and Treatment scheme of COVID-19, and close contacts were defined by the Prevention and Control Scheme of COVID-19. These two schemes were released by the National Health Commission of China (Supplementary materials) [16,17].

Identification and quarantine of contacts

Once a suspected or confirmed COVID-19 case was identified, the case would be reported as an index case and isolated, and the Center for Diseases Control and Prevention (CDC) will conduct a field investigation. Information of index cases was collected by clinical workers, including demographic information, exposure history, clinical symptoms, date of symptom onset, laboratory test results, and the severity. This information was directly reported to the National Internet-Based Infectious Diseases Reporting System. Information of contacts was collected by CDC using a standardized questionnaire, including general demographic characteristics, relationships with the index case, and patterns and frequency of contract. Meanwhile, their throat swabs were collected and detected by real-time reverse transcriptase polymerase chain reaction assay (RT-PCR). During the quarantine, health status of all contacts was monitored, and their throat swabs were collected every several days to test their infection status. Once they were identified with positive of severe

acute respiratory syndrome coronavirus 2 (SARS-COV-2), they would be transferred to a designated hospital for diagnosis and treatment. Clinical symptoms and severity of these infected contacts were followed up and recorded by clinical workers. After 14 days' quarantine, contacts with negative SARS-COV-2 were released.

Statistical analysis

Categorical variables were described using percentage (%), and a Chi-square test was used to test the differences in distributions of categorical variables between index and secondary cases. If conditions for Chi-square test were not satisfied, Fisher's exact test was used.

Attack rate was calculated as the percentage of contacts who were later confirmed to be infected with SARS-COV-2. We estimated the attack rates of contacts by gender, age, relationships to index cases (household members, relatives, social activities, etc.), transportations (flight, train, public transportation, private car, and the Dream Cruise) where infection occurred, course of disease (incubation period, symptomatic period, and different days from symptom onset) of index cases when the contact occurred, severity of index cases (mild, moderate, severe, and critically severe), and clinical symptoms of index cases. These attack rates were calculated only using sub-datasets of the index cases and contacts with detailed information because some cases had no complete information for estimate. Logistic regression was also conducted to estimate the risk factors of COVID-19. All data analyses were conducted by R software (version 3.5.0, R Foundation for Statistical Computing).

Results

General characteristics of contacts

As of March 15, 2020, a total of 11,686 contacts were traced and quarantined. The first contact was

identified on January 10, 2020. Contacts (n=106) without key formation were excluded, and 11,580 contacts were finally included in the analysis. Figure 1 showed the daily number of quarantined contacts, which peaked (n=574) on January 31. Of total contacts, 6,183 (53.4%) were males; 8,419 (72.7%) were adults aged 20-59 years, and 9,725 (84.0%) contacts were quarantined in centralized stations. The number of contacts occurred at home, in social activities, on transportations, and in health care settings were 4,893 (41.0%), 2,016 (16.9%), 3,198 (26.5%) and 1,348 (11.3%), respectively. Many contacts were from family members of index cases (4,707, 40.7%), social activity contacts (3,344, 28.9%), transportation contacts (2,778, 24.0%), and health care workers (573, 4.9%) (Table 1). All contacts were linked to 1,158 index cases, with a mean of 7.8 (95%CI: 7.0-8.7) close contacts per index case. The average contacts per index case varied with contact circumstances and relationships to the index cases (Table S1). The average period from exposure to quarantine was 6.4 days, and the average duration of quarantine was 9.7 days.

Attack rates of COVID-19

Until March 15, 515 (4.4%) contacts were identified to be infected with SARS-COV-2. The attack rates varied by age groups with the highest for the group aged 60-69 years (11.1%), and the lowest for the group of 20-29 years (2.3%). The attack rate of children < 10 years was 5.7%, and the attack rates were higher in children whose index cases aged 30-39 years (8.5%), and 50-59 years (7.0%) (Table S2).

We also observed a higher attack rate in females (5.6%) than in males (3.5%). In addition, contacts having close relationship with index cases had higher attack rate (attack rate: 23.3% for spouse; 10.6% for non-spouse family members; 7.0% for close relatives; 4.1% for other relatives, 1.3% for social activity contacts, etc.). Different attack rates also occurred in various transportations where

infection occurred. Attack rates were 0.8% on flight, 1.2% on train, 2.1% on public transportation, 4.2% on private car and 9.4% on the Dream Cruise (9.4%).

When considering the time contacting with the index cases, attack rates were 3.3% and 7.0% when contacts occurred in the index cases' incubation period and symptomatic period. In detail, attack rate increased from five days prior to the symptom onset of index cases (1.7%), to a peak during 3-4 days (10.1%) after onset, and then decreased to 4.0% after 17 days of the onset. In addition, attack rates increased from 4.6% for the contacts of mild cases to 7.5% for the contacts of critically severe cases. Table S3 shows attack rates for the contacts of index cases with different clinical symptoms, and higher attack rates were observed in index cases with dyspnea (11.2%), dizzy (10.6%), muscle soreness (10.4%), and shortness of breath (10.0%).

Risk of infection for COVID-19

Compared with people aged 20-29 years, children <10 years (RR: 2.59, 95%CI: 1.79-3.76) and children aged 10-19 (RR: 1.81, 95%CI: 1.17-2.81) had higher risk to be infected with COVID-19 (Figure 2A). The risks were also higher in people aged 30-39 years (RR: 1.96, 95%CI: 1.41-2.71), 50-59 years (RR: 2.30, 95%CI: 1.65-3.27), 60-69 years (RR: 5.29, 95%CI: 3.76-7.46) and 70-79 years (RR: 3.03, 95%CI: 1.81-5.08). Moreover, young adults (aged 30-39 years), whose index cases aged <20 years, 30-39 years, and 50-69 years, had higher infected risk (Table S4). We also observed a higher risk in females than in males (RR: 1.66, 95%CI: 1.39-2.00) (Figure 2B). In addition, people having close relationship with index cases encountered higher risk to be infected (RR and 95%CI: 20.68 [14.28-29.95] for spouse; 9.55 [6.73-13.55] for non-spouse family members; 5.90 [4.06-8.59] for close relatives; 3.37 [2.15-5.28] for other relatives) (Figure 2C). In terms of the infected risk in transportations, we did not observe significant difference across various transportations except in

the Dream Cruises (RR: 4.19, 95%CI: 1.21-14.50) (Figure 2D).

