

THE UNIVERSITY of EDINBURGH

Edinburgh Research Explorer

Understanding the potential drivers for respiratory syncytial virus rebound during the COVID-19 pandemic

Citation for published version:

Li, Y, Wang, X, Cong, B, University, S, Feikin, DR & Nair, H 2022, 'Understanding the potential drivers for respiratory syncytial virus rebound during the COVID-19 pandemic', Journal of Infectious Diseases. https://doi.org/10.1093/infdis/jiab606

Digital Object Identifier (DOI):

10.1093/infdis/jiab606

Link: Link to publication record in Edinburgh Research Explorer

Document Version: Publisher's PDF, also known as Version of record

Published In: Journal of Infectious Diseases

General rights

Copyright for the publications made accessible via the Edinburgh Research Explorer is retained by the author(s) and / or other copyright owners and it is a condition of accessing these publications that users recognise and abide by the legal requirements associated with these rights.

Take down policy The University of Edinburgh has made every reasonable effort to ensure that Edinburgh Research Explorer content complies with UK legislation. If you believe that the public display of this file breaches copyright please contact openaccess@ed.ac.uk providing details, and we will remove access to the work immediately and investigate your claim.



Understanding the potential drivers for respiratory syncytial virus rebound during the COVID-19 pandemic

Prof. You Li, PhD^{1,2*}; Xin Wang, PhD^{1,2}; Bingbing Cong, BMed¹; Shuyu Deng, BMed¹; Daniel R Feikin, MD³; Prof. Harish Nair, PhD²

1 School of Public Health, Nanjing Medical University; Nanjing, China

2 Centre for Global Health, Usher Institute, University of Edinburgh; Scotland, United Kingdom

3 Department of Immunizations, Vaccines, and Biologicals, WHO, Geneva, Switzerland

*Corresponding author: Prof. You Li (You.Li@njmu.edu.cn), Department of Epidemiology, School of Public Health, Nanjing Medical University, 101 Longmian Rd., Nanjing, China 211166

Summary: Increasing population susceptibility and full (re)-opening of schools have driven respiratory syncytial virus rebound during the COVID-19 pandemic, overriding the effect of temperature.

© World Health Organization, 2022. All rights reserved. The World Health Organization has granted the Publisher permission for the reproduction of this article.

This is an Open Access article distributed under the terms of the Creative Commons Attribution 3.0 IGO License (https://creativecommons.org/licenses/by/3.0/igo/) which permits unrestricted reuse, distribution, and reproduction in any medium, provided the original work is properly cited.

zce

Non-pharmaceutical interventions (NPIs) were widely introduced to combat the COVID-19 pandemic. These interventions also likely led to substantially reduced activity of respiratory syncytial virus (RSV). From late 2020, some countries observed out-of-season RSV epidemics. Here, we analyzed the role of NPIs, population mobility, climate, and SARS-CoV-2 circulation in RSV rebound through a time-to-event analysis across 18 countries. Full (re)-opening of schools was associated with an increased risk for RSV rebound (HR = 23.29 [95% CI: 1.09–495.84]); every 5°C increase in temperature was associated with a decreased risk (HR = 0.63 [0.40–0.99]). There was an increasing trend in the risk for RSV rebound over time, highlighting the role of increased population susceptibility. No other factors were found statistically significant. Further analysis suggests increasing population susceptibility and full (re)-opening of schools could both override the counter-effect of high temperatures, which explains the out-of-season RSV epidemics during the COVID-19 pandemic.

Key words: Respiratory syncytial virus; pandemic; seasonality; COVID-19; non-pharmaceutical intervention; temperature; humidity; wind speed; school; susceptibility

