

Transmissibility and geographic spread of the 1889 influenza pandemic

Alain-Jacques Valleron^{a,b,c,1}, Anne Cori^{a,b}, Sophie Valtat^{a,b}, Sofia Meurisse^c, Fabrice Carrat^{a,b,c}, and Pierre-Yves Boëlle^{a,b}

^aInstitut National de la Santé et de la Recherche Médicale, U 707, F-75012 Paris, France; ^bUniversité Pierre et Marie Curie-Paris 6, UMR-S 707, F-75012 Paris, France; ^cAssistance Publique-Hôpitaux de Paris, Hôpital Saint Antoine, Unité de Santé Publique, F-75012 Paris, France

Edited by Barry R. Bloom, Harvard School of Public Health, Boston, MA, and approved March 26, 2010 (received for review January 30, 2010)

Until now, mortality and spreading mechanisms of influenza pandemics have been studied only for the 1918, 1957, and 1968 pandemics; none have concerned the 19th century. Herein, we examined the 1889 “Russian” pandemic. Clinical attack rates were retrieved for 408 geographic entities in 14 European countries and in the United States. Case fatality ratios were estimated from datasets in the French, British and German armies, and morbidity and mortality records of Swiss cities. Weekly all-cause mortality was analyzed in 96 European and American cities. The pandemic spread rapidly, taking only 4 months to circumnavigate the planet, peaking in the United States 70 days after the original peak in St. Petersburg. The median and interquartile range of clinical attack rates was 60% (45–70%). The case fatality ratios ranged from 0.1% to 0.28%, which is comparable to those of 1957 and 1968, and 10-fold lower than in 1918. The median basic reproduction number (R_0) was 2.1, which is comparable to the values found for the other pandemics, despite the different viruses and contact networks. R_0 values varied widely from one city to another, and only a small minority of those values was within the range in which modelers’ mitigation scenarios predicted effectiveness. The 1889 and 1918 R_0 correlated for the subset of cities for which both values were available. Social and geographic factors probably shape the local R_0 , and they could be identified to design optimal mitigation scenarios tailored to each city.

epidemics | history | reproduction number | attack rate | epidemiology

The ongoing A/H1N1 global influenza epidemic was qualified as a “pandemic” on 11 June 2009, when the World Health Organization raised its alert to phase 6. This new episode adds to a series of 11 pandemics that occurred in the 18th, 19th, and 20th century (1). Interpandemic intervals have ranged from 8 (between 1781 and 1789) to a maximum of 42 years (between 1847 and 1889). This maximum value of 42 was just 1 year more than the time elapsed between the last pandemic of the 20th century (1968) and the first pandemic of the 21st century (2009).

During this long period, the threat of the pandemic—and the need for preparedness—increased with the years passed, which fueled the investigation of mitigation scenarios based on social distancing, vaccination, and/or antiviral treatment (2–7). These studies have relied on mathematical modeling for which “plausible” parameters could be derived only from analysis of past pandemics (2, 6). Most studies focused on 1918, which became de facto the stereotype of the feared pandemic to come. Many researchers extracted quantitative information, especially the value of the basic reproduction number R_0 (the mean number of persons infected by a single infectious individual introduced into a totally susceptible population) for that pandemic.

Comparatively, the two other pandemics of the 20th century (1957 and 1968) were much less studied, and no quantitative analysis at all was undertaken for any of the 19th century pandemics, most likely because it was wrongly assumed that not enough data existed to support such analyses. We focused here on the 1889–1890 season when the last pandemic of the 19th century, known as the “Russian flu,” occurred. The 1889 Russian flu, possibly caused by an H3N8 virus (8), was the first pandemic

to occur in a highly connected world: at that time, the 19 largest European countries, including Russia, had 202,887 km of railroads (9), which is more than now (10) (Appendix S1, section 1.2 showing the 19th century railway network). Transatlantic travel by boat took less than 6 days at that time (11), instead of less than 1 day now (which is not a substantial difference, given the time scale of the global spread of a pandemic).

We searched all available information on the 1889 pandemic, and analyzed its spread, the sizes and dynamics of the local epidemics, and the associated mortality experience. Our goals were to extend the knowledge on influenza pandemics, and to verify that the study of global pandemics of the 19th century was feasible.