When considering time contacting with index cases, the risk of exposure to index cases in the symptomatic period was higher than in the incubation period (RR: 2.15, 95%CI: 1.67-2.79) (Figure 2E). More specifically, the infected risk increased from five plus days prior to the symptom onset of index cases (RR: 0.30, 95%CI: 0.15-0.60), to a peak during 3-4 days (RR: 1.87, 95%CI: 1.33-2.61) after onset, and then decreased to 0.30 (95%CI: 0.12-0.77) after 17 days of the onset (Figure 2F). Moreover, contact with index cases with critically severe symptoms was associated with a higher infected risk (RR: 1.61, 95%CI: 1.00-2.57) (Figure 2G). Figure 2H shows the infected risk for the contacts of index cases with different clinical symptoms compared to fever, and there were higher risks in index cases with dizzy (RR: 1.58, 95%CI: 1.08-2.30), myalgia (RR: 1.49, 95%CI: 1.15-1.94), and chill (RR: 1.42, 95%CI: 1.05-1.92).

Discussion

After reporting the first case on January 15, 2020, Guangdong Provincial government mobilized enormous resources to respond to the COVID-19 epidemic. More than 11,000 close contacts of COVID-19 were traced and quarantined. One third of the total cases reported in Guangdong Province were identified from these contacts, which indicate that contact tracing strategy have played an important role in containing the spreading of COVID-19. The analysis of index cases and their close contacts provides insight into the attack rates and risk factors of infection for COVID-19.

We found that attack rates were higher in the elderly with the highest in the group aged 60-69 years, and logistic regression demonstrated the statistical significance. These findings are consistent with the results for SARS in Beijing [7]. Recent studies also reported that elderly contacts were more likely to encounter COVID-19 infection [13,14]. However, another recent article in Taiwan did not observe significant higher infected risk of elderly contacts, which may ascribe its insufficient sample size [15]. Our findings thus confirmed greater vulnerability of the elderly. Those contacts aged 60-69 years could have more physical activities than older people, which may cause closer contact with

index case for a longer period [7]. Meanwhile, the immunity of the age may be weaker than younger adults, making them more susceptible to infection. Therefore, more efforts are needed to protect the elderly from the infection of COVID-19.

The susceptibility of children to COVID-19 is controversial [8,18]. Clinical data of COVID-19 showed much lower percentage of children aged <10 years [19,20]. A recent systematic review considering literatures of COVID-19 in children pointed out that children cases are usually less severe than adult cases, and more children cases are asymptomatic infection, which makes them less opportunity to be tested and identified [21]. However, we found higher infected risk of COVID-19 in children <10 years that their RR were larger than contacts aged 10-59 years, which indicates that children were also susceptible to COVID-19. Furthermore, we observed a higher attack rate in children whose index cases aged 30-39 and 50-59 years. Although limited sample size may cause insignificant RR, our results still implicated that the children may be mainly infected by their parents and grandparents. Two recent studies reported consistent results with our study [13,22]. For instance, Dong et al. analyzed 2,143 pediatric COVID-19 patients across China, and found that children were susceptible to COVID-19 [22]. Additionally, young adults (30-39 years) were more likely to be infected by children aged < 20 years, their peers aged 30-39 years, as well as people aged 50-69 years. These findings may be attributed to the status that young adults are the primary caregivers once their children and parents got sick, and they are also the individuals who have many social activities with their peers. These findings suggested that people should performed strict personal protection both at home and in public places. Compared with previous studies, our study prospectively collected data based on contacts tracing, which had explicit temporality for causal inference and reduced recall bias, and therefore provide more reliable evidence. Our finding is helpful for preventing people from being infected with COVID-19.

We observed that female contacts were more likely to be infected by SARS-CoV-2 than male contacts, which is consistent with previous studies [13,14]. For example, a recent study conducted in Guangzhou also found higher attack rates in females than in males [14]. This difference in attack rate between sex may be due to several reasons: (1) females play predominant roles as caregivers within family and may have closer contact and longer contact period with the index cases [23]; (2) females comprise a large proportion of health-care workers [24]. Therefore, our findings suggest more prevention measures specifically implemented to protect females from infection during the

epidemic of COVID-19.

We observed that the relationships between contacts and index cases significantly affected the infected risks. Compared with the social activity contacts, the risk of being infected was more than 20 times higher among the spouse and more than 9 times higher among other family members, which was consistent with previous studies on SARS and H1N1 [7,25]. A newly published research also found that more infections were acquired in household [15]. Family members are more likely to have closer contact with index case for a longer contact period with shorter distance. Another possible reason is that family members may have some certain linkage with index cases in living habits which may cause higher predisposition in infection than other close contacts. Unfortunately, individuals commonly take protective measures in public place like washing hands and wearing mask, but neglect personal protection at home. This indicates the necessity for public to pay attention to personal protective at home especially when family members develop symptom or have travel history of epidemic areas.

We also compared attack rates occurred on different transportations, and found lower attack rates occurred on trains or flights. This result indicates that the possibility of transmission of SARS-COV-2 on flight and train was low, which may be related to the advanced air purification system and sanitation in these transportations. However, after controlling for age and sex, the results of logistic regression did not find significant difference across various transportations except in the Dream Cruises. The insignificance may be attributed to the limited sample size and the risk difference may actually exist. Future studies with larger sample should be conducted to explore this issue and provide evidence to guide the development of prevention in transportations.

Although previous studies reported that both asymptomatic and symptomatic cases could infect other persons [26-28], the differences in contagiousness at different phases of COVID-19 remain unclear. Our study shows the contagiousness peaked during 3-4 days after symptom onset, which is consistent with previous studies, which showed higher virus shedding during several days after the onset of symptoms [29-31]. For example, To et al. found that salivary viral load in COVID-19 cases was highest during the first week after symptom onset, and the viral RNA was detected 25 days after symptom onset [31]. In addition, we found contacts before the symptom onset could also lead to infection, which indicates the transmission of COVID-19 in incubation period. Although viral shedding before symptom onset is still limited, Zou et al. reported an asymptomatic patient who had

similar amount of virus to those symptomatic cases [30]. Another study conducted in children also detected positive virus before onset of symptoms in several children cases [20]. These findings suggested COVID-19 could be transmitted before onset of symptoms.

The present study found that severe index cases could cause higher attack rates than mild cases. In addition, compared with cases with fever, dizzy, myalgia and chill caused higher infected risks to their contacts, while cases with rhinorrhea, expectoration and chest tightness caused lower infected risks. To et al.'s study showed higher virus load in specimens of severe patients than mild patients [31], which verified our findings. However, studies are needed to detect the virus load in cases with different clinical symptoms for assessing their contagiousness.