Introduction

Respiratory syncytial virus (RSV) is the most common pathogen that causes hospitalisation for pneumonia and bronchiolitis among young children globally [1-3]. RSV seasonal epidemics occur annually in most parts of the world and typically in autumn/winter in temperate regions [4]. Understanding RSV seasonality has important implications for health-care services planning, immunisation strategies, as well as recruitment for clinical trials of RSV prevention and treatment. Following the onset of the COVID-19 pandemic in early 2020, non-pharmaceutical interventions (NPIs) were widely enforced by countries to reduce the spread of the SARS-CoV-2 virus. These interventions also likely resulted in substantial reduction in the circulation of RSV during its typical autumn/winter season in both the northern [5-11] and southern hemispheres [12-15] in 2020. Interestingly, some countries observed delayed out-of-season RSV rebound since late 2020, while other countries have not yet observed any RSV epidemics [5-7, 10, 11, 13, 15-17]. The underlying drivers for RSV rebound in some settings remain unknown. While the relaxation of NPIs can be an important driver[18], other factors such as climate [4, 19] and the possible viral interactions [20-22] could have also played a role in both RSV suppression and subsequent rebound. In this study, we sought to disentangle the role of these factors in RSV rebound through a time-to-event analysis among 18 countries.

Methods

Study design

Overview

This was a multi-country longitudinal observational study. The outcome of interest was the occurrence of RSV rebound since the onset of COVID-19 pandemic. The exposures of interest included school opening status, population mobility, ban on international arrivals, COVID-19 notification rate and meteorological factors. Eighteen countries (Australia, Belgium, Canada, Chile, Denmark, England, France, Iceland, Ireland, Japan, Netherlands, New Zealand, Paraguay, Portugal, Slovenia, South Korea, Spain, Sweden) with available data on both RSV activity (between 2019 and

2021) and exposures of interest were selected (appendix pp 2–3). We followed the STROBE guidelines for the reporting of our study (appendix pp 10–11).

Outcome

Data on RSV weekly activity between 2019 and 2021 were accessed from national/regional viral surveillance reports identified through several previous works on RSV seasonality [4, 5, 20]; detailed data sources for RSV are available in the appendix (pp 2–3). For each RSV season, the season onset was defined based on whether an increasing trend in weekly reported RSV cases was observed. An increasing trend was confirmed when the number of increasing weeks exceeded the number of nonincreasing weeks by five (i.e. five net increasing weeks) in any given intervals. (Figure 1, Panel A) For example, if five consecutive increasing weeks were observed, then the RSV onset would be defined as the fifth increasing week; if one non-increasing week was observed among several increasing weeks, then the RSV onset would not be confirmed until the sixth increasing week (so that the number of net increasing week is 6-1=5). The method for defining RSV season onset used in this study had several advantages compared with other existing methods. First, this method was based on short term trend in positive tests and therefore was relatively less affected by varying testing practice over time (e.g., before and during the COVID-19 pandemic) and among countries. Second, this method was not dependent on annual RSV data and could be used prospectively for detecting RSV season onset for timely response. Third, this method did not require the number of negative tests that were not available in some countries.

RSV rebound was defined as the first RSV onset that occurred after the expected RSV season onset during the COVID-19 pandemic. For each country, the period at risk for RSV rebound started from the expected onset of the first RSV season since the beginning of the COVID-19 pandemic, denoted as week 0, based on the timing of its last pre-pandemic RSV season onset (e.g. if RSV onset was week 40 in 2019, then the expected RSV onset after the beginning of the COVID-19 was week 40 in 2020). The period at risk for RSV rebound ended either when RSV rebound occurred or when the observation ended (the last week of available RSV data by 8-September-2021), whichever came earlier. (**Figure 1**, Panel B)

Exposures

We considered several time-dependent exposures that were perceived to be associated with RSV rebound and had available data. Briefly, we considered the Google retail and recreation community mobility metric as an objective measure for NPI stringency; we included climate factors, daily average temperature, relative humidity, and wind speed; we included a binary indicator of whether countries banned international arrivals from any countries; we included COVID-19 14-day cumulative notification rate (available on a weekly basis); and we included school opening status. For school opening status, three levels were included: fully open, partially open (defined as: [a] open/closed in certain regions only; and/or [b] open/closed for some grade levels/age groups only; and/or [c] open but with reduced in-person class time, combined with distance learning), and closure (as reference). Detailed description of these exposures is in the appendix (p3).

