- 1 Early epidemiological assessment of the transmission
- 2 potential and virulence of coronavirus disease 2019
- **3 (COVID-19) in Wuhan City: China, January-February,**
- **4 2020**
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Abstract

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Background: Since the first cluster of cases was identified in Wuhan City, China, in December, 2019, coronavirus disease 2019 (COVID-19) has rapidly spread across China, causing multiple introductions in 109 countries/territories/areas as of March 10th. Despite the scarcity of publicly available data, scientists around the world have made strides in estimating the magnitude of the epidemic, the basic reproduction number, and transmission patterns. Recently more evidence suggests that a substantial fraction of the infected individuals with the novel coronavirus show little if any symptoms, which suggest the need to reassess the transmission potential of this emerging disease. In this study, we derive estimates of the transmissibility and virulence of COVID-19 in Wuhan City, China, by reconstructing the underlying transmission dynamics using multiple data sources. **Methods:** We employ statistical methods and publicly available epidemiological datasets to jointly derive estimates of transmissibility and severity associated with the novel coronavirus. For this purpose, the daily series of laboratory-confirmed COVID-19 cases and deaths in Wuhan City and epidemiological data of Japanese evacuees from Wuhan City on

board government-chartered flights were integrated into our analysis. 43 **Results:** 44 Our posterior estimates of basic reproduction number (R) in Wuhan City, China in 45 2019–2020 reached values as high as 5.20 (95% CrI: 5.04–5.47) and the enhanced public 46 health intervention after January 23rd in 2020 was associated with a declined R at 0.58 47 48 (95% CrI: 0.51–0.64), with the total number of infections (i.e. cumulative infections) estimated at 1905526 (95%CrI: 1350283-2655936) in Wuhan City, raising the 49 proportion of infected individuals to 19.1% (95%CrI: 13.5-26.6%). We also found that 50 51 most recent crude infection fatality ratio (IFR) and time-delay adjusted IFR is estimated 52to be 0.04% (95% CrI: 0.03%–0.06%) and 0.12% (95% CrI: 0.08–0.17%), which is 53 several orders of magnitude smaller than the crude CFR estimated at 4.19% 54 **Conclusions:** 55 We have estimated key epidemiological parameters of the transmissibility and virulence 56 of COVID-19 in Wuhan, China during January-February, 2020 using an ecological 57 modelling approach. The power of our approach lies in the ability to infer epidemiological parameters with quantified uncertainty from partial observations 58 collected by surveillance systems. 59

Keywords: epidemic; transmissibility; mathematical model; COVID-19; China

Background

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The novel coronavirus (COVID-19) emerging from China is a deadly respiratory pathogen that belongs to the same family as the coronavirus responsible for the 2002-2003 Severe Acute Respiratory Syndrome (SARS) outbreaks [1]. Since the first cluster of cases was identified in Wuhan City, China, in December, 2019, COVID-19 has rapidly spread across China as well as caused multiple introductions in 109 countries/territories/areas as of March 10th, 2020 [2]. Nevertheless, China has been hit hard by this emerging infectious disease, especially the city of Wuhan in Hubei Province, where the first cluster of severe pneumonia caused by the novel virus was identified. Meanwhile, the cumulative number of laboratory and clinically confirmed cases and deaths in mainland China has reached 80778 and 3158, respectively, as of March 10th, 2020 [3]. Because the morbidity and mortality burden associated with the novel coronavirus has disproportionally affected the city of Wuhan, the center of the epidemic in China, the central government of the People's Republic of China imposed a lockdown and social distancing measures in this city and surrounding areas starting on January 23rd 2020. Indeed, out of the 80778 COVID-19 cases reported in China, 49978 cases (61.9%) are from Wuhan City. In terms of the death count, a total of 2423 deaths (76.7%) have been recorded in Wuhan city out of the 3158 deaths reported throughout China. To guide the effectiveness of interventions, it is crucial to gauge the uncertainty relating to key epidemiological parameters characterizing the transmissibility and the severity of the disease. Despite the scarcity of publicly available data, scientists around the world have made strides in estimating the magnitude of the epidemic, the basic reproduction number, and transmission patterns [4-5]. Moreover, accumulating evidence suggests that a substantial fraction of the infected individuals with the novel coronavirus show little if any symptoms, which suggest the need to reassess the transmission potential of this emerging disease [6]. For this purpose, in this study we employ statistical methods and publicly available epidemiological datasets to jointly derive estimates of transmissibility and severity associated with the novel coronavirus.

