

# SARS-CoV-2 vaccines: will they arrive too late?

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## Abstract

The COVID-19 pandemic has already ravaged many countries and lockdowns at various levels have been in place for some time. The indirect effect on the economy is devastating and decisions on reopening activities and vaccine acquisition and deployment must be taken. There are two main questions: what are the signals we need to watch to decide the places, the activities and the level at which these can be restarted? what should be the vaccination policies? Here we suggest how to use the observed deaths by COVID-19 in an arbitrary population as a surrogate for the progression of the epidemic with the purpose of decision making. The main result is that if the number of deaths in a region is close to 2 per thousand individuals, the fraction of remaining susceptible may be too small for the vaccine to make a difference.

*Keywords:* COVID-19, SARS-CoV-2, IFR, Asymptomatic, Immunity, Total Infections, Lethality, Vaccines

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## 1. Introduction

COVID-19 pandemic has shown to have a low lethality, nevertheless, the burden of the disease so far is huge. Economical activities are suspended or reduced and there is pressure for reopening of schools which requires more in-  
5 formation, not only available to policymakers but also to the general public,

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to avoid civic unrest. The competition for developing a vaccine is keen, with about 321 vaccine candidates with 32 in clinical trials in progress (Le et al., 2020), raising safety concerns (Harrison and Wu, 2020; Peeples, 2020), and the additional economic cost for the purchase and deployment of vaccines has not  
10 been yet added to the burden of the disease.

At the moment, without vaccines or effective pharmaceutical treatments available, the decision on whether reopening activities or not depends mostly on the number of additional infections/deaths that will be caused because of some policy, say, opening schools, allowing public gatherings, opening touristic places,  
15 increasing the current density allowed in some places as cinemas, restaurants, buses, and planes, etc. For a highly infectious virus as SARS-CoV-2, the decision must be rooted in the amount of remaining susceptible in the region affected for some change in policy.

Using the number of individuals that access private or public hospitals as a  
20 surrogate of the current number of infections, or even the number of confirmed cases, is not accurate, because those quantities strongly depend on the availability of medical services that are not always accessible to the bulk of a population in many countries, or simply because of lack of resources for test deployment or because of policies that disregard testing. The number of deaths is less dubious.  
25 if we manage to calculate the average number of infections that will result in the death of the average individual, then, we can use this number to estimate the number of infections that were required to observe the current number of deaths, and from here, the share of susceptible available in a well-defined population. Even although deaths were not reported as COVID-19 related, records  
30 may exist that shed light on the likelihood that death was or not caused by SARS-CoV-2 infection, which is part of the policy adopted by some countries like Belgium, which is, without doubt, one of the reasons why this country keeps one of the highest number of deaths per capita in Europe.

Before proceeding, we need to deploy two facts. The first one attempts to  
35 establish that, with few exceptions, it is almost impossible to stop the pandemic by mitigation and control measures, and the most we can do is reducing the in-

fection rate (at a huge economic and social cost) which is known as *flattening the curve*. The second establishes that we are in the condition now to have reliable lower bound estimates for IFR, the Infection Fatality Risk of SARS-CoV-2. It  
40 is the confluence of these two facts that allow us to establish a surrogate for the progression of the epidemic in a region and thus, for the calculation of the share of susceptible available, which is the basis to decide on reopening activities and vaccine deployment.

**Fact 1: SARS-CoV-2 is highly contagious and will infect most of the**  
45 **population**

Several studies have shown that the basic reproductive number  $R_0$  is high (Sanche et al., 2020; D'Arienzo and Coniglio, 2020; Najafimehr et al., 2020; Liu et al., 2020; Alimohamadi et al., 2020), and it has been calculated as high as 5.7. It is important to remember that  $R_0$  is the potential of disease transmission  
50 in the absence of any control or mitigation measures, and it is the potential that comes into play as soon as control or mitigation measures are suspended. We calculated the  $R_0$  for 40 countries using only the first two weeks of data, analyzing the progression in the number of deaths. These 40 countries were the first countries affected by pandemics except for China. The  $R_0$  calculated  
55 is shown in Figure 1. The estimation of the true potential of the disease is more evident for those countries that were affected first (e.g. Italy and Spain), where our estimate of  $R_0$  is about the same as the reported by Sanche et al. (2020). The learning curve is evident in the reduction of  $R_0$  when the pandemic advances. Estimates of  $R_0$  in countries where the pandemic arrived at later  
60 stages will not reflect the true potential of the virus alone, but the preparedness of a population.

