

# DECOMPOSITION TECHNIQUES IN POPULATION HEALTH RESEARCH

José Manuel Aburto & Serena Vigizzi



EUROPEAN DOCTORAL  
SCHOOL OF  
DEMOGRAPHY



# Recap

To avoid the interaction term, Kitagawa suggests

$$\Delta CDR = \underbrace{\sum_x \left( \frac{M_x(t_2) + M_x(t_1)}{2} \right) \left( \frac{N_x(t_2)}{N(t_2)} - \frac{N_x(t_1)}{N(t_1)} \right)}_{\text{Changes in x-composition}} + \underbrace{\sum_x \left( \frac{\frac{N_x(t_2)}{N(t_2)} + \frac{N_x(t_1)}{N(t_1)}}{2} \right) (M_x(t_2) - M_x(t_1))}_{\text{Changes in rates}} \quad (1)$$

# Recap

$$\dot{\bar{v}} = \underbrace{\bar{\dot{v}}}_{\text{Direct component}} + \underbrace{\text{Cov}(v, \dot{w})}_{\text{Structural or compositional component}} \quad (2)$$

Main result in Vaupel & Canudas-Romo (2002)

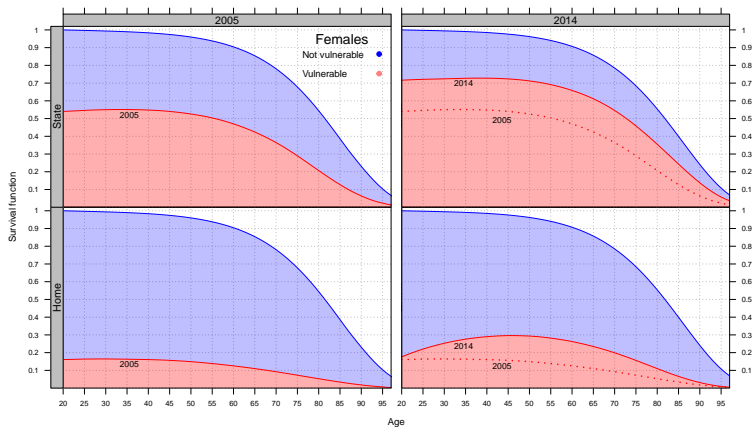
# Recap

Life expectancy at age  $x$  is defined as

$$e(x) = \frac{\int_x^{\infty} \ell(a) da}{\ell(x)}$$

Then we can define disability-free life expectancy as

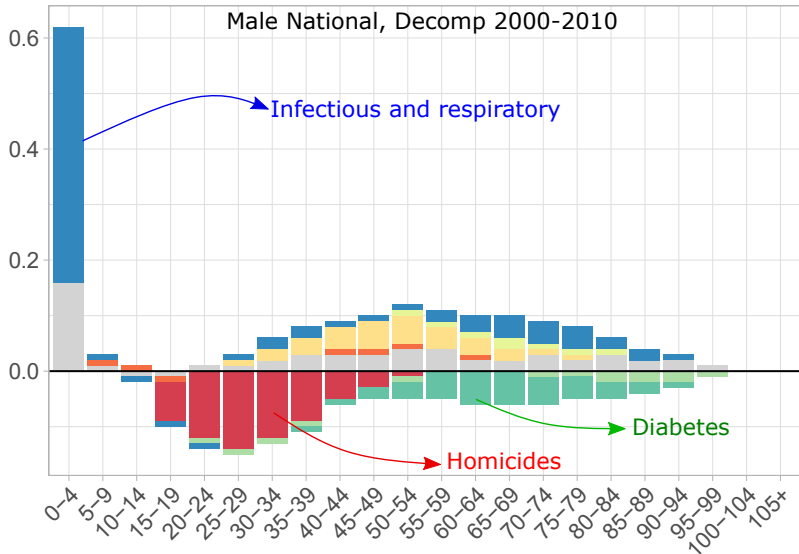
$$e^{DF}(x) = \frac{\int_x^{\infty} [1 - \pi(a)] \ell(a) da}{\ell(x)}$$



# Recap

Arriaga (1984) Effects of mortality change by age groups on life expectancies ( $\sum_n \Delta_x = \text{Total change}$ ):

$${}_n\Delta_x = \underbrace{\frac{\ell_x^1}{\ell_0^1} \left( \frac{{}_nL_x^2}{\ell_x^2} - \frac{{}_nL_x^1}{\ell_x^1} \right)}_{\text{Direct effect}} + \underbrace{\frac{T_{x+n}^2}{\ell_0^1} \left( \frac{\ell_x^1}{\ell_x^2} - \frac{\ell_{x+n}^1}{\ell_{x+n}^2} \right)}_{\text{Indirect and interaction effects}}$$



# Outline

- ▶ Cause-deleted life tables
- ▶ Measures of variation in ages at death
- ▶ Linear integral decomposition



## Cause-deleted life tables (based on Chapter 4, Preston et al 2001)

Often we are interested in knowing what a life table would look if only one cause of death were operating to diminish a cohort.

