Effectiveness of Isolation Policies in Schools: Evidence from a Mathematical Model of Influenza and COVID-19

Supplementary Article S1 - Expanded Methods

The text below describes the modeling methodology used in the study. In addition to this text, we provide the source code for the model, the associated data, and the expanded results (<u>https://github.com/sashagutfraind/feverfighter</u>). A live configurable dashboard for applying the model is also available (<u>https://epi1.shinyapps.io/FeverFighter/</u>).

Modeling Methodology

We use the methodology of a multi-stage SEIR model^{1,2}, that uses daily time steps combined with viral shedding and symptom data for influenza and COVID-19. In order to model the disease precisely, the model stratifies the infected population along three dimensions: the student cohort, the degree of isolation, and the stage of disease (Table S1 and Eqn A.1). The model divides the student population into cohorts representing the school grades (the school staff and the community could also be represented as their own cohort(s)). Each cohort is indexed by *i* and includes its own variables: the susceptibles, S_i , infected (explained below), and recovered, R_i . Contact rates across cohorts and within them are described by a contact matrix, giving the model the ability to simulate a diverse set of institutions and communities. The model further divides the cohorts into isolated and unisolated (*H* and *I*).

The model generalizes the classical SEIR model in which there is a sharp division between the exposed but not infectious population, and the infectious population (the E and the I in SEIR). In our model, each stage of the infection has a certain rate of infectiousness and symptoms (which can be zero, for the classical E compartment).

Table S1. Variables of the models. Index <i>i</i> indicates the cohort and <i>d</i> the day of infection (up to 9 and 32 for
influenza and COVID-19, respectively)

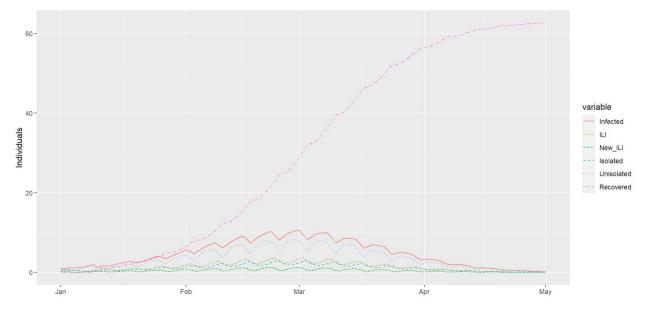
Variable	Interpretation
S_i	Naïve persons in cohort <i>i</i> (for immunity, $\tilde{S}_i(0) = S_i(0)(1 - vv_e)$, see below)
I _{i,d}	Infected persons in cohort i on day d of infection who are not isolated
$H_{i,d}$	Similarly, but who are <i>isolated</i> (e.g. at home)
R _i	Recovered in cohort <i>i</i>

More precisely, let the force of infection λ_i experienced by persons in cohort *i* depend on the number of infected at each stage, the rate of viral shedding (s_d) , and the contact matrix between all the cohorts $(c_{i,i})$. See Eqn A.1.

Eqn A.1. Our modeling framework - the multi-state discrete time SEIR infection model stratified by cohort and day of infection. Variables are dependent on time *t*. Return rates from isolation at day $r_d(p)$ depend on symptom isolation policy p, $\delta_{1,d}$ is the Kronecker delta function (= 1 if d is 1, and otherwise zero). d_f is the final stage of infection. The parameters are listed in Table S2.

$$\begin{aligned} \Delta S_{i} &= -bS_{i}\lambda_{i} \\ H_{i,d} &= (1 - r_{d}(p))(H_{i,d-1} + I_{i,d-1})(1 - \delta_{1,d}) \\ I_{i,d} &= r_{d}(p)(H_{i,d-1} + I_{i,d-1})(1 - \delta_{1,d}) + \delta_{1,d}bS_{i}\lambda_{i} \\ \Delta R_{i} &= H_{i,d_{f}} + I_{i,d_{f}} \\ \lambda_{i} &= \sum_{j}\sum_{d} c_{i,j}s_{d}I_{j,d} \end{aligned}$$

We model vaccination as a reduction in the initially susceptible population for each cohort, which depends on the vaccination rate and the vaccine efficacy: $\tilde{S}_i(0) = S_i(0)(1 - vv_e)$. This transformation could be used to account for any resistance to infection, whether induced, or arising from genetics or prior exposure to similar pathogens.



