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**Research** Special Paper

# Effect of school closures on mortality from coronavirus disease 2019: old and new predictions

BMJ 2020; 371 doi: <https://doi.org/10.1136/bmj.m3588> (Published 07 October 2020) Cite this as: BMJ 2020;371:m3588

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Accepted 15 September 2020

## Abstract

**Objective** To replicate and analyse the information available to UK policymakers when the lockdown decision was taken in March 2020 in the United Kingdom.

**Design** Independent calculations using the CovidSim code, which implements Imperial College London's individual based model, with data available in March 2020 applied to the coronavirus disease 2019 (covid-19) epidemic.

**Setting** Simulations considering the spread of covid-19 in Great Britain and Northern Ireland.

**Population** About 70 million simulated people matched as closely as possible to actual UK demographics, geography, and social behaviours.

**Main outcome measures** Replication of summary data on the covid-19 epidemic reported to the UK government Scientific Advisory Group for Emergencies (SAGE), and a detailed study of unpublished results, especially the effect of school closures.

**Results** The CovidSim model would have produced a good forecast of the subsequent data if initialised with a reproduction number of about 3.5 for covid-19. The model predicted that school closures and isolation of younger people would increase the total number of deaths, albeit postponed to a second and subsequent waves. The findings of this study suggest that prompt interventions were shown to be highly effective at reducing peak demand for intensive care unit (ICU) beds but also prolong the epidemic, in some cases resulting in more deaths long term. This happens because covid-19 related mortality is highly skewed towards older age groups. In the absence of an effective vaccination programme, none of the proposed mitigation strategies in the UK would reduce the predicted total number of deaths below 200 000.

**Conclusions** It was predicted in March 2020 that in response to covid-19 a broad lockdown, as opposed to a focus on shielding the most vulnerable members of society, would reduce immediate demand for ICU beds at the cost of more deaths long term. The optimal strategy for saving lives in a covid-19 epidemic is different from that anticipated for an influenza epidemic with a different mortality age profile.

## Introduction

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IBMIC was developed from an influenza pandemic model.<sup>156</sup> The original code used for Report 9 has not been released. However, the team at Imperial College London, headed by epidemiologist Neil Ferguson, collaborated with Microsoft, GitHub, and the Royal Society Rapid Assistance in Modelling the Pandemic (RAMP) initiative to recreate the model in the CovidSim code: this version has been stringently externally validated.<sup>7</sup> We used GitHub tagged version 0.14.0 plus additional patches dated before 3 June 2020, the full technical details of which are published elsewhere.<sup>8</sup> Ferguson et al<sup>9</sup> supplied the input files relevant to Report 9 that were included in the GitHub release.

CovidSim performs simulations of the UK at a detailed level without requiring personal data. The model includes millions of individual “people” going about their daily business—for example, within communities and at home, schools, universities, places of work, and hospitals. The geographical representation of the UK is taken from census data, so the distribution of age, health, wealth, and household size for simulated people in each area is appropriate. The model also includes appropriate numbers, age distribution, and commuting distances of people in the simulated schools and workplaces, each in line with national averages. The network of interactions is age dependent: people interact mainly with their own age group and with family, teachers, and carers. The virus (severe acute respiratory syndrome coronavirus 2) initially infects random members of this network of interacting coworkers, strangers, friends, and family. Whenever an infected person interacts with a non-infected person, there is a probability that the virus will spread. This probability depends on the time and proximity of the interaction and the infectiousness of the person according to the stage of the disease. Infected people might be admitted to hospital and might die, with the probability dependent on age, pre-existing conditions, and stage of the disease. This extremely detailed model is then parameterised using the best available expert clinical and behavioural evidence,<sup>5</sup> with coronavirus specific features being updated as more coronavirus specific data become available from the worldwide pandemic.<sup>8</sup> Therefore, the model has the required complexity to consider non-pharmaceutical interventions, which would reduce the number of interactions between simulated people in the model ([table 1](#)). To predict policymaking, it is assumed that these interventions are implemented when demand for ICU beds reaches a particular “trigger” level. As the model contains far more realistic detail than the data available, the results are averages over many runs with different starting conditions, all of which are consistent with known data. The real epidemic is just one of these possibilities, so the code determines the range of scenarios for which plans should be made. This is particularly important when the numbers of localised outbreaks are low: the prediction that local spikes will occur somewhere is reliable, and the most likely places can be identified, but predicting exactly when and where is not possible with the level of data available. All interventions reduce the reproduction number and slow the spread of the disease. However, a counterintuitive result presented in Report 9 ([table 3](#) and [table A1](#) in that report) is the prediction that, once all other considered interventions are in place, the additional closure of schools and universities would increase the total number of deaths. Similarly, adding general social distancing to a scenario involving household isolation of suspected cases (case isolation) and household quarantine of family members, with appropriate estimates for compliance, was also projected to increase the total number of deaths.

