

# Evidence for transmission of COVID-19 prior to symptom onset

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## Abstract

We collated contact tracing data from COVID-19 clusters in Singapore and Tianjin, China and estimated the extent of pre-symptomatic transmission by estimating incubation periods and serial intervals. The mean incubation periods accounting for intermediate cases were 4.91 days (95%CI 4.35, 5.69) and 7.54 (95%CI 6.76, 8.56) days for Singapore and Tianjin, respectively. The mean serial interval was 4.17 (95%CI 2.44, 5.89) and 4.31 (95%CI 2.91, 5.72) days (Singapore, Tianjin). The serial intervals are shorter than incubation periods, suggesting that pre-symptomatic transmission may occur in a large proportion of transmission events (0.4-0.5 in Singapore and 0.6-0.8 in Tianjin, in our analysis with intermediate cases, and more without intermediates). Given the evidence for pre-symptomatic transmission it is vital that even individuals who appear healthy abide by public health measures to control COVID-19.

## Introduction

The novel coronavirus disease, COVID-19, was first identified in Wuhan, Hubei Province, China in December 2019 (*Li Q et al., 2020; Huang C et al., 2020*). The virus causing the disease was soon

26 named severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) (*Hui DS et al., 2020*) and  
 27 quickly spread to other regions of China and then across the globe, causing a pandemic with over  
 28 5 million cases and 300,000 deaths at the time of writing (*Johns Hopkins University, 2020*). In Tianjin,  
 29 a metropolis located at the north of China, the first case was confirmed on January 21, 2020 (*Tianjin*  
 30 *Health Commission, 2020*). Two days later, the first case was confirmed in Singapore (*Ministry of*  
 31 *Health Singapore, 2020*), a city country in Southeast Asia. As of February 28, 2020, 93 and 135 cases  
 32 had been confirmed in Singapore and Tianjin (*Ministry of Health Singapore, 2020; Tianjin Health*  
 33 *Commission, 2020*). The first Singapore COVID-19 case was confirmed as an individual who had  
 34 travelled to Singapore from Wuhan. Many of the initial cases were imported from Wuhan, with  
 35 later cases being caused by local transmission. Singaporean officials worked to identify potential  
 36 contacts of confirmed cases; close contacts were monitored and quarantined for 14 days from  
 37 their last exposure to the patient, and other low risk contacts were put under active surveillance  
 38 and contacted daily to monitor their health status. These early outbreaks continue to provide the  
 39 opportunity to estimate key parameters to understand COVID-19 transmission dynamics.

40 We screened publicly available data to identify datasets for two COVID-19 clusters that could be  
 41 used to estimate transmission dynamics. In both Singapore and Tianjin, the COVID-19 outbreak  
 42 occurred within a relatively closed system where immediate public health responses were imple-  
 43 mented, contacts were identified and quarantined, and key infection dates were tracked and up-  
 44 dated daily. With its experiences in control of the SARS outbreak, the Singaporean government  
 45 had been adopting a case-by-case control policy from January 2, 2020. Only close contacts of a con-  
 46 firmed case were monitored and quarantined for 14 days. In Tianjin, a number of COVID-19 cases  
 47 were traced to a department store, where numerous customers and sales associates were likely  
 48 infected. Additional customers who had potential contact were asked to come forward through  
 49 state news and social media, as well as asked if they had visited the department store at various  
 50 checkpoints in the city. All individuals identified as having visited the store in late January were  
 51 quarantined and sections of the Baodi District where the store is located were sealed and put  
 52 under security patrol.

53 We estimate the serial interval and incubation period of COVID-19 from clusters of cases in Singa-  
 54 pore and Tianjin. The serial interval is defined as the length of time between symptom onset in a  
 55 primary case (infecter) and symptom onset in a secondary case (infectee), whereas the incubation  
 56 period is defined as the length of time between an infectee's exposure to a virus and their symp-  
 57 tom onset. Both are important parameters that are widely used in modelling in infectious disease,  
 58 as they impact model dynamics and hence fits of models to data. While the pandemic has pro-  
 59 gressed far beyond these early outbreaks, it remains the case that mathematical modelling, using  
 60 parameters derived from estimates like these, is widely used in forecasting and policy.

61 The serial interval and incubation period distributions, in particular, can be used to identify the ex-  
 62 tent of pre-symptomatic transmission (i.e. viral transmission from an individual that occurs prior  
 63 to symptom onset). There is evidence that pre-symptomatic transmission accounts for a consider-  
 64 able portion of COVID-19 spread (*Arons, MM et al., 2020; Baggett TP et al., 2020; Li et al., 2020*) and  
 65 it is important to determine the degree to which this is occurring (*Peak et al., 2020*). Early COVID-19  
 66 estimates borrowed parameters from SARS (*Wu JT et al., 2020; Abbott S et al., 2020; Jiang X et al.,*  
 67 *2020*), but more recent estimates have been made using information from early clusters of COVID-  
 68 19 cases, primarily in Wuhan. Depending on the population used, estimates for incubation periods  
 69 have ranged from 3.6–6.4 days and serial intervals have ranged from 4.0–7.5 days (*Li Q et al., 2020;*  
 70 *Ki M, 2020; Backer JA et al., 2020; Linton NM et al., 2020; Nishiura H et al., 2020*); however, it is cru-  
 71 cial that the estimates of incubation period and serial interval are based on the same outbreak, and  
 72 are compared to those obtained from outbreaks in other populations. Distinct outbreak clusters  
 73 are ideal for understanding how COVID-19 can spread through a population with no prior expo-  
 74 sure to the virus. Here, we estimate the portion of transmission that is pre-symptomatic based  
 75 on estimates of the incubation period and serial interval. We estimate both quantities under two  
 76 frameworks: first, we use samples as directly as is feasible from the data, for example assuming  
 77 that the health authorities' epidemiological inferences regarding who exposed whom and who was  
 78 exposed at which times are correct. Second, we use estimation methods that allow for unknown  
 79 intermediate cases, such that the presumed exposure and infection events may not be complete.  
 80 We also separate the analysis of incubation period according to earlier and later phases of the  
 81 outbreaks, since measures were introduced during the time frame of the data.

## 82 Results

### 83 Descriptive Analyses

84 Figures 1 and 2 show the daily counts, putative origin of the exposure and individual time courses  
 85 for the Singapore and Tianjin data. In the Singapore dataset, new hospitalization and discharge  
 86 cases were documented daily from January 23 to February 26, 2020. 66.7% (62/93) of the confirmed  
 87 cases recovered and were discharged from the hospital by the end of the study period (Figure 1(a)).  
 88 The disease progression timeline of the 93 documented cases in Figure 1(c) indicates that symptom  
 89 onset occurred  $1.71 \pm 3.01$  (mean  $\pm$  SD) days after the end of possible viral exposure window and  
 90 cases were confirmed  $7.43 \pm 5.28$  days after symptom onset. The mean length of hospital stay was  
 91  $13.3 \pm 6.01$  days before individuals recovered and were discharged.

92 In the Tianjin dataset, new confirmed cases were documented daily from January 21 to February 22,  
 93 2020. 48.1% (65/135) recovered and 2.2% (3/135) had died by the end of the study period (Figure

2(a)). The timeline of the 135 cases is shown in Figure 2(c). Symptom onset occurred  $4.98 \pm 4.83$  (mean  $\pm$  SD) days after the end of the possible viral exposure window. Cases were confirmed  $5.23 \pm 4.15$  days after symptom onset. The duration of hospital stay of the Tianjin cases is unknown as the discharge date of each case was not available. In both datasets, daily counts decline over time, which is likely a combination of delays to symptom onset and between symptom onset and reporting, combined with the effects of strong social distancing and contact tracing.

## Incubation period

In the Singapore dataset, we find that the median incubation period in our direct analysis (without accounting for intermediate cases) is 5.32 days with the gamma distribution; shape 3.05 (95%CI 2.0, 3.84); and scale 1.95 (1.23, 2.34). The mean incubation period is 5.99 (95%CI 4.97, 7.14) days. In Tianjin we find a median 8.06 days; shape 4.74 (3.35, 5.72); scale 1.83 (1.29, 2.04). The mean is 8.68 (7.72, 9.7) days. These results are summarised in Table 1, and we also fitted Weibull and log normal distributions; see Appendix 1 Table 1. These are consistent with, or slightly longer than, previous estimates, see Appendix 1 Table 5 for comparison.

In Singapore, these estimates are based on a combination of cases for whom last possible exposure is given by travel, and later cases (for whom the presumed infector was used). In Tianjin, social distancing measures were implemented during the outbreak. We find that the estimated incubation period is different, particularly in Tianjin, for cases with symptom onset on or prior to January 31st: see Figure 3 and Figure 4. The estimated median incubation period for pre-Feb 1 cases in Tianjin is 6.48 days; the  $q = (0.025, 0.975)$  quantiles are (2.5, 13.3) days. In contrast, post-Jan 31 the median is 12.13 days with  $q = (0.025, 0.975)$  quantiles (7.3, 18.7) days. The means are 6.88 (5.97, 7.87) days for early cases and 12.4 (11.1, 13.7) days for later cases. Social distancing seems unlikely to change the natural course of infection, but these results might be explained if exposure occurred during group quarantine or otherwise later than the last time individuals thought they could have been exposed. Pre-symptomatic transmission would enable this, if an individual was thought to have been exposed before group quarantine, but in actuality was exposed during quarantine by a pre-symptomatic individual. The time interval in the data would then not be a sample of the incubation period, instead it would be a sample of one or more generation times plus an incubation period.

