DECOMPOSITION TECHNIQUES IN POPULATION HEALTH RESEARCH

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To avoid the interaction term, Kitagawa suggests

$$\Delta CDR = \underbrace{\sum_{x} \left(\frac{M_{x}(t_{2}) + M_{x}(t_{1})}{2} \right) \left(\frac{N_{x}(t_{2})}{N(t_{2})} - \frac{N_{x}(t_{1})}{N(t_{1})} \right)}_{\text{Changes in x-composition}} + \underbrace{\sum_{x} \left(\frac{N_{x}(t_{2})}{N(t_{2})} + \frac{N_{x}(t_{1})}{N(t_{1})} \right) \left(M_{x}(t_{2}) - M_{x}(t_{1}) \right)}_{\text{Changes in rates}}$$
(1)

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

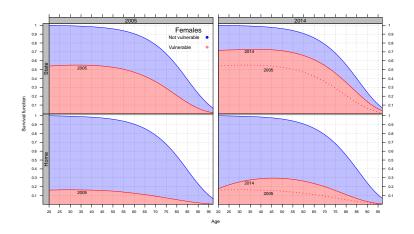
Main result in Vaupel & Canudas-Romo (2002)

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

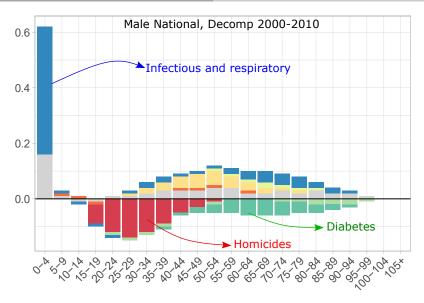
$$e^{DF}(x) = \frac{\int_x^{\infty} [1 - \pi(a)] \ell(a) da}{\ell(x)}$$



Arriaga (1984) Effects of mortality change by age groups on life expectancies $(\sum_{n} \Delta_{x} = \text{Total change})$:

$${}_{n}\Delta_{x} = \underbrace{\frac{\ell_{x}^{1}}{\ell_{0}^{1}} \left(\frac{{}_{n}L_{x}^{2}}{\ell_{x}^{2}} - \frac{{}_{n}L_{x}^{1}}{\ell_{x}^{1}} \right)}_{\text{Direct effect}} + \underbrace{\frac{T_{x+n}^{2}}{\ell_{0}^{1}} \left(\frac{\ell_{x}^{1}}{\ell_{x}^{2}} - \frac{\ell_{x+n}^{1}}{\ell_{x+n}^{2}} \right)}_{\text{Indirect and interaction effect}}$$

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)

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'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 $(m^i(x))$.

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

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► Since, by assupmtion,

$$_{n}^{*}p_{x}^{i}=e^{-\int_{x}^{x+n}\mu^{i}(a)da}=e^{-\int_{x}^{x+n}R_{i}\cdot\mu(a)da}$$

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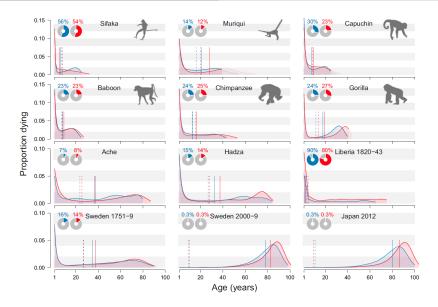
$$_{n}^{\ast}p_{x}^{i}=[_{n}p_{x}]^{R_{i}}$$

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$$= [{}_{n}p_{x}]^{\frac{nD_{x}^{\prime}}{nD_{x}}}$$

Now with this simple relation we can create hypothetical scenarios.

https://population-health.shinyapps.io/saudi-arabia-health-profile/



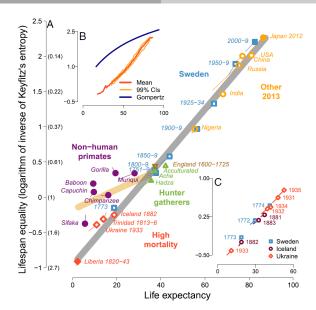
Lifespan variation indicators

Indicators	•	Conventional life table notation
Life disparity	e^{\dagger}	$\sum_{y=0}^{\omega} d_y e(y)$
Gini coefficient	G	$1-rac{1}{e_0}\sum_{y=0}^{\omega}\ell_{y+1}^2$
Theil's index	Т	$\sum_{y=0}^{\omega} d_y \left[rac{ar{x}_y}{e_0} \ln rac{ar{x}_y}{e_0} ight]$
Mean logarithmic deviation	MLD	$\sum_{y=0}^{\omega} d_y (\ln(e_0/ar{x}_y)$
Variance	V	$\sum_{y=0}^{\omega} d_y (\bar{x}_y - e_0)^2$
Standard deviation	SD	\sqrt{V}
Interