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# SARS-CoV-2 infection incidence during the first and second COVID-19 waves in Italy

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# ABSTRACT

We assessed the relation between COVID-19 waves in Italy, which was severely affected during the pandemic. We evaluated the hypothesis that a larger impact from the first wave (February–May 2020) predicts a smaller peak during the second wave (September–October 2020), in the absence of local changes in public health interventions and area-specific differences in time trends of environmental parameters. Based on publicly available data on province-specific SARS-CoV-2 infections and both crude and multivariable cubic spline regression models, we found that for provinces with the lowest incidence rates in the first wave, the incidence in the second wave increased roughly in proportion with the incidence in the first wave until an incidence of about 500–600 cases/100,000 in the first wave. Above that value, provinces with higher incidences in the first wave experienced lower incidences in the second wave. It appears that a comparatively high cumulative incidence of infection, even if far below theoretical thresholds required for herd immunity, may provide noticeable protection during the second wave. We speculate that, if real, the mechanism for this pattern could be depletion of most susceptible individuals and of superspreaders in the first wave. A population learning effect regarding cautious behavior could have also contributed. Since no area-specific variation of the national policy against the SARS-CoV-2 outbreak was allowed until early November 2020, neither individual behaviors nor established or purported environmental risk factors of COVID-19, such as air pollution and meteorological factors, are likely to have confounded the inverse trends we observed in infection incidence over time.

# **1. Introduction**

Coronavirus disease (COVID-19) is a severe and potentially lifethreatening disease due to infection with the severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2), which spread globally during 2020 after the initial outbreak in late 2019 in China [\(Docea et al., 2020](#page-4-0)). Its transmission occurs predominantly through air [\(Wiersinga et al.,](#page-5-0)  [2020\)](#page-5-0). Therapeutic options to treat the infection and its sequelae are limited [\(CORIST, 2020](#page-4-0); [Tsatsakis et al., 2020a](#page-5-0); [Wiersinga et al., 2020](#page-5-0)). Before a vaccine was available, and in the absence of a highly effective therapy, attempts to control the SARS-CoV-2 pandemic during 2020 relied on public health measures such as donning protective gear, social distancing, and shutdowns that limited population mobility and interaction ([Ma and Lipsitch, 2021;](#page-4-0) [Vinceti et al., 2020\)](#page-5-0). Reduced social interaction lowers  $R_t$ , and can stem the spread of the virus, but lockdowns cannot be sustained indefinitely. With normal social intercourse, protection from herd immunity is thought to be attainable only at high prevalences of immunity, on the order of at least 50% and likely considerably above that for this virus ([Fontanet and Cauchemez, 2020](#page-4-0); [Omer et al., 2020;](#page-4-0) [Randolph and Barreiro, 2020\)](#page-5-0).

Italy offers a unique opportunity to investigate how successive waves of the infection may interact [\(Lipsitch et al., 2020\)](#page-4-0). Italy was the first nation with widespread population involvement, with a high mortality from COVID-19 in the spring of 2020, followed by a summer with

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relatively low infection incidence after easing of the tight lockdown that was in effect from March 8 through May 2, 2020 ([Goumenou et al.,](#page-4-0)  [2020;](#page-4-0) [Vinceti et al., 2020](#page-5-0)). The second wave arrived in late August and progressed to a peak in late October 2020. A characteristic of the response to the pandemic in Italy is that the non-pharmacological interventions, including social distancing, mask use and the closing of schools and retail establishments, were evenly adopted throughout the country. There were no important local differences in attempted control until November 6, 2020, when area-specific policies driven by the local spread of the outbreak began ([DPCM, 2020](#page-4-0)). This history allows a relatively unconfounded comparison of the first and second waves of COVID-19 epidemiology during the 2020 February–October period between Italian communities, assuming that environmental and behavioral risk factors for COVID-19 such as outdoor air pollution, meteorological effects, and social interaction are close to uniform across Italy ([Ambika et al., 2021;](#page-4-0) [Calina et al., 2021;](#page-4-0) [Copat et al., 2020](#page-4-0); [Domingo and Rovira, 2020](#page-4-0); [Filippini et al., 2020, 2021](#page-4-0); [Marques et al.,](#page-4-0)  [2021; Mele et al., 2021;](#page-4-0) [Sidiropoulou et al., 2021](#page-5-0); [Sunyer et al., 2021](#page-5-0); [Tsatsakis et al., 2020b\)](#page-5-0). The vaccination campaign did not begin in Italy until December 27, 2020 ([Ministry of Health, 2021](#page-4-0)). Here we consider how the two waves of infection interacted within Italian communities during the 2020 February–October period.