In Singapore we find the same effect, though much less pronounced. The estimated median incubation time is 5.26, with (0.025, 0.975) quantiles of (1.30, 13.8) days for early cases (also defined as cases with symptom onset on or prior to January 31st) and 5.35 (quantiles (1.22, 14.6)) days for late-arising cases. The means are 5.91 (4.50, 7.64) days for early cases and 6.06 (4.70, 7.67) days for later

cases. Fits of gamma and log-normal distributions are similar; see Appendix 1 Table 2. Changes in perception of exposure times after control measures were introduced (i.e. people may assume that they must have been exposed prior to control measures), together with pre-symptomatic transmission, could result in missing intermediate transmission events and hence lengthened incubation period estimates. This in part motivates our analysis with intermediate cases.

Our estimates of the incubation period with intermediates are similar, under the assumption that intermediates are relatively rare. Results are shown in Figure 5 and Table 1. We find that the median of the bootstrapped mean incubation periods for Singapore with a low (0.05 per day) rate of unknown intermediates is 4.91 days (4.35, 5.69 95% bootstrap CI), compared to a generation time of 3.71 (2.36, 4.91) days. The Tianjin bootstrapped mean incubation period is 7.54 (6.76, 8.56 95%CI) days and the generation time is only 2.82 (1.83, 3.52) days. The estimates are lower when the assumed probability of unknown intermediates is higher. Indeed, if intermediates were present between assumed exposure and onset, naturally the generation time would be shorter than if they were not. The mean generation times are consistently shorter than the mean incubation periods, indicating that infection can occur prior to symptom onset. The difference is particularly pronounced in Tianjin, where long intervals were observed.

However, this approach makes a number of assumptions, and is limited by the fact that if we do not know the true infectors then we are also unlikely to know the true exposure. The data we have is well suited to this method in the sense that there were particular events where exposure is thought to have occurred, and so we can account for intermediates in the manner we have done, but we do not have information for the alternative scenario in which the true exposures were prior to those given in the data. This could happen if, for example, individuals were exposed before attending an event or before known contact, and developed symptoms well after it. Exposure would thus be wrongly attributed to the event or contact. We have accommodated this with uncertainty in the exposure intervals, in particular not insisting that individuals who are likely to be the index case for a cluster (e.g., who developed symptoms on the same day as an event) *must* have been exposed then, but instead allowing the possibility that they were exposed earlier.

## Serial intervals

Figure 6 represents the empirical serial intervals between all potential transmission case-pairs as noted in the data and represented in Figure 7, split into groups based on date of first symptom onset for each case-pair. The empirical mean serial intervals shorten in the 'late' group in both Singapore and Tianjin; however, the empirically-derived 95% confidence intervals overlap (Singapore early 4.44 (-2.81, 11.7) vs. late 3.18 (-1.52, 7.88); Tianjin early 5.48 (-0.968, 11.9) vs. late 4.18 (-2.33,

160 10.7)). Negative lower bounds are due to the high standard deviation.

161 Shortening serial intervals are expected as increased quarantine measures are enacted during the  
 162 course of an outbreak and can be an indication of improved control through successful contact  
 163 tracing, as seen in SARS *Lipsitch et al. (2003)*. Our results suggest that serial intervals shortened as  
 164 the outbreak progressed in both clusters, but they could also be due to right truncation. Accounting  
 165 for this, we found that the mean serial intervals were 4 and 5 days (Singapore, Tianjin); a Cox  
 166 regression found no significant difference between the early and late groups' serial intervals. This  
 167 estimate is made directly from case pairs in the data without accounting for intermediate infectors  
 168 and co-primary infection, as in the ICC analysis.

169 Table 1 shows our ICC estimates of the mean and standard deviation for the serial intervals, with  
 170 comparison to other analyses and assumptions in Appendix 1 Table 5. The ICC method finds the  
 171 mean serial interval to be 4.17 (2.44, 5.89 95% bootstrap CI) days (0.882 bootstrap standard devia-  
 172 tion) for Singapore and 4.31 (2.91, 5.72) days (0.716 bootstrap sd) for Tianjin, using the first 4 cases  
 173 in each cluster. This is consistent with the results with right truncation.

## 174 Pre-symptomatic transmission

175 We estimated incubation periods and serial intervals with and without accounting for intermedi-  
 176 ate unknown cases. To estimate the portion of transmission that occurs before symptom onset,  
 177 we compare the "direct" (no intermediate) estimates of each, and the "indirect" (accounting for in-  
 178 termediates) estimates of each. We estimate consistently shorter serial intervals than incubation  
 179 period, suggesting that there is pre-symptomatic transmission.

180 We took the covariation of incubation periods and serial intervals (and of generation times and  
 181 incubation periods) into account by sampling the intervals jointly before estimating the fraction of  
 182 the relevant differences that are negative. Even accounting for correlation, the estimated fraction  
 183 of pre-symptomatic transmission for Singapore is 0.74 (regardless of early/late split) and for Tianjin  
 184 is 0.72, 0.96, 0.81 (early, late, all), based on the direct estimates of the incubation periods and serial  
 185 intervals (see also Figure 8). When we use the incubation period estimates that account for interme-  
 186 diates, the portions pre-symptomatic transmission are 0.53 in Singapore and 0.79 in Tianjin, when  
 187 the assumed "rate of appearance" of intermediates  $r$  is 0.05 (i.e. when we assume a relatively low  
 188 rate of unknown intermediates). If this rate  $r$  is increased, the portion of pre-symptomatic trans-  
 189 mission decreases, but even for  $r = 0.2$  we estimate the pre-symptomatic portion to be 0.38 in  
 190 Singapore and 0.64 in Tianjin.

191 These results were obtained under an estimated correlation between the incubation period and

192 serial interval of 0.289 in Tianjin. If instead the correlation were 0.1, the portion of pre-symptomatic  
 193 transmission in Tianjin under  $r = 0.05, 0.1, 0.15$  and  $0.2$ , respectively, is estimated as 0.783, 0.725,  
 194 0.663 and 0.62. With correlation 0.8, the equivalent portions are 0.849, 0.781, 0.704 and 0.660.  
 195 We therefore find that the degree of positive correlation does not greatly impact our estimates  
 196 of pre-symptomatic transmission. We retain high estimates of the fraction pre-symptomatic in  
 197 Tianjin, due to the long apparent incubation periods. It seems likely that these are an artifact of  
 198 either pre-symptomatic transmission during quarantine/lockdown, or of other assumptions made  
 199 about exposures in the creation of the original dataset. We conclude that overall for this data and  
 200 under reasonable assumptions, we see evidence of at least 65% of transmission occurring before  
 201 symptom onset.

202 In our direct analysis, we estimate that infection occurred on average 1.99 and 3.68 days before  
 203 symptom onset of the infector (Singapore, Tianjin). Because the incubation period is different for  
 204 early- and late-occurring cases in our data, on average transmission for early-occurring cases is  
 205 1.91 and 2.06 days before symptom onset (Singapore, Tianjin) and 1.88, 7.4 days before (Singapore,  
 206 Tianjin) for late-occurring cases. Taking a low rate ( $r = 0.05$ ) of potential unknown intermediate  
 207 cases into account, the mean difference reduces to 0.77 and 3.23 days (Singapore, Tianjin), though  
 208 we still estimate a significant portion of pre-symptomatic transmission (0.53, 0.79), as above.

209 Overall, serial intervals are robustly shorter than incubation periods in our analyses (Table 1).  
 210 These estimates are strengthened by the fact that we have estimated both incubation period and  
 211 serial interval in the same populations and by the fact that we obtain the same result in two dis-  
 212 tinct datasets. In both sets of estimates, samples of the incubation period minus serial interval are  
 213 negative with probability 0.38 or higher (Singapore) and 0.64 or higher (Tianjin), and these lower  
 214 bounds require a high rate of unknown intermediates early in the outbreak. This indicates that  
 215 a substantial portion of transmission may occur before symptom onset (see Appendix 1 and Fig-  
 216 ure 8), consistent with the clinical observations reported by *Rothe C et al. (2020)* and *Bai Y et al.*  
 217 *(2020)*.

218 Shorter serial intervals yield lower reproduction number estimates. For example, if the epidemic  
 219 grows at a rate of 0.15 (doubling time of 4.6 days *Jung S et al. (2020)*, scenario 1), an estimated  
 220 reproduction number using the mean of the bootstrapped estimates is  $R = 1.76$  (1.30, 2.17) with  
 221 a serial interval of 4.17 days (Singapore) and  $R = 1.95$  (1.72, 2.47) with a serial interval of 4.3 days  
 222 (Tianjin). In contrast, if a longer serial interval (7.5 days (*Jung S et al., 2020; Li Q et al., 2020*)) is used,  
 223 the estimate is  $R = 3.05$ . This is based on the relationship between  $R_0$ , serial interval, and growth  
 224 rate, and is a simple estimate that does not take into account a complex and variable natural history  
 225 of infection (*Wallinga J and Lipsitch M (2007)*). It serves primarily to illustrate how our estimated  
 226 serial intervals impact  $R$  in simple models for COVID-19 dynamics.

## Discussion

Here we use transmission clusters in two locations where cases have reported links, exposure and symptom onset times to estimate both the incubation period and serial interval of COVID-19. We make these datasets available in a convenient spreadsheet form; they were available publicly but the Singapore dataset was presented in free text updates and the Tianjin cluster was described on multiple sites and in graphical form, in Chinese. We anticipate that the datasets themselves will remain useful for understanding COVID-19's early spread in these well-documented outbreaks.

The incubation period and serial interval are key parameters for transmission modelling and for informing public health interventions; modelling remains one of the primary policy aids in use in planning local and global COVID-19 responses. Serial intervals, together with  $R_0$ , control the shape and distribution of the epidemic curve (*Anderson RM et al., 2004*). They influence the disease's incidence and prevalence, how quickly an epidemic grows, and how quickly intervention methods need to be implemented by public health officials to control the disease (*Anderson RM et al., 2004; Fraser C et al., 2004*). In particular, the portion of transmission events that occur before symptom onset is a central quantity for infection control (*Fraser C et al., 2004*), and will impact the efficacy of contact tracing and case finding efforts *Peak et al. (2020)*.

Singapore and Tianjin officials both reacted quickly when COVID-19 cases appeared and started implementing contact tracing and containment measures, however there was a dramatic difference in the severity of the measures taken. The first case was identified in Singapore on Jan 23, 2020 and in Tianjin on Jan 21. By Feb 9, Singapore had identified 989 close contacts and implemented a travel advisory to defer all travel to Hubei Province and all non-essential travel to Mainland China, asked travellers to monitor their health closely for two weeks upon return to Singapore, and asked the public to adopt precautions including avoiding close contact with people who are unwell, practicing good hygiene and hand washing, and wearing a mask if you had respiratory symptoms (*Ministry of Health Singapore, 2020*). Comparatively, by Feb 9 in Tianjin, 11,700 contacts were under observations and the Baodi district of almost 1 million people was placed under lockdown with restrictions including: one person per household could leave every two days to purchase basic needs, public gatherings were banned, no one could leave their homes between 10PM and 6AM without an exemption, entrances to Tianjin were put under control, and all the buses linking nearby provinces and cities were halted ([www.chinadaily.com](http://www.chinadaily.com)). While Singapore contained the virus spread relatively well until mid-March, they reached 500 confirmed cases on March 23, 1000 cases on April 1, 10,000 cases on April 22, and 25,000 cases on May 13 (*Ministry of Health Singapore, 2020*); Tianjin province began to flatten their epidemic curve by mid- to late-February and had plateaued at 192 confirmed cases as of May 19 ([github.com/CSSEGISandData/COVID-19](https://github.com/CSSEGISandData/COVID-19)).



261 In Singapore and Tianjin we estimated relatively short serial intervals. Of particular note, early  
 262 estimates of  $R_0$  for COVID-19 used the SARS serial interval of 8.4 days (*Wu JT et al., 2020; Abbott S*  
 263 *et al., 2020; Majumder M and Mandl KD, 2020*). Our serial interval findings from two populations  
 264 mirror those of *Zhao S et al. (2020)* and *Nishiura H et al. (2020)*, who estimated a serial interval  
 265 of 4.4 and 4.0 days. Du et al (*Du Z et al., 2020*) obtain a similar estimate for the serial interval  
 266 (3.96 days with 95%CI: 3.53-4.39) but with standard deviation 4.75 days, based on 468 cases in 18  
 267 provinces. Furthermore, we estimate the serial interval to be shorter than the incubation period in  
 268 both clusters, which suggests pre-symptomatic transmission. This indicates that spread of SARS-  
 269 CoV-2 is likely to be difficult to stop by isolation of detected cases alone. However, shorter serial  
 270 intervals also lead to lower estimates of  $R_0$ , and our serial intervals support  $R_0$  values just below  
 271 2; if correct this means that half of the transmissions need to be prevented to contain outbreaks.

272 We stratified the incubation period analysis for Tianjin by time of symptom onset (pre- or post-  
 273 Jan 31, 2020; motivated by quarantine/social distancing measures) and found that the apparent  
 274 incubation period was longer for those with post-quarantine symptom onset. The reason for this  
 275 is unclear, but one possible explanation is that there were (unknown, therefore unreported) expo-  
 276 sures during the quarantine period. If people are quarantined in groups of (presumed) uninfected  
 277 cases, pre-symptomatic transmission in quarantine would result in true exposure times that are  
 278 more recent than reported last possible exposure times.

279 Although it may seem contradictory that e.g., Singapore's efforts were able to keep the epidemic  
 280 under control using mainly case-based controls if pre-symptomatic transmission is common, it re-  
 281 mains the case that detailed contact tracing combined with case finding may be key to limiting both  
 282 symptomatic and pre-symptomatic spread. In Singapore, symptom-free close contacts of known  
 283 cases were quarantined preemptively for 14 days, and other less high risk contacts were placed  
 284 under phone surveillance *Lee et al. (2020)*. In addition, if case finding is able to prevent a large  
 285 portion of symptomatic transmission, it seems logical that the remaining observed transmission  
 286 may be pre-symptomatic. The large extent of pre-symptomatic spread that is occurring, however,  
 287 may be one reason that the spread of COVID-19 in Singapore was ultimately only delayed and not  
 288 prevented.

289 There are several limitations to this work. First, the times of exposure and the presumed infectors  
 290 are uncertain, and the incubation period is variable. We have not incorporated uncertainty in the  
 291 dates of symptom onset. We have used the mixture model approach for serial intervals to avoid  
 292 assuming that the presumed infector is always the true infector, but the mixture does not capture  
 293 all possible transmission configurations. Our  $R_0$  estimates are simple, based on a doubling time of  
 294 4.6 days, and could be refined with more sophisticated modelling in combination with case count  
 295 data. We have not adjusted for truncation (e.g. shorter serial intervals are likely to be observed

first) or the growth curve of the epidemic. However, the serial interval estimates are consistent between the two datasets, are robust to the parameter choices, and are consistently shorter than the estimated incubation times.

We identified both the incubation period and the serial interval in Singapore and Tianjin COVID-19 clusters. Our results suggest that there is substantial transmission prior to onset of symptoms, as the serial interval is shorter than incubation period by 2-4 days. We find differences in estimated incubation period between early and later cases; this may be due to pre-symptomatic transmission or differences in reporting and/or in perceived exposure as the outbreak progressed, in the context of social distancing measures. Evidence of transmission from apparently healthy individuals makes broad-scale social distancing measures particularly important in controlling the spread of the disease.

## Materials and Methods

### Data

All datasets and R code are available on GitHub ([github.com/carolinecolijn/ClustersCOVID19](https://github.com/carolinecolijn/ClustersCOVID19)).

Singapore data was obtained from the Ministry of Health Singapore (*Ministry of Health Singapore, 2020*) online press releases. The Singapore dataset comprised 93 confirmed cases from the date of the initial case on January 23, 2020 until February 26, 2020. Tianjin data was obtained from the Tianjin Health Commission (*Tianjin Health Commission, 2020*) online press releases. The Tianjin dataset comprises 135 cases confirmed from January 21 to February 22, 2020. The symptom onsets were available on the official website for all but a few patients who had not had symptoms before being diagnosed at a quarantine center. Both datasets contained mainly information on exposure times, contacts among cases, time of symptom onset (See Appendix 1 for column descriptions and data processing).

### Statistical analysis

All statistical analyses were performed using R (*R Core Team, 2013*).

**Incubation periods: not accounting for intermediate cases.** The daily incidence of hospitalization and mortality was plotted with the cumulative number of cases confirmed and discharged. The daily incidence was also visualized by date of symptom onset. For the symptom onset plots,

any cases that did not have information on date of onset of symptoms were removed. Cases were then grouped based on their assumed source of infection (see Appendix 1 for full details).

Incubation periods were estimated in two ways: directly from the exposure to symptom onset times, and using a model allowing for unknown intermediate cases to have been the true source of infection (see below). The direct estimates were based on the earliest and latest possible exposure times, and on the reported times of symptom onset. It is impossible to confirm the exact times of exposure and thus we used interval censoring, which uses the likelihood of a time falling in a defined window, (R package *icenReg* (*Anderson-Bergman, 2017*)) to make parametric estimates of the incubation period distribution. For cases without a known earliest possible exposure time, we assume that the case must have been exposed within the 20 days prior to their symptom onset. For cases without a known latest possible exposure time, we assume that exposure had to have occurred before symptom onset. Some cases had a travel history or contact with a known location or presumed source of the virus and this defined their window for exposure. In the Singapore dataset, other individuals had estimated exposure times based on the symptom times for their presumed infector. For these, we define an exposure window using the symptoms of their presumed infector  $-7/ + 4$  days. Having defined exposure windows, we proceed with interval censoring. In both datasets we stratified the data according to whether symptom onset occurred early or late, and estimated incubation periods separately. We define 'early' cases as those with symptom onset on or prior to January 31st.

**Incubation periods: accounting for intermediate cases.** Standard estimates of the incubation period from exposure and symptom data require knowledge of the true exposure event. In our data, exposures were frequently attributed to attendance at events or locations where there had been known COVID-19 cases. It is conceivable that some cases were not in fact exposed at the event, but subsequent to it, by an unknown (perhaps asymptomatic) case who also attended the event or was otherwise linked. We developed the following approach to account for possible unknown intermediates. Suppose the data suggest that case  $i$  was exposed at an event by individual  $A$ , but in fact, there is an unknown intermediate  $x$  who was infected at the event and who subsequently infected  $i$ . In this case, the time between  $i$ 's apparent exposure and  $i$ 's symptom onset is not a sample of the incubation period. Instead, it is one generation time (the time between  $A$  infecting  $x$  and  $x$  infecting  $i$ ) followed by one incubation period (from  $x$  infecting  $i$  to  $i$ 's symptoms). Similarly, if  $x$  infects a second unknown intermediate  $y$ , and  $y$  infects  $i$ , then the time elapsed is two generation intervals followed by an incubation period. Under the simplifying assumption that the generation time and the incubation period follow a gamma distribution with the same scale parameter, we can explicitly write the density for the elapsed time, given  $k$  intermediates. We model the assumption that longer times between (presumed) exposure and symptom onset have more

room for undetected intermediate cases. To describe this with likelihoods, we model unknown intermediate cases occurring with a probability proportional to the length of the apparent incubation period, using a Poisson process (see Appendix 1). We estimate the mean incubation period and generation interval with this approach, also accounting for right truncation (which is not available in the interval censoring estimator in `icenReg`). If  $f(t)$ ,  $g(t)$  are the densities for the incubation period and generation time respectively, then with  $k$  intermediates, the time elapsed has density  $h_k(t) = g * \dots * g * f = g^{(k)} * f$ , where  $*$  denotes convolution, i.e.,  $g * f = \int_0^t g(s)f(t-s)ds$ . The right truncation time  $T_i$  is the time between  $i$ 's exposure and the end of the observation period (because if the symptom onset does not happen after  $T_i$  has elapsed it will not be observed, and this can bias estimates). Let the time from symptom onset to the beginning of the exposure window be  $t_{max}^i$ , and to the end of the window  $t_{min}^i$ . The incubation period is then somewhere in the interval  $(t_{min}^i, t_{max}^i)$ . The likelihood of observing a time in this interval, conditional on  $k$  intermediates, is  $L_k^i = \frac{H_k(t_{max}^i) - H_k(t_{min}^i)}{H_k(T_i)}$ . We use a Poisson process with rate  $r$  to model the probability that there are  $k$  intermediates. This means that the likelihood for the  $i$ 'th observation is  $L^i = \sum_{k=0}^3 p_k L_k^i$ . The complete likelihood is the product over all cases,  $L = \prod_i L^i$ . To compute this, note that if  $g$  and  $f$  are both gamma densities with shapes  $a_g$  and  $a_i$ , and if they have the same scale parameter  $b$ , then the convolution  $g * f(t) = \text{Gamma}(a_g + a_i, b)$ . We can extend this to  $k$  intermediates: the density is  $g^{(k)} * f = \text{Gamma}(ka_g + a_i, b)$ . We truncate the number of possible intermediates at  $n$ , so we condition the usual Poisson probability for  $k$  arrivals,  $\rho_{k,r} = r^k e^{-r} / k!$ , accordingly. Let  $C_{n,r} = \sum_{i=0}^n p_i(r)$ , and use

$$p_k = \begin{cases} \rho_{k,r} / C_{n,r}, & k \leq n \\ 0 & \text{otherwise} \end{cases} \quad (1)$$

We use maximum likelihood to estimate the shape parameters  $a_g$  and  $a_i$  of the generation and incubation periods under a range of intermediate "arrival rates"  $r$  and we use bootstrapping to estimate the credible intervals. We refer to this analysis as the "incubation period with intermediates" analysis.

**Serial intervals: not accounting for intermediates** We illustrate the empirical serial intervals implied by contact links reported in the data. We compute the mean and standard deviation of these in entirety, and separated into early- and late-occurring cases, calculating summary statistics of possible transmission pairs in 'early' (i.e., first date of symptom onset on or before Jan 31, 2020) vs. 'late' portions of both clusters. We estimate the mean serial intervals using these "directly reported" contacts, accounting for right truncation (R package `SurvTrunc`) and using Cox proportional hazards to determine whether there is a significant early vs. late difference. We use the non-parametric survival curves to estimate the mean serial interval for both datasets.

**Serial intervals: accounting for intermediates** As with incubation periods, reported serial inter-

vals may miss unknown intermediates, and co-infectors of two cases presumed to be a transmission pair. We used the expectation-maximization approach described in Vink et al (Vink, MA et al., 2014), which not only takes unknown intermediates into account but also explicitly models a fixed set of possible mis-allocation of infector-infectee pairs in contact data. Briefly, this approach assigns the case with earliest symptom onset in a cluster a "putative index" (PI) status, and uses the time difference between symptom onset of subsequent cases in the cluster and the putative index as "index case to case" (ICC) intervals for putative index cases in small, closely-linked sets of cases ("small clusters"). The ICC intervals are the time differences between the symptom onset time  $t_{pi}$  of the putative index (PI) case and the other members' symptom onset (call these times  $t_j$ , where  $j$  is another case in the same small cluster as this PI). These intervals are not samples of the serial interval distribution, because it need not be the case that the PI infected the others. Vink et al Vink, MA et al. (2014) used a mixture model in which ICC intervals  $t_j - t_{pi}$  can arise in four ways: (1) an outside case infects PI and  $j$ ; (2) PI infects  $j$ ; (3) PI infects an unknown who infects  $j$  and (4) PI infects unknown 1 who infects unknown 2 who infects  $j$ . Accordingly, if the serial interval  $x \sim \mathcal{N}(\mu, \sigma^2)$ , the density for the ICC intervals is

$$f(x; \mu, \sigma^2) = \sum_i w_i f_i(\mu, \sigma^2)$$

where  $w_i$  are weights of the  $i$ 'th component density and  $f_i$  are the component densities for the  $i$ 'th transmission route. Expectation-maximization is used to determine  $\mu$  and  $\sigma$  (See Vink, MA et al. (2014) for more details).

For each dataset we create a network, with individuals represented by nodes. The network's edges are the reported direct contacts between individuals. Every such network (or graph) consists of one or more components – sets of nodes that are connected by edges. We use the components of the network to define transmission clusters. Since the four models in the mixture are likely insufficient to model the transmission in large clusters, we restrict the analysis to only the first 4 cases per cluster (or the first 3, 5, or 6 cases per cluster to determine impact of altering number of cases per cluster; see Appendix 1). We defined the first case within the cluster as the case with the earliest date of symptom onset within the cluster; however we also examined the impact of using the earliest end exposure time if the first symptomatic case was not the index case for the cluster (See Appendix 1). Given the serial interval, we calculate an approximate reproduction number using the empirical growth rate ((Wallinga J and Lipsitch M, 2007):  $R = \exp r\mu - 1/2r^2\sigma^2$ , where  $r$ ,  $\mu$  and  $\sigma$  are the exponential growth rate, the mean serial interval and the standard deviation of the serial interval, respectively). To obtain confidence intervals for  $R$  we resample  $\mu$  and  $\sigma$  using bootstrapping.

**Pre-symptomatic transmission** We estimate the portion of transmission that occurs before symptoms as the fraction of samples where serial interval minus incubation period is negative. We in-

427 introduce an approach to take covariation between the two variables into account, as follows. The  
428 mean difference between two random variables is the difference between the means. Therefore,  
429 the mean serial interval minus the mean incubation periods gives an estimate of the mean time  
430 before symptoms that transmission occurs according to our data. However, the distribution of the  
431 difference depends on the covariance between the incubation period and the serial interval. Un-  
432 fortunately it is challenging to obtain a good estimate of the covariance between these quantities.  
433 We estimated the covariance (and correlation) using case pairs; each pair is associated with two  
434 numbers: a serial interval estimate and an incubation period for the infectee. The covariance be-  
435 tween these is a (somewhat crude) estimate of the covariance in the incubation period and serial  
436 intervals. We sampled incubation periods and serial intervals from our estimated distributions, en-  
437 suring that we respected the observed correlation, and used the serial interval - incubation period  
438 differences to estimate the portion of transmission that is pre-symptomatic. Further details of this  
439 approach are given in Appendix 1.

440 In estimating pre-symptomatic transmission, we compare "direct" (not accounting for intermedi-  
441 ates) incubation periods and serial intervals, and we compare the two *with* accounting for interme-  
442 diates. We take the covariation into account throughout.