This study has several strengths. First, our study includes the largest number of close contacts of COVID-19 to date. Second, our study is a retrospective cohort study, which provides information with explicit temporality for causal inference and the recall bias was reduced. Third, we estimated the attack rates and infected risks for different contacts, which is helpful for identifying susceptible groups to develop specific protection. Fourth, we estimated the contagiousness across the course of COVID-19.

Some limitations also need to be noted. First, although we used a large dataset with more than ten thousand of contacts, the sample size of cases was limited in some subgroups, which may lead to insufficient power to identify the statistical significance. Second, a number of asymptomatic infections may be missed and their close contacts cannot be identified. Third, since the imperfect sensitivity of the RT-PCR test, some potential infections among close contacts may be missed. Fourth, the data were collected by a variety of epidemiological investigation groups across Guangdong Province. Despite using the same protocol, the implementation may have inconsistency and some noise may be introduced.

Conclusions

Children, old people, females and family members are susceptible to be infected with COVID-19, while index cases in incubation period had lower contagiousness. Our findings will be helpful for developing targeted prevention and control strategies to combat the worldwide pandemic.

Ethical approval and consent to participate

Data collection and analysis of cases and close contacts were determined by the National Health

Commission of the People's Republic of China to be part of a continuing public health outbreak investigation and were thus considered exempt from institutional review board approval.

No study participants were involved in the preparation of this article. The results of the article will be summarised in media press releases from the Guangdong Provincial Center for Disease Control and Prevention.

Consent for publication

Not applicable.

Competing interests

The authors declare that they have no competing interests.

Funding

This study was supported by [the National Key Research and Development Program of China#1] under Grant [number 2018YFA0606200, 2018YFA0606202]; [the Science and Technology Program of Guangdong Province#2] under Grant [number 2018B020207006, 2019B020208005, 2019B111103001]; and [Guangzhou Science and technology Plan Project#3] under Grant [number 201804010383].

Author contributions

TL, WJL, HJZ, JFH, ZHC and GHH contributed equally to this article. WJL, HJZ and WJM conceptualized the paper. TL, JFH, ZHC and GHH analyzed the data, with input from TS, SWC, PW, JLL, YHL, MJC, JXH, JWN, LX, JPX, JXH, LFL, QH, ZHR, APD, WLZ, JSL, XL, XHT, MK, LCG, ZHZ, DXG, GMC, and MRD. TL, WJL, HJZ, TS, JFH, ZHC, GHH and WJM wrote the initial draft with all authors providing critical feedback and edits to subsequent revisions. All authors approved the final draft of the manuscript. WJM is the guarantor. The corresponding author attests

that all listed authors meet authorship criteria and that no others meeting the criteria have been omitted.

Data sharing

No additional data available.

Acknowledgements

We thank all members from health departments and CDCs in Guangdong Province for their contribution in data collection, COVID-19 control and prevention.

References

1. Li Q, Guan X, Wu P, et al. Early Transmission Dynamics in Wuhan, China, of Novel Coronavirus–Infected Pneumonia. *N Engl J Med*, 2020; 382(13):1199-1207. doi: 10.1056/NEJMoa2001316.
2. World Health Organization. Coronavirus disease (COVID-19) Situation Report – 152. 2020. Available at: https://www.who.int/docs/default-source/coronaviruse/situation-reports/20200620-covid-19-sitrep-152.pdf?sfvrsn=83aff8ee_2
3. World Health Organization. Coronavirus disease (COVID-19) Situation Report – 52. 2020. Available at: https://www.who.int/docs/default-source/coronaviruse/situation-reports/20200312-sitrep-52-covid-19.pdf?sfvrsn=e2bfc9c0_4
4. Glasser JW, Hupert N, McCauley MM, et al. Modeling and public health emergency responses: lessons from SARS. *Epidemics*, 2011; 3(1):32-37. doi: 10.1016/j.epidem.2011.01.001.
5. Swanson KC, Altare C, Wesseh CS, et al. Contact tracing performance during the Ebola epidemic in Liberia, 2014-2015. *PLoS Negl Trop Dis*, 2018; 12(9): e0006762. doi: 10.1371/journal.pntd.0006762.
6. Kang M, Song T, Zhong H, et al. Contact Tracing for Imported Case of Middle East Respiratory

Syndrome, China, 2015. *Emerg Infect Dis*, 2016; 22(9):1644-1646. doi: 10.3201/eid2209.152116.

7. Pang X, Zhu Z, Xu F, et al. Evaluation of Control Measures Implemented in the Severe Acute Respiratory Syndrome Outbreak in Beijing, 2003. *JAMA*, 2003; 290(24):3215-3221. doi: 10.1001/jama.290.24.3215.

8. WHO-China Joint Mission Members. Report of the WHO-China Joint Mission on Coronavirus Disease 2019 (COVID-19). WHO; 2020. Available at [https://www.who.int/publications-detail/report-of-the-who-china-joint-mission-on-coronavirus-disease-2019-\(covid-19\)](https://www.who.int/publications-detail/report-of-the-who-china-joint-mission-on-coronavirus-disease-2019-(covid-19)). Accessed 1 March 2020.

9. Hellewell J, Abbott S, Gimma A, et al. Feasibility of controlling COVID-19 outbreaks by isolation of cases and contacts. *Lancet Glob Health*, 2020; 8(4): e488-e496. doi: 10.1016/S2214-109X(20)30074-7.

10. Keeling MJ, Hollingsworth TD, Read JM. The Efficacy of Contact Tracing for the Containment of the 2019 Novel Coronavirus (COVID-19). *medRxiv*. 2020. Available from: <https://doi.org/10.1101/2020.02.14.20023036>.

11. Du RH, Liang LR, Yang CQ, et al. Predictors of mortality for patients with COVID-19 pneumonia caused by SARS-CoV-2: a prospective cohort study. *Eur Respir J*. 2020. 55(5). doi: 10.1183/13993003.00524-2020.

12. Zhou F, Yu T, Du R, et al. Clinical course and risk factors for mortality of adult inpatients with COVID-19 in Wuhan, China: a retrospective cohort study. *Lancet*, 2020; 395(10229):1054-1062. doi: 10.1016/S0140-6736(20)30566-3.

13. Bi Q, Wu Y, Mei S, et al. Epidemiology and transmission of COVID-19 in 391 cases and 1286 of their close contacts in Shenzhen, China: a retrospective cohort study. *Lancet Infect Dis*, 2020; S1473-3099(20):30287-5. doi: [https://doi.org/10.1016/S1473-3099\(20\)30287-5](https://doi.org/10.1016/S1473-3099(20)30287-5).