Table (1) in Appendix contains the main statistics derived from the analysis of  $\lambda$  and  $R_0$  for the countries analyzed. From the estimated  $\lambda$ 's, we can see that the first countries to be affected had an  $R_0$  larger than 5 whereas this was  
65 decreasing to achieve values just above 2.5, this implies that at the beginning

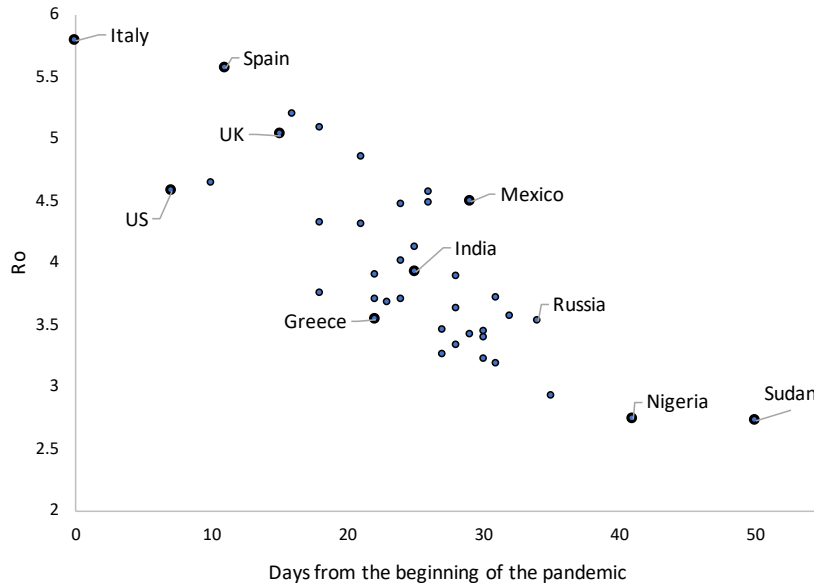


Figure 1: Age of the pandemic vs.  $R_0$ . For each country, the baseline was taken as the date when the accumulated number of deaths was at least 5. Day zero is 02/24/2020. See Table 1 for expanded data.

$\lambda$  was as high as one effective contact every three days to one effective contact every 5 days, on average.

There are some facts we need to consider when a virus spreads with that strength: first, forecasting the size of the epidemics (total number of infected) is simpler, since social contact structures are bypassed and become irrelevant, and thus the contact network resembles more a random mixing pattern. For this random mixing, the estimated fraction of infected  $f$ , can be approximated (Kermack and McKendrick, 1927) with:

$$f = 1 - e^{-R_0 f} \quad (1)$$

(for a probabilistic derivation, see Hernandez-Suarez and Mendoza-Cano, 2009).

For instance, if  $R_0 = 5.7$  the fraction of infected is  $f = 0.99$ . However, if we manage to reduce  $R_0$  by half,  $f = 0.93$ , which is not a big difference in the size

of the epidemic. Nevertheless, reducing  $R_0$  by half is an incredible task that involves mainly lockdowns and face-masks, and the former has a huge economic cost that can not hold for long except for rich countries or countries whose political or cultural organization allows its implementation. Nevertheless, even if a country manages to implement actions to reduce  $R_0$  to a value smaller to one, and can support citizens economically to maintain the lockdown for long periods, a handful of infected individuals that enter the country from abroad can restart the infection process, if lockdowns are lifted. From here, the observed waves (see Figure 2 for examples in Japan, Cuba, S. Korea and New Zealand). This is particularly true for a disease where the number of asymptomatic individuals is by far, larger than that of symptomatic and thus, difficult to control. As a consequence, equation (1) can be seen as a quota that must be fulfilled, as long as the fraction of remaining susceptible is larger than  $f$ .

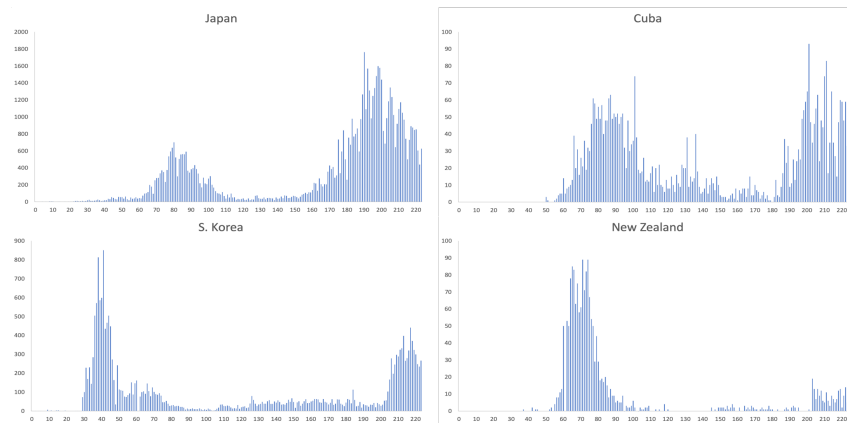


Figure 2: **The appearance of a second wave.** Confirmed cases per day. The current IFR\* (per million inhabitants) is Japan: 10; Cuba: 6; S. Korea: 9; New Zealand: 4

90 **Fact 2: the overall IFR of SARS-CoV-2 is somewhere between 0.2  
and 0.3 %**

All calculations in this section are done with the data on deaths reported to Aug 3, 2020 (Dong et al., 2020). Also, in this section we use IFR to indicate the true overall Infection Fatality Risk of SARS-CoV-2, that is, it is the fraction of  
95 individuals that die if infected with SARS-CoV-2, whereas IFR\* indicates some estimate of IFR for a region of interest. Many attempts have been made to estimate this important parameter that would be a remarkable surrogate to measure the progress of the epidemics in an arbitrarily defined region or population, but the main inconvenient is that while the number of deaths from SARS-CoV-2  
100 can be approximated, the denominator, the total number of infected from the disease is elusive, especially considering that a large fraction of individuals that are infected are asymptomatic. Besides, the efficacy of immunity tests to detect who has been infected and recovered has been challenged with the findings that the ability to detect antibodies is reduced in a few days, especially in those with  
105 mild or no symptoms (Long et al., 2020; Ibarondo et al.). Recently, Eyre et al. (2020) reported that a high fraction of individuals with none or mild symptoms may be undiagnosed mainly due to the calibration strategies of some standardized tests and concluded that samples from individuals with mild/asymptomatic infection should be included in SARS-CoV-2 immunoassay evaluations.

110 Some studies to estimate the IFR have been recently released for Iceland (Gudbjartsson et al.), India (Mukhopadhyay and Chakraborty, 2020), Germany (Streeck et al., 2020) and Denmark (Erikstrup et al., 2020) among others. In a review of 23 studies where the IFR was estimated Ioannidis (2020) a median of 0.25% is reported, which is about 2.5 deaths for each 1000 individuals. The study  
115 in Iceland, (Gudbjartsson et al.), the most comprehensive to date, proposes an IFR of 0.3% (95% CI, 0.2 to 0.6). Special care must be taken with this later estimate, since Iceland has a population of 321,641, with only 29 deaths so far.

If the disease will affect most of the population, as suggested previously, then, the most effective way to estimate the overall IFR is by measuring the

120 fraction of deaths in a population in which the epidemic has evolved for a long  
time and it is over or near the end. It is evident that we need to focus in regions  
with: *a)* high observed IFR\* and *b)* lack of new waves. Such is the case of  
regions as New Jersey and New York states (see figure 3). The current IFR\*  
calculated for these states is respectively 1.809 and 1.699.