**Table 1**  
Peak demand for UK-wide intensive care unit beds (in 000s) for different intervention scenarios and different intensive care unit (ICU) triggers during the coronavirus disease 2019 epidemic

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**Table 2**  
Definition of interventions from Report 9 considered in CovidSim

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**Patient and public involvement**

Patients or the public were not involved in the design, conduct, reporting, or dissemination plans of our research. All data used were retrieved from existing public sources, as referenced. We plan to share this on social media, Twitter, and blogs.

**Results**

To reproduce the result tables for the scenarios presented in Report 9, we averaged over 10 simulation runs with the same random number seeds as used in the original report. The simulations are run for 800 days, with day 1 being 1 January 2020. The simulated intervention period lasts for three months (91 days), with some interventions extended for an additional 30 days. In reality, interventions were in place for rather longer, which delayed the second wave but had little effect on deaths. The mitigation scenarios in Report 9 considered reproduction numbers of  $R_0=2.2$  and  $R_0=2.4$ . As highlighted by Ferguson et al,<sup>8</sup> the results we obtain here are not precisely identical to those in Report 9 because they are an average of 10 stochastic realisations, the population dataset has changed to an open source one, and the algorithm used to assign individuals from households to other places such as schools, universities, and workplaces has been modified to be deterministic. We also count deaths in all waves, not just the first. The stochasticity gives a variance of around 5% in total number of deaths and ICU bed demand between different realisations using different random numbers. More important is the uncertainty of the timing of the peak of the infections between realisations, which is around five days. We compared these predictions to the death rates from the actual trajectory of covid-19.<sup>1011</sup> NHS England stopped publishing data on critical bed occupancy in March 2020,<sup>12</sup> so it was not possible to compare ICU data from the model with real world data.

[Table 1](#) shows the demand for ICU beds and [table 3](#) shows the total number of deaths; in both, the same mitigation scenarios as presented in Report 9 were used. As in Report 9, for each mitigation scenario we considered a range of ICU triggers. In [table 1](#) we report the peak ICU bed demand across the full simulation for each trigger, as was presented in Report 9, but we also include the peak demand for ICU beds during the period of the intervention (first wave). The latter we define as the period during