14. Jing QL, Liu MJ, Zhang ZB, et al. Household secondary attack rate of COVID-19 and associated determinants in Guangzhou, China: a retrospective cohort study. *Lancet Infect Dis*. 2020. doi: [https://doi.org/10.1016/S1473-3099\(20\)30471-0](https://doi.org/10.1016/S1473-3099(20)30471-0).
15. Cheng HY, Jian SW, Liu DP, et al. Contact Tracing Assessment of COVID-19 Transmission Dynamics in Taiwan and Risk at Different Exposure Periods Before and After Symptom Onset. *JAMA Intern Med*. 2020. doi: 10.1001/jamainternmed.2020.2020.
16. National Health Commission of the People's Republic of China. Diagnosis and Treatment Procedure of COVID-19. 2020. Available at: <http://www.nhc.gov.cn/yzygj/s7653p/202003/46c9294a7dfe4cef80dc7f5912eb1989.shtml>. Accessed 5 March 2020.
17. National Health Commission of the People's Republic of China. Prevention and control Scheme of COVID-19. 2020. Available at: <http://www.nhc.gov.cn/jkj/s3577/202003/4856d5b0458141fa9f376853224d41d7.shtml>.
18. Lee P, Hu Y, Chen P, et al. Are children less susceptible to COVID-19? *J Microbiol Immunol Infect*, 2020; S1684-1182(20)30039-6. doi: 10.1016/j.jmii.2020.02.011.
19. Wu Z, McGoogan JM. Characteristics of and important lessons from the coronavirus disease 2019 (COVID-19) outbreak in China: summary of a report of 72314 cases from the Chinese Center for Disease Control and Prevention. *JAMA*, 2020. doi: 10.1001/jama.2020.2648.
20. Xu Y, Li X, Zhu B, et al. Characteristics of pediatric SARS-CoV-2 infection and potential evidence for persistent fecal viral shedding. *Nat Med*, 2020; 26(4):502-505. doi: 10.1038/s41591-020-0817-4.
21. Mehta NS, Mytton OT, Mullins EWS, et al. SARS-CoV-2 (COVID-19): What do we know about children? A systematic review. *Clin Infect Dis*. 2020. doi: 10.1093/cid/ciaa556.

22. Dong Y, Mo X, HuY, et al. Epidemiological Characteristics of 2143 Pediatric Patients With 2019 Coronavirus Disease in China. *Pediatrics*, 2020. doi: 10.1542/peds.2020-0702.
23. Wenham C, Smith J, Morgan R. COVID-19: the gendered impacts of the outbreak. *Lancet*. 2020. 395(10227):846-848. doi: 10.1016/S0140-6736(20)30526-2.24. Boniol M, McIsaac M, Xu L, et al. Gender equity in the health workforce: Analysis of 104 countries Geneva: World Health Organization. 2019. Available from: https://www.who.int/hrh/resources/gender_equity-health_workforce_analysis/en/
25. Pang X, Yang P, Li S, et al. Pandemic (H1N1) 2009 among quarantined close contacts, Beijing, People's Republic of China. *Emerg Infect Dis*, 2011; 17(10):1824-1830. doi: 10.3201/eid1710.101344.
26. Rothe C, Schunk M, Sothmann P, et al. Transmission of 2019-nCoV infection from an asymptomatic contact in Germany. *N Engl J Med*, 2020; 382(10):970-971. doi: 10.1056/NEJMc2001468.
27. Bai Y, Yao L, Wei T, et al. Presumed Asymptomatic Carrier Transmission of COVID-19. *JAMA*, 2020; 323(14):1406-1407. doi: 10.1001/jama.2020.2565.
28. Chan JF-W, Yuan S, Kok K-H, et al. A familial cluster of pneumonia associated with the 2019 novel coronavirus indicating person-to-person transmission: a study of a family cluster. *Lancet*, 2020; 395(10223):514-523. doi: 10.1016/S0140-6736(20)30154-9.
29. Woelfel R, Corman VM, Guggemos W, et al. Clinical presentation and virological assessment of hospitalized cases of coronavirus disease 2019 in a travel-associated transmission cluster. *medRxiv*, 2020. Available from: <https://doi.org/10.1101/2020.03.05.20030502>.
30. Zou L, Ruan F, Huang M, et al. SARS-CoV-2 Viral Load in Upper Respiratory Specimens of Infected Patients. *N Engl J Med*, 2020; 382(12):1177-1179. doi: 10.1056/NEJMc2001737.
31. To KK-W, Tsang OT-Y, Leung W-S, et al. Temporal profiles of viral load in posterior oropharyngeal

saliva samples and serum antibody responses during infection by SARS-CoV-2: an observational cohort study. *Lancet Infect Dis*, 2020; 20(5):565-574.doi: [https://doi.org/10.1016/S1473-3099\(20\)30196-1](https://doi.org/10.1016/S1473-3099(20)30196-1).

Table 1. General characteristics of contacts to COVID-19 cases in Guangdong Province

	n	%
Sex		
Male	6183	53.4
Female	5397	46.6
Age (years)		
0-9	1048	9.0
10-19	819	7.1
20-29	2420	20.9
30-39	2601	22.5
40-49	1878	16.2
50-59	1520	13.1
60-69	831	7.2
70-79	314	2.7
≥80	149	1.3
Places of quarantine		
At home	1855	16.0
Centralized stations	9725	84.0
Contact circumstances		
Family	4893	40.9
Social activities	2016	16.8
Transportation	3198	26.7
Flight	695	5.8
Train	902	7.5
Public transportation*	229	1.9
Private car	213	1.8
The Dream Cruises	64	0.5
Unknown	1095	9.2
Health care institutes	1348	11.3
Others	519	4.3
Relationship with index cases		
Family members	4707	40.7
Spouse	563	4.9
Family members (non-spouse)	1878	16.2
Close relatives	1341	11.6
Other relatives	925	8.0
Social activity contacts	3344	28.9
Transportation contacts	2778	24.0
Health care workers	573	4.9
Others	178	1.5
Infection spectrum of contacts		
No infection	11065	95.6
Asymptomatic infections	66	0.6
Mild confirmed cases	104	0.9
Moderate confirmed cases	300	2.6
Severe confirmed cases	31	0.2
Critically severe confirmed cases	12	0.1
Dead cases	2	<0.01

*: Indicate other public transportations mainly including bus, taxi, subway, ferry, etc.