Results

Global Spread. The dissemination of the 1889 pandemic was extremely rapid (Fig. 1 and Movie S1). The mortality peaks occurred during the weeks ending 1 December in St. Petersburg, 22 December in Germany, 5 January in Paris, and 12 January in the United States. The median delay between the notification of the first case and the mortality peak was 5.0 weeks [interquartile range (IQR) = 4.0–6.5] (Appendix S1, section 1.5). The mean (\pm SD) speed of the pandemic was estimated at 394 (\pm 255) km/week in continental Europe and 1015 (\pm 727) km/week in the United States.

Clinical Attack Rate and Mortality Burden. The weighted mean of the 408 clinical attack rates (CAR) found was 50%. The median was 60% and the IQR was 45–70%. The 1888–1889 all-causes mortality time course was studied for 96 studied cities of Europe and the United States. The peak size [defined as (maximum mortality – baseline mortality) / baseline mortality in %] varied widely from city to city (Fig. 2A): the mean peak size was 142%, the median was 124%, and the IQR was 87–186%. The mortality burden was estimated in each of the 96 studied cities. The mean was 0.17%, the median was 0.16% of the population, and the IQR was 0.13–0.25%. The 1889 case fatality ratio (CFR) was directly obtained from data in the French and British armies, and was indirectly estimated from surveys performed in seven Swiss cities and in the German army. The CFR ranged between 0.1% and 0.28% (Appendix S1, section 1.7).

Determinants of Local Epidemic Dynamics. The R_0 were estimated in 96 cities studied using a discrete time Susceptible–Exposed (Latent)–Infected–Removed (SEIR) model as described by Mills et al. (12). The median R_0 was 2.1 (IQR = 1.9–2.4) (Fig. 2B). In all, 7% of the R_0 values were less than 1.8 and 22% were less than

Author contributions: A.-J.V. designed research; A.-J.V., A.C., S.V., F.C., and P.-Y.B. performed research; A.C., S.M., and P.-Y.B. analyzed data; and A.-J.V., A.C., S.V., and P.-Y.B. wrote the paper.

The authors declare no conflict of interest.

This article is a PNAS Direct Submission.

¹To whom correspondence should be addressed. E-mail: alain-jacques.valleron@upmc.fr.

This article contains supporting information online at www.pnas.org/lookup/suppl/doi:10.1073/pnas.1000886107/-DCSupplemental.