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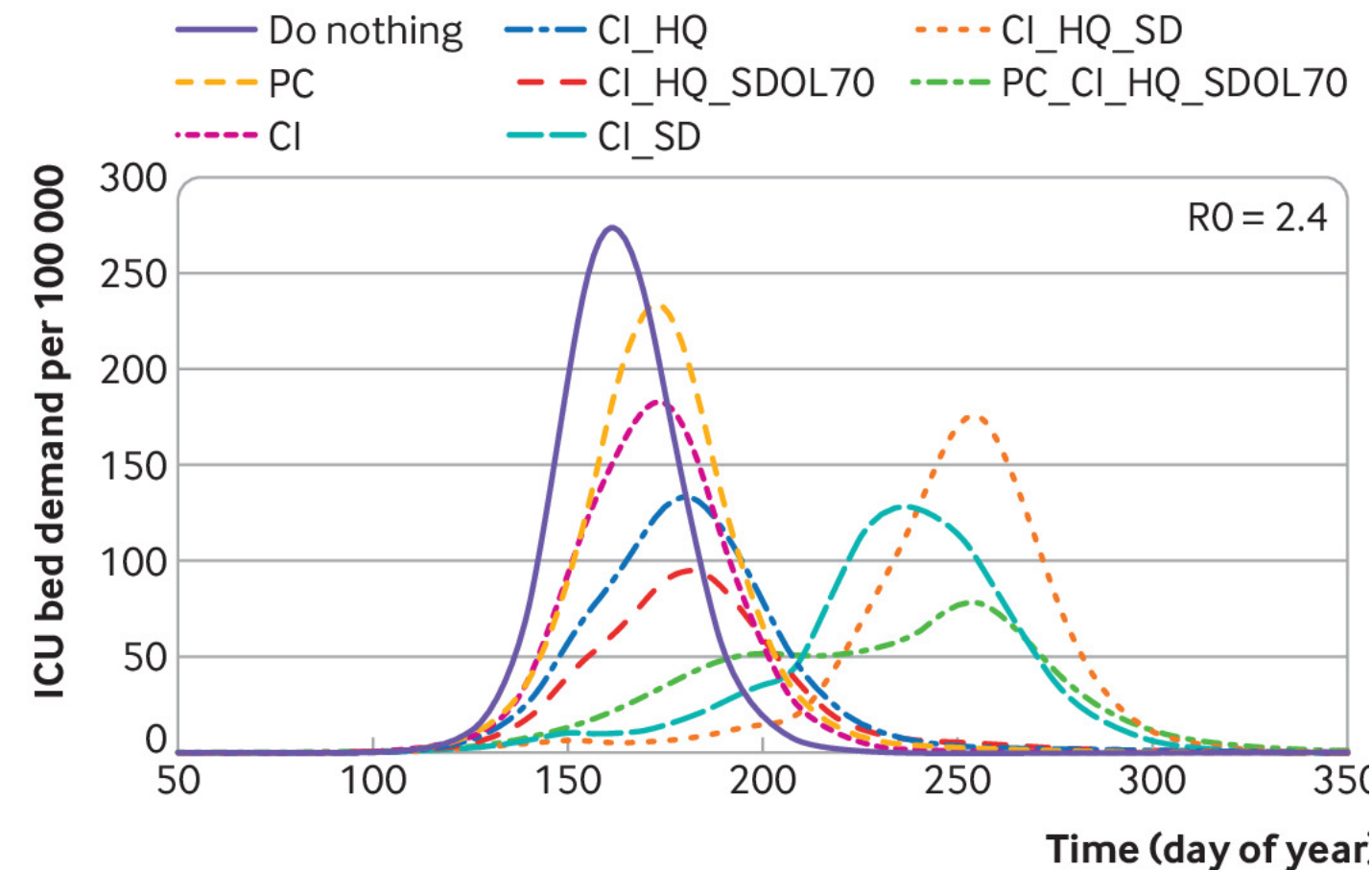
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Table 1 and table 3 show that in some mitigation scenarios the peak demand for ICU beds and most deaths occur during the period when the intervention is in place. There are, however, other scenarios when the opposite is true. The reason for this is illustrated in figure 1. The mitigation scenarios of “do nothing,” place closures, case isolation, case isolation with household quarantine, and case isolation, household quarantine, and social distancing of over 70s are as presented in figure 2 of Report 9. We also show some additional scenarios (case isolation and social distancing; case isolation, household quarantine, and general social distancing; and place closures, case isolation, household quarantine, and social distancing of over 70s) that are not shown in figure 2 of Report 9 but are included in table 1 and table 3 and in the tables in Report 9.



**Fig 1**  
Flattening the curve. The first five curves are the same scenarios as presented in figure 2 of report 9. Three additional scenarios are also shown (summarised in table 1 and table 3). The scenario of place closures, case isolation, household quarantine, and social distancing of over 70s would minimise peak demand for intensive care but prolong the epidemic, resulting in more people needing intensive care and more deaths. These findings illustrate why adding place closures to a scenario with case isolation, household quarantine, and social distancing of over 70s can lead to more deaths than the equivalent scenario without place closures. Doing so suppresses the infection when the interventions are present but leads to a second wave when interventions are lifted. In the model this happened in July 2020, after a 91 day lockdown: in practice the first lockdown was extended into August, so the second wave was postponed to September. The total number of deaths in the scenario of case isolation, household quarantine, and social distancing of over 70s is 260 000, whereas when place closures are included the total number is 350 000. Similarly, comparing general social distancing with equivalent scenarios without social distancing, the second wave peak in the case isolation, household quarantine, and general social distancing scenario is higher than the first wave peak in the case isolation and household quarantine scenario. ICU=intensive care unit; PC=place closures; CI=case isolation; HQ=household quarantine; SDOL70=social distancing of over 70s; SD=general social distancing

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In the simulations presented here, the main interventions are in place for three months and end on about day 200 (some interventions are extended for an additional 30 days). Figure 1 shows that weaker intervention scenarios lead to a single wave that occurs during the period in which the interventions are in place. Hence the peak demand for ICU beds occurs during this period, as do most deaths.

Stronger interventions, however, are associated with suppression of the infection such that a second wave is observed once the interventions are lifted. For example,