Table 2. Attack rates of COVID-19 in contacts with different characteristics

Characteristics	Total contacts	Total infections	Attack Rate (%)
Age of contacts (years)			
0-9	1048	60	5.7
10-19	819	33	4.0
20-29	2420	56	2.3
30-39	2601	113	4.4
40-49	1878	56	3.0
50-59	1520	76	5.0
60-69	831	92	11.1
70-79	314	21	6.7
≥80	149	7	4.7
Sex			
Male	6183	213	3.4
Female	5397	302	5.6
Relationship to the index case			
Spouse	563	131	23.3
Family members (non-spouse)	1878	199	10.6
Close relatives	1341	94	7.0
Other relatives	925	38	4.1
Social activity contacts	3344	41	1.3
Transportation contacts	2778	10	0.3
Health care workers	573	2	0.3
Others	178	0	0.0
Contacts on different transportations			
Flight	695	6	0.8
Train	901	11	1.2
Public transportation*	229	5	2.1
Private car	213	9	4.2
The Dream Cruises	63	6	9.5
Unknown	1104	14	1.3

*: Indicate other public transportations mainly including bus, taxi, subway, ferry, etc.

Table 2. Attack rates of COVID-19 in contacts with different characteristics (*continue*)

Characteristics	Total contacts	Total infections	Attack Rate (%)
Disease history of confirmed index cases#			
Incubation period	2211	72	3.3
Symptomatic period	5904	411	7.0
Contacts to the index cases at different time (days to the symptom onset)*			
≤ -5	522	9	1.7
-4 to -3	283	6	2.1
-2 to -1	974	25	2.5
0	1020	61	5.6
1 to 2	1036	81	7.3
3 to 4	865	97	10.1
5 to 6	702	61	8.0
7 to 8	371	31	7.7
9 to 10	223	16	6.7
11 to 12	106	6	5.4
13 to 14	109	4	3.5
15 to 16	188	10	5.1
≥ 17	265	11	4.0
Clinical severity of index case			
Mild	1244	57	4.6
Moderate	5637	344	6.1
Severe	812	52	6.4
Critically severe	371	28	7.5

*: Minus number indicates days before the symptom onset, plus number indicates the days

after the symptom onset in confirmed cases, and zero indicates the day of symptom onset. In

order to precisely estimate the contacting time, only the pairs with only one index case and

one secondary case were included.

Figure legends

Figure 1. Daily numbers of quarantined contacts, and confirmed cases or asymptomatic infections identified from the quarantined contacts in Guangdong Province.

Figure 2. Infected risks of COVID-19 in contacts with different characteristics.

Panel A: In contacts with different ages;

Panel B: In males and females;

Panel C: In contacts who had different relationships to the index case;

Panel D: In contacts exposed to the index cases on different transportations;

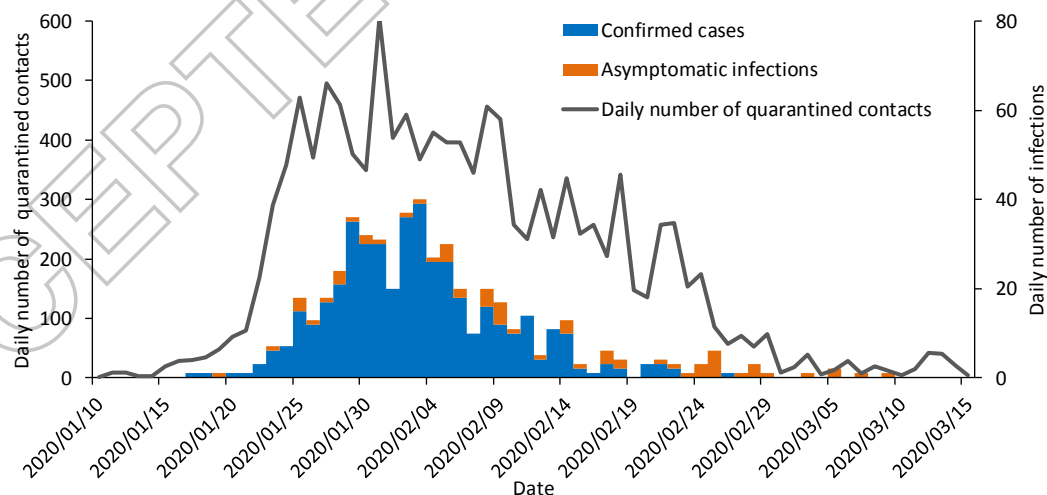
Panel E: In contacts exposed to the index cases at different time;

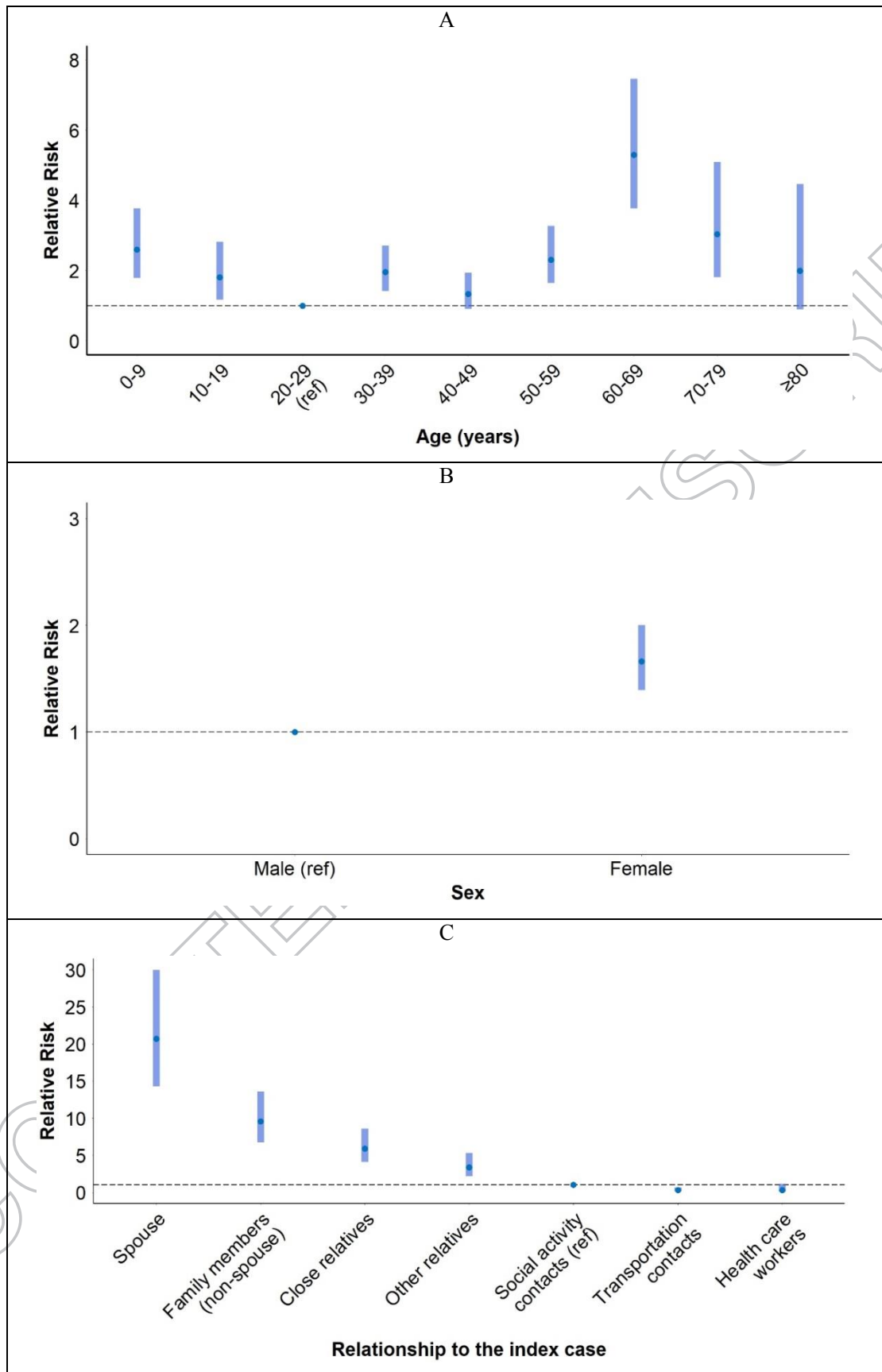
Panel F: In contacts exposed to the index cases in different course of disease;