Typical outbreak curves are shown in Figure S1.

Figure S1. The forecasted epidemic curve of an influenza outbreak in a median school size based on US Department of Education data. Simulation is for the case with no formal isolation policy (where isolation occurs due to voluntary choice alone). Transmission is reduced during weekends, resulting in visible ripples in the outbreak curves every 7 days. ILI, influenza-like illness.

Contact Rates

We assumed that persons have the most physical contact with others in their cohort. The rate of contact with other cohorts is controlled by a parameter which was varied in our sensitivity analysis. To adjust for generally higher contact rates during the winter months, we use a seasonal term cf.^{3,4} which multiplies the baseline contact rate, b_0 , by a factor that peaks on January 1st. :

 $b = b_0(1 + b_s cos(2\pi D/365))$ Eqn A.2

Here, D is the day of the year counting from January 1st. The value b is multiplied by b_h if the day falls on a weekend or by b_c if the school is in closure or vacation (see Table S2).

Sensitivity Analysis

In sensitivity analysis, we assigned the parameters to truncated normal distributions with the mean, standard deviation (SD) and range obtained from previous research or calibration. For influenza, we then adjusted the transmission parameter of the model to give a 25% attack rate (thus matching the typical rate ⁵, after accounting for symptomatic rate⁶). For COVID-19, we adjusted the transmission parameter to give an 11.3% attack rate observed in a representative school outbreak (see Calibration section below). Experiments with alternative attack rates give qualitatively similar results (results not shown).

Table S2. Summary of parameters and their interpretation. All time rate constants are in units of day $^{-1}$. Parameters were obtained from previous studies or, when not available, by calibrating the model to an outbreak (see calibration section below). SD, standard deviation; CI, confidence interval.

Parameter	Interpretation	Range (SD)	Sources, if available
<i>b</i> ₀	Core transmission rate	Influenza: 0.000402 (0) COVID-19: 0.000511 (0)	Calibrated. See section on model calibration
b _h	Relative contact rate during weekends	0.28014 (0.00875)	Calibrated. Comparable to school closure in ⁷
<i>b</i> _c	Relative contact rate during closures, holidays or vacations	0.141479 (0.001822)	Calibrated. Comparable to school closure in ⁷
b_s	Relative seasonal amplitude of transmission	0.478733 (0.001187)	Calibrated
$C_{i,j}$	Relative contact rate for cohorts <i>i</i> and <i>j</i> where	1 if $i = j$ else0.605565 (0.000672)	Calibrated. In some settings, this could be estimated from the degree of epidemiological separations of the cohorts
1	Symptomatic rate	Influenza: 84% (CI: 81-87%) COVID-19: 18.1% (CI:13.9-22.9)	For influenza, based on meta-analysis of asymptomatic rate ⁶ ; for COVID-19 ⁸
р	Fraction of persons complying with isolation	0.158385 (0.199545)	Calibrated. Quantifies compliance with the policy
r _d	Fraction of persons returning to cohort on day <i>d</i> after infection	Based on symptom data	See Table S3 and Table S4 below
S _d	Shedding at day d	Table S3 and Table S4	
t _{start}	Day of first infected case	Input data or calibration	Often reported approximately in post-outbreak investigations
V	Vaccination rate (if vaccine available)	0.8 (0.1)	Seasonal flu: 60% [0.5, 0.7] ⁹ Novel flu strains: 0%
Ve	Vaccine efficacy (if vaccine available)	Influenza 0.5 (0.1) Covid-19: 0.7 (0.2)	For influenza ^{10,11} . Higher mean for COVID-19 based on novel mRNA vaccines. For novel infections with no immune response, 0%.
X _i	Persons in cohort <i>i</i>	Normal: 70 Large: 140	For a typical K-5 for the US ¹² . Six cohorts per school ¹³
у	Fever attention	0.698836 (0.001567)	Calibrated. Fraction of persons who would return from home, of those having symptoms on the previous day

