Child survival for mothers: mortality change and related measures

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Abstract

Background

The expected number of descendants for mothers is an important relation in kinship theory. Child survival (and child death) affect mothers in a variety of ways, yet it is not clear how changes in mortality schedules, given a fertility pattern, affect the number of surviving children from the standpoint of mothers.

Objective

Existing formal demographic tools are used within a stable population framework to characterize child survival from the point of view of a mother, considering both absolute and relative changes in mortality patterns.

Results

The expected change in child survival given an absolute change in mortality depends on the difference between the age of a mother and the age structure of her surviving children. The effects of relative changes in mortality are explained using a fertility-weighted measure of entropy.

Contribution

The paper proposes new expressions to characterize the effects of changes in mortality on child survival. The different measures it introduces can be applied to populations at different stages of the demographic transition.

1. Relationship

 CS_a is the expected number of surviving children to a woman aged a in a stable female population with fertility rates m_x and force of mortality μ_x , as proposed by Goodman, Keyfitz, and Pullum (1974):

$$CS_a = \int_{\alpha}^{a} m_x l_{a-x} dx. \tag{1}$$

We now consider the consequences of a change δ in mortality in the range $[0, a - \alpha]$, where α is the start age of fertility risk.

A discrete approximate of the effect on CS_a given an absolute change in mortality $m_{x,\delta} = m_x + \delta$ is (Wrycza and Baudisch (2012)):

$$\frac{\Delta CS_a}{CS_a} \approx -(a - \overline{x}_{(CS)})\Delta\delta. \tag{2}$$

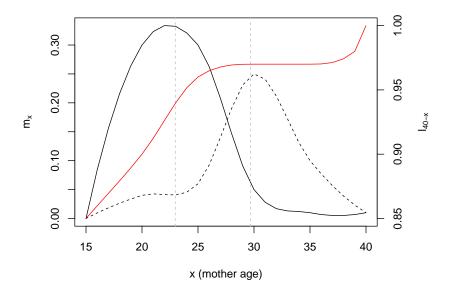


Figure 1: Two different fertility schedules (black lines) and a big hump mortality case (right axis, line in red). In dashed grey lines are the mean age at birth for each pattern.

In this equation, the expected change in child survival is inversely proportional to the difference between maternal age a and the mean age of the mother at the birth of her surviving daughters $\overline{x}_{(CS)}$. In other words, the magnitude of the change depends negatively on the age distribution of the surviving offspring (as younger children experience longer periods of exposure to risk).

The relationship is less clear for proportional changes in mortality $m_{x,\delta} = m_x(1+\delta)$ given the interaction of birth and mortality rates (Keyfitz and Caswell (2005)). Neverthless we can say that, in discrete terms:

$$\frac{\Delta CS_a}{CS_a} \approx -H_{CS_a} \Delta \delta. \tag{3}$$

Where H_{CS_a} can be interpreted as a temporary entropy constant until age $a-\alpha$. Depending the age structure of the survival daughters $C_{CS_a}(x) = \frac{m_x l_{a-x}}{CS_a}$, is the way that is taken each age contribution to the disparity measure $H_{CS_a} = \int_{\alpha}^{a} C_{CS_a}(x) H(x)$ (Aburto et al. (2019)).

The effect of a relative change δ will be bigger when the mortality is more disperse in ages where the fertility is more concentrated. For example in 1, with a big hump in first adult ages, the youngest schedule would have a bigger change in than the older one in C_{CS_a} .

2. Proofs

2.1. Absolute change

Considering that $m_{x,\delta} = m_x + \delta$ (Wrycza and Baudisch (2012)), and $l_{a-x} = e^{-\int_0^{a-x} (\mu_t + \delta)}$:

$$CS_{(a)}^{\delta} = \int_{\alpha}^{a} m_x l_{a-x} e^{-\delta(a-x)} dx. \tag{4}$$

To find the effects on daughter survival of adding δ to the age-specific death rates of daughters, we get the derivative of $dCS_{(a)}^{\delta}/d\delta$, evaluated near zero (Keyfitz and Caswell (2005), section 4.3):

$$= -a \int_{\alpha}^{a} m_x l_{a-x} dx + \int_{\alpha}^{a} m_x l_{a-x} x dx. \tag{5}$$

Since $CS_a = \int_{\alpha}^{a} m_x l_{a-x} dx$, we can rewrite equation (5) as

$$\frac{dCS^{\delta}}{d\delta} = -aCS_a + \int_{\alpha}^{a} x m_x l_{a-x} dx. \tag{6}$$

Dividing both sides by CS_a and multiplying by $d\delta$, we get

$$\frac{dCS_a}{CS_a} = -(a - \overline{x}_{(CS)})d\delta \tag{7}$$

where $\overline{x}_{(CS)}$ is the mean age of women at the birth of their surviving daughters in a stationary population.