Methods

Epidemiological data

We linked our model to two different datasets. First, the daily series of laboratory–confirmed nCov cases and deaths in Wuhan City were extracted according to date of symptoms onset or reporting date from several sources [3, 7-8]. Our analysis relies on epidemiological data reported prior to February 11th, 2020 because of the change in case definition that was announced on February 12th, 2020 [9]. As of February 11th, 2020, a total of 19559 confirmed cases including 820 deaths were reported in Wuhan City. Second, epidemiological data of Japanese evacuees from Wuhan City on board government–chartered flights were obtained from the Japanese government. After arriving in Japan, all of the Japanese evacuees were kept in isolation for about 14 days and examined for infection using polymerase chain reaction (PCR) tests [7]. As of February 11th, a total of four flights with the Japanese evacuees left Wuhan City. We collected information on the timing of the evacuee fights that left Wuhan City as well as the number of passengers that tested positive for COVID-19 in order to calibrate our model (Table S1).

Statistical analysis

- Using the following integral equation model, we estimate the reproduction
- number of COVID-19. Here, infected and reported cases are denoted by i and c,
- 112 respectively.

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- We connected the daily incidence series with a discrete—time integral equation
- to describe the epidemic dynamics. Let g_s denote the probability mass function of the
- serial interval, e.g., the time from illness onset in a primary case to illness onset in the
- secondary case, of length s days, which is given by

$$g_{s} = G(s) - G(s-1)) ,$$

- For s > 0 where G(.) represents the cumulative distribution function of the gamma
- distribution. Mathematically, we describe the expected number of new cases with day t,
- 119 E[c(t)] as follows,

$$E[c(t)] = \sum_{s=1}^{\infty} E[c(t-s)]R,$$

- where E[c(t)] represents the expected number of new cases with onset day t, where R
- represents the average number of secondary cases per case.
- Subsequently, we also employed the time-dependent variation in R to take into
- account the impact of enhanced interventions on the transmission potential. This time
- dependence was modelled by introducing a parameter δ_I , which is given by

$$\delta_1 = \begin{cases} 1 & otherwise \\ \beta_1 & \text{if } t = period_1 \\ \beta_2 & \text{if } t = period_1 \end{cases}$$

- where period₁ and period₂ represent the corresponding period from January 23rd to
- February 2nd 2020 and from February 3rd to February 11th, 2020, respectively.
- January 23rd 2020 is the date when the central government of the People's Republic of

- China imposed a lockdown in Wuhan and other cities in Hubei in an effort to quarantine the epicentre of the coronavirus (COVID-19) to mitigate transmission. Furthermore, we evenly divide the interval into two periods to incorporate the time-dependent effects on R using the parameters β_I and β_I which scale the extent of the intervention, taking
- values smaller than 1[10].
- To account for the probability of occurrence, θ [11] ,we assume that the number
- of observed cases on day t, h(t), occurred according to a Bernoulli sampling process,
- with the expected values $E(c_i; H_{t-1})$, where $E(c_i; H_{t-1})$ denotes the conditional expected
- incidence on day t, given the history of observed data from day 1 to day (t-1), denoted
- by H_{t-1} . Thus, the number of expected newly observed cases is written as follows:

$$\mathbf{E}[h(t);H_{t-1}] = \{ \begin{aligned} (1-\theta) + \theta \mathbf{E}[c(t);H_{t-1}], & \text{if } h = 0, \\ \theta \mathbf{E}[c;H_{t-1}], & \text{otherwise,} \end{aligned}$$