To estimate the rate of symptomatic COVID-19 cases, our priority was to use only studies that had unbiased sampling of all at-risk persons, since surveillance-based methods tend to capture individuals with higher expression of symptoms⁶. Therefore, we adopted the estimate of Poletti et al.⁸ which tested all household members of known COVID-19 cases and is unique in reporting the symptom rate for ages 0 to 19, instead of all age ranges. It used a symptom definition of upper or lower respiratory tract symptoms, or fever \geq 37.5 °C and a sample size of N=692.

Their estimate is consistent with a large random survey from Spain¹⁴ that reported a rate of 34.53% but that included adults that have a higher rate of symptomatic infections^{8,15}.

Modeling of isolation behaviors and the effect of isolation policy

We model the complex interaction between the factors that affect the population. As quantified below, the rate of returning to school is decreased at the more severe stages of the disease, when the disease has a higher symptomatic rate, or when the patients are more aware of their symptoms. As described below, the symptom-based isolation policy is modeled as a decrease in the rate at which students return to school.

To be exact, let the values r_d express the fraction of persons returning from home on day d after they become infected when no policy is in place. It is modeled as a function of four factors: the symptoms in the previous days (f_s) , the symptom propensity (i.e. rate of symptomatic infections) $l \in [0, 1]$, the attention to symptoms $y \in [0, 1]$, and the compliance to policy (if any) $p \in [0, 1]$. To be precise, in the simplest case (no isolation policy), increasing f_{d-1} , l or y would decrease the return rate: $r_d = 1 - lyf_{d-1}$. The introduction of the isolation policy decreases the return to school rate at day d by making the persons more attentive to the recent days of symptoms. Namely, under a one-day isolation policy, the rate is modified to:

$$r_d = 1 - [l(y + (1 - y)p)]f_{d-1}$$
 on day d; Eqn A.3

This model ensures that the rate of symptoms lf_{d-1} sets an upper bound on the effectiveness of the symptom-isolation policy. In the case of 100% compliance and 100% symptom attention, the return rate is $1 - lf_{d-1}$ and not lower. Under a two-day isolation policy, the rate on day *d* is given by replacing f_{d-1} in Eqn A.3 with $max(f_{d-1}, f_{d-2})$, and in general, $max_{i=1, d-1}(f_i)$ for longer isolation policies (see Table S3 and Table S4).

Viral Shedding and Symptom Burden

Influenza viral shedding appears to vary by subtype. However, the patterns of shedding were similar in both children and adults^{16,17}, and between the seasonal and p(H1N1) outbreaks ^{17–19}. We allowed for a proportion *l* of the infections to be asymptomatic^{16,20}, which in our model increases the rates of return to the community and transmission, while also reducing the effectiveness of control policies. Meta-analysis of influenza studies²¹ was used to determine shedding and symptom rates by disease state (see Table S3, Table S4).

Table S3. Daily symptom and shedding rates for influenza from ²¹. Viral shedding rate is based on log10 of titers, and is multiplied by the transmissibility parameter in the model. Symptom scores are relative to their peak. New influenza-like illness (ILI) rates is the probability of newly reporting ILI on a given day. Predicted return rates when symptom propensity is l = 0.84 and symptom attention is y = 0.5. Return rates: A = no policy, B = policy of one day of isolation with 0.5 compliance, C = policy of one day of isolation with 100% compliance. As the rate of compliance rises, the return rate generally decreases since students self-isolate at home. Rows are days from the time of infection. Total Score is an overall indicator of illness reported in ²¹.