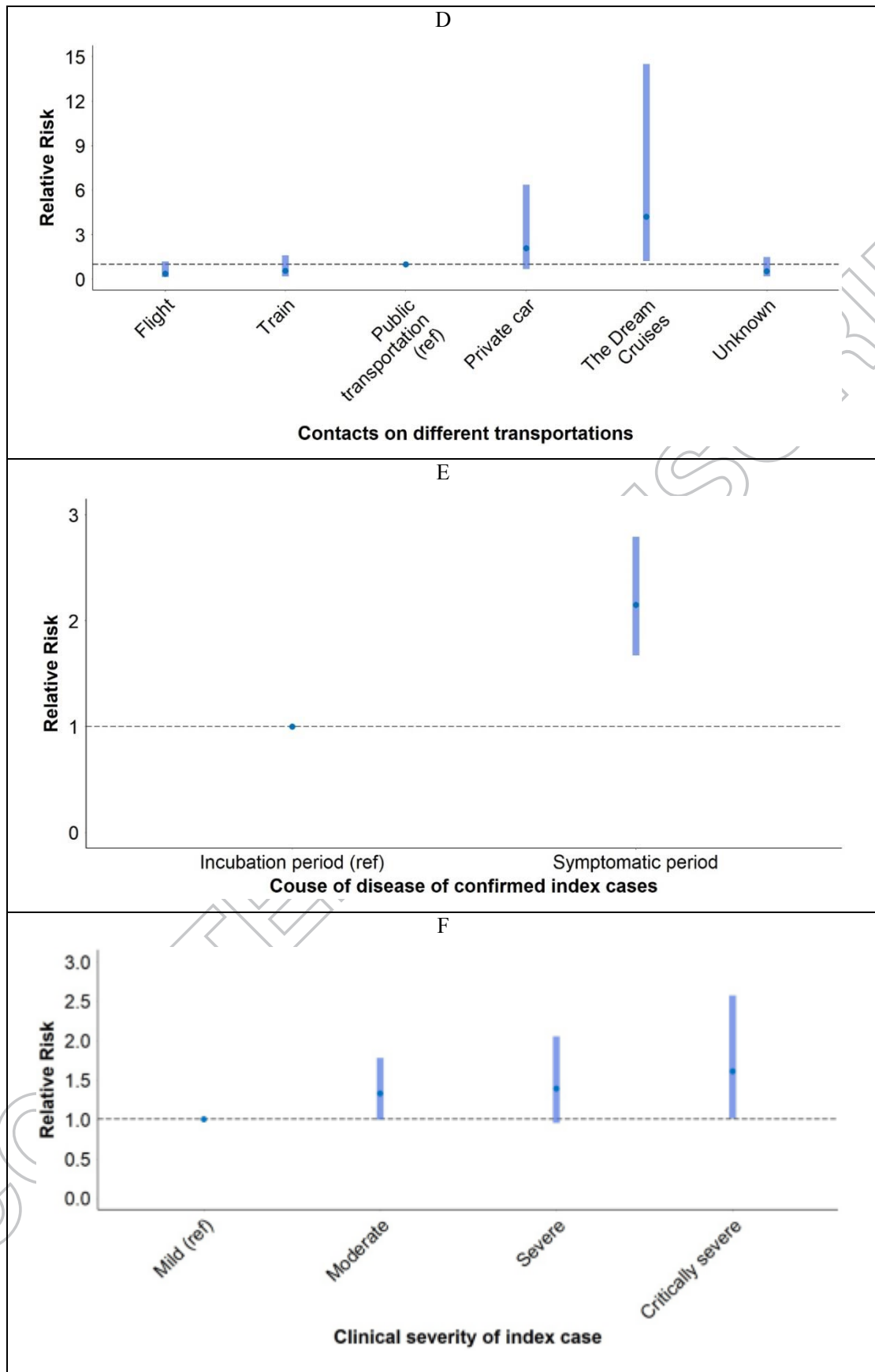
Panel G: In contacts exposed to the index cases with different clinical severity;