Data analysis

We used a piece-wise additive mixed model (PAMM) for the time-to-event analysis.[23] Briefly, in PAMM, the observation period is broken down into a finite number of intervals and one assumes that hazard rates are piece-wise constant in each of these intervals; then a generalized additive model is applied to estimate the baseline hazard as well as other time-varying covariates semi-parametrically. This was done using the R package "pammtools" [24] and "mgcv" [25]. We first considered a complete model with all exposures included, and then the main model was determined through a stepwise backwards variable elimination process from the complete model by comparing model Akaike information criterion (Appendix pp 5–6). This was to maintain the balance between goodness of fit and parsimony. The complete model is given by:

$$\log(\lambda_i(t; x_{ji})) = \beta_0 + \sum_{j=1}^J \beta_j x_{ji} + f(t),$$

where λ denotes hazard rates; *i* denotes each country; *t* denotes time at risk; x_j denotes the exposure of interest, *j*; *f* denotes a spline smooth function that will be estimated through restricted maximum likelihood (REML).

As the definition for RSV onset was based on the history of RSV activity for five or more weeks, we selected to average the exposures using a five-week time window before fitting the data into the model. We also applied a time lag of two weeks between exposures and outcome considering the possible time lag between RSV infection and reporting. We conducted a series of sensitivity analyses that assessed different RSV definitions, time windows for averaging exposures and time lags between exposures and outcome; we also conducted an ad-hoc sensitivity analysis that used a dichotomous school opening status, school open vs closure (details of all sensitivity analyses in Appendix p4). Furthermore, we conducted an ad-hoc exploratory analysis that allowed for time-varying effects and non-linear effect of temperature (through a spline smooth function); based on this model, we predicted the risk for RSV rebound for the first ten weeks after schools fully reopens or closes at different time (relative to the expected RSV onset) as well as the risk for RSV rebound when schools remain fully open or closed, with varied temperatures.

All statistical analyses and visualisations were conducted using the R software (version 4.0.5).

Results

Countries included

All 18 countries included in the analysis experienced delayed RSV onset. Eleven countries (61%) observed RSV rebound based on data available by 8-September-2021; compared with the expected RSV onset, RSV rebound was delayed by a range of 5 to 54 weeks (**Table 1**). Detailed country-specific data on changes in the exposures of interest over time are in the appendix (pp 7–8).

Drivers of RSV rebound

From the complete model that included all exposures of interest, we found that both partial and full (re)-opening of schools might increase the risk for RSV rebound although the hazard ratio (HR) was not statistically significant for neither. As an independent factor, increased temperatures could reduce the risk (HR = 0.58 for every 5°C increase [95% CI: 0.36-0.95). Other factors did not apparently have an effect on the risk for RSV rebound. (**Figure 2**, Panel A)

Our main model, selected through the backwards model selection process, showed that full (re)opening of schools was associated with an increased risk for RSV rebound (HR = 23.29 [1.09-495.84]) and that every 5°C increase in temperature was associated with a decreased risk for RSV rebound (HR = 0.63, 95% CI: 0.40-0.99). Partial (re)-opening of schools was not found to be associated with the risk for RSV rebound. (**Figure 2**, Panel A) Moreover, there was an increasing trend in the risk for RSV rebound over time since the expected RSV onset in 2020 or 2020/21 season, from both the complete and main models. (**Figure 2**, Panel B)

Sensitivity analysis

Results from predefined sensitivity analyses generally confirmed the findings above. Notably, the findings were sensitive to a less specific definition for RSV onset (i.e. 4 net increasing weeks rather than 5 in the main analysis, which resulted in three more countries having RSV rebound, Bolivia, Denmark, and Portugal). The results from the ad-hoc sensitivity analysis that used a dichotomous school opening status showed statistically non-significant HR estimates for school reopening. (Appendix p9)

Different scenarios on school opening and risk for RSV rebound

Furthermore, we assessed the risks for RSV rebound in the first ten weeks following fully (re)opening of schools or school closures. Full (re)-opening of schools could substantially increase the risk for RSV rebound particularly with decreased temperatures and even at high temperatures (**Figure 3**, Panel A). Closing schools (from fully open) could gradually decrease the risk for RSV rebound although to a lesser extent when temperatures decrease (**Figure 3**, Panel B). The risk for RSV rebound also increased over time (since the expected typical RSV onset) even at high temperatures when schools remain closed (**Figure 3**, Panel C1) or fully open (**Figure 3**, Panel C2).