		Symptoms	Return scenar	у					
Day after infection	Viral Shedding	Total Score	Fever (Systemic)	Respiratory	Nasal	New ILI Rates	А	В	C
1	1.89	0.25	0.12	0.15	0.18	0.25	1.00	1.00	1.00

2	3.00	0.67	0.93	0.66	0.79	0.50	0.919	0.911	0.903
3	2.63	0.85	0.75	0.95	0.94	0.25	0.347	0.281	0.215
4	2.16	0.67	0.60	0.91	0.91	0.00	0.478	0.425	0.372
5	1.54	0.47	0.30	0.56	0.70	0.00	0.58	0.538	0.496
6	1.07	0.18	0.17	0.56	0.47	0.00	0.789	0.767	0.746
7	0.74	0.06	0.08	0.49	0.17	0.00	0.881	0.869	0.857
8	0.30	0.06	0.07	0.36	0.02	0.00	0.941	0.935	0.929
9	0.35	0.00	0.08	0.14	0.00	0.00	0.951	0.946	0.941

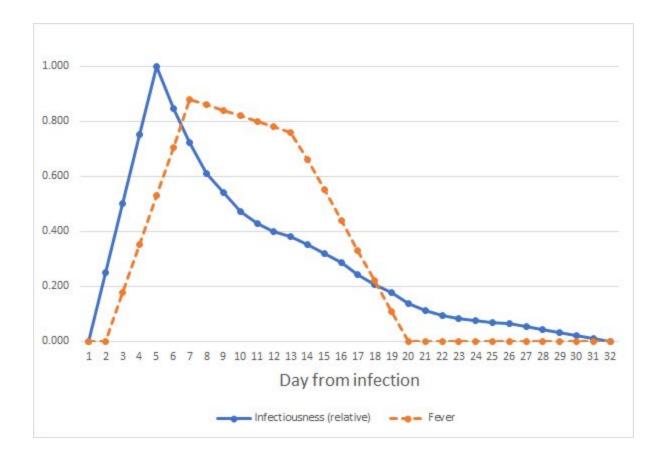


Figure S2. Estimated SARS-CoV-2 infectiousness and symptom propensity in symptomatic cases. Infectiousness is based on 22 with linear interpolation added before day 5 and after day 28^{23-27} . Fever scores for patients with fever are based on collections of case reports²⁸.

Table S4. Estimated symptoms, shedding rates, and return rates for symptomatic persons infected with SARS-CoV-2. Asymptomatic infections are accounted for through a separate parameter. Return rates calculated for when attention is focused on fever, symptom propensity rate is l = 0.1809, and symptom attention is y = 0.5. Return rates: A = no policy, B = policy of one day of isolation with 0.5 compliance, C = policy of one day of isolation with 100% compliance. As the rate of compliance rises, the return rate generally decreases since students self-isolate at home. Rows are days from the time of infection.

		Symptoms	Return rate by scenario					Symp toms	Return	rate by so	cenario
Day after infection	Viral Shedding	Fever (Systemic)	A	В	С	Day after infection	Viral Shed ding	Fever (Syste mic)	A	В	С
1	0.000	0	1	1	1	17	0.244	0.330	0.934	0.927	0.92
2	0.250	0	1	1	1	18	0.207	0.220	0.95	0.945	0.94
3	0.500	0.176	1	1	1	19	0.177	0.110	0.967	0.964	0.96
4	0.750	0.352	0.973	0.971	0.968	20	0.139	0	0.983	0.982	0.98
5	1.000	0.528	0.947	0.942	0.936	21	0.112	0	1	1	1
6	0.845	0.704	0.92	0.912	0.904	22	0.095	0	1	1	1
7	0.721	0.880	0.894	0.883	0.873	23	0.082	0	1	1	1
8	0.611	0.860	0.867	0.854	0.841	24	0.077	0	1	1	1
9	0.540	0.840	0.871	0.857	0.844	25	0.070	0	1	1	1
10	0.473	0.820	0.874	0.861	0.848	26	0.067	0	1	1	1
11	0.427	0.800	0.877	0.864	0.852	27	0.056	0	1	1	1
12	0.400	0.780	0.88	0.867	0.855	28	0.044	0	1	1	1