## Cause-deleted life tables (based on Chapter 4, Preston et al 2001)

Often we are interested in knowing what a life table would look if only one cause of death were operating to diminish a cohort.

The activity of a particular decrement  $i$  will be observed as it works itself out within a multiple decrement process.

## Cause-deleted life tables (based on Chapter 4, Preston et al 2001)

Often we are interested in knowing what a life table would look if only one cause of death were operating to diminish a cohort.

The activity of a particular decrement  $i$  will be observed as it works itself out within a multiple decrement process.

**'What would happen if...'**

- The rate of decrement from  $\mu^i(x)$  if  $i$  were the only decrement ( ${}^*m^i(x)$ ) differs from what it would be if  $i$  were working in the presence of other decrements ( $m^i(x)$ ).

- Assume the force of decrement of cause  $i$  is proportional all-cause force of decrement:

$$\mu^i(a) = R_i \cdot \mu(a) \quad \text{for } x \leq a \leq x + n$$

- Assume the force of decrement of cause  $i$  is proportional all-cause force of decrement:

$$\mu^i(a) = R_i \cdot \mu(a) \quad \text{for } x \leq a \leq x + n \quad (3)$$

- Since, by assumption,

$${}_n p_x^i = e^{-\int_x^{x+n} \mu^i(a) da} = e^{-\int_x^{x+n} R_i \cdot \mu(a) da}$$

- Assume the force of decrement of cause  $i$  is proportional all-cause force of decrement:

$$\mu^i(a) = R_i \cdot \mu(a) \quad \text{for } x \leq a \leq x + n \quad (3)$$

- Since, by assumption,

$${}_n p_x^i = e^{-\int_x^{x+n} \mu^i(a) da} = e^{-\int_x^{x+n} R_i \cdot \mu(a) da}$$

$${}_n p_x^i = e^{-R_i \int_x^{x+n} \mu(a) da} = [e^{-\int_x^{x+n} \mu(a) da}]^{R_i}$$

- Assume the force of decrement of cause  $i$  is proportional all-cause force of decrement:

$$\mu^i(a) = R_i \cdot \mu(a) \quad \text{for } x \leq a \leq x + n \quad (3)$$

- Since, by assumption,

$${}_n^*p_x^i = e^{-\int_x^{x+n} \mu^i(a) da} = e^{-\int_x^{x+n} R_i \cdot \mu(a) da}$$

$${}_n^*p_x^i = e^{-R_i \int_x^{x+n} \mu(a) da} = [e^{-\int_x^{x+n} \mu(a) da}]^{R_i}$$

$${}_n^*p_x^i = [{}_np_x]^{R_i}$$



$${}^*{}_np_x^i = [{}_np_x]^{R_i}$$

$$= [{}_np_x]^{\frac{{}_nD_x^i}{{}_nD_x}}$$

Now with this simple relation we can create  
hypothetical scenarios.

<https://life-expectancy.org>

## Lifespan variation indicators

Indicators		Conventional life table notation
Life disparity	$e^\dagger$	$\sum_{y=0}^{\omega} d_y e(y)$
Gini coefficient	G	$1 - \frac{1}{e_0} \sum_{y=0}^{\omega} \ell_{y+1}^2$
Theil's index	T	$\sum_{y=0}^{\omega} d_y \left[ \frac{\bar{x}_y}{e_0} \ln \frac{\bar{x}_y}{e_0} \right]$
Mean logarithmic deviation	MLD	$\sum_{y=0}^{\omega} d_y (\ln(e_0/\bar{x}_y))$
Variance	V	$\sum_{y=0}^{\omega} d_y (\bar{x}_y - e_0)^2$
Standard deviation	SD	$\sqrt{V}$
Interquartile range	IQR	

**Table 1** Pearson correlation coefficients between pairs of indices, calculated from birth (ages 0–110+) in the top panel and calculated conditional on survival to age 10 (ages 10–110+) in the bottom panel, for all female and male life tables in the Human Mortality Database (7,516 in total)

	$e^{\dagger}$	$G$	$T$	$MLD$	$S$	$V$	$IQR$
$e^{\dagger}$	1.000						
$G$	.978	1.000					
$T$	.947	.991	1.000				
$MLD$	.965	.991	.992	1.000			
$S$	.981	.933	.893	.930	1.000		
$V$	.987	.945	.911	.944	.996	1.000	
$IQR$	.967	.966	.948	.956	.920	.944	1.000