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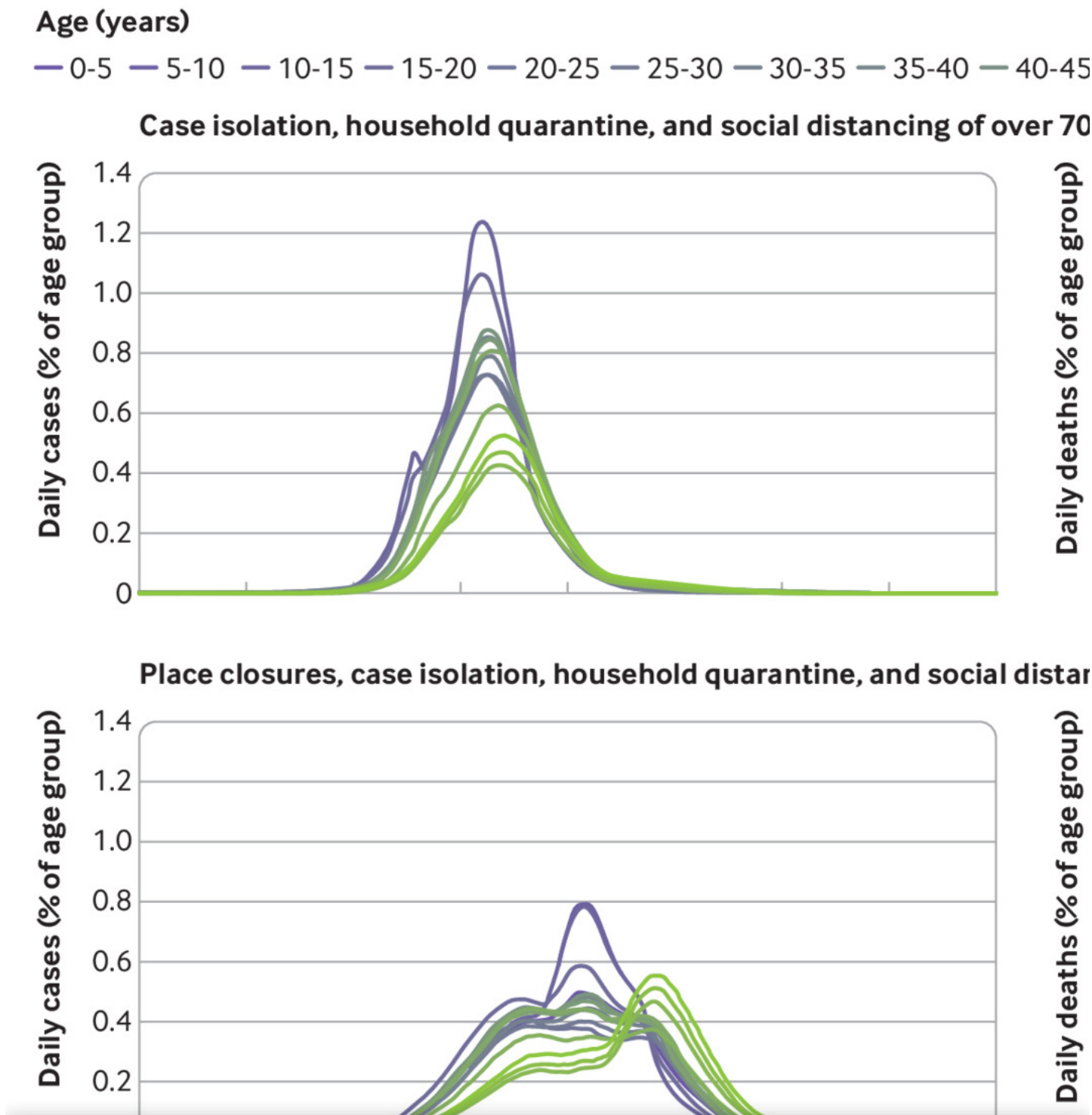
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cases is seen. The data are broken down into age groups, showing that younger people contribute most to the total number of cases, but that deaths are primarily in older age groups. Adding the place closure intervention (and keeping all other things constant) gives the behaviour shown in the second row of plots. The initial peak is greatly suppressed, but the end of place closures while other social distancing is in place prompts a second peak of cases among younger people. This then leads to a third, more deadly, peak of cases affecting elderly people when social distancing of over 70s is removed. Postponing the spread of covid-19 means that more people are still infectious and are available to infect older age groups, of whom a much larger fraction then die.