13	0.379	0.760	0.883	0.871	0.859	29	0.033	0	1	1	1
14	0.351	0.660	0.886	0.874	0.863	30	0.022	0	1	1	1
15	0.318	0.550	0.901	0.891	0.881	31	0.011	0	1	1	1
16	0.287	0.440	0.917	0.909	0.901	32	0.000	0	1	1	1

Model Validation and Calibration

The model has been validated by comparing it to empirical data on three influenza outbreaks and one outbreak of COVID-19. The available data generally indicates the daily number of new ILI cases or the daily number of new absences, and both were also calculated from the model. In all calibrations, we found the model to give a tight fit to the data matching the attack rate, the peak date, and the overall shape, as illustrated in **Figure S3**, **Figure S4**, **and Figure S5**. For COVID-19, the fit is reported in Table S5.

For validation, we needed to estimate from the model the number of individuals with new influenza-like illness (ILI) and the number of newly absent. For ILI, the number of individuals with ILI at a given time *t* is computed by considering $J_s(t)$ - the number of infected individuals on date *t* (including isolated and unisolated) who are in stage *s* and the rate of new ILI cases at stage *s*, l_s . The quantity is multiplied by l_s - the fraction of individuals that

develop ILI (set at 84%²⁰): $ILI(t) = l \sum_{s} l_s J_s(t)$.

The symptom data indicates that ILI symptoms first arise 24 hours after influenza infection and peak around 2.5 days after infection. Accordingly, we assume that, of the population reporting ILI symptoms, $\frac{1}{4}$, $\frac{1}{2}$, and $\frac{1}{4}$ first reports on day 1, 2, and 3, respectively.

We calculated the new absentees per day by considering the number of infected and the return rate. The number of

isolated persons is given by: $\sum_{s} (1 - r_s) I_s(t)$. Furthermore, the number of newly-absent (i.e. isolated) students is given

by $\sum_{s} \prod_{q=1}^{s-1} r_q (1-r_s) I_s(t)$. We will use this formula in calibrating the model to absenteeism data below.

Unknown parameter values and ranges were estimated using a genetic algorithm²⁹. The algorithm adjusted the values of the unknown parameters to match the well-characterized A(H1N1)v influenza outbreak in a boarding school³⁰. Parameter ranges were estimated from the distribution of parameters in the top 10% of the solutions. A separate set of simulations with seasonal influenza calibrated to an outbreak of seasonal influenza ³¹.