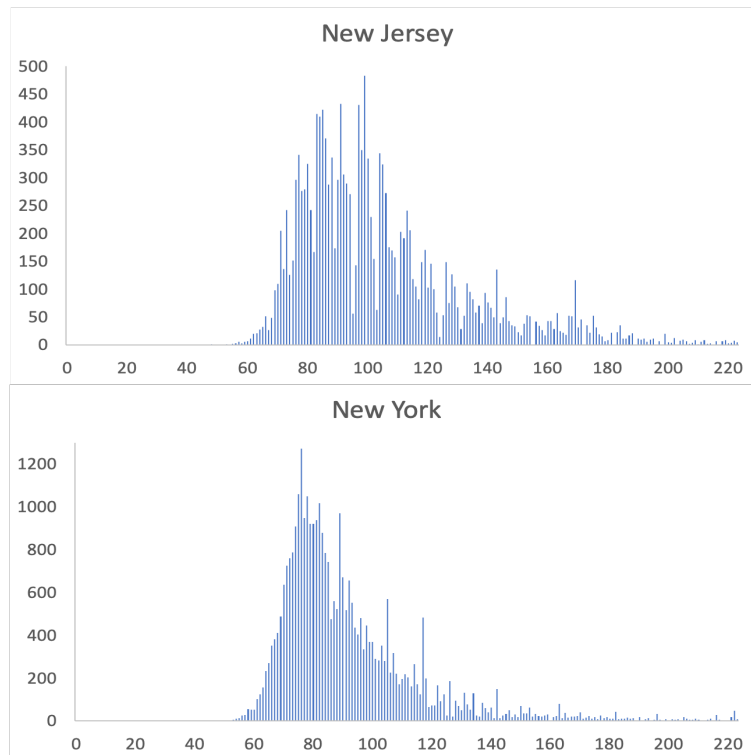


Figure 3: **Absence of second waves in the states of New Jersey and New York.** Confirmed cases per day. The current IFR\* (per million inhabitants) is New Jersey: 1.809; New York: 1.699

125 The lack of new waves in these two states indicates that the virus has infected  
and killed a fraction of individuals close to the IFR, and thus the epidemic  
rampaged those states and it stopped due to what is commonly called *herd*  
*immunity*, that actually conveys no immunity at all, but a lack of a significant  
mass of susceptible for the  $R_0$  to be effective. Thus, the progression of the

130 epidemic in a region can be estimated with  $IFR^*/IFR$ .

## 2. Consequences from the previous facts for vaccines and the return to normality

As a consequence of the two previous facts, we can conclude that it is possible to monitor the development of the epidemic in a region by following the number of deaths. One can estimate the fraction of total infected at time  $t$ ,  
135 asymptomatic or not, with:

$$\theta_t = \frac{x_t}{N \times IFR}$$

where  $x_t$  is the number of observed deaths at time  $t$  and  $N$  is the population size. The estimated fraction of susceptible as a function of the IFR in New Jersey and New York is shown in figure 4. We can see that if  $IFR=2/1000$ , the fraction of remaining susceptible in New Jersey and New York is 9.5% and 15%  
140 respectively, nevertheless, if it is as high as 2.3, the fraction of susceptible raises to 20% and 25% for those states.

## 3. Discussion

Because of the aforementioned facts, it is very important to monitor the COVID-19 related deaths and every effort has to be made to improve current  
145 estimates of the IFR. Hernandez-Suarez et al. (2020) suggested that in light of the high contagiousness of SARS-CoV-2, the secondary cases resulting in fatalities in a household with at least one confirmed infected, may be useful to estimate the IFR, which would provide a large amount of data and minimal testing requirements. Antibody testing in New York shows to date, that out  
150 of 2'089,089 tests, 535,031 were positive for antibodies (NYC-Health) which implies 25.6% of the population may be immune, contradicting our findings that for an IFR as high as 2.3 per thousand the percentage of infected should be close to 75% already. Nevertheless, the fraction of people with antibodies is



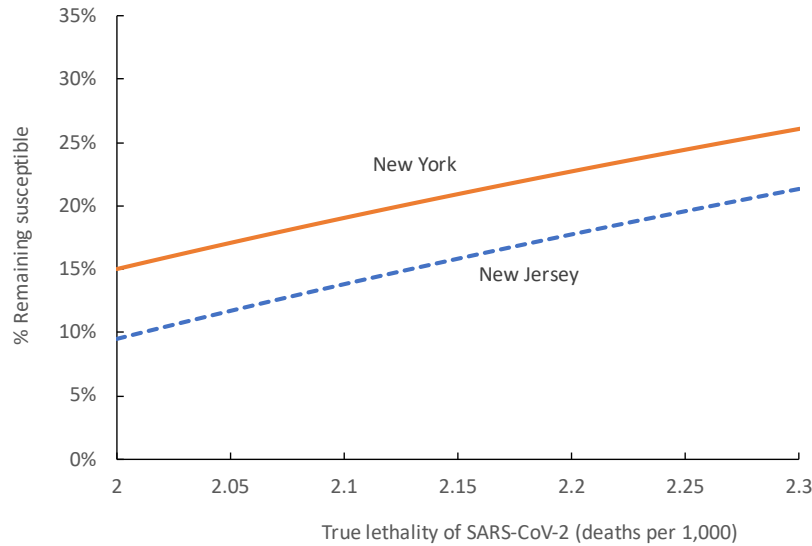


Figure 4: The fraction of remaining susceptible in the states of New Jersey and New York as a function of the lethality of SARS-CoV-2.