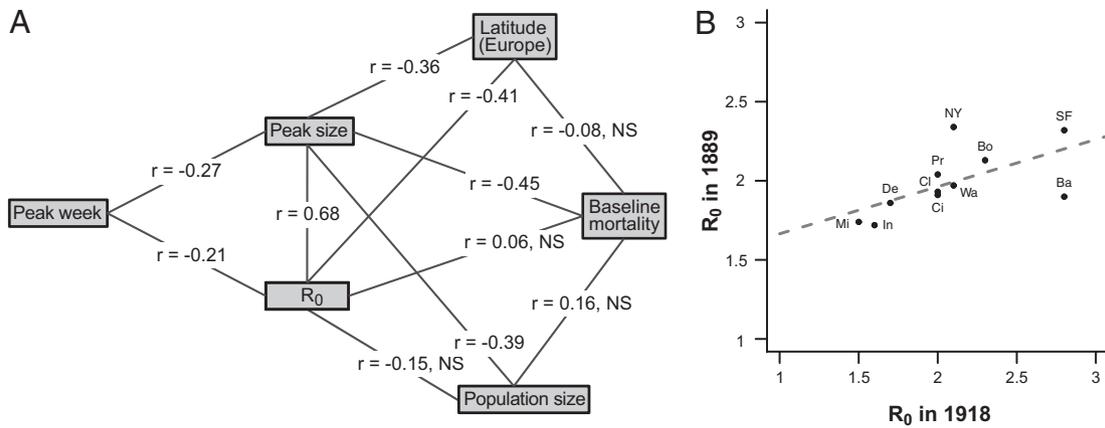


Fig. 3. (A) Spearman rank correlations between R_0 estimations, population size, peak sizes, peak times (for all cities), and latitudes (for European cities). Significance levels: $P < 10^{-5}$ for $|r| > 0.45$; $P < 10^{-4}$ for $|r| > 0.40$; $P < 10^{-3}$ for $|r| > 0.35$, $P < 0.05$ for $|r| > 0.20$. (B) Correlation between 1918 (12) and 1889 R_0 values (this study) for the 11 US cities studied in common (Ba, Baltimore; Bo, Boston; Ci, Cincinnati; Cl, Cleveland; De, Detroit; In, Indianapolis; Mi, Minneapolis; NY, New York; Pr, Providence; SF, San Francisco; Wa, Washington). Pearson's correlation = 0.62 ($P < 0.05$). Dotted line corresponds to linear regression.

would have little, if any, effect (15, 16). One possible hypothesis is that the important predictor of the speed of the pandemic is not the absolute numbers of passengers traveling between cities but the “connectedness” of the network of cities: (i.e., the facility to connect any city A with any city B with railway routes).

The negative correlation that we found between the excess mortality (as measured by the peak value) and the R_0 , on one hand, and the latitude in Europe, on the other hand, is in line with what was observed in 1918, when the mortality burden was higher in the southern region of Europe than in the northern part (17). The correlation analysis showed that there were characteristics of the cities, such as the number of inhabitants per dwelling, that predicted the R_0 , hence the strength of the local epidemic. The fact that the R_0 values were positively correlated almost 30 years apart indicates that there are also intrinsic geographic, weather, and/or sociological characteristics of the cities that could be used to predict which cities are at risk for high R_0 . The identification of such predictors would be of practical interest, as they could guide local mitigation policies.

In conclusion, adding the 1889 pandemic to the short list of the three pandemics studied previously led to the emergence of some similarities among them: R_0 values were consistently ~2

and CAR within the range 30–60% (18–25). The only striking difference concerned mortality: 1889 was comparable to 1957, 1968, and to the “pseudo-pandemics” of 1947 and 1977–1978 (26), thereby making the 1918 pandemic even more exceptional. Finally, the 2009 A/H1N1, at least so far, seems to be another “mild” pandemic, similar to what our analysis revealed the 1889 Russian flu to be.

Methods

Data. We retrieved information on the 1889–1890 all-cause mortality time series in government reports and journals for 172 cities in Europe and the United States, from which the 96 cities with populations exceeding 35,000 inhabitants and a unique mortality peak during the period were selected for analysis. The date of notification of the first case was known in 23 of the 96 cities (Appendix S1, section 1, data sources description, and Dataset S1).

The case fatality ratio (CFR), which is the proportion of infected patients who die from the disease, was estimated in young adults based on statistics collected at the time on several hundreds of thousands of soldiers in the French, British, and German armies. We derived CFR estimations from the ratio of mortality to morbidity statistics obtained in a survey made in seven Swiss cities (total population ~170,000 inhabitants) (Appendix S1, section 1.7).

Data on the clinical attack rate (CAR), which is the proportion of the population with clinical signs during the epidemic, were retrieved from 408