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## 448 **Competing Interests**

449 Authors have no competing interests to disclose.

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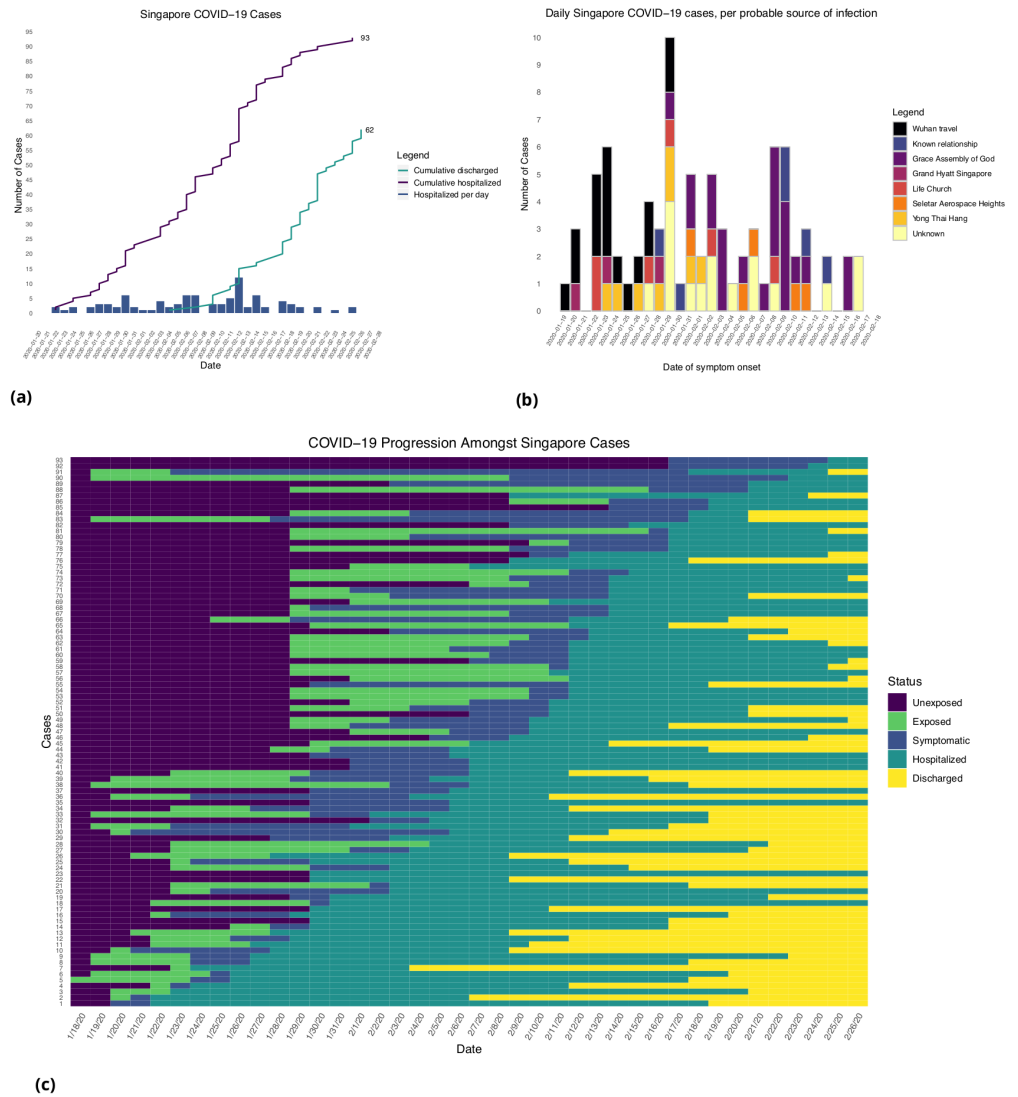
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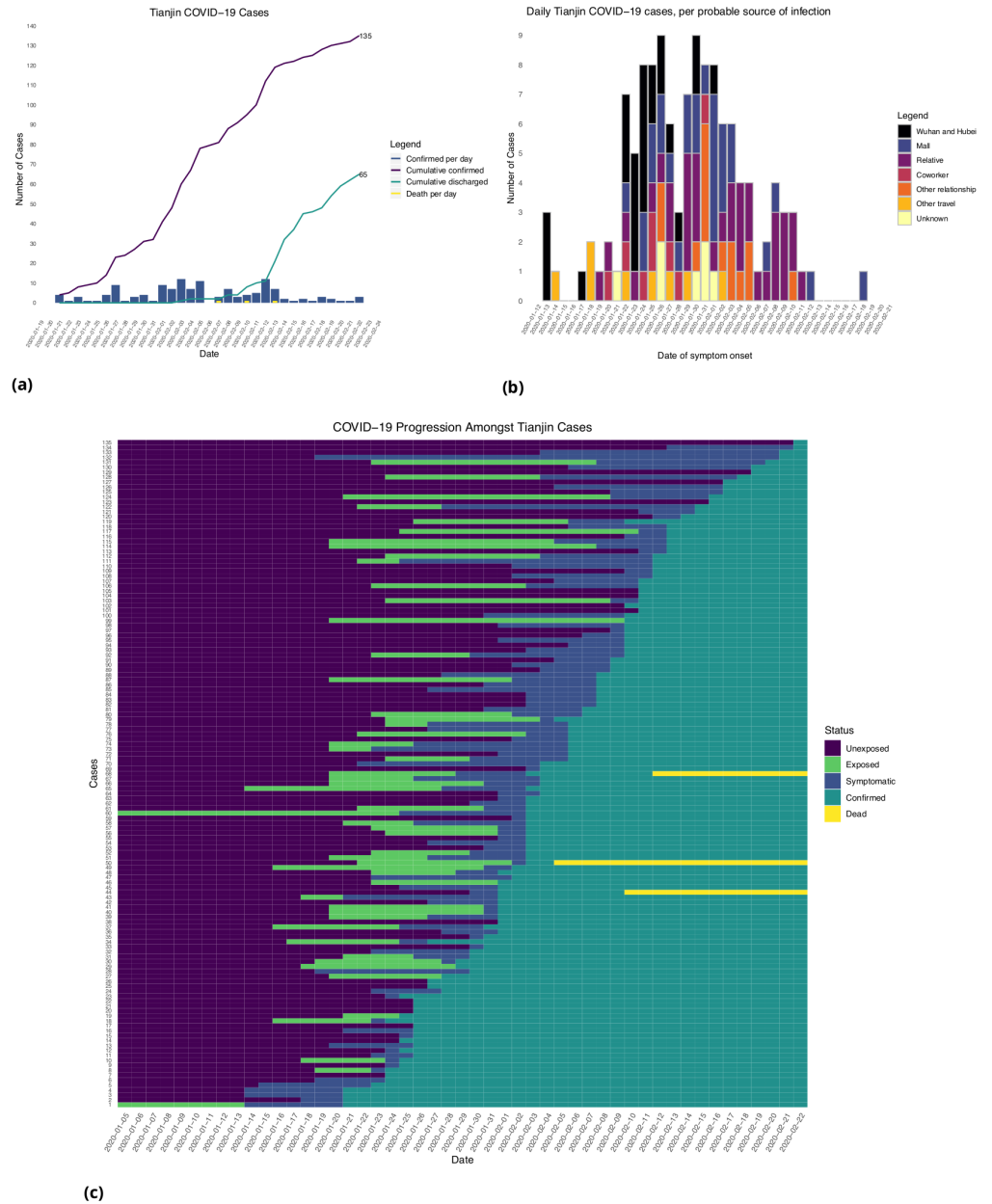
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**Table 1.** Mean incubation period, serial interval and pre-symptomatic transmission. Incubation periods are based on the gamma estimates because these are the most convenient for taking the covariation of serial intervals and incubation periods into account (done throughout the table). 95% CIs are provided in brackets.

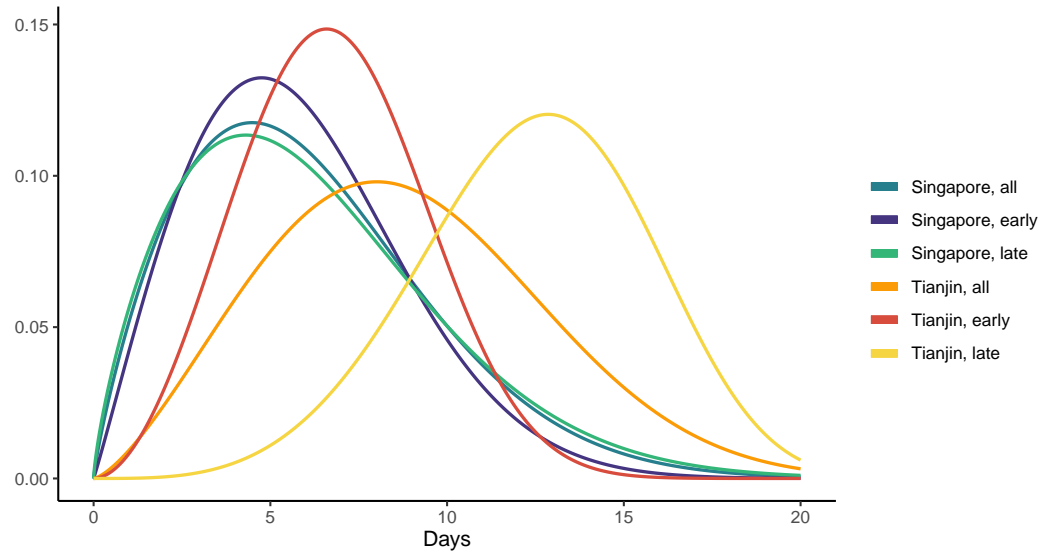
<b>Without intermediates</b>	Incubation (days)	Serial interval (days)	Mean difference (days)	Portion pre-symptomatic (-)
Singapore (all)	5.99 (4.97, 7.14)	4.0 (2.73, 5.57)	1.99	0.74
Singapore (early)	5.91 (4.50, 7.64)		1.91	0.742
Singapore (late)	6.06 (4.70, 7.67)		2.06	0.744
Tianjin (all)	8.68 (7.72, 9.7)	5.0 (3.82, 6.12)	3.68	0.81
Tianjin (early)	6.88 (5.97, 7.87)		1.88	0.72
Tianjin (late)	12.4 (11.1, 13.7)		7.4	0.96
<b>Account for intermediates</b>				
Singapore $r = 0.05$	4.91	4.17 (2.44, 5.89)	0.77	0.53
Singapore $r = 0.1$	4.43		0.26	0.46
Singapore $r = 0.15$	4.12		-0.05	0.41
Singapore $r = 0.2$	3.89		-0.28	0.38
Tianjin $r = 0.05$	7.54	4.31 (2.91, 5.72)	3.23	0.79
Tianjin $r = 0.1$	6.89		2.58	0.74
Tianjin $r = 0.15$	6.30		1.99	0.67
Tianjin $r = 0.2$	5.91		1.6	0.64