155 not a good surrogate of the fraction of infected with SARS-CoV-2 because it underestimates the number of infected for two reasons: first, antibody testing is voluntary and it is natural to expect that individuals with no symptoms are less compelled to be tested than those symptomatic, so, a large fraction of infected is not tested, and second, we already mentioned some failures in testing reported, 160 apparently due to calibration procedures that tend to fail in individuals with none or mild symptoms (Eyre et al., 2020).

The idea that the share of susceptible in New Jersey is relatively small, is supported on the fact that the number of active cases reported to date is over 19,000 and unless they are fully quarantined the infectious pressure from these 165 individuals and those unreported must be huge, nevertheless, no new peak is observed. The same happens in New York, although the share of susceptible is higher. The progression of the pandemics in San Marino (see Figure 5) also supports this idea, since despite of being the country at the top of the mortality per individual (1.237 per thousand) and having only two deaths since the end

170 of April, it appears that the saturation required for the epidemic to halt has not been reached, and a second wave has started in the last two weeks.

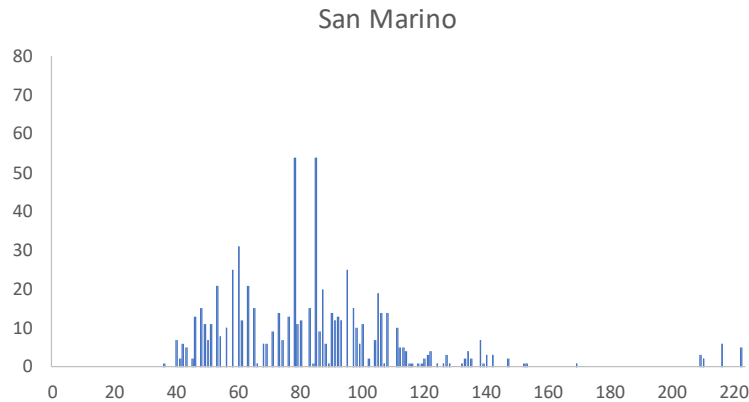


Figure 5: **Second wave in San Marino.** Confirmed cases per day. From April 25 to September 3, there has been only two deaths.

Everything leads to the following question: what will be the purpose of a vaccine in a region where the IFR\* is already large enough to suggest there is only a small fraction of susceptible with limited chance of infection? This question is especially valid from the comprehensive study in Iceland that strongly suggests the existence of immunity due to infection (Gudbjartsson et al.). Unless it is proven that immunity wanes beyond a protective level after some time, everything seems to indicate that those regions where the share of estimated susceptible is already low, should have less priority in the distribution of vaccines. If facts 1 and 2 are not taken into account, a vaccination campaign in a region where the proportion of deaths among the population is close to the IFR, will just give the impression of being effective but will not play a role, regardless of the efficacy of the vaccine. In these cases, the false sense of being protected would be the only thing countries will be paying for.