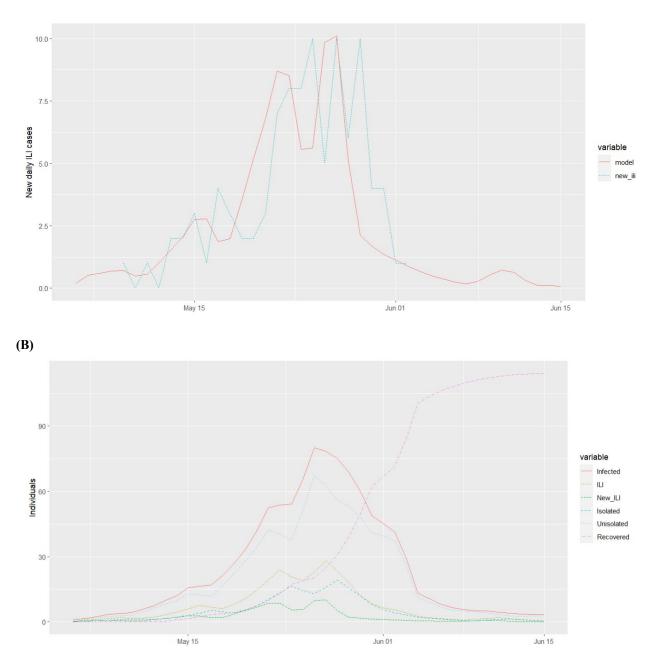


Figure S3. Calibration to data from a boarding school in South East England during the A(H1N1)v outbreak from May 9, 2009 through June 2, 2009, infecting 101 of the 1307 students³⁰. (A) model predictions and actual new cases of influenza-like illness (ILI) and (B) estimated course of the epidemic from the model.

(A)

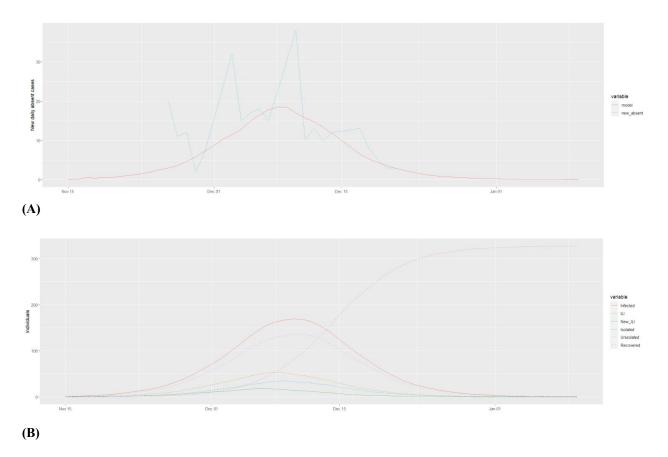


Figure S4. Calibration to absenteeism for a primary school in Thames Valley, UK during the 2012/2013 influenza season abstracted from ³¹. The data shows spikes of new absences every Monday due to lack of data collection on Saturdays and Sundays. (A) model predictions and match to actual new cases of absence from school and (B) estimated course of the epidemic from the model.

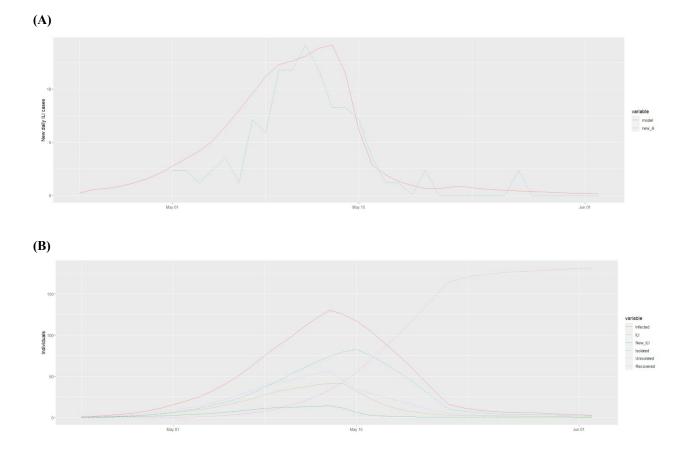


Figure S5. Calibration to data from an elementary school in Pennsylvania affected by the 2009 Pandemic Influenza $A(H1N1)^{32}$. (A) model predictions and actual new cases of influenza-like illness (ILI) and (B) estimated course of the epidemic from the model.

Table S5: Outbreak of COVID-19 in the Rehavia Gymnasia (Jerusalem, Israel) - high school with grades 7-12 containing 1200 students and 180 staff. The school was closed starting 5/28 and complete serosurvey found 159 infections³³.