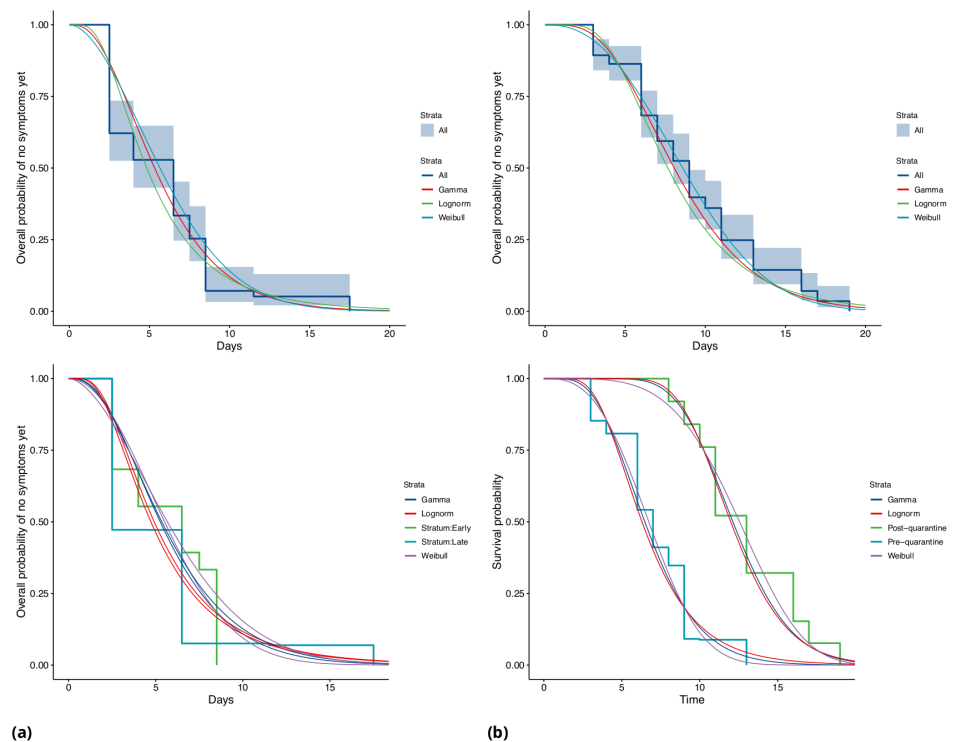
**Figure 1.** Singapore COVID-19 cases. (a) Daily hospitalized cases and cumulative hospitalized and discharged cases. (b) Daily incidence with probable source of infection. (c) Disease timeline, including dates at which each case is unexposed, exposed, symptomatic, hospitalized, and discharged. Not all cases go through each status as a result of missing dates for some cases.



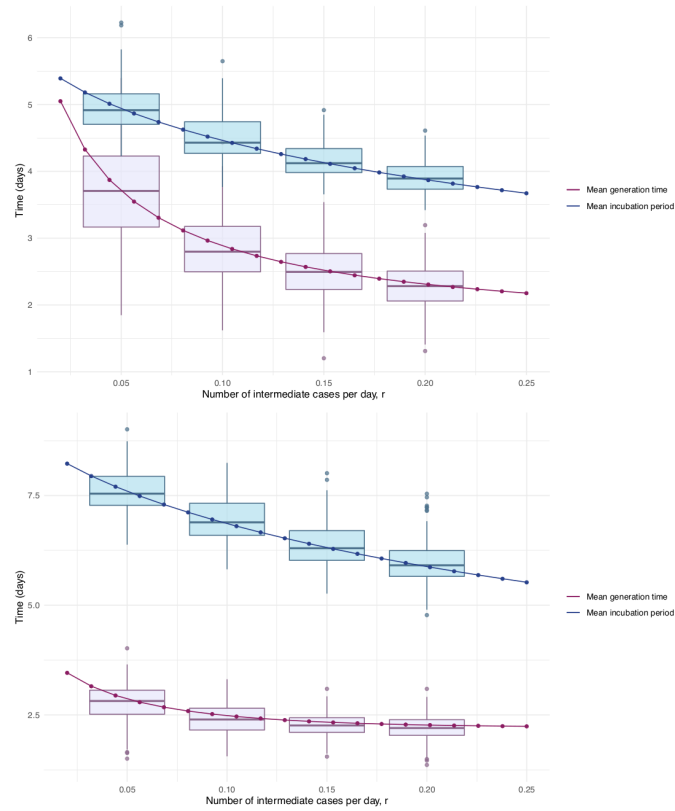
**Figure 2.** Tianjin COVID-19 cases. (a) Daily and cumulative confirmed cases, cumulative discharges and daily death cases. (b) Daily incidence with probable source of infection. (c) Disease progression timeline; not all cases go through each status as a result of missing dates for some cases.



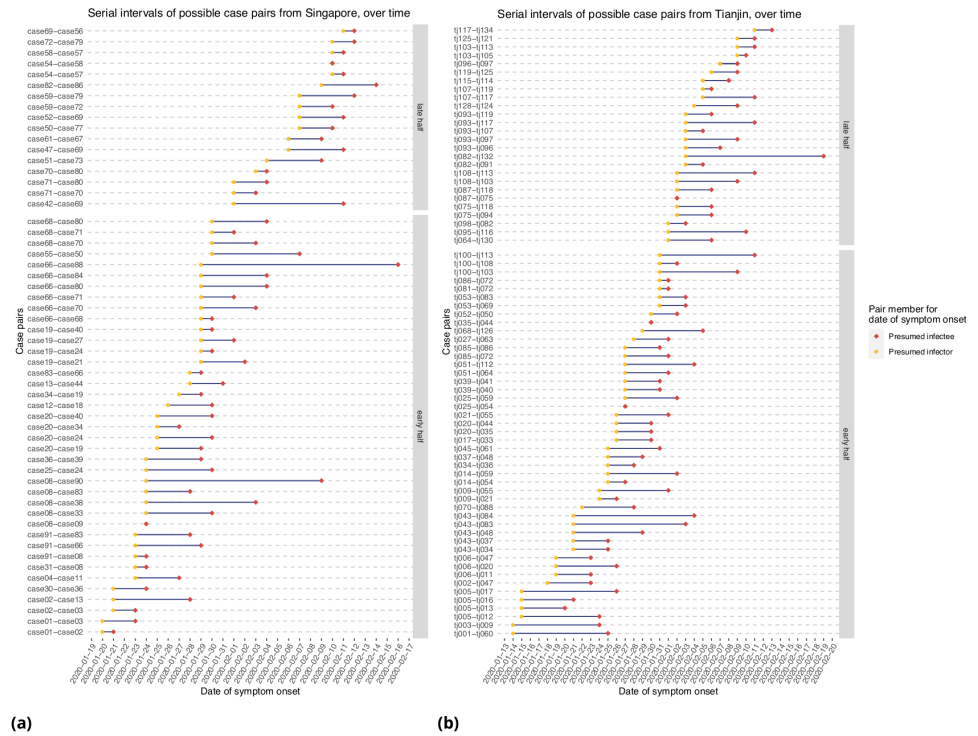
**Figure 3.** Fitted gamma COVID-19 incubation period distributions (without intermediates). Cases are defined as 'early' if they have symptom onset on or prior to January 31, and are classified 'late' otherwise.



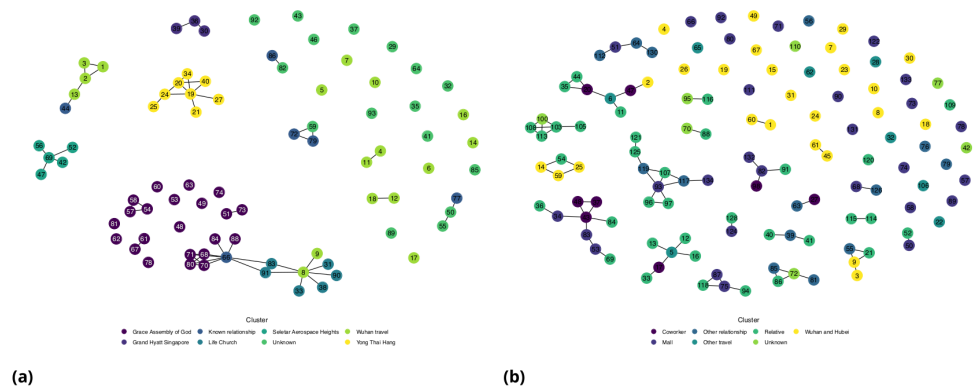
**Figure 4.** COVID-19 incubation period Kaplan-Meier curves for (a) Singapore and (b) Tianjin. Top panels show unstratified data (all cases with symptom onset given). Bottom panels show 'early' and 'late' cases, where early cases are defined as those with symptom onset on or prior to January 31, and late otherwise.



