

Research proposal: Mortality exposure in times of Covid-19

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Introduction

The subject of death has, once again, come to the forefront of public debate as the world grapples with a global pandemic of the Covid-19 disease. Much attention has been given to estimating age-specific case fatality rates and mortality rates in general [ref]. The degree to which the pandemic will increase the exposure to mortality in the population as a whole has received little or no attention in the demographic literature. Studies in sociology and public health have consistently shown the negative and long-term consequences of bereavement on mental, physical, and emotional health, especially for women [ref]. Yet, no study to date has considered the degree to which covid-19 excess mortality will increase exposure to death for people around the world.

Given the age-gradient of the Covid-19 case fatality rates [ref], we expect that the pandemic will result in the death of thousands of elderly relatives, such as parents, grandparents, great-grandparents, aunts and uncles. More worryingly, it might result in the death of adult children for even older parents who depend on the crucial support given by their offspring or grandchildren. For many in the Global North this will be their first close encounter with death, as historically low mortality rates have implied a reduction in the exposure to mortality at all ages in high-income countries (Murphy 2011, @alburez_gutierrez_womens_2019).

Project description

This study will combine existing and future data on Covid-19 excess mortality [ref] with powerful methods from formal demography (Goodman, Keyfitz, and Pullum 1974) to estimate the potential burden of covid-19 bereavement. As such, we are not interested in the total number of people who will die from Covid-19, but in the number of people who will be affected by the death of a relative because of the disease. In other words, this is a study on ‘excess bereavement’. Given the range of covid-19 mortality rates in different countries, we ask: (1) How many people will lose a relative to Covid-19, either in the nuclear or in the extended family? (2) What is the age distribution of the bereaved population?

The project comprises a methodological and an empirical component. In terms of methodology, we seek to generalize the Kin-Cohort Method, which is itself an extension of the Goodman, Keyfitz Pullum kinship equations (Goodman, Keyfitz, and Pullum 1974), to estimate the potential increase in the exposure to the death of relatives. This means deriving the mathematical expressions needed to model the effect of changes in mortality on the population-level prevalence of excess bereavement.

We exemplify this for the case of parental bereavement, which we understand as the probability of losing a child. Let CS_a be the expected number of surviving children to a mother alive at aged a in a female stable population with fertility rates m_x , mortality hazard μ_x and survival function $l_x = e^{-\int_0^x \mu_t dt}$ (with unit radix $l_0 = 1$), as proposed by Goodman, Keyfitz, and Pullum (1974):

$$CS_a = \int_0^a m_x l_{a-x} dx.$$

Now consider an absolute increase in mortality δ in mortality in the range $[0, a - \alpha]$, where α is the start age of fertility risk. We can think of this hypothetical increase in mortality as the excess mortality derived from

the Covid-19 disease so that $m_{x,\delta} = m_x + \delta$ (Wrycza and Baudisch (2012)) and $l_{a-x}^\delta = e^{-\int_0^{a-x} (\mu_t + \delta) dt}$.

$$CS_a^\delta = \int_a^\infty m_x l_{a-x}^\delta e^{-\delta(a-x)} dx.$$

As we have shown elsewhere [ref], we can conceptualize the effects of this absolute change δ in mortality as the difference between maternal age a and the mean age of the mother at the birth of her surviving daughters k_a :

$$\frac{\Delta CS_a}{CS_a} \approx -(a - k_a) \Delta \delta.$$

This is a first step towards developing a formal understanding the consequences of an increase in age-specific mortality on child survival but further development is needed, especially for the case in which mortality does not increase by the same factor at all ages. This is, of course, the case of Covid-19, which is known to affect people in older ages [ref]. Further work is also needed to generalize these relationships to other types of kin, such as mothers, grandmothers, cousins and aunts. A final strand of methodological component of the project will seek to develop the introduce heterogeneity in the Kin-Cohort models. In their current state, these models ignore the role of heterogeneity in the experience of kin death, a particular limitation when considering that deaths from the infectious disease are likely to be clustered in the population.

The empirical componente of the project will apply the mathematical models to month-specific mortality and fertility rates (interpolated, if needed, from the available data taking the seasonality of mortality and fertility into account) to estimate the excess burden of kin death derived from Covid-19. Combining this with data on the age and sex structure of the population will allow us to estimate excess bereavement, which we will compare to a baseline counter-factual scenario using rates that exclude the excess mortality from Covid-19. The study will produce a range of estimates, given the uncertainty inherent to the Covid-19 mortality statistics that are required as input for our mathematical models.

As an initial step, the analysis will focus on Germany and Italy, for which quality data on covid-19 death rates [ref] is already available at the sub-national level and where more quality data is expected in the future [ref]. Our methodology is country-agnostic, meaning that it can be applied to other settings once quality data on Covid-19 excess mortality is becomes available.

Expected main results

- A set of academic publications describing the methodology and the main results of the study
- A flexible and scalable methodology for estimating the excess burden of covid-19 bereavement. The code to produce the estimates will be made freely available on the code-sharing platform Github.
- The first set of estimates of how the Covid-19 disease will affect the exposure to mortality at a population-level
- A online platform for scientists and policy makers to access and download the data easily (see this interactive app)

Work plan

References

- Alburez-Gutierrez, Diego, Martin Kolk, and Emilio Zagheni. 2019. “Women’s Experience of Child Death over the Life Course: A Global Demographic Perspective.” Preprint. SocArXiv. <https://doi.org/10.31235/osf.io/s69fz>.
- Goodman, Leo A, Nathan Keyfitz, and Thomas W Pullum. 1974. “Family Formation and the Frequency of Various Kinship Relationships.” *Theoretical Population Biology* 5 (1): 1–27.

Murphy, Michael. 2011. “Long-Term Effects of the Demographic Transition on Family and Kinship Networks in Britain.” *Population and Development Review* 37 (January): 55–80. <https://doi.org/10.1111/j.1728-4457.2011.00378.x>.

Wrycza, Tomasz, and Annette Baudisch. 2012. “How Life Expectancy Varies with Perturbations in Age-Specific Mortality.” *Demographic Research* 27: 365–76.