Discussion

Our findings suggest that full (re)-opening of schools is the predominant risk factor for RSV rebound, increasing the risk for RSV rebound by as much as 23 fold (95% CI: 1.09–459.84). High temperature decreases the risk for RSV rebound, with every 5°C increase reducing the risk by 37% (95% CI: 1–

60). The risk for RSV rebound also increases over time since the expected typical RSV onset, highlighting the role of the increased susceptible population. Our scenario analysis suggests that full (re)-opening of schools can substantially increase the risk for RSV rebound when temperature drops and still increase the risk even at high temperatures. Growing susceptibility and full (re)-opening of schools could both override the counter-effect of high temperatures, which explains the out-of-season RSV epidemics during the COVID-19 pandemic. Based on empirical data, these findings provide timely evidence-based recommendation for the prevention and control of RSV epidemics in the context of COVID-19 pandemic.

The predominant role of full (re)-opening of schools in RSV rebound highlighted in our study is consistent with the findings from a household cohort study in Kenya [26], which suggests that school-age children play an important role in the spread of RSV, especially to infants within the family who are most vulnerable to developing severe RSV disease. Second to full (re)-opening of schools, high temperature could decrease the risk for RSV rebound, which aligns well with its typical season in most temperate countries [4]. Moreover, our findings reveal a continuously increasing trend in the risk for RSV rebound over time. This is likely a result of the increase in RSV susceptible population over time, due to the growing number of newborns after the COVID-19 pandemic who remain naïve to RSV as well as the buildup of the number of older children who were not infected by RSV in early infancy, including the school-age children who play an important role in RSV transmission.

Our scenario analysis suggests that countries in the northern hemisphere that have not observed RSV rebound and therefore have a larger than normal susceptible population might expect RSV rebound soon if schools fully reopen in fall 2021. Health systems in these countries should prepare for a surge in RSV cases that might happen even earlier than their typical RSV season. Our scenario analysis also suggests that school reopening could substantially increase the risk for RSV rebound even at high temperatures. This could help explain the delay in out-of-season RSV rebound observed in some countries, such as UK.

We acknowledge several caveats to interpreting these results. First, while we identified school opening and temperature as important drivers, we might lack the statistical power to rule out other exposures of interest as important risk/protective factors. One example is ban on international arrivals. International travels declined substantially following the COVID-19 pandemic, which slowed the global seeding of RSV and might have delayed the normal RSV season. This is supported by a recent study in Australia by Eden and colleagues [17], which revealed a significant reduction in RSV genetic diversity following the COVID-19 pandemic. Another example is concurrent SARS-CoV-2 activity. Viral interference could play a role in the delayed RSV onset. A recent systematic analysis [20] showed that the 2009 influenza pandemic, in which widescale NPI was not employed, delayed RSV onset on average by 0.58 months, suggesting possible viral interference between the pandemic influenza strain and RSV. More generally, viral interference could also explain why some viruses such as rhinoviruses [10, 27, 28] restored circulation early after NPIs were relaxed whereas the activity of other viruses such as influenza virus [15, 29] remained low. However, we were unable to include these viruses in our model due to the absence of accessible data.

Second, we focused our analysis on the timing of RSV rebound; due to data scarcity, we were unable to evaluate how different factors could affect the magnitude or severity profile of RSV rebound. A modelling study using pre-COVID-19-pandemic RSV data by Baker and colleagues predicted that future RSV rebound would occur with higher-than-usual magnitude [18]. However, RSV rebound with lower-than-usual magnitude was observed in countries such as US and France [30]. A better understanding of how future RSV epidemics would evolve requires the continuation of RSV surveillance, which was interrupted in multiple sites during the COVID-19 pandemic. We were also unable to stratify our analysis by age group due to data scarcity; studies from Australia [31] and France [32] both suggest that compared with the pre-pandemic period, children hospitalised for RSV were significantly older during the COVID-19 pandemic. For school opening, due to the absence of relevant data, we could not further assess the effect of opening of different grades (e.g. primary vs secondary) that are expected to drive RSV transmission differently. We were also unable to consider the opening of pre-school facilities (e.g., day-care centres) due to the absence of relevant data.

Third, in addition to school closure and international travel bans as individual NPIs, we attempted to use the Google retail and recreation mobility as an objective measure for other NPIs (e.g. limits on visits to restaurants, cinemas, shopping malls, etc.) considering that the contexts of these NPIs were often fully comparable among different countries. As a result, we were unable to separate the NPIs out in our study. We were also unable to fully account for several NPIs such as wearing of facecovering and social distancing that could not be captured well by the mobility data.