- Further, we model the time–dependent variation in the reporting probability.
- This time dependence was modelled by introducing a parameter δ_2 , which is given by

$$\delta_2 = \begin{cases} \alpha_1, & \text{if } t = period_3, \\ \alpha_2, & \text{if } t = period_4, \\ 1, & \text{otherwise,} \end{cases}$$

- where period₃ and period₄ represent the corresponding periods from the start of our
- study period to Jan 16 and from Jan 17 to Jan 22, respectively, while α_1 and α_2 scale the
- extent of the reporting probability (where α_1 and α_2 is expected to be smaller than 1).
- We evenly divide the duration before the lockdown was put in place into two to
- incorporate the time dependency of the reporting probability. The number of expected
- newly observed cases should be updated as

$$\begin{split} \mathbf{E}[h(t);H_{t-1}] = \{ & (1-\theta) + q\delta\theta \mathbf{E}[c(t);H_{t-1}], & \text{if } h_a = 0 \;, \\ & q\delta\theta \mathbf{E}[c(t);H_{t-1}], & \text{otherwise,} \end{split}$$

We assume the incidence, h(t) is the result of the Binomial sampling process with the

expectation E[h]. The likelihood function for the time series of observed cases that we employ to estimate the effective reproduction number and other relevant parameters is given by:

$$L_1(U;c) = \prod_{t=1}^T \binom{E(h(t);H(t-1))}{c(t)} q^{c(t)} (1-q)^{E(h(t);H(t-1)-c(t))} ,$$

where *U* indicates parameter sets that are estimated from this likelihood.

Subsequently, the conditional probability of non–infection given residents in Wuhan City at the time point of t_i , p_{ti} , was assumed to follow a binomial distribution, and the likelihood function is given by:

$$L_{2}(p_{t_{i}}; M_{t_{i}}, m_{t_{i}}) = \binom{M_{t_{i}}}{m_{t_{i}}} p_{t_{i}}^{m_{t_{i}}} (1 - p_{t_{i}})^{M_{t_{i}} - m_{t_{i}}},$$

Where M_{ti} and m_{ti} is the number of government charted flight passengers and non–infected passengers at the date of t_i , respectively, and p_{ti} is the proportion of the estimated non–infected population in Wuhan at the date of t_i , calculated from the h(t) and catchment population in Wuhan City [3,13].

Serial interval estimates of COVID-19 were derived from previous studies of nCov, indicating that it follows a gamma distribution with the mean and SD at 7.5 and 3.4 days, respectively, based on ref. [14]. The maximum value of the serial interval was fixed at 28 days as the cumulative probability distribution of the gamma distribution up to 28 days reaches 0.999.

Infection fatality ratio

Crude CFR and crude IFR is defined as the number of cumulative deaths divided by the number of cumulative cases or infections at a specific point in time

without adjusting the time delay from illness onset or hospitalization to death. Next, we employed an integral equation model in order to estimate the real–time IFR. First, we estimated the real–time CFR as described elsewhere [15-17]. For the estimation, we employ the delay from hospitalization to death, f_s , which is assumed to be given by $f_s = F(s) - F(s-1)$ for s>0 where H(s) follows a gamma distribution with mean 10.1 days and SD 5.4 days, obtained from the available observed data [18].

$$\begin{split} L_{3}(\pi;c_{t},\theta) &= \prod_{t_{i}} \left(\sum_{t=1}^{t_{i}} c_{t} \right), \left(\pi \frac{\sum_{t=2}^{t_{i}} \sum_{s=1}^{t-1} c_{t-s} f_{s}}{\sum_{t=1}^{t_{i}} c_{t}} \right)^{D_{t_{i}}} \left(1 \right. \\ &- \pi \frac{\sum_{t=2}^{t_{i}} \sum_{s=1}^{t-1} c_{t-s} f_{s}}{\sum_{t=1}^{t_{i}} c_{t}} \right)^{\sum_{t=1}^{t_{i}} c_{t} - D_{t_{i}}} \end{split}$$

where c_t represents the number of new cases with reported day t, and D_{ti} is the number of new deaths with reported day t_i [16-18]. We assume that the cumulative number of observed deaths, D_t is the result of the binomial sampling process with probability π . Subsequently, crude IFR and time—delay adjusted IFR are calculated using the estimated π and h_t .