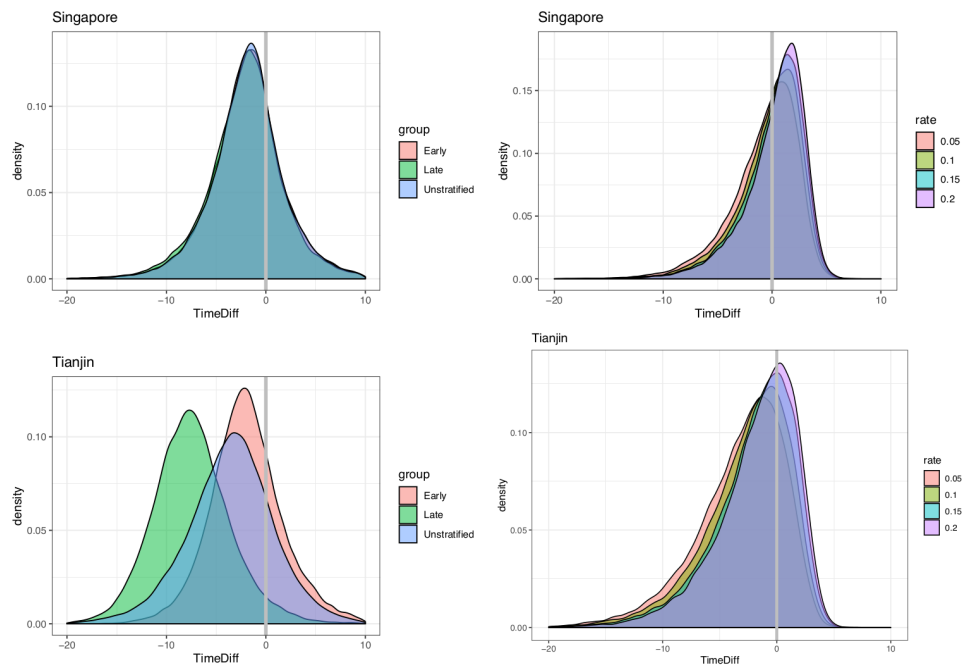
**Figure 5.** Mean incubation period and generation time estimates from the incubation period intermediates analysis, under the assumption that the scale parameter for both distributions is equal, shown with dependence on the mean number of unknown intermediate cases per day of the empirical time elapsed between exposure and symptom onset. The incubation period is longer than the generation time, so this analysis suggests that symptom onset occurs after infectiousness begins. Top: Singapore. Bottom: Tianjin. The means are the scale times the shape, which is fixed at 2.1 in Singapore and 2.2 in Tianjin. Varying this fixed value for the shape parameter was not found to significantly impact the results.



**Figure 6.** Serial intervals of possible case pairs in (a) Singapore and (b) Tianjin. Pairs represent a presumed infector and their presumed infectee plotted by date of symptom onset. Cases are defined as 'early' if they have symptom onset on or prior to January 31st.



**Figure 7.** Network diagram for (a) Singapore (b) Tianjin



**Figure 8.** Pre-symptomatic infection as estimated by samples of (serial interval - incubation period), accounting for covariation. Top: Singapore. Bottom: Tianjin. Left: without intermediates. Right: accounting for intermediates. Grey vertical line: 0. Samples below zero indicate pre-symptomatic transmission. In all cases there is substantial pre-symptomatic transmission.



## Appendix 1

### Details of the Singapore and Tianjin datasets

#### The Singapore dataset

In the Singapore dataset: "related cases" are direct known contacts between cases; "cluster links" are cases that are linked together through an identified cluster event; "presumed infected date" and "presumed reason" are the earliest known date and the reason that each case was known to likely be infected; "last poss exposure" and "contact based exposure" are sub-classifications of "presumed infected date," representing either the last date that each case could have been infected – the date of arrival in Singapore for travellers from Wuhan – or the date that each case was likely infected during a local transmission event in Singapore, respectively; "cluster" is the Ministry of Health Singapore's classification of cases into transmission cluster events.

These data inform the "start\_source" and "end\_source" columns which encode the earliest and latest possible dates of case exposure. For example, we assume that those travelling from Wuhan were exposed before travel (due to evidence of lack of community transmission in Singapore at the time), and that those cases associated with a particular event or location (e.g., Grace Assembly gatherings, the visit to the Yong Thai store by a tour group from Wuhan) were not exposed prior to that event. For the latter, we set "end\_source" to the date of the event + 4 days, to allow for some uncertainty and the possibility of an intermediate infector. For cluster cases thought to originate from a particular index case but lacking information on dates of contact, "start\_source" is set to the first symptom onset in the cluster - 7 days. The "end\_source" is set assuming that once a case in a cluster was identified, people were well aware of this and ceased mixing within the group; thus "end\_source" is the minimum of the earliest quarantine, hospitalization or symptom onset in the cluster, and the symptom onset date of the case in question.

In the absence of other information, we set the "start\_source" of a case to their symptom onset date - 20 days (to allow for a wide range of epidemiologically feasible incubation periods), and the "end\_source" to their symptom onset (since all cases must be exposed before they show symptoms).

All cases in the Singapore dataset were categorized into an infection source group based on information provided in the "presumed reason" column without conflict. The group designations were not used in the statistical estimates.

## The Tianjin dataset

In the Tianjin cases summary spreadsheet, the main columns are: gender, age, symptom onset, symptom type, confirmation date, severity and death date (*Tianjin Health Commission (2020)*); detailed information from daily reports for the first 80 patients provided travel or exposure history and contact information, from which we obtained exposure windows (start source, end source). For backup and to complete missing information for later cases we also referred to *Jinyun News (2020)*, Tianjin official local media, who used Baodi local government reports (*Baodi district of Tianjin (2020)*). They reported detailed activity for those confirmed cases when their corresponding epidemiological history investigation was finished.

The “start\_source” and “end\_source” columns were defined similarly to the Singapore dataset where possible, with reasoning provided in the “Infection\_source” column and further explanation in “recorrection for start and end source”. In most cases, start and end times in the Tianjin dataset were defined by known windows of contact with other individuals with confirmed COVID-19 infections, the Baodi shopping mall or travel to areas with higher levels of infection such as Wuhan. Again, in the absence of other information, we set the “start\_source” of a case to their symptom onset date - 20 days and the “end\_source” to their symptom onset.

Cases in the Tianjin dataset were categorized into an infection source group based on information provided in the the “Infection source” column. There were a small number of cases ( $n = 12$ ) that could be classified into two possible infection source groups (e.g. from Wuhan and has a close relationship with another known case). These cases were assigned their infection source groups based on the following hierarchy of possible sources: (highest priority) Wuhan or Hubei origin > Mall (for shoppers, workers, or individuals living near to the Baodi mall outbreak) > Family relationship > Work relationship > Other known relationship > Other travel > Unknown (lowest priority).

## Statistical methods

### Incubation period

The “start\_source” and “end\_source” columns in each dataset are used to define the maximum and minimum possible incubation periods for each case. We additionally assume that incubation times have to be at least 1 day in length, and that the maximum incubation times are at least 3 days to take into account some uncertainty on symptom onset reporting.

We explored several distributions for the incubation period: gamma, Weibull and log normal. As shown in Figure 4, once fit the resulting distributions all provide very similar results. Appendix

1 Table 1 summarizes the parameter estimates for these three distributions. Appendix 1 Table 2 gives the parameters for the incubation period for early- and late-occurring cases in both datasets.

### **Serial interval**

We used bootstrapping to explore the range for the point estimates of  $\mu$  and  $\sigma$  from the mixture model. Appendix 1 Figure 1 shows the results. The mean of the bootstrapped mean estimates is  $4.49 \pm 0.716$  for Tianjin and  $3.83 \pm 0.882$  days in Singapore. Bootstrap values are consistent with a serial interval that is considerably shorter than the incubation periods in both datasets. Appendix 1 Table 3 shows the sensitivity analysis; we varied the the number of cases per cluster to include in the ICC interval data and we explored sorting the cases in the clusters according to the time of last exposure (i.e., the putative index status assigned to the individual with the earliest end to their exposure window, instead of the first symptomatic individual).

The primary analysis removes all cases that are missing dates of symptom onset. To explore the potential impact of removing cases we repeated the serial interval estimates—when accounting for intermediates—by including these missing cases with imputed dates of symptom onset. There are 10 cases with missing date of symptom onset in both Tianjin and Singapore datasets. All cases missing date of symptom onset have a date of confirmation for infection with SARS-CoV-2; therefore, imputed dates were calculated by: (date of confirmation for case) - (average difference between date of symptom onset and date of confirmation, for all cases used in main analysis). This average difference between date of symptom onset and date of confirmation is 5.23 days in Tianjin and 7.43 days in Singapore. Imputing dates in this manner assumes that dates of symptom onset are missing completely at random. This assumption seems reasonable as the range of date of confirmation for the 10 imputed cases covers the majority of the range of date of confirmation for cases in the main analysis, in both datasets (Feb 1 to Feb 22, 2020 for imputed cases vs. Jan 21 to Feb 22, 2020 for main analysis cases in Tianjin and Jan 31 to Feb 21, 2020 vs. Jan 23 to Feb 26, 2020 in Singapore). Appendix 1 Table 4 contains the results of serial interval estimates including cases with imputed date of symptom onset and demonstrates that there is no substantial difference compared to serial interval estimates from the main analysis where missing cases are removed (Appendix 1 Table 3).