Fourth, we only selected 18 countries that had available RSV surveillance data and data on all exposures of interest. Tropical countries were underrepresented in our analysis. A recent study from Thailand observed a delay of about two months in the RSV season [10]. Lastly, we were unable to account for any changes in health-care seeking behaviours and health-care practices since the COVID-19 pandemic, which might contribute to a short-term delay in RSV reporting.

By breaking the longstanding periodicity of RSV activity, the ongoing COVID-19 pandemic as well as the public health responses to it, offer a unique opportunity to disentangle different factors that could affect RSV transmission dynamics. Our study highlights full (re)-opening of schools and growing population susceptibility as the predominant drivers for RSV rebound that could override the counter-effect of high temperatures. Our findings could help explain the seasonal RSV epidemics observed in every fall (when schools are opened and temperature drops) in most temperate countries. These findings have important implications for countries' preparedness for RSV rebound and shed light on the mystery of the mechanism of RSV seasonality. Although it remains unknown whether RSV will return to its pre-COVID-19 pandemic seasonality, experience from the previous 2009 influenza pandemic suggests that RSV season restored to normality one year after the pandemic [20]. It will be important to continue, or in some cases re-establish, surveillance for RSV at this stage of the COVID-19 pandemic to both better understand the epidemiology of RSV transmission, as well as prepare for the burden of RSV rebound on the public health system.

Acknowledgments YL, XW and HN are members of the Respiratory Syncytial Virus Consortium in Europe (RESCEU). RESCEU has received funding from the Innovative Medicines Initiatives (IMI) 2 Joint Undertaking under grant agreement number 116019. This Joint Undertaking receives support from the EU's Horizon 2020 Research and Innovation programme and the European Federation of Pharmaceutical Industries and Associations.

Contributors YL and XW conceptualised the study. YL led data acquisition with substantial contribution from BC and SD. YL led data analysis and visualisation with substantial contribution from XW. YL wrote the draft report, and all other authors revised the report critically for important intellectual content. All authors have read and approved the final version of the report. YL, BC and SD had access to the study data, and the corresponding author (YL) had final responsibility for the decision to submit for publication.

Potential conflicts of interest YL and HN report grants from WHO relating to the current study. YL reports grants from Wellcome Trust outside the submitted work. HN reports grants from the Innovative Medicines Initiative, consulting fees from BMGF, Pfizer, and Sanofi, honoraria from Abbvie, support from Sanofi for attending meetings, and participation on advisory boards from Sanofi, Janssen, Novavax, Reviral, Resvinet, and WHO, outside the submitted work. All other authors declare no competing interests. The authors alone are responsible for the views expressed in this article and they do not necessarily represent the views, decisions, or policies of the institutions with which they are affiliated.

Data sharing All data used in this study are publicly available and have been cited in the article or its supplementary appendix.

Funding The Bill & Melinda Gates Foundation, World Health Organization.

Address for correspondence You Li (you.li@njmu.edu.cn), School of Public Health, Nanjing Medical University, 101 Longmian Avenue, Jiangning District, Nanjing 211166, China

References

1. O'Brien KL, Baggett HC, Brooks WA, et al. Causes of severe pneumonia requiring hospital admission in children without HIV infection from Africa and Asia: the PERCH multi-country case-control study. Lancet **2019**; 394:757-79.

2. Li Y, Johnson EK, Shi T, et al. National burden estimates of hospitalisations for acute lower respiratory infections due to respiratory syncytial virus in young children in 2019 among 58 countries: a modelling study. Lancet Respir Med **2021**; 9:175-85.

3. Shi T, McAllister DA, O'Brien KL, et al. Global, regional, and national disease burden estimates of acute lower respiratory infections due to respiratory syncytial virus in young children in 2015: a systematic review and modelling study. Lancet **2017**; 390:946-58.

4. Li Y, Reeves RM, Wang X, et al. Global patterns in monthly activity of influenza virus, respiratory syncytial virus, parainfluenza virus, and metapneumovirus: a systematic analysis. Lancet Glob Health
2019; 7:e1031-e45.