Date	Reported new cases	Predicted Infected	Predicted daily new symptom atic	Date	Reported new cases	Predicted Infected	Predicted daily new symptom atic
5/11/2020		1	0	5/21/2020		18	0
5/12/2020		1	0	5/22/2020		19	0
5/13/2020		1	0	5/23/2020		20	0
5/14/2020		2	0	5/24/2020		34	0
5/15/2020		2	0	5/25/2020		50	0
5/16/2020		2	0	5/26/2020	1st known case	71	1
5/17/2020		4	0	5/27/2020	2 new cases	98	1
5/18/2020		6	0	5/28/2020	Last day of classes	136	1
5/19/2020		9	0	5/29/2020	School shut down	142	2
5/20/2020		12	0	Final serological survey of the school	Actual infected 159	Predicted infected 159	-

REFERENCES

- 1. Andreasen V, Frommelt T. A School-Oriented, Age-Structured Epidemic Model. *SIAM J Appl Math.* 2005;65(6):1870–1887.
- 2. Pellis L, Ferguson NM, Fraser C. Epidemic growth rate and household reproduction number in communities of households, schools and workplaces. *J Math Biol*. 2011;63(4):691–734.
- 3. Keeling MJ, Rohani P, Grenfell BT. Seasonally forced disease dynamics explored as switching between attractors. *Physica D*. 2001;148(3-4):317–335.
- 4. Grassly NC, Fraser C. Seasonal infectious disease epidemiology. Proc Biol Sci. 2006;273(1600):2541–2550.
- 5. Somes MP, Turner RM, Dwyer LJ, et al. Estimating the annual attack rate of seasonal influenza among unvaccinated individuals: A systematic review and meta-analysis. *Vaccine*. 2018;36(23):3199–3207.
- 6. Leung NHL, Xu C, Ip DKM, et al. The fraction of influenza virus infections that are asymptomatic: a systematic review and meta-analysis. *Epidemiology*. 2015;26(6):862.
- 7. Halloran ME, Ferguson NM, Eubank S, et al. Modeling targeted layered containment of an influenza pandemic in the United States. *Proc Natl Acad Sci U S A*. 2008;105(12):4639–4644.
- 8. Poletti P, Tirani M, Cereda D, et al. Probability of symptoms and critical disease after SARS-CoV-2 infection. *arXiv* [*q*-bioPE]. 2020. http://arxiv.org/abs/2006.08471.
- 9. Tammy A. Santibanez Centers for Disease Control and Prevention. *Flu Vaccination Coverage*. CDC; 2016. https://www.cdc.gov/flu/fluvaxview/coverage-1415estimates.htm. Accessed October 1, 2017.
- 10. Tricco AC, Chit A, Soobiah C, et al. Comparing influenza vaccine efficacy against mismatched and matched strains: a systematic review and meta-analysis. *BMC Med.* 2013;11:153.
- 11. Belongia EA, Simpson MD, King JP, et al. Variable influenza vaccine effectiveness by subtype: a systematic review and meta-analysis of test-negative design studies. *Lancet Infect Dis.* 2016;16(8):942–951.
- 12. US Department of Education. Common Core of Data (CCD). https://nces.ed.gov/ccd/pubschuniv.asp. Accessed October 2, 2020.
- 13. Contributors to Wikimedia projects. Educational stage. https://en.wikipedia.org/wiki/Educational_stage. Published December 6, 2005. Accessed November 16, 2020.
- 14. Pollán M, Pérez-Gómez B, Pastor-Barriuso R, et al. Prevalence of SARS-CoV-2 in Spain (ENE-COVID): a nationwide, population-based seroepidemiological study. *Lancet*. July 2020.
- 15. Ludvigsson JF. Systematic review of COVID-19 in children shows milder cases and a better prognosis than adults. *Acta Paediatr*. 2020;109(6):1088–1095.
- 16. Loeb M, Singh PK, Fox J, et al. Longitudinal study of influenza molecular viral shedding in Hutterite communities. *J Infect Dis.* 2012;206(7):1078–1084.
- 17. Fielding JE, Kelly HA, Mercer GN, et al. Systematic review of influenza A(H1N1)pdm09 virus shedding: duration is affected by severity, but not age. *Influenza Other Respi Viruses*. 2014;8(2):142–150.
- 18. Biggerstaff M, Cauchemez S, Reed C, et al. Estimates of the reproduction number for seasonal, pandemic, and zoonotic influenza: a systematic review of the literature. *BMC Infect Dis.* 2014;14:480.
- 19. Boëlle P-Y, Ansart S, Cori A, et al. Transmission parameters of the A/H1N1 (2009) influenza virus pandemic:

a review. Influenza Other Respi Viruses. 2011;5(5):306-316.

- 20. Call SA, Vollenweider MA, Hornung CA, et al. Does this patient have influenza? *JAMA*. 2005;293(8):987–997.
- 21. Carrat F, Vergu E, Ferguson NM, et al. Time lines of infection and disease in human influenza: a review of volunteer challenge studies. *Am J Epidemiol*. 2008;167(7):775–785.
- 22. He X, Lau EHY, Wu P, et al. Temporal dynamics in viral shedding and transmissibility of COVID-19. *Nature Medicine*. 2000;26:672–675.
- 23. Wyllie AL, Fournier J, Casanovas-Massana A, et al. Saliva is more sensitive for SARS-CoV-2 detection in COVID-19 patients than nasopharyngeal swabs. *Infectious Diseases (except HIV/AIDS)*. April 2020.
- 24. Cheng H-Y, Jian S-W, Liu D-P, et al. High transmissibility of COVID-19 near symptom onset. *Infectious Diseases (except HIV/AIDS)*. March 2020.
- 25. Wölfel R, Corman VM, Guggemos W, et al. Virological assessment of hospitalized patients with COVID-2019. *Nature*. 2020;581(7809):465–469.
- 26. Benefield AE, Skrip LA, Clement A, et al. SARS-CoV-2 viral load peaks prior to symptom onset: a systematic review and individual-pooled analysis of coronavirus viral load from 66 studies. *Infectious Diseases (except HIV/AIDS)*. September 2020.
- van Kampen JJA, van de Vijver DAMC, Fraaij PLA, et al. Shedding of infectious virus in hospitalized patients with coronavirus disease-2019 (COVID-19): duration and key determinants. *Infectious Diseases (except HIV/AIDS)*. June 2020.
- Chan JF-W, Yuan S, Kok K-H, et al. A familial cluster of pneumonia associated with the 2019 novel coronavirus indicating person-to-person transmission: a study of a family cluster. *Lancet*. 2020;395(10223):514–523.
- 29. genalg: R Based Genetic Algorithm. https://CRAN.R-project.org/package=genalg. Accessed October 13, 2020.
- 30. Smith A, Coles S, Johnson S, et al. An outbreak of influenza A(H1N1)v in a boarding school in South East England, May-June 2009. *Eurosurveillance*. 2009;14(27).
- McCann LJ, Suchanek O, McCarthy ND, et al. Descriptive epidemiology of school outbreaks of seasonal influenza B during 2012/2013 in the Thames Valley, United Kingdom. *Public Health*. 2014;128(12):1121–1124.
- Bhattarai A, Villanueva J, Palekar RS, et al. Viral Shedding Duration of Pandemic Influenza A H1N1 Virus during an Elementary School Outbreak—Pennsylvania, May–June 2009. *Clin Infect Dis*. 2011;52(suppl_1):S102–S108.
- 33. ToI Staff. 3 teachers in shuttered Jerusalem school have virus; pupils, staff to be tested. *Times of Israel*. https://www.timesofisrael.com/3-teachers-in-shuttered-jerusalem-school-have-virus-pupils-staff-to-be-tested/. Published May 28, 2020. Accessed July 19, 2020.