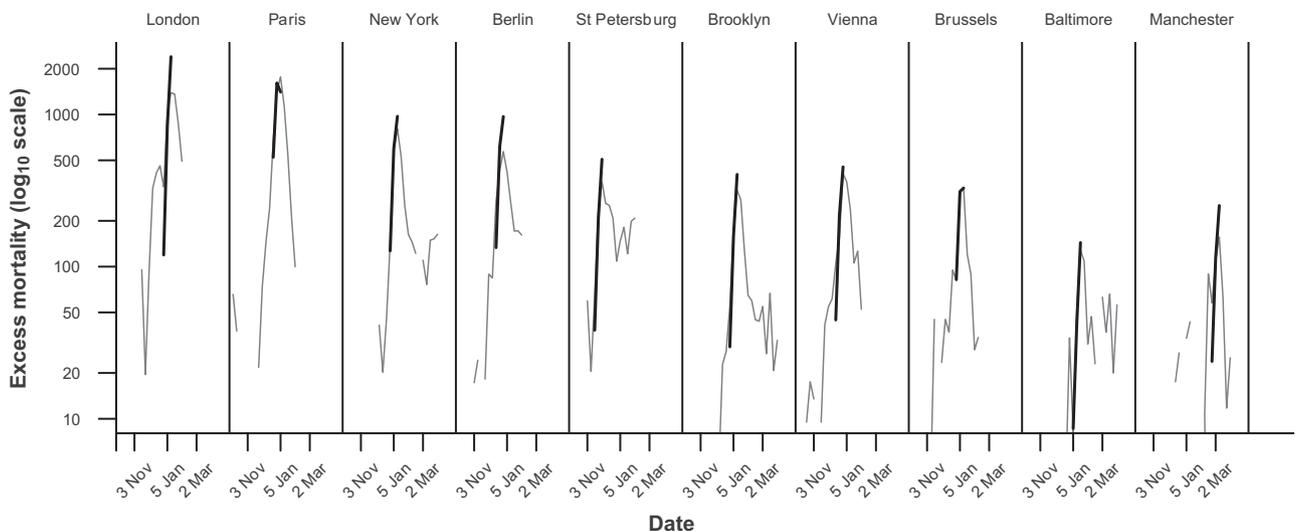


Fig. 4. Fit of influenza mortality rates obtained with the SEIR model for the 10 cities with the largest population sizes. Thin lines indicate raw data; bold lines indicate model fits.

administrative units located in 14 European countries and in the United States (Appendix S1, section 1.6, and Dataset S2). Of these administration units, 331 were from Switzerland, 52 from Germany, and the 31 remaining from 13 countries (Appendix S1, section 1.6 clinical attack rate per country). The weighted mean of CAR was computed for each country, with weights proportional to the countries' population sizes.

Descriptive Analysis of the Mortality Patterns in the 96 Cities. The speed of the epidemic was estimated from the ratios of the distance to the times elapsed between mortality peaks observed in all combinations of two cities.

Baseline mortality was defined by averaging mortality rates over the first 4 weeks of the datasets. This estimation of baseline was found highly correlated to those performed by considering control periods in 1888, or longer, when available (Appendix S1, section 1.4.2). The variation found in the estimates of the baseline guided the sensitivity analyses (Appendix S1, section 3.3). The quality of the excess mortality estimations was assessed via comparison with the mortality rate by respiratory diseases, which was known directly in eight US cities in this study (Appendix S1, section 1.4.2).

The peak size, defined as the relative increase of mortality at the peak's highest amplitude compared with baseline mortality, was chosen as an indicator of the pandemic impact on the mortality in each studied city.

The mortality burden in each city was estimated as the difference between all-cause and baseline mortalities over the period (peak week -4 to peak week +4). In eight cities with sufficiently long datasets, an alternative estimate of the

mortality burden was obtained by using a baseline computed with a periodic regression model (27). The two estimates differed by less than 25% on average.

SEIR Model. We analyzed all-cause-mortality kinetics for the 96 selected cities; their R_0 were estimated using a discrete time SEIR model describing the dynamics of local epidemics after introduction of one infectious case (Appendix S1, section 2). The estimation of the generation time and infectious period was based on the data accrued in a comprehensive review of virus-excretion profiles obtained after experimental influenza infection of 1,280 volunteers (28). That review did not find significant differences between the virus-load kinetics of the groups infected with different influenza strains (B, H1N1, and H3N2). The average generation time (i.e., the time for a primary infected individual to transmit the infection to a secondary case) was 2.6 days. The model that we developed (Appendix S1, section 2.1) found latent and infectious periods lasting respectively 1.6 ± 0.3 and 1.0 ± 1.0 days (means \pm SD). We then fitted the observed excess mortality to the SEIR model, assuming a CFR of 0.2% (Fig. 4 and Appendix S1, section 2.2).

ACKNOWLEDGMENTS. We thank Dr. Mills and Dr. Lipsitch for providing the individual 1918 R_0 values that they estimated, and Dr. Lipsitch for critical comments on an earlier version of this paper. We also thank Camille Pelat and Amandine Lacombe for help in database management and programming, and Georges Marion and Virginie Hoggchoko for help in translation of the German reports. This work was supported by European Union contracts InfTrans (6th PCRD) and FluModCont (7th PCRD).

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