### **Pre-symptomatic transmission: methods details**

We estimated the portion pre-symptomatic transmission taking into account that the serial interval and incubation period are not independent, as described in the main text. In Singapore, we found that the covariance is 5.88, the Pearson correlation is 0.43 ( $p = 0.001$ ) and the Spearman ( $\rho=0.174$ )

and Kendall ( $\tau=0.134$ ) correlations were not significant ( $p = 0.2$ ); this is an intermediate signal of covariation. In Tianjin the covariance was 2.63, the correlation 0.29 and the statistical signal similar. We used both our "direct" and "intermediate" incubation period analysis to determine the portion pre-symptomatic transmission, accounting for the covariation.

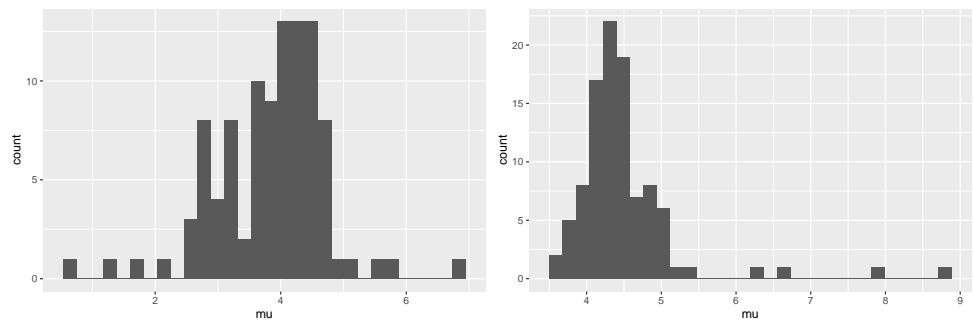
To do this, we first sampled the incubation period parameters using the fits to data in the main text. These fits include a variance estimate between the shape and scale parameters, so we sample the shape and scale accordingly (using the gamma distribution). We created 100 incubation period (shape, scale) pairs (i.e., 100 samples). There are samplers in R for multivariate distributions whose margins are both gamma (`rmvgamma` in the `lcmix` package) and of course multivariate normal samplers, but we do not have a sampler for jointly distributed random variables with a normal distribution on one margin and a gamma on the other. Therefore, we use a gamma distribution for the serial interval, with the same mean and variance as the normal distribution estimated directly from the case-pair data. We obtain 100 serial interval gamma (shape, scale) pairs with the appropriate mean and variance. For each of these 100 distributions, sample *jointly* 500 incubation periods and serial intervals, with correlation of approximately 0.3. We therefore have  $100 \times 500 = 50,000$  joint samples of incubation period and serial interval. The fraction of the (serial interval minus incubation period) samples is an estimate of the fraction of transmission that is pre-symptomatic, accounting for covariation.

We take the same approach for the estimates that account for intermediates; therefore, we sample from the gamma distribution for the incubation period as estimated with intermediates, from the ICC estimate of the serial intervals (i.e., we sample 100 incubation period shape, scale pairs, and 100 generation time shape, scale pairs, and for each we create 500 samples of the incubation period and generation time, accounting for covariance). This yields the estimates in Table 1 and the density plots in Figure 8.

### **Additional published estimations**

Estimates of incubation period and serial interval from other studies are shown in Appendix 1 Table 5. Of note, the majority of studies do not estimate both incubation period and serial interval in the same population.

## Appendix 1 Figure 1



**Appendix 1 - Figure 1.** Bootstrap values of the mean serial interval for (left) Singapore and (right) Tianjin, based on 100 replicates using the first 4 cases in each cluster.

## Appendix 1 Tables 1–5

**Appendix 1 - Table 1.** Incubation period estimates (without intermediates) using gamma, Weibull and log normal distributions. 95% confidence intervals for the shape and scale (log mean and sd for log normal) parameters are shown in brackets.

<b>Gamma</b>	Median	Shape	Scale
Singapore Cluster	5.32	3.05 (2.0, 3.84)	1.95 (1.23, 2.34)
Tianjin Cluster	8.06	4.74 (3.35, 5.72)	1.83 (1.29, 2.04)

<b>Weibull</b>	Median	Shape	Scale
Singapore Cluster	5.66	1.83 (1.45, 2.30)	6.91 (5.77, 8.29)
Tianjin Cluster	8.59	2.41 (1.99, 2.90)	10.01 (8.94, 11.20)

<b>Log normal</b>	Median	log Mean	Standard Deviation
Singapore Cluster	4.83	1.57 (1.38, 1.81) (mean 4.81)	0.60 (0.47, 0.76)
Tianjin Cluster	7.66	2.04 (1.92, 2.22) (mean 7.69)	0.47 (0.39, 0.56)

**Appendix 1 - Table 2.** Incubation period estimates (without intermediates) using stratified data

<b>Tianjin</b>			
Gamma	Median	Shape	Scale
Early	6.48	6.01 (3.61, 7.26)	1.140 (0.66,1.276)
Late	12.1	17.78 (9.52, 21.47)	0.695 (0.379,0.778)
Weibull	Median	Shape	Scale
Early	6.73	2.88 (2.16, 3.48)	7.643 (6.735, 8.553)
Late	12.6	4.34 (3.10, 5.24)	13.661 (12.245, 15.289)
Log normal	Median	log mean	standard deviation
Early	6.30	1.84 (1.70,2.03)	0.426 (0.331,0.547)
Late	12.0	2.48 (2.38,2.67)	0.233 (0.172,0.315)
<b>Singapore</b>			
Gamma	Median	Shape	Scale
Early	5.26	3.22 (1.67, 4.05)	1.818 (0.847,2.18)
Late	5.35	2.96 (1.68,3.72)	2.034 (1.132,2.439)
Weibull	Median	Shape	Scale
Early	5.51	2.05 (1.34,2.58)	6.587 (5.077,7.897)
Late	5.67	1.75 (1.29,2.21)	6.989 (5.408,8.38)
Log normal	Median	log mean	standard deviation
Early	4.91	1.59 (1.33,1.82)	0.598 (0.421,0.848)
Late	4.72	1.55 (1.25,1.78)	0.606 (0.441,0.834)

**Appendix 1 - Table 3.** Serial interval estimates: accounting for intermediates

ordering	Number cases per cluster	$\mu$ (Tianjin)	$\sigma$ (Tianjin)	$\mu$ (Singapore)	$\sigma$ (Singapore)
Onset	3	4.17	0.998	4.03	1.06
Onset	4	4.31	0.935	4.17	1.06
Onset	5	4.43	0.999	4.43	1.09
Onset	6	4.54	1.05	4.76	1.15
Last Exposure	4	5.09	1.27	4.26	1.17
Bootstrap	4	4.49 (sd 0.716)	0.995 (sd 0.307)	3.83 (sd 0.882)	1.24 (sd 0.538)



**Appendix 1 - Table 4.** Serial interval estimates: accounting for intermediates and using imputed dates of symptom onset

ordering	Number cases per cluster	$\mu$ (Tianjin)	$\sigma$ (Tianjin)	$\mu$ (Singapore)	$\sigma$ (Singapore)
Onset	3	4.35	0.907	4.18	1.05
Onset	4	4.40	0.864	4.27	1.04
Onset	5	4.48	0.909	4.41	0.981
Onset	6	4.55	0.948	4.71	1.08
Last Exposure	4	4.81	0.948	4.62	2.11
Bootstrap	4	4.53 (sd 0.585)	0.941 (sd 0.358)	4.31 (sd 1.03)	1.50 (sd 0.629)

**Appendix 1 - Table 5.** Mean incubation period and mean serial interval estimates for COVID-19 generated by other studies.

<b>Data</b>	<b>Number of Cases</b>	<b>Mean Incubation Period (days)</b>	<b>Mean Serial Interval (days)</b>	<b>Reference</b>
Wuhan first cases	425	5.2 (95CI 4.1-7.0)	7.5 (95CI 5.3-19)	<i>Li Q et al. (2020)</i>
South Korea first cases	24	3.6	4.6	<i>Ki M (2020)</i>
Travellers from Wuhan	88	6.4 (95CI 5.6-7.7)	-	<i>Backer JA et al. (2020)</i>
Diagnosis outside Wuhan (excluding Wuhan residents)	52	5.0 (95CI 4.2-6.0)	-	<i>Linton NM et al. (2020)</i>
Diagnosis outside Wuhan (including Wuhan residents)	158	5.6 (95CI 5.0-6.3)	-	<i>Linton NM et al. (2020)</i>
Transmission chains				
in Hong Kong	21 chains	-	4.4 (95CI 2.9-6.7)	<i>Zhao S et al. (2020)</i>
Infector-infectee pairs*	28 pairs	-	4.0 (95CI 3.1-4.9)	<i>Nishiura H et al. (2020)</i>

\*Note: included 3 infector-infectee pairs from this Singapore cluster. Remainder from Vietnam (4), South Korea (7), Germany (4), Taiwan (1) and China (9)