The total likelihood is calculated as $L=L_1L_2L_3$ and model parameters were estimated using a Monte Carlo Markov Chain (MCMC) method in a Bayesian framework. Posterior distributions of the model parameters were estimated based on sampling from the three Markov chains. For each chain, we drew 100,000 samples from the posterior distribution after a burn–in of 20,000 iterations. Convergence of MCMC chains were evaluated using the potential scale reduction statistic [19-20]. Estimates and 95% credibility intervals for these estimates are based on the posterior probability

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distribution of each parameter and based on the samples drawn from the posterior distributions. All statistical analyses were conducted in R version 3.5.2 (R Foundation for Statistical Computing, Vienna, Austria) using the 'rstan' package. **Results** The daily series of COVID-19 laboratory-confirmed incidence and cumulative incidence in Wuhan in 2019–2020 are displayed in Figure 1. Overall, our dynamical models yield a good fit to the temporal dynamics (i.e. incidence, cumulative incidence) including an early exponential growth pattern in Wuhan. In incidence data, a few fluctuations are seen, probably indicating that the surveillance system likely missed many cases during the early transmission phase (Figure 1). Our posterior estimates of basic reproduction number (R) in Wuhan City, China in 2019–2020 was estimated to be as high as 5.20 (95% CrI: 5.04–5.47). The time-dependent scaling factors quantifying the extent of enhanced public health intervention on R is 0.99 (95%CrI: 0.94–1.00), declining R to 5.12 (95%CrI: 4.98–5.26) from January 23rd to February 1st and 0.11 (95% CrI: 0.10–0.13), declining R to 0.58 (95% CrI: 0.51–0.64) from February 2nd to February 11th, 2020. Other parameter estimates for the probability of occurrence and reporting rate are 0.97 (95% CrI: 0.84–1.00) and 0.010 (95% CrI: 0.007–0.014), respectively. Moreover, the time-dependent scaling factor quantifying the extent of reporting rate, α , is estimated to be 0.08 (95% CrI: 0.03–0.19) before January 16th and to be 0.99 (95% CrI: 0.96–1.00) from January 17th to January 22nd. The total number of estimated laboratory–confirmed cases (i.e. cumulative

cases) is 18913 (95% CrI: 16444–19705) while the actual numbers of reported

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laboratory-confirmed cases during our study period is 19559 as of February 11th, 2020. Moreover, we inferred the total number of COVID-19 infections (Figure S1). Our results indicate that the total number of infections (i.e. cumulative infections) is 1905526 (95%CrI: 1350283-2655936). The Observed and posterior estimates of the cumulative number of deaths from COVID-19 in Wuhan are displayed in Figure 2, and model-based posterior estimates of the cumulative number of deaths is 820 (95% CrI: 744–900), while actual number of reported deaths is 820. The estimated temporal variation in the death risk caused by COVID-19 in Wuhan, China, 2019–2020 is shown in Figure 3 and Figure S2. Observed and posterior estimated of crude CFR in Wuhan City is presented in Figure 2A, while observed and posterior estimates of time-delay adjusted CFR is shown in Figure 2B. Furthermore, Figure 3A and 3B illustrates time-delay no-adjusted IFR and time-delay adjusted IFR, respectively. The latest estimate of the crude CFR and time-delay adjusted CFR in Wuhan appeared to be 4.2% (95% CrI: 3.9–4.9%) and 12.2% (95% CrI: 11.4–13.0%), respectively, whereas the latest model-based posterior estimates of time-delay not adjusted IFR and adjusted IFR, presented in Figure 3 C and D, are 0.04%(95% CrI: 0.03%-0.06%) and 0.12% (95% CrI: 0.08-0.17%), respectively, while the observed crude CFR is calculated to be 4.19% (Table 1). **Discussion** In this study we derived estimates of the transmissibility and virulence of COVID-19 in Wuhan City, China, by reconstructing the underlying transmission

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dynamics using multiple data sources. Applying dynamic modeling, the reproduction number and death risks as well as probabilities of occurrence and reporting rate were estimated. Our posterior estimates of basic reproduction number (R) in Wuhan City, China in 2019–2020 is calculated to be as high as 5.20 (95%CrI: 5.04–5.47). The time-dependent scaling factor quantifying the extent of enhanced public health intervention on R is 0.99 (95%CrI: 0.94–1.00), declining R to 5.12 (95%CrI: 4.98–5.26) from January 23rd to February 1st and 0.11 (95% CrI: 0.10–0.13), declining R to 0.58 (95%CrI: 0.51–0.64) for February 2nd to February 11th, 2020. These R estimates capturing the underlying transmission dynamics modify the impact of COVID-19, with the total number of infections (i.e. cumulative infections) estimated at 1905526 (95%CrI: 1350283–2655936) in Wuhan City, raising the proportion of infected individuals to 19.1% (95%CrI: 13.5–26.6%) with a catchment population in Wuhan City of 10 million people. Our estimates of mean reproduction number reached values as high as 5.20, an estimate that is slightly higher than previous mean estimates in the range 2.2-3.8 derived by fitting epidemic models to the initial growth phase of the observed case incidence [21-23]. By comparison, the R estimate for the Diamond Princess cruise ship in Japan reached values as high as ~11 [24]. Further, these estimates are higher than recent mean R estimates derived from the growth rates of the COVID-19 outbreaks in Singapore $(R\sim1.1)$ [25] and Korea $(R\sim1.5)$ [26]. The sustained high R values in Wuhan City even after the lockdown and mobility restrictions suggests that transmission is occurring inside the household or amplified in healthcare settings [18], which is a landmark of past SARS and MERS outbreaks

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[27-28]. Considering the potent transmissibility of COVID-19 in confined settings, as illustrated by COVID-19 outbreaks aboard cruise ships, including the Diamond Princess cruise ship, where the total number of secondary or tertiary infections reached 705 among more than 3,700 passengers as of February 28th, 2020 and also by the COVID-19 outbreak tied to the Shincheonji religious sect in South Korea where church members appear to have infected from seven to 10 people. [29-31], it is crucial to prevent transmission in confined settings including hospital-based transmission by strengthening infection control measures as well as transmission stemming from large social gatherings. Our most recent estimates of the crude CFR and time-delay adjusted CFR are at 4.2% (95% CrI: 3.9–4.9%) and 12.23% (95% CrI: 11.4–13.0%), respectively. In contrast, our most recent crude IFR and time-delay adjusted IFR is estimated to be 0.04% (95% CrI: 0.03%–0.06%) and 0.12% (95% CrI: 0.08–0.17%), which is several orders of magnitude smaller than the crude CFR at 4.19%. These findings indicate that the death risk in Wuhan is estimated to be much higher than those in other areas, which is likely explained by hospital-based transmission [32]. Indeed, past nosocomial outbreaks have been reported to elevate the CFR associated with MERS and SARS outbreaks, where inpatients affected by underlying disease or seniors infected in the hospital setting have raised the CFR to values as high as 20% for a MERS outbreak [33-34]. Public health authorities are interested in quantifying R and CFR to measure the transmission potential and virulence of an infectious disease, especially when emerging/re–emerging epidemics occur in order to decide the intensity of the public health response. In the context of a substantial fraction of unobserved infections due to

COVID-19, R estimates derived from the trajectory of infections and the IFR are more realistic indicators compared to estimates derived from observed cases alone [18, 35-36]. Our analysis also revealed a high probability of occurrence and quite low reporting probabilities in Wuhan City. High probability of occurrence in the above equation suggests that zero observed cases at some point is not due to the absence of those infected, but more likely due to a low reporting rate. A very low reporting probability suggests that it is difficult to diagnose COVID-19 cases or a breakdown in medical care delivery. Moreover, we also identified a remarkable change in the reporting rate, estimated to be 12-fold lower in the 1st period (-Jan 16, 2020) and about the same during the 2^{nd} period (January $17^{th} - 22^{nd}$, 2020), relative to the that estimated after January 23rd 2020. Our results are not free from the limitations. First, our methodology aims to capture the underlying transmission dynamics using multiple data sources. By implementing mass screening in certain populations is a useful approach to ascertain the real proportion of those infected and a way of adding credibility to the estimated values. Second, it is worth noting that the data of Japanese evacuee employed in our analysis is not a random sample from the Wuhan catchment population. Indeed, it also plausible that their risk of infection in this sample is not as high as that of local residents in Wuhan, underestimating the fatality risk.

Conclusion

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In summary, we have estimated key epidemiological parameters of the

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transmissibility and virulence of COVID-19 in Wuhan, China, January-February, 2020 using an ecological modelling approach and several epidemiological datasets. The power of our approach lies in the ability to infer epidemiological parameters with quantified uncertainty from partial observations collected by surveillance systems. List of abbreviations CFR: Case fatality ratio, IFR: Infection Fatality ratio, SARS: Severe Acute Respiratory Syndrome, MERS: Middle East Respiratory Syndrome **Additional files** Additional file 1: Appendix. Table S1. Information related to Japanese evacuees from Wuhan City on board government-chartered flights **Declarations** Ethics approval and consent to participate Not applicable.

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program.

Consent for publication Not applicable. Availability of data and materials The present study relies on published data and access information to essential components of the data are available from the corresponding author. **Competing interests** The authors declare that they have no competing interests. **Funding** KM acknowledges support from the Japan Society for the Promotion of Science (JSPS) KAKENHI Grant Number 18K17368 and from the Leading Initiative for Excellent Young Researchers from the Ministry of Education, Culture, Sport, Science & Technology of Japan. KK acknowledges support from the JSPS KAKENHI Grant Number 18K19336 and 19H05330. GC acknowledges support from NSF grant 1414374 as part of the joint NSF-NIH-USDA Ecology and Evolution of Infectious Diseases

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Authors' contributions KM and GC conceived the early study idea. KM and KK built the model. KM implemented statistical analysis and wrote the first full draft. GC advised on and helped shape the research. All authors contributed to the interpretation of the results and edited and commented on several earlier versions of the manuscript. **Acknowledgements** Not applicable. **REFERENCES** 1. Jon Cohen. Mining coronavirus genomes for clues to the outbreak's origins. **Science**. Jan 31, 2020. https://www.sciencemag.org/news/2020/01/mining-coronavirus-genomes-cluesoutbreak-s-origins. Accessed, 3 Feb, 2020 2. World Health Organization (WHO). Novel Coronavirus (2019-nCoV) situation reports. Available from: https://www.who.int/emergencies/diseases/novel-coronavirus-2019/situation-rep orts Accessed, [cited 2020 Mar 10]

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Figures Figure 1. Observed and posterior estimates of the daily new cases and cumulative cases of the COVID-19 cases in Wuhan, China, 2019–2020 Observed and posterior estimates of laboratory-confirmed reported cases (A) and cumulative reported cases (B) are presented. Observed data are presented in the dot, while dashed line indicates 50 percentile, and areas surrounded by light grey and deep grey indicates 95% and 50% credible intervals (CrI) for posterior estimates, respectively. Epidemic day 1 corresponds to the day that starts at January 1st, 2020. Figure 2. Observed and posterior estimates of the cumulative deaths of the **COVID-19 in Wuhan, China, 2019–2020** Observed and posterior estimates of the cumulative deaths of the COVID-19 in Wuhan is presented. Observed data are presented in the dot, while dashed line indicates 50 percentile, and areas surrounded by light grey and deep grey indicates 95% and 50% credible intervals (CrI) for posterior estimates, respectively. Epidemic day 1 corresponds to the day that starts at January 1st, 2020. Figure 3.Temporal variation of the infection fatality risks caused by COVID-19 in Wuhan, China, 2019-2020

(A) Posterior estimates of crude infection fatality ratio in Wuhan City. (B) Posterior estimates of time–delay adjusted infection fatality ratio in Wuhan City.