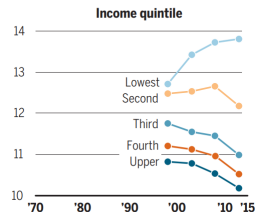
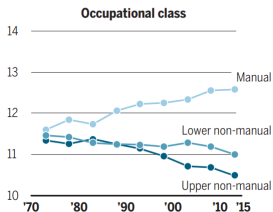
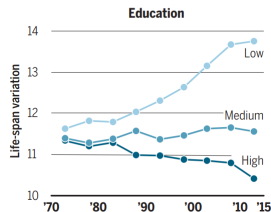
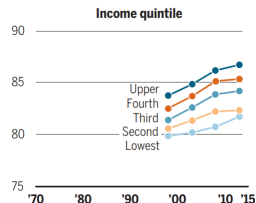
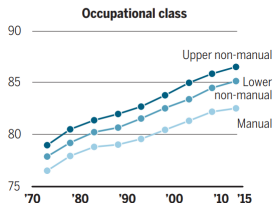
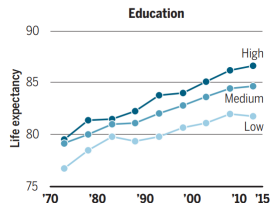
  

	$e_{10}^{\dagger}$	$G_{10}$	$T_{10}$	$MLD_{10}$	$S_{10}$	$V_{10}$	$IQR_{10}$
$e_{10}^{\dagger}$	1.000						
$G_{10}$	.986	1.000					
$T_{10}$	.978	.995	1.000				
$MLD_{10}$	.979	.990	.995	1.000			
$S_{10}$	.986	.962	.961	.973	1.000		
$V_{10}$	.984	.964	.971	.980	.998	1.000	
$IQR_{10}$	.981	.978	.977	.976	.958	.966	1.000

Source: van Raalte & Caswell (2013)

## Trends in life expectancy and life-span variation for Finnish females, 1971–1975 to 2011–2014

Life expectancy is the average age at death, and life-span variation is the standard deviation, conditional upon survival to age 30, with age-specific death rates frozen at those observed in the given year. See supplementary materials for data and methods, including trends for males (which are qualitatively similar), and robustness checks using alternative measures of life-span variation.



How can we decompose by age  
and cause of death these (any)  
function?

## Linear integral decomposition (Horiuchi et al 2008)

- Relies on the assumption that values of the covariates change **continuously**, or gradually, along an actual or hypothetical dimension

## Linear integral decomposition (Horiuchi et al 2008)

- ▶ Relies on the assumption that values of the covariates change **continuously**, or gradually, along an actual or hypothetical dimension
- ▶ Applied when difference in a dependent variable is expressed as a sum of the effects of differences in its covariates. (TFR, mean completed parity)

- Let  $y$  be a demographic function (e.g.  $e^{\dagger}$ ), which is differentiable, of  $n$  covariates (e.g. rates) denoted by  $\mathbf{x} = [x_1, x_2, \dots, x_n]$ .



- ▶ Let  $y$  be a demographic function (e.g.  $e^{\dagger}$ ), which is differentiable, of  $n$  covariates (e.g. rates) denoted by  $\mathbf{x} = [x_1, x_2, \dots, x_n]$ .
- ▶ Assume they both depend on an underlying dimension  $t$  (e.g. time).

- ▶ Let  $y$  be a demographic function (e.g.  $e^{\dagger}$ ), which is differentiable, of  $n$  covariates (e.g. rates) denoted by  $\mathbf{x} = [x_1, x_2, \dots, x_n]$ .
- ▶ Assume they both depend on an underlying dimension  $t$  (e.g. time).
- ▶ We have observations at two time points and  $\mathbf{x}$  is a differentiable vector function of  $t$  between  $t_1$  and  $t_2$ .

- ▶ Let  $y$  be a demographic function (e.g.  $e^{\dagger}$ ), which is differentiable, of  $n$  covariates (e.g. rates) denoted by  $\mathbf{x} = [x_1, x_2, \dots, x_n]$ .
- ▶ Assume they both depend on an underlying dimension  $t$  (e.g. time).
- ▶ We have observations at two time points and  $\mathbf{x}$  is a differentiable vector function of  $t$  between  $t_1$  and  $t_2$ .