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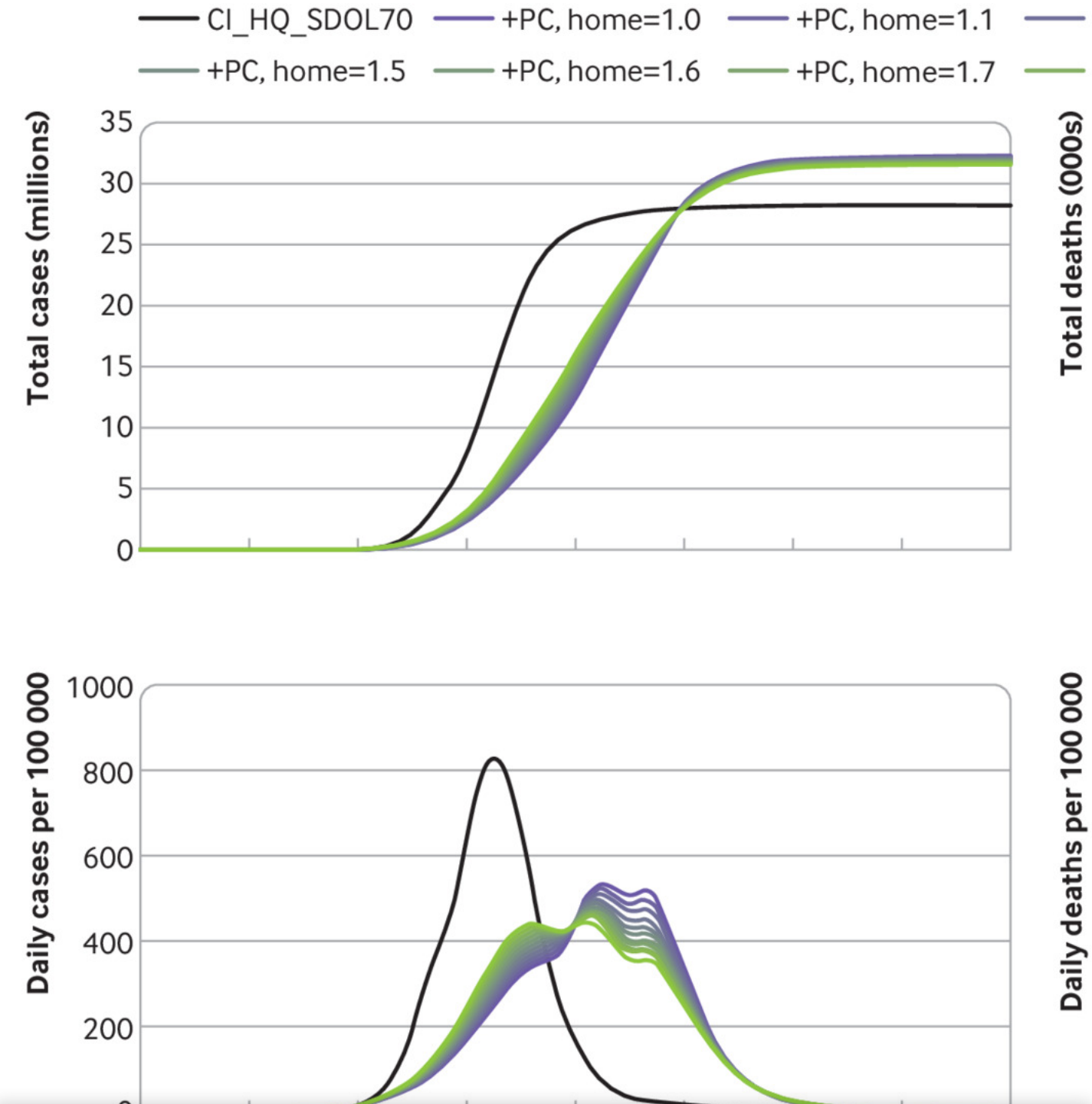
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One criticism of school closure is that reduced contact at school leads to increased contact at home, meaning that children infect high risk adults rather than low risk children. We investigated this by increasing the infection rate at home to an extremely high level. **Figure 3** shows that this makes an insignificant difference compared with the overall effect of adding school closures (despite the description of place closure interventions in table 2 of Report 9, university closures are not included in the scenario parameter file representing place closure, case isolation, household quarantine, and social distancing of over 70s<sup>9</sup>) to the other interventions.



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Although Report 9 discusses the possibility that relaxing the interventions could lead to a second peak later in the year, we wanted to explore this in more detail using the newer CovidSim code and latest set of parameter files.<sup>8</sup>

The interventions we consider are place closures, case isolation, household quarantine, and general social distancing, which are implemented using the PC\_CI\_HQ\_SD parameter file. Specifically, we use the parameter file available in the data/param\_files subdirectory of the GitHub repository. The only modification was to change the duration of the interventions to 91 days.

These interventions start in late March (day 83) and last for three months (91 days). These simulations are also initialised so that about 15 600 deaths occur by day 100 (9 April) in all scenarios, mostly in people infected before the interventions were implemented. Initialisation is done by modifying the “number of deaths accumulated before alert” parameter in the preUK\_2.0.txt parameter file. This compares with how the Report 9 simulations were initialised, which used the reported deaths to 14 March. The results are presented in [figure 4](#). The top panel shows the cumulative number of deaths, using data from National Records of Scotland<sup>11</sup> and Connors and Fordham,<sup>13</sup> whereas the bottom panel shows ICU bed demand per 100 000 people. Although our simulations include Northern Ireland, the available reported data do not. Therefore, the simulation results and data presented in [figure 4](#) are only for England, Wales, and Scotland. We also consider a range of reproduction numbers and find that values higher than those considered in Report 9 best reproduce the data, with a value between 3.0 and 3.5 probably providing the best fit. This is consistent with the analysis presented in Flaxman et al,<sup>14</sup> but we acknowledge that the data could also be fitted by changes to the other scenario parameters. In both panels we also show the “do nothing” scenario for a reproduction number of 3.0.