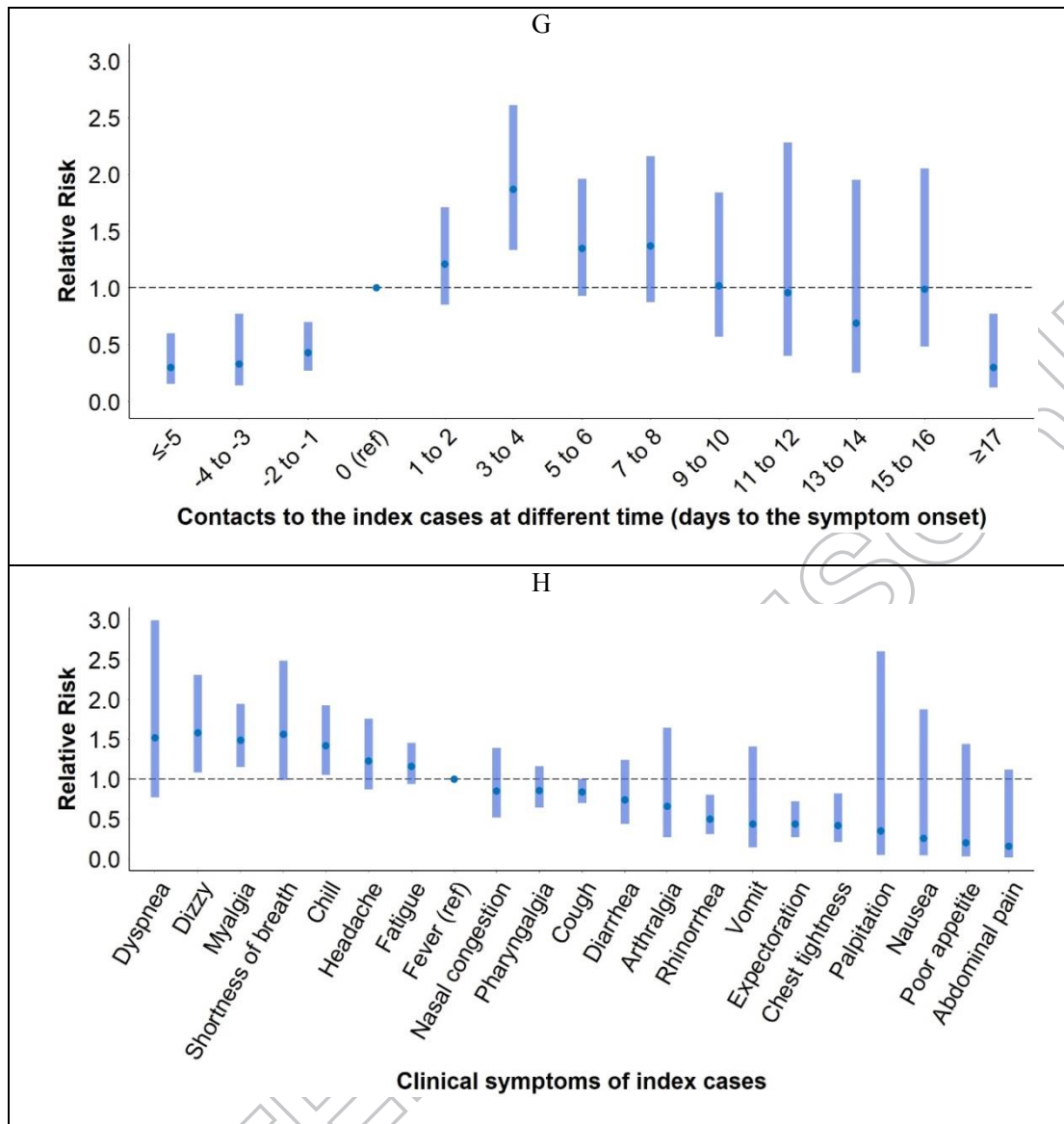
Panel H: In contacts exposed to the index cases with different clinical symptoms.

Adjusted for age and/or sex









Supplementary materials

Risk factors associated with COVID-19 infection: a retrospective cohort study based on contacts tracing

Tao Liu^{1*}, Wenjia Liang^{2*}, Haojie Zhong^{2*}, Jianfeng He^{2*}, Zihui Chen^{1*}, Guanhao He^{1*}, Tie Song², Shaowei Chen¹, Ping Wang¹, Jialing Li², Yunhua Lan², Mingji Cheng², Jinxu Huang², Jiwei Niu², Liang Xia², Jianpeng Xiao¹, Jianxiong Hu¹, Lifeng Lin², Qiong Huang², Zuhua Rong¹, Aiping Deng², Weilin Zeng¹, Jiansen Li², Xing Li¹, Xiaohua Tan², Min Kang², Lingchuan Guo¹, Zhihua Zhu¹, Dexin Gong¹, Guimin Chen¹, Moran Dong¹, Wenjun Ma^{1†}

1. Guangdong Provincial Institute of Public Health, Guangdong Provincial Center for Disease Control and Prevention, Guangzhou, China.

2. Guangdong Provincial Center for Disease Control and Prevention, Guangzhou, China.

*These authors contributed equally to this work

†The author was corresponding author.

Correspondence to:

Dr. Wenjun Ma, Guangdong Provincial Institute of Public Health, Guangdong Provincial Center for Disease Control and Prevention, Guangzhou 511430, China

mawj@gdiph.org.cn

1.1 Definitions of suspect and confirmed cases

Considering both the following epidemiological history and clinical manifestations:

Epidemiological history

- (1) History of travel to or residence in Wuhan and its surrounding areas, or in other communities where cases have been reported within 14 days prior to the onset of the disease;
- (2) In contact with novel coronavirus infected people (with positive results for the nucleic acid test) within 14 days prior to the onset of the disease;
- (3) In contact with patients who have fever or respiratory symptoms from Wuhan and its surrounding area, or from communities where confirmed cases have been reported within 14 days before the onset of the disease; or
- (4) Clustered cases (2 or more cases with fever and/or respiratory symptoms in a small area such families, offices, schools etc. within 2 weeks).

Clinical manifestations

- (1) Fever and/or respiratory symptoms;
- (2) The aforementioned imaging characteristics of novel coronavirus pneumonia (SARS-COV-2);
- (3) Normal or decreased WBC count, normal or decreased lymphocyte count in the early stage of onset.

A suspect case has any of the epidemiological history plus any two clinical manifestations or all three clinical manifestations if there is no clear epidemiological history.

Confirmed cases

Suspect cases with one of the following etiological or serological evidences:

- (1) Real-time fluorescent RT-PCR indicates positive for new coronavirus nucleic acid;
- (2) Viral gene sequence is highly homologous to known new coronaviruses.
- (3) SARS-COV-2 virus specific IgM and IgG are detectable in serum; SARS-COV-2 virus specific

IgG is detectable or reaches a titration of at least 4-fold increase during convalescence compared with the acute phase.

1.2 Definition of close contacts to the index case

A contact is a person who experienced any one of the following exposures during the 2 days before and the 14 days after the onset of symptoms of a probable or confirmed case:

- (1) Face-to-face contact with a probable or confirmed case within 1 meter and for more than 15 minutes.
- (2) Direct physical contact with a probable or confirmed case.
- (3) Direct care for a patient with probable or confirmed COVID-19 disease without using proper personal protective equipment. or
- (4) Other situations as indicated by local risk assessments.

Note: for confirmed asymptomatic cases, the period of contact is measured as the 2 days before through the 14 days after the date on which the sample was taken which led to confirmation.

1.3 Definition of asymptomatic infection

Asymptomatic infection was defined as those whose specimens are detected with positive of SARS-COV-2 virus, but have no clinical symptoms including fever, cough, etc.