#### **2. Methods**

# *2.1. Study population and outcome*

We used data available without cost from public sources. The number of newly diagnosed infections with SARS-CoV-2 was reported on the website of the Civil Protection Agency ([CPD - Italian Civil Protection](#page-4-0)  [Department, 2020](#page-4-0)) for two time frames: from February 24-May 31, 2020 (first wave), and from September 1-October 31, 2020 (second wave). These newly diagnosed infections correspond to the new positive tests of infection based on quantitative reverse transcription polymerase chain reaction reported by regional Health Services. Daily data flow from all regions of Italy was mandatory. Based on these data and the population data available at the Italian National Institute of Statistics website at January 1, 2020 [\(ISTAT, 2020b\)](#page-4-0), we computed wave-specific incidence of SARS-CoV-2 infection during the first and second waves. We also used the regional anti-SARS-CoV-2 antibody seroprevalence data made available by the National Institute of Statistics ([ISTAT, 2020a](#page-4-0)).

# *2.2. Data analysis*

We assessed the association between first-wave incidence and regional seroprevalence data made available in the national survey with a weighted linear regression analysis, weighting observations by the ratio of provincial or regional population size to average size for all Italian provinces or regions. To investigate the relation between first and second wave province-specific incidence rates, we used linear regression to fit a restricted cubic spline model that weighted provincial data by population size and based on three knots at fixed percentiles of firstwave distribution. We also calculated a pointwise 95% confidence interval (CI) for the forecasted second-wave incidence. We used Stata software (Version-16.1 Stata Inc., College Station-TX, 2021) for all analyses, based on a code developed by NO and TF and available on request.

#### **3. Results**

[Fig. 1](#page-3-0) shows SARS-CoV-2 incidence in the investigated periods. In the first wave, the number of diagnosed cases was 233,019, while in the shorter period of the second wave it was 409,241. Accordingly, the cumulative incidence during the first wave was 387/100,000 as a national average, ranging from 60/100,000 in Calabria region to 943/ 100,000 in Aosta Valley region. Corresponding figures for the slightly shorter period of the second wave were 679/100,000 for the national average, ranging from 171 (Calabria) to 1571/100,000 (Aosta Valley). The region-specific cumulative incidence and seroprevalence were, as expected, strongly correlated (Supplemental Fig. S1).

In the spline model we found an inverted U-shaped relation between first-wave incidence and the estimated predicted mean of the secondwave incidence computed on residents uninfected during the first wave, i.e. the susceptible population [\(Fig. 2](#page-3-0)). If the first-wave incidence was relatively low, the incidence was positively correlated with the incidence in the second wave. If the first-wave incidence, however, was greater than about 500–600 cases/100,000, the incidence was negatively correlated with the second-wave severity. These findings were similar when we computed the incidence in the second wave among all residents, independently of having been infected during the first wave, rather than just in the susceptible population (Supplemental Fig. S2). Adding potential confounders to the regression models such as proportion of elderly people, residents living alone, and an indicator of mobility, also did not appreciably change the results (Supplemental Fig. S<sub>3</sub>). Based on the relation between estimated seroprevalence and firstwave incidence, and the downward trend on the right side of the inverted U distribution in [Fig. 2](#page-3-0), we estimated that the seroprevalence threshold for a protective effect in the second wave is around approximately 2.8%–3.6%. Values above this cutpoint, especially much higher levels, were associated with more limited spread during the second wave.

### **4. Discussion**

We saw an inverted U-shaped relation between first-wave and second-wave incidence. Above 500–600 infection cases/100,000, greater first-wave incidence predicted progressively smaller secondwave incidence. This inflection point corresponded with a populationbased anti-SARS-CoV-2 antibody prevalence of approximately 3%. The official case numbers from which we computed first and second wave infection rates underestimated the actual incidence, since they mostly comprised symptomatic cases, particularly during the first wave. Nonetheless, these figures were highly correlated with the seroprevalence data at the regional level.