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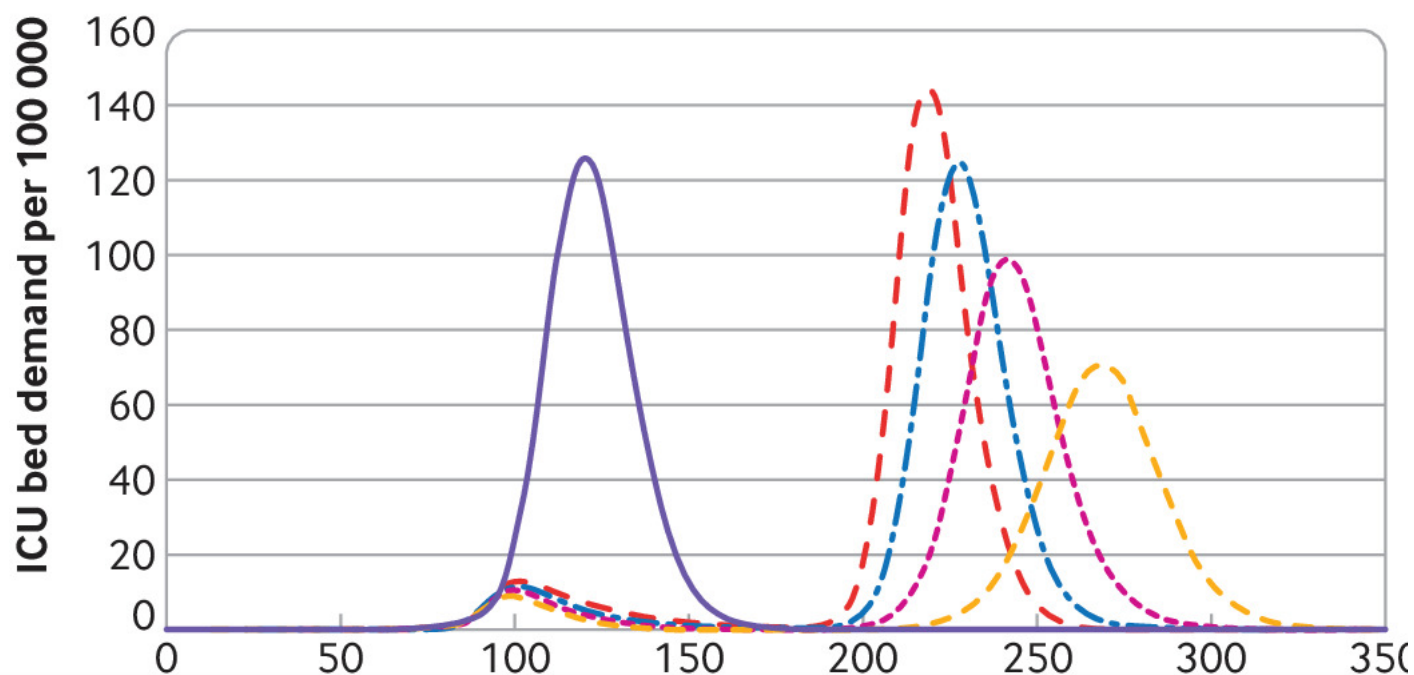
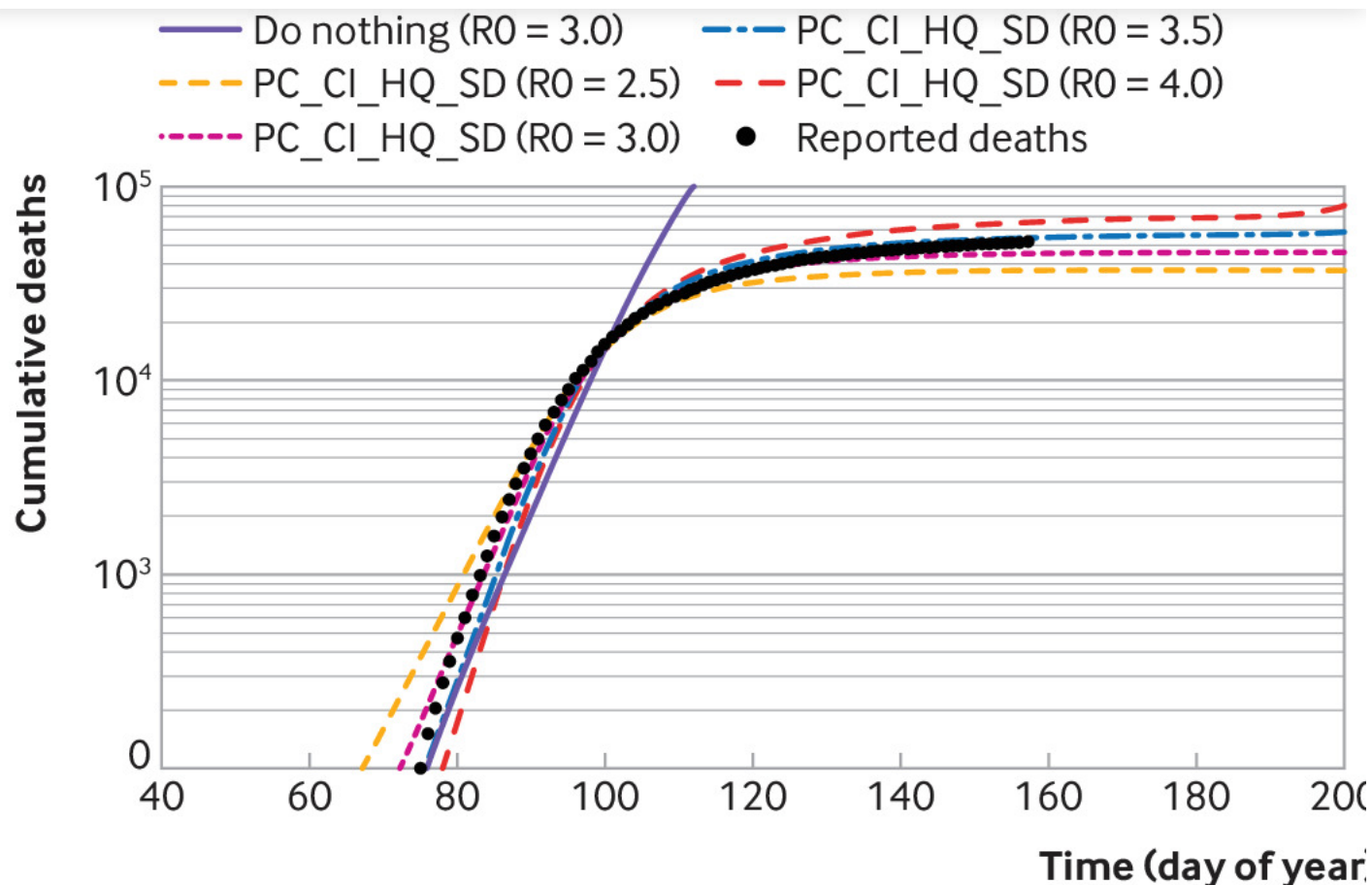
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the over 70s age group. (An editorial published in *The BMJ* on 19 September 2020 suggests this 5% could be an underestimate because IgA antibodies and T cell immunity were overlooked.<sup>16</sup>) With only 5-10% immunity after the lockdown, the epidemiological situation at the outset of the second wave is similar to that of March. Consequently, the number of second wave infections is predicted to be similar to that of the first wave, with a somewhat lower death rate.