Table S1. Characteristics of contact tracing and quarantine of COVID-19 in Guangdong Province

	Parameter (95%CI)
Average number of contacts linked to one index case*	7.8 (7.0, 8.7)
Contacts divided by contact circumstances	
Family	5.3 (4.7, 6.0)
Social activity	12.5 (10.2, 14.7)
Transportation	10.8 (9.3, 12.2)
Health care institute	22.8 (17.4, 28.2)
Others	8.0 (1.9, 14.1)
Contacts divided by relationship to index cases	
Household member	5.1 (4.2, 5.9)
Social activity contact	14.6 (12.3, 16.8)
Transportation contact	12.7 (10.9, 14.5)
Health care worker	15.9 (11.9, 19.9)
Others	11.2 (-2.0, 24.4)
Average days from exposure to quarantine	6.4 (6.2, 6.6)
Average days of quarantine	9.7 (9.6, 9.8)

*: The parameter was estimated in the total contacts.

Table S2. Attack rates of COVID-19 in contacts of index cases with different
clinical symptoms

Clinical symptoms of index cases	Number of contacts	Total infections	Attack Rate (%)
Total	8115	483	6.0
Dyspnea	89	10	11.2
Dizzy	322	34	10.6
Myalgia	760	79	10.4
Shortness of breath	219	22	10.0
Chill	598	54	9.0
Headache	469	39	8.3
Fatigue	1552	122	7.9
Fever	4989	335	6.7
Nasal congestion	305	18	5.9
Pharyngalgia	945	55	5.8
Cough	3996	225	5.6
Diarrhea	320	16	5.0
Arthralgia	117	5	4.3
Rhinorrhea	525	19	3.6
Vomit	85	3	3.5
Expectoration	595	18	3.0
Chest tightness	315	9	2.9
Palpitation	38	1	2.6
Nausea	65	1	1.5
Poor appetite	79	1	1.3
Abdominal pain	121	1	0.8
Chest pain	7	0	0.0

Adjusted for age and sex.

Table S3. Attack rates (%) of COVID-19 in contacts who contacted to the index cases with different ages

Age of contact (years)	Age (years) of the index case (number of infections/number of contacts, %)								
	0-9	10-19	20-29	30-39	40-49	50-59	60-69	70-79	≥80
0-9	0 (0.0)	0 (0.0)	1/88 (1.1)	17/201 (8.5)	3/94 (3.2)	8/114 (7.0)	3/106 (2.8)	0 (0.0)	0 (0.0)
10-19	0 (0.0)	1/24 (4.2)	0 (0.0)	5/101 (5.0)	9/113 (8.0)	2/53 (3.8)	1/69 (1.4)	2/29 (6.9)	0 (0.0)
20-29	0 (0.0)	0 (0.0)	6/357 (1.7)	4/249 (1.6)	9/266 (3.4)	13/259 (5.0)	1/149 (0.7)	1/71 (1.4)	2/26 (7.7)
30-39	4/39 (10.3)	2/27 (7.4)	4/254 (1.6)	25/361 (6.9)	2/259 (0.8)	21/284 (7.4)	19/273 (7.0)	2/54 (3.7)	0 (0.0)
40-49	1/11 (9.1)	2/42 (4.8)	1/185 (0.5)	2/197 (1.0)	12/260 (4.6)	3/199 (1.5)	5/167 (3.0)	11/61 (18.0)	1/27 (3.7)
50-59	0 (0.0)	0 (0.0)	4/197 (2.0)	9/144 (6.3)	4/114 (3.5)	24/271 (8.9)	7/152 (4.6)	4/45 (8.9)	1/27 (3.7)
60-69	0 (0.0)	2/12 (16.7)	1/47 (2.1)	13/103 (12.6)	5/59 (8.5)	12/86 (14.0)	33/168 (19.6)	7/40 (17.5)	0 (0.0)
70-79	0 (0.0)	0 (0.0)	0 (0.0)	2/19 (10.5)	4/38 (10.5)	1/41 (2.4)	3/47 (6.4)	5/39 (12.8)	2/6 (33.3)
≥80	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)	1/25 (4.2)	3/25 (12.0)	3/13 (23.1)	0 (0.0)	0 (0.0)

Table S4. Relative risk (RR) of COVID-19 in contacts who contacted to the index cases with different ages

Age of contact (years)	Age of the index case (years)								
	0-9	10-19	20-29	30-39	40-49	50-59	60-69	70-79	≥80
0-9	-	-	0.36 (0.04, 3.52)	2.81 (0.80, 9.84)	Reference	2.31 (0.59, 8.97)	0.89 (0.17, 4.52)	-	-
10-19	-	0.51 (0.06, 4.23)	-	0.60 (0.19, 1.86)	Reference	0.45 (0.09, 2.17)	0.17 (0.02, 1.37)	0.86 (0.17, 4.21)	-
20-29	-	-	0.51 (0.18, 1.45)	0.49 (0.15, 1.62)	Reference	1.61 (0.67, 3.88)	0.21 (0.03, 1.67)	0.44 (0.05, 3.55)	2.52 (0.51, 12.42)
30-39	12.77 (2.23, 72.87)	10.25 (1.37, 76.71)	2.11 (0.38, 11.66)	8.77 (2.05, 37.45)	Reference	10.28 (2.38, 44.39)	8.39 (1.93, 36.50)	3.97 (0.54, 28.96)	-
40-49	2.05 (0.24, 17.41)	1.07 (0.23, 4.97)	0.11 (0.01, 0.85)	0.21 (0.05, 0.97)	Reference	0.31 (0.09, 1.11)	0.63 (0.22, 1.83)	4.37 (1.82, 10.51)	0.75 (0.09, 6.05)
50-59	-	-	0.51 (0.12, 2.08)	1.65 (0.49, 5.55)	Reference	2.47 (0.83, 7.31)	1.17 (0.33, 4.11)	2.44 (0.58, 10.31)	0.90 (0.10, 8.47)
60-69	-	2.26 (0.38, 13.48)	0.25 (0.03, 2.27)	1.53 (0.51, 4.54)	Reference	1.85 (0.61, 5.61)	2.56 (0.94, 6.93)	2.24 (0.65, 7.68)	-
70-79	-	-	-	1.05 (0.17, 6.39)	Reference	0.21 (0.02, 2.00)	0.57 (0.12, 2.71)	1.19 (0.29, 4.85)	3.95 (0.53, 29.25)

≥ 80	-	-	-	-	Reference	2.78 (0.26, 30.21)	5.71 (0.48, 68.55)	-	-
-----------	---	---	---	---	-----------	--------------------	--------------------	---	---

Adjusted for sex.