The seroprevalence levels corresponding to these first-wave incidences are much lower than those expected to bring about herd immunity, which for COVID-19 is calculated to be above 50% ([Fontanet](#page-4-0)  and Cauchemez,  $2020$ ). The R<sub>t</sub> and the corresponding level for herd immunity could be affected by heterogeneity of behavior, if high-risk people are more likely to have contracted and spread the disease in the first wave (such as superspreaders, frequent travelers, nursing home and hospital personnel) ([Calo et al., 2020](#page-4-0); [Kault, 2020;](#page-4-0) [Kochanczyk](#page-4-0)  [et al., 2020](#page-4-0); [Lewis, 2021; Ma and Lipsitch, 2021](#page-4-0); [Signorelli et al., 2020\)](#page-5-0) and therefore play a smaller role in the spread of the virus during a second wave. In fact, individuals most likely to contract and to spread the virus were also most likely to be infected early and after that to show immunity, diminishing  $R_t$  during later waves on the presumption that immunity persists during the interval between waves (Braun et al., [2020; Long et al., 2020](#page-4-0); [Wajnberg et al., 2020](#page-5-0)). This presumption seems reasonable, as a study from China suggested that the duration of neutralizing antibodies against SARS-CoV-2 persisted for at least nine months [\(He et al., 2021\)](#page-4-0). In addition, recent evidence indicates that seroprevalence estimates, usually based on anti-SARS-CoV-2 antibody prevalence as was the case in Italy and other countries [\(Deeks et al.,](#page-4-0)  [2020;](#page-4-0) [ISTAT, 2020a;](#page-4-0) [Pollan et al., 2020](#page-5-0)), considerably underestimate the prevalence of effective immunity. T-cell mediated immunity may occur even in anti-SARS-CoV-2 antibody negative individuals ([Braun](#page-4-0)  [et al., 2020;](#page-4-0) [Sekine et al., 2020\)](#page-5-0), perhaps from cross-reactivity from other human coronavirus infections [\(Calina et al., 2020](#page-4-0); [Kostoff et al.,](#page-4-0)  [2020\)](#page-4-0), such as those accounting for 20% of upper respiratory tract infections, and might last for years [\(Braun et al., 2020](#page-4-0); [Callow et al., 1990](#page-4-0); [Hope and Bradley, 2021;](#page-4-0) [Jarjour et al., 2021;](#page-4-0) [Ledford, 2021;](#page-4-0) [Neagu](#page-4-0) 

<span id="page-3-0"></span>

**Fig. 1.** Incidence of officially diagnosed SARS-CoV-2 infections in the Italian provinces in the two COVID-19 waves.

[et al., 2021;](#page-4-0) [Ng et al., 2016](#page-4-0)). The effect of this unmeasured immunity could depress the  $R_t$  closer to 1 in populations with high first-wave infection incidence.



**Fig. 2.** Population-weighted cubic spline regression analysis of the relation between first and second wave incidence of SARS-CoV-2 infection in the Italian provinces based on susceptible population (removing subjects with a previous positive swab test up to August 31, 2020). Shadow area, forecasted interval of the mean incidence point estimate; circles, province-specific values weighted for population size.

Populations of the provinces more severely hit by the epidemic during the first wave may have undergone a more profound behavioral change during the summer and the second wave, by voluntarily adhering to prudent behaviors protective against contracting COVID-19. On the other hand, this theory is unlikely to explain the association we observed, because all of Italy engaged in the same set of public health measures adopted to curb the COVID-19 outbreak, including the light and tight lockdowns after the beginning of the outbreak until October 2020, as well as the prescriptions for public health measures such as systematic mask wearing ([Vinceti et al., 2020\)](#page-5-0). It was not until November 6, 2020 that the Italian government activated area-specific differential approaches for COVID-19 prevention and control, including bar, restaurant, shop and school closure, and mobility restrictions, adapted to the spread of the COVID-19 outbreak in each region. Furthermore, neither the SARS-CoV-2 variant B.1.1.7, a variant first detected in the UK in late 2021, nor the South African and Brazilian variants, were detected in Italy during the period investigated in this study ([ISS, 2021\)](#page-4-0), making it unlikely that there was any confounding related to lineage-specific transmission dynamics of the virus [\(Davies](#page-4-0)  [et al., 2021\)](#page-4-0). It appears that the severity of the first wave of the COVID-19 outbreak may have been a determinant of the severity of the subsequent wave, even with comparatively low levels of immunity at the population level.

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# **Credit author statement**

MV and TF designed the original study, and with KJR analyzed and interpreted the data, and wrote the manuscript. NO designed and carried out data analysis with TF and SDF. All authors read and approved the final manuscript.

#### **Declaration of competing interest**

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

# **Appendix A. Supplementary data**

Supplementary data to this article can be found online at [https://doi.](https://doi.org/10.1016/j.envres.2021.111097)  [org/10.1016/j.envres.2021.111097.](https://doi.org/10.1016/j.envres.2021.111097)

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