#### 185 4. Bibliography

- Alimohamadi, Y., Taghdir, M., Sepandi, M., 2020. The estimate of the basic reproduction number for novel coronavirus disease (COVID-19): a systematic review and meta-analysis. *Journal of Preventive Medicine and Public Health* .
- 190 D'Arienzo, M., Coniglio, A., 2020. Assessment of the sars-cov-2 basic reproduction number,  $r_0$ , based on the early phase of covid-19 outbreak in italy. *Biosafety and Health* 2, 57 – 59. doi:10.1016/j.bsheal.2020.03.004.
- Dong, E., Du, H., Gardner, L., 2020. An interactive web-based dashboard to track covid-19 in real time. *The Lancet infectious diseases* 20, 533–534.
- 195 Erikstrup, C., Hother, C.E., Pedersen, O.B.V., Mølbak, K., Skov, R.L., Holm, D.K., Sækmose, S., Nilsson, A.C., Brooks, P.T., Boldsen, J.K., Mikkelsen, C., Gybel-Brask, M., Sørensen, E., Dinh, K.M., Mikkelsen, S., Møller, B.K., Haunstrup, T., Harritshøj, L., Jensen, B.A., Hjalgrim, H., Lillevang, S.T., Ullum, H., 2020. Estimation of SARS-CoV-2 infection fatality rate by real-time antibody screening of blood donors. *medRxiv* doi:10.1101/2020.04.24.20075291.
- 200 Eyre, D.W., Lumley, S.F., O'Donnell, D., Stoesser, N.E., Matthews, P.C., Howarth, A., Hatch, S.B., Marsden, B.D., Cox, S., James, T., et al., 2020. Stringent thresholds for sars-cov-2 igg assays result in under-detection of cases reporting loss of taste/smell. *medRxiv* .
- Gudbjartsson, D.F., Norddahl, G.L., Melsted, P., Gunnarsdottir, K., Holm, H., Eythorsson, E., Arnthorsson, A.O., Helgason, D., Bjarnadottir, K., Ingvarsson, R.F., Thorsteinsdottir, B., Kristjansdottir, S., Birgisdottir, K., Kristinsdottir, A.M., Sigurdsson, M.I., Arnadottir, G.A., Ivarsdottir, E.V., 210 Andresdottir, M., Jonsson, F., Agustsdottir, A.B., Berglund, J., Eiriksdottir, B., Fridriksdottir, R., Gardarsdottir, E.E., Gottfredsson, M., Gretarsdottir,

O.S., Gudmundsdottir, S., Gudmundsson, K.R., Gunnarsdottir, T.R., Gylfason, A., Helgason, A., Jensson, B.O., Jonasdottir, A., Jonsson, H., Kristjansson, T., Kristinsson, K.G., Magnusdottir, D.N., Magnusson, O.T., Olafsdottir, L.B., Rognvaldsson, S., le Roux, L., Sigmundsdottir, G., Sigurdsson, A., Sveinbjornsson, G., Sveinsdottir, K.E., Sveinsdottir, M., Thorarensen, E.A., Thorbjornsson, B., Thordardottir, M., Saemundsdottir, J., Kristjansson, S.H., Josefsdottir, K.S., Masson, G., Georgsson, G., Kristjansson, M., Moller, A., Palsson, R., Gudnason, T., Thorsteinsdottir, U., Jonsdottir, I., Sulem, P., Stefansson, K., . Humoral immune response to sars-cov-2 in iceland. *New England Journal of Medicine* doi:10.1056/NEJMoA2026116.

Harrison, E.A., Wu, J.W., 2020. Vaccine confidence in the time of covid-19. *European Journal of Epidemiology* 35, 325–330. URL: <https://doi.org/10.1007/s10654-020-00634-3>, doi:10.1007/s10654-020-00634-3.

Hernandez-Suarez, C., Mendoza-Cano, O., 2009. Applications of occupancy urn models to epidemiology. *Mathematical Biosciences and Engineering* 6, 509–520.

Hernandez-Suarez, C., Verme, P., Murillo-Zamora, E., 2020. Estimation of the infection fatality rate and the total number of sars-cov-2 infections. *medRxiv* doi:10.1101/2020.04.23.20077446.

Ibarrondo, F.J., Fulcher, J.A., Goodman-Meza, D., Elliott, J., Hofmann, C., Hausner, M.A., Ferbas, K.G., Tobin, N.H., Aldrovandi, G.M., Yang, O.O., . Rapid decay of anti-SARS-CoV-2 antibodies in persons with mild COVID-19. *New England Journal of Medicine* doi:10.1056/NEJMc2025179.

Ioannidis, J., 2020. The infection fatality rate of COVID-19 inferred from seroprevalence data. *medRxiv* doi:10.1101/2020.05.13.20101253.

Kermack, W.O., McKendrick, A.G., 1927. A contribution to the mathematical theory of epidemics. *Proceedings of the Royal Society of London. Series A* 115, 700–721.