$$y(t) = f(\mathbf{x}(t)) = f(x_1(t), x_2(t), \dots, x_n(t)) \quad (4)$$

By the fundamental theorem of calculus

$$y(t_2) - y(t_1) = \int_{t_1}^{t_2} \frac{d}{dt} y(t) dt$$

By the fundamental theorem of calculus

$$y(t_2) - y(t_1) = \int_{t_1}^{t_2} \frac{d}{dt} y(t) dt \quad (5)$$

Applying the chain rule for partial derivatives of a composite function

$$y(t_2) - y(t_1) = \int_{t_1}^{t_2} \left[ \sum_{i=1}^n \frac{\partial}{\partial x_i(t)} y(t) \cdot \frac{d}{dt} x_i(t) \right] dt \quad (6)$$

Exchange of integration and summation, and  
applying the substitution rule for definite integrals

$$y(t_2) - y(t_1) = \sum_{i=1}^n \int_{x_i(t_1)}^{x_i(t_2)} \frac{\partial}{\partial x_i(t)} y(t) dx_i(t)$$

Exchange of integration and summation, and  
applying the substitution rule for definite integrals

$$y(t_2) - y(t_1) = \sum_{i=1}^n \int_{x_i(t_1)}^{x_i(t_2)} \frac{\partial}{\partial x_i(t)} y(t) dx_i(t) \quad (7)$$

(Simplifying notation)

$$y_2 - y_1 = \sum_{i=1}^n \int_{x_{i1}}^{x_{i2}} \frac{\partial y}{\partial x_{i1}} dx_i = \sum_{i=1}^n c_i \quad (8)$$

$$y_2 - y_1 = \sum_{i=1}^n c_i \quad (9)$$

$c_i$  is the total change in  $y$  produced by changes in the  $i$ -th covariate,  $x_i$ .

Important: Theoretical foundation for decomp analysis: implies that even if a dependent variable is not an additive function of its covariates, a change in the dependent variable can be expressed as a sum of effects of the covariates.

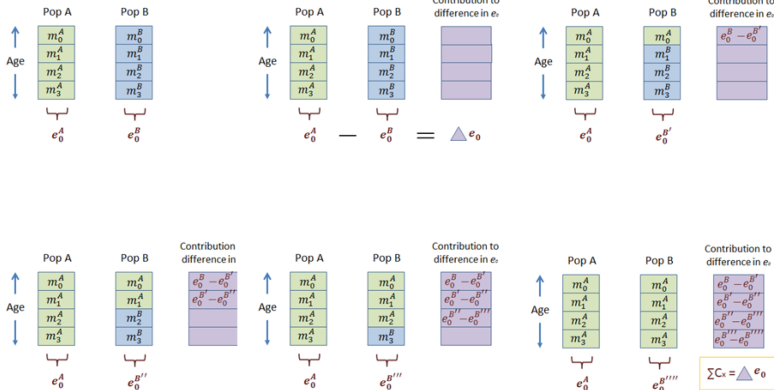


## Stepwise decomposition (Andreev et al 2002)

- ▶ To decompose a function for which there is no analytic solution (or for which the analytic solution is complicated).
- ▶ Step-wise decomposition alters the rates one element at a time (the element usually age), and recalculates the index function (i.e. life expectancy or any other summary measure).

# Stepwise decomposition

## Step-wise decomposition method

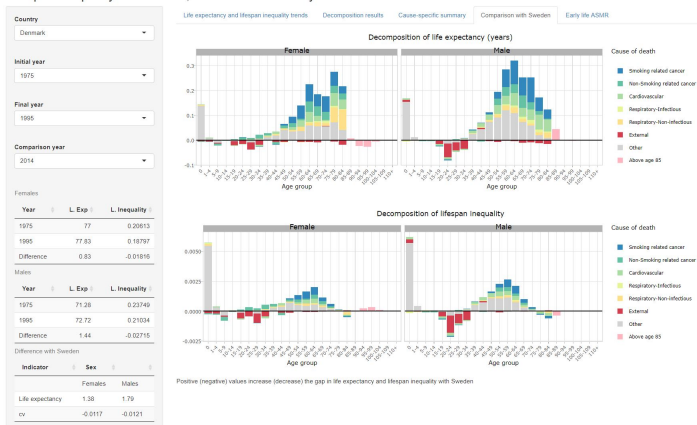


\*NB: Usually replacements are also done in reverse age contributions averaged.

## Exercise 1: Mexico's example

# Extending to causes of death

Lifespan inequality in Denmark, Sweden and Norway



[https://jmaburto.shinyapps.io/DK\\_App/](https://jmaburto.shinyapps.io/DK_App/)

## Exercise 2: by cause of death

Challenge 1 Use the linear integral model to decompose the change in the standard deviation of the age-at-death distribution and life expectancy by age and cause of death for 3 countries you might be interested in (over time or between them). Interpret the results of life expectancy alongside standard deviation. Make it interesting. You can use data from HCoD, HMD, WHO, GBD.