2.2. Relative change

Considering that $m_{x,\delta} = m_x(1+\delta)$ and $l_{a-x,\delta} = e^{-\int_0^{a-x} \mu_t(1+\delta)dt} = l_{a-x}^{(1+\delta)}$:

$$CS_a^{\delta} = \int_{\alpha}^a m_x l_{a-x}^{(1+\delta)} dx. \tag{8}$$

Using the derivative of $dCS_{(a)}^{\delta}/d\delta = log(l_{a-x})l_{a-x}^{(1+\delta)}$ and evaluating δ near zero we get the final expression:

$$\frac{dCS_a}{CS_a} = \frac{\int_{\alpha}^{a} m_x l_{a-x} \log(l_{a-x}) dx}{CS_a} d\delta \tag{9}$$

Calling $C_{CS_a}(x) = \frac{m_x l_a - x}{CS_a}$ the weighting function of the successful fertility pattern (onset distribution of survival daughters), and making an analogy with the classical entropy function H where the weights are just the survival function (Keyfitz and Caswell (2005)), we can define this information function $H_{CS_a} = \int_{\alpha}^{a} C_{CS_a} H(x)$, and get:

$$\frac{dCS_a}{CS_a} = -H_{CS_a}d\delta. \tag{10}$$

3. History

Kin count estimation has a long pedigree in mathematical demography. Lotka (1931) proposed to model the orphanhood in a theorical population, comparing the effect across demographic regimes. Equation 1 was originally proposed by Brass (1953) to estimate child mortality. The notion of estimating the expected number of living daughters in a stable population was generalized by Goodman, Keyfitz, and Pullum (1974) for other kin relations (granddaughters, cousins, etc.). The "counting method" approach, being the net reproduction rate R_0 limited to a, or the (not complete) ratio between generations (multiplying and dividing by the mother generation B), was further popularized by Keyfitz and Caswell (2005). Bongaarts (1987) used a similar approach to estimate descendants in his 'Family Status Model'. More recently, Wrycza and Baudisch (2012) has looked at the formal effect of different kind of changes in mortality by age in a paper published in this series.

4. Related measures

This section proposes two additional measures related to the child survival measure for a mother aged a.

4.1. Time with children lost

Mean Time Spent with Lost (MTSL) is an absolute measure of the expected total time with a death daughter that lives a mother, which can be expressed in terms of a temporary expected lost years measure e^{\dagger} :

$$MTSL_a = \int_{\alpha}^{a} m_x \int_{0}^{a-x} d_t e_{0,a-x-t} dt dx = \int_{\alpha}^{a} m_x e_{o,a-x}^{\dagger} dx.$$
 (11)

Where d_t is the death distribution from birth, and $e_{0,a-x-t}$ is the life expectancy at birth until age a-x-t.

It is reasonable to think that mothers relates these lost years with the expected time with live daughters, getting the *intensity* of her losts ($Intensity\ Time\ Lost\ (ITL)$). This makes possible also to compare between fertility regimes:

$$ITL_a = \frac{\int_{\alpha}^{a} m_x e_{o,a-x}^{\dagger} dx}{\int_{\alpha}^{a} m_x e_{o,a-x} dx}$$
 (12)

4.2. Mean Age at Child Loss

Different populations could have different mean age at lost (MAL_a) for mothers aged a with at least a death child:

$$MAL_a = K_a + \int_{\alpha}^{a} f_{x,a} MAD_{a-x}$$

$$\tag{13}$$

 K_a is the mean age at birth for womens aged a, MAD_{a-x} refers to the mean age at death for those newborns who dies before a-x, TFR_a is the cumulated fertility for a women aged a, and $f_{x,a}$ is the cumulative age distribution of fertility until age a.

Equation 13 can be derived by starting with the mother age x+t at each death child age at death t, weighted by the fertility and survival function:

$$MAL_{a} = \frac{\int_{\alpha}^{a} m_{x} \int_{0}^{a-x} \frac{d_{t}(x+t)dt}{\int_{0}^{a-x} d_{t}dt}}{\int_{\alpha}^{a} m_{x}}$$

$$MAL_{a} = \frac{\int_{\alpha}^{a} m_{x}x}{TFR_{a}} + \frac{\int_{\alpha}^{a} m_{x}MAD_{a-x}}{TFR_{a}}$$

$$MAL_{a} = K_{a} + \frac{\int_{\alpha}^{a} m_{x}MAD_{a-x}}{TFR_{a}}$$

$$(14)$$

Arriving to the equation 13.

5. Applications

• The proposed measures are applied to two different demographic regimes.

[PENDING]

• A computation approach is used to evaluate the relative contribution of each component given a relative change in δ .

[PENDING]

5. Bibliography

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