- 240 Le, T.T., Cramer, J., Chen, R., Mayhew, S., 2020. Evolution of the covid-19 vaccine development landscape. *Nat Rev Drug Discov* .
- Liu, Y., Gayle, A.A., Wilder-Smith, A., Rocklöv, J., 2020. The reproductive number of COVID-19 is higher compared to SARS coronavirus. *Journal of Travel Medicine* 27. doi:10.1093/jtm/taaa021. taaa021.
- 245 Long, Q.X., Tang, X.J., Shi, Q.L., Li, Q., Deng, H.J., Yuan, J., Hu, J.L., Xu, W., Zhang, Y., Lv, F.J., Su, K., Zhang, F., Gong, J., Wu, B., Liu, X.M., Li, J.J., Qiu, J.F., Chen, J., Huang, A.L., 2020. Clinical and immunological assessment of asymptomatic SARS-CoV-2 infections. *Nature Medicine* doi:10.1038/s41591-020-0965-6.
- 250 Mukhopadhyay, S., Chakraborty, D., 2020. Estimation of undetected covid-19 infections in india. medRxiv doi:10.1101/2020.04.20.20072892.
- Najafimehr, H., Mohamed Ali, K., Safari, S., Yousefifard, M., Hosseini, M., 2020. Estimation of basic reproduction number for covid-19 and the reasons for its differences. *International Journal of Clinical Practice* 74. doi:10.1111/ijcp.13518.
- 255 NYC-Health, . COVID-19: Data. <https://www1.nyc.gov/site/doh/covid/covid-19-data.page>. Accessed: 2020-09-03.
- Peeples, L., 2020. News feature: Avoiding pitfalls in the pursuit of a covid-19 vaccine. *Proceedings of the National Academy of Sciences* 117, 8218–8221. URL: <https://www.pnas.org/content/117/15/8218>, doi:10.1073/pnas.2005456117, arXiv:<https://www.pnas.org/content/117/15/8218.full.pdf>.
- 260 Sanche, S., Lin, Y.T., Xu, C., Romero-Severson, E., Hengartner, N., Ke, R., 2020. High contagiousness and rapid spread of severe acute respiratory syndrome coronavirus 2. *Emerging Infectious Diseases* 26, 1470.
- 265 Streeck, H., Schulte, B., Kuemmerer, B., Richter, E., Hoeller, T., Fuhrmann, C., Bartok, E., Dolscheid, R., Berger, M., Wessendorf, L., Eschbach-Bludau,

M., Kellings, A., Schwaiger, A., Coenen, M., Hoffmann, P., Noethen, M., Eis-  
Huebinger, A.M., Exner, M., Schmithausen, R., Schmid, M., Kuemmerer, B.,  
270 2020. Infection fatality rate of SARS-CoV-2 infection in a german community  
with a super-spreading event. medRxiv doi:10.1101/2020.05.04.20090076.

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280 disseminating results for the use of practitioners and policy makers. This work  
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### **5. Appendix**

Table 1: Contact rate  $\lambda$  and  $R_0$  for the first 40 countries with diagnosed SARS-CoV-2.  $R_0$  was calculated at the beginning of the epidemic in each country.

Country	$\lambda$	$R_0$	$R^2$	Country	$\lambda$	$R_0$	$R^2$
Italy	0.366	5.79	0.997	Philippines	0.237	3.748	0.988
Spain	0.352	5.569	0.995	Colombia	0.235	3.708	0.996
Netherlands	0.329	5.196	0.998	Denmark	0.234	3.696	0.994
Germany	0.322	5.083	0.999	Poland	0.234	3.694	0.992
United Kingdom	0.319	5.036	0.995	Norway	0.233	3.681	0.997
Belgium	0.306	4.844	0.996	Romania	0.229	3.627	0.987
France	0.294	4.645	0.979	Chile	0.226	3.569	0.996
US	0.29	4.579	0.993	Greece	0.224	3.545	0.994
Peru	0.289	4.571	0.996	Russia	0.223	3.528	0.996
Mexico	0.284	4.494	0.998	Pakistan	0.218	3.45	0.993
Turkey	0.283	4.48	0.998	Israel	0.218	3.444	0.995
Brazil	0.282	4.461	0.997	Morocco	0.216	3.418	0.993
Switzerland	0.273	4.323	0.995	Ukraine	0.215	3.393	0.995
Sweden	0.273	4.31	0.982	Panama	0.211	3.333	0.991
Ecuador	0.26	4.116	0.997	Hungary	0.206	3.252	0.994
Austria	0.254	4.009	0.992	Slovenia	0.204	3.222	0.998
India	0.248	3.925	0.99	Finland	0.201	3.18	0.994
Portugal	0.248	3.922	0.992	UA Emirates	0.185	2.926	0.993
Canada	0.247	3.903	0.996	Nigeria	0.173	2.732	0.995
Ireland	0.246	3.888	0.992	Sudan	0.173	2.73	0.995