5. van Summeren J, Meijer A, Aspelund G, et al. Low levels of respiratory syncytial virus activity in Europe during the 2020/21 season: what can we expect in the coming summer and autumn/winter? Euro Surveill **2021**; 26:2100639.

6. Olsen SJ, Winn AK, Budd AP, et al. Changes in Influenza and Other Respiratory Virus Activity During the COVID-19 Pandemic - United States, 2020-2021. MMWR Morb Mortal Wkly Rep 2021; 70:1013-9.

7. Agha R, Avner JR. Delayed Seasonal RSV Surge Observed During the COVID-19 Pandemic.
 Pediatrics 2021:e2021052089.

8. Kim JH, Roh YH, Ahn JG, et al. Respiratory syncytial virus and influenza epidemics disappearance in Korea during the 2020-2021 season of COVID-19. Int J Infect Dis **2021**; 110:29-35.

9. Nolen LD, Seeman S, Bruden D, et al. Impact of Social Distancing and Travel Restrictions on Non-Coronavirus Disease 2019 (Non-COVID-19) Respiratory Hospital Admissions in Young Children in Rural Alaska. Clin Infect Dis **2021**; 72:2196-8. 10. Thongpan I, Vichaiwattana P, Vongpunsawad S, Poovorawan Y. Upsurge of human rhinovirus infection followed by a delayed seasonal respiratory syncytial virus infection in Thai children during the coronavirus pandemic. Influenza Other Respir Viruses **2021**.

11. Weinberger Opek M, Yeshayahu Y, Glatman-Freedman A, Kaufman Z, Sorek N, Brosh-Nissimov T. Delayed respiratory syncytial virus epidemic in children after relaxation of COVID-19 physical distancing measures, Ashdod, Israel, 2021. Euro Surveill **2021**; 26.

12. Huang QS, Wood T, Jelley L, et al. Impact of the COVID-19 nonpharmaceutical interventions on influenza and other respiratory viral infections in New Zealand. Nat Commun **2021**; 12:1001.

Foley DA, Yeoh DK, Minney-Smith CA, et al. The Interseasonal Resurgence of Respiratory
 Syncytial Virus in Australian Children Following the Reduction of Coronavirus Disease 2019-Related
 Public Health Measures. Clin Infect Dis 2021:ciaa1906.

14. Varela FH, Scotta MC, Polese-Bonatto M, et al. Absence of detection of RSV and influenza during the COVID-19 pandemic in a Brazilian cohort: Likely role of lower transmission in the community. J Glob Health **2021**; 11:05007.

15. Tempia S, Walaza S, Bhiman JN, et al. Decline of influenza and respiratory syncytial virus detection in facility-based surveillance during the COVID-19 pandemic, South Africa, January to October 2020. Euro Surveill **2021**; 26.

16. Casalegno J-S, Ploin D, Cantais A, et al. Characteristics of the delayed respiratory syncytial virus epidemic, 2020/2021, Rhône Loire, France. Euro Surveill **2021**; 26:2100630.

17. Eden J-S, Sikazwe C, Xie R, et al. Off-season RSV epidemics in Australia after easing of COVID-19 restrictions. medRxiv **2021**:2021.07.21.21260810.

Baker RE, Park SW, Yang W, Vecchi GA, Metcalf CJE, Grenfell BT. The impact of COVID-19
 nonpharmaceutical interventions on the future dynamics of endemic infections. Proc Natl Acad Sci U
 S A 2020; 117:30547-53.

19. Baker RE, Mahmud AS, Wagner CE, et al. Epidemic dynamics of respiratory syncytial virus in current and future climates. Nat Commun **2019**; 10:5512.

20. Li Y, Wang X, Msosa T, de Wit F, Murdock J, Nair H. The impact of the 2009 influenza pandemic on the seasonality of human respiratory syncytial virus: A systematic analysis. Influenza Other Respir Viruses **2021**.

21. Nickbakhsh S, Mair C, Matthews L, et al. Virus-virus interactions impact the population dynamics of influenza and the common cold. Proc Natl Acad Sci U S A **2019**; 116:27142-50.

22. Dee K, Goldfarb DM, Haney J, et al. Human Rhinovirus Infection Blocks Severe Acute Respiratory Syndrome Coronavirus 2 Replication Within the Respiratory Epithelium: Implications for COVID-19 Epidemiology. J Infect Dis **2021**; 224:31-8.

23. Bender A, Groll A, Scheipl F. A generalized additive model approach to time-to-event analysis. Stat Modelling **2018**; 18:299-321.

24. Bender A, Scheipl F. pammtools: Piece-wise exponential additive mixed modeling tools. arXiv **2018**:1806.01042.

25. Wood SN. Stable and efficient multiple smoothing parameter estimation for generalized additive models. J Am Stat Assoc **2004**; 99:673-86.

26. Munywoki PK, Koech DC, Agoti CN, et al. The Source of Respiratory Syncytial Virus Infection In Infants: A Household Cohort Study In Rural Kenya. J Infect Dis **2013**; 209:1685-92.

27. Takashita E, Kawakami C, Momoki T, et al. Increased risk of rhinovirus infection in children during the coronavirus disease-19 pandemic. Influenza Other Respir Viruses **2021**; 15:488-94.

28. Kuitunen I, Artama M, Haapanen M, Renko M. Rhinovirus spread in children during the COVID-19 pandemic despite social restrictions-A nationwide register study in Finland. J Med Virol **2021**.

29. FluCov. FluCov Epi-Bulletin – June 2021. Available at:

https://www.nivel.nl/sites/default/files/algemene-

content/FluCov%20EpiBulletin_June2021_06072021.pdf. Accessed 10-August 2021.

30. Williams TC, Sinha I, Barr IG, Zambon M. Transmission of paediatric respiratory syncytial virus and influenza in the wake of the COVID-19 pandemic. Euro Surveill **2021**; 26:2100186.

31. Foley DA, Phuong LK, Peplinski J, et al. Examining the interseasonal resurgence of respiratory syncytial virus in Western Australia. Arch Dis Child **2021**.

32. Fourgeaud J, Toubiana J, Chappuy H, et al. Impact of public health measures on the post-COVID-19 respiratory syncytial virus epidemics in France. Eur J Clin Microbiol Infect Dis **2021**:1-7.

nusci Reedice Reedice

Country	Start of period at	RSV	End of period at	Duration of period at
	risk for RSV	rebound	risk for RSV	risk for RSV rebound
	rebound (T1)		rebound (T2)	(T2-T1; weeks)
Australia	Week 15, 2020*	Yes	Week 38, 2020	23
Belgium	Week 45, 2020	Yes	Week 8, 2021	16
Canada	Week 42, 2020	Yes	Week 28, 2021	27
Chile	Week 25, 2020	No	Week 27, 2021	62
Denmark	Week 46, 2020	No	Week 20, 2021	34
England	Week 43, 2020	Yes	Week 22, 2021	32
France	Week 45, 2020	Yes	Week 4, 2021	12
Iceland	Week 4, 2021	Yes	Week 9, 2021	5
Ireland	Week 44, 2020	No	Week 30, 2021	39
Japan	Week 26, 2020	Yes	Week 6, 2021	33
Netherlands	Week 50, 2020	Yes	Week 26, 2021	29
New Zealand	Week 24, 2020	Yes	Week 25, 2021	54
Paraguay	Week 21, 2020	No	Week 22, 2021	62
Portugal	Week 49, 2020	No	Week 20, 2021	30
Slovenia	Week 49, 2020	Yes	Week 20, 2021	32
South Korea	Week 41, 2020	No	Week 29, 2021	49
Spain	Week 49, 2020	Yes	Week 20, 2021	30

Table 1. Overview of countries included in the analysis.

Country	Start of period at	RSV	End of period at	Duration of period at
	risk for RSV	rebound	risk for RSV	risk for RSV rebound
	rebound (T1)		rebound (T2)	(T2-T1; weeks)

RSV = respiratory syncytial virus. The period at risk for RSV rebound started at the expected week of RSV onset based on the 2019 data and ended at the week of RSV rebound or the last week of the latest RSV reports (that were available by 8-September-2021), whichever came earlier. *RSV season had already started in Australia in the beginning of 2020 until being interrupted by the COVID-19 pandemic; we selected the week when RSV season was interrupted as the start of the period at risk for Australia.

k certe

Figure 1. Schematic figure of the study design. RSV = respiratory syncytial virus. (A) Definition of RSV onset. The numbers next to the dots denote the difference in weeks between the number of weeks with increased RSV activity and the number of weeks with non-increased RSV activity (i.e. "net increasing weeks"). (B) Definition of period at risk for RSV rebound.

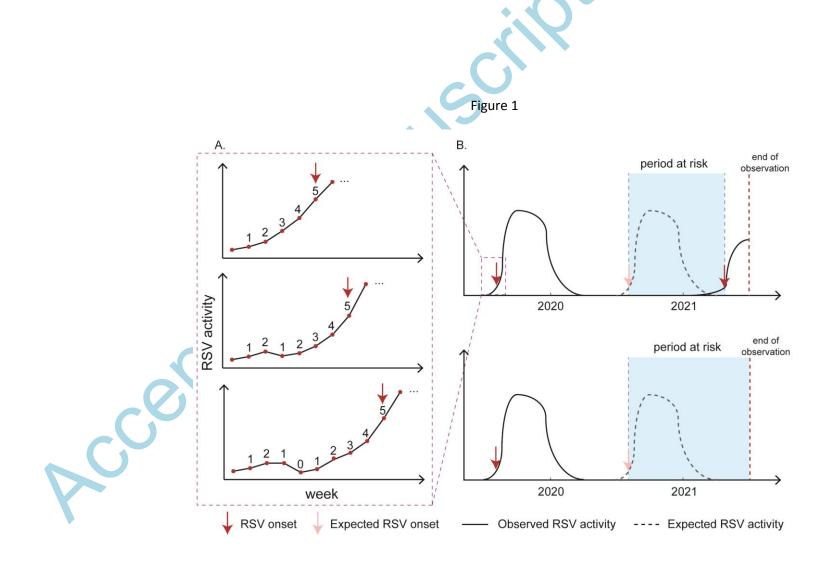
Figure 2. Effect of (A) time-dependent exposures and (B) time length at risk on respiratory

syncytial virus rebound. (A) Dots denote the point estimates and error bars denote the corresponding 95% confidence intervals. (B) the reference is the starting week of observation (i.e. week 0); lines in the middle denote the point estimates and upper and lower lines denote the corresponding 95% confidence intervals.

Figure 3. Predicted risk for RSV rebound under different scenarios on school opening status. A

two-week time lag in the effect of school opening / closing was assumed. For all comparisons, reference temperature was set as 10° C (the median temperature when a typical RSV season occurs in the 18 countries) and reference week was week 0 (i.e. the week when school opening status changes for panels A and B, and the week of typical RSV onset for panel C.) HR = hazard ratio.

Recei



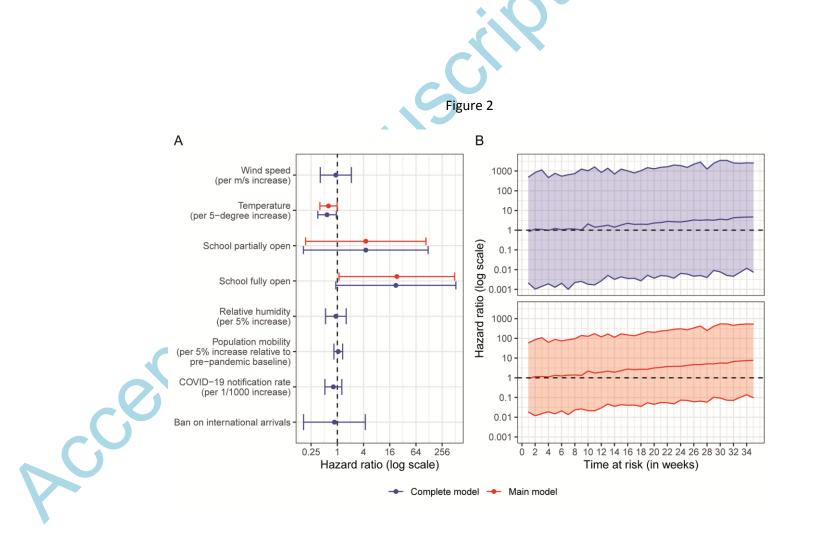


Figure 3