Black dots shows observed data, and light and dark indicates 95% and 50% credible intervals for posterior estimates, respectively. Epidemic day 1 corresponds to the day that starts at January 1st, 2020.

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Figure S1. Observed daily new cases and posterior estimates of the daily new infections of the COVID-19 in Wuhan, China, 2019-2020 Observed daily new cases and posterior estimates of infections of the COVID-19 are presented. Observed data are presented in the dot, while dashed line indicates 50 percentile, and areas surrounded by light grey and deep grey indicates 95% and 50% credible intervals (CrI) for posterior estimates, respectively. Epidemic day 1 corresponds to the day that starts at January 1st, 2020. Figure S2.Temporal variation of the case fatality risks caused by COVID-19 in Wuhan, China, 2019-2020 (A) Observed and posterior estimates of crude case fatality ratio in Wuhan City, (B) Observed crude case fatality ratio and posterior estimates of time-delay adjusted CFR in Wuhan City. This figure is submitted to the ref [18]. The purpose of the study is to compare the case fatality ration (CFR. Not IFR) in three different areas (Wuhan City, in Hubei Province excluding Wuhan City and in China excluding Hubei Province) to interpret the current severity of the epidemic in China, and the purpose is different from this study.

Tables

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Table 1 – Death risk by COVID-19 in Wuhan City, China, 2020 (As of February 12, 2020)

Death Risk	Latest estimate	Range of median
	Latest estimate	estimates
Crude CFR (Observed)	4.19%	2.0 - 9.0%
Crude CFR (Estimated)	$4.2\%~(95\%\mathrm{CrI}^{\ \ddagger}\colon 3.9-4.9\%)$	3.4 - 7.2%
Time delay adjusted CFR	12.2% (95% CrI: $11.4 - 13.0%$)	4.1 - 34.8%
Crud IFR	0.04% (95%CrI: 0.03 – 0.06%)	0.02 - 0.07%
Time delay adjusted IFR	0.12% (95% CrI: 0.08 - 0.17%)	0.04 - 0.33%

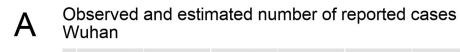
517 CrI: Credibility intervals, CFR: Case fatality ratio, IFR: Infection fatality

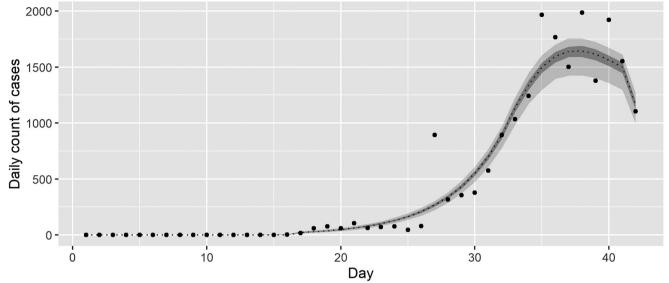
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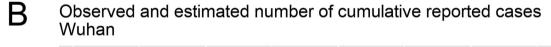
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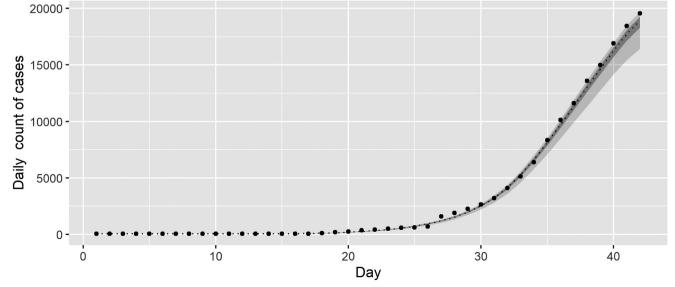
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[‡]Upper and lower 95% credibility interval









Observed and estimated number of reported cumulative death Wuhan