In practice, it seems that mandatory and voluntary interventions short of a full lockdown will continue and maintain the reproduction number closer to 1. This will mean slower exponential growth of the second wave and keep the peak demand for ICU beds manageable, although since the epidemic is prolonged, the effect on total deaths is smaller. It is worth noting that a reproduction number of 1 is also the value that prolongs the need for interventions for the longest. At this level, the inhomogeneity of transmissions, particularly the unpredictability of superspreading events, becomes critical. Despite the level of detail in the model, the data are insufficient to model real people: we observed that for a major national epidemic, insufficient data introduce an uncertainty of about five days in the predictions. At a local level, and with a lower reproduction number, this uncertainty in the timing of the epidemic is greatly increased: it is impossible to predict when a particular town will experience an outbreak (specifically, different towns experience outbreaks in different runs of the code).

## Discussion

In this paper we used the recently released CovidSim code<sup>8</sup> to reinvestigate the mitigation scenarios for covid-19 from IBMIC presented in mid-March 2020 in Report 9.<sup>2</sup> The motivation behind this was that some of the results presented in the report suggested that the addition of interventions restricting younger people might actually increase the total number of deaths from covid-19.

We find that the CovidSim code reliably reproduces the results from Report 9 and that the IBMIC can accurately track the data on death rates in the UK. Reproducing the real data does require an adjustment to the parameters and a slightly higher reproduction number than considered in Report 9 and implies an earlier start to the epidemic than suggested by the report. We emphasise that the unavailability of these parameters in early March 2020 is not a failure of the IBMIC model.

We confirm that adding school and university closures to case isolation, household quarantine, and social distancing of over 70s would lead to more deaths compared with the equivalent scenario without the closures of schools and universities. Similarly, general social distancing was also projected to reduce the number of cases but increase the total number of deaths compared with social distancing of over 70s only. We note that in assessing the impact of school closures, UK policy advice has concentrated on reducing total number of cases and not the number of deaths.<sup>17</sup>

The qualitative explanation is that, within all mitigation scenarios in the model, the epidemic ends with widespread immunity, with a large fraction of the population infected. Strategies that minimise deaths involve the infected fraction primarily being in the low risk younger age groups—for example, focusing stricter social distancing measures on care homes where people are likely to die rather than schools where they are not. Optimal death reduction strategies are different from those aimed at reducing the burden on ICUs, and different again from those that lower the overall case rate. It is therefore impossible to optimise a strategy for dealing with covid-19 unless these three desirable outcomes are prioritised.

We find that scenarios that are very effective when the interventions are in place, can then lead to subsequent waves during which most of the infections, and deaths, occur. Our comparison of updated model results with the published death data suggests that a similar second wave will occur later this year if interventions are fully lifted. More realistically, if the case isolation, household quarantine, and social distancing of over 70s strategy is followed, alongside other non-pharmaceutical intervention measures such as non-mandatory social distancing and improved medical outcomes, the second wave will grow more slowly than the first, with more cases but lower mortality.

Since this paper was first written (June 2020), UK policy has moved to more local interventions. CovidSim models the geography of all towns, but only the simulated people are representative of the true population. This uncertainty means that the model cannot reliably predict which town will experience an outbreak. Specifically, whereas the timing of the national outbreak is uncertain by days, the timing of an outbreak in a town is uncertain by months. IBMIC is the most precise model available, but substantially more personal data would be needed to obtain reliable local predictions.

Finally, we re-emphasise that the results in this work are not intended to be detailed predictions for the second wave of covid-19. Rather, we re-examined the evidence available at the start of the epidemic. More accurate information is now available about the compliance with lockdown rules and age dependent mortality. The difficulty in shielding care home residents is a particularly important set of health data that was not available to modellers at the outset.

Nevertheless, in all mitigation scenarios, epidemics modelled using CovidSim eventually finish with widespread infection and immunity, and the final death toll depends primarily on the age distribution of those infected and not the total number.

### **What is already known on this topic**

Detailed models of individual interactions, which can take many hours of supercomputer time to run, are a reliable way to predict the course of an epidemic and investigate counterfactual scenarios

The IBMIC model is the most detailed individual based model of the United Kingdom appropriate for simulation of the spread of an epidemic

The UK-wide lockdown as a result of coronavirus disease 2019 (covid-19) was implemented as a highly effective way of reducing the spread of the epidemic

### **What this study adds**

The model used for Report 9 was independently validated and verified, and predicts that, in the absence of an effective vaccine for covid-19, school closures would result in more overall deaths than no school closures

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- Funding: This paper was supported by a UK Research and Innovation grant ST/V00221X/1 under covid-19 initiative. This work was undertaken, in part, as a contribution to the Rapid Assistance in Modelling the Pandemic (RAMP) initiative, coordinated by the Royal Society. The funders had no role in considering the study design or in the collection, analysis, interpretation of data, writing of the report, or decision to submit the article for publication.
- Competing interests: All authors have completed the ICMJE uniform disclosure form at [www.icmje.org/coi\\_disclosure.pdf](http://www.icmje.org/coi_disclosure.pdf) and declare: support from UK Research and Innovation for the submitted work; no financial relationships with any organisations that might have an interest in the submitted work in the previous three years; no other relationships or activities that could appear to have influenced the submitted work.
- Ethical approval: Not required.
- Data sharing: The full simulation and datasets can be accessed and run from GitHub using the SHA1 hash code 92d414769c6387a08ab65d9830f7f9775fdd3a71. Code examples and raw data sufficient to reproduce all results in this research are available at <https://doi.org/10.7488/ds/2912>.
- The lead author (GJA) affirms that the manuscript is an honest, accurate, and transparent account of the study being reported; that no important aspects of the study have been omitted; and that any discrepancies from the study as planned have been explained.
- Dissemination to participants and related patient and public communities: Since this research uses public demographic data for the whole of the UK, there are no plans for dissemination of this research to specific participants, beyond publishing it.
- Provenance and peer review: Not commissioned; externally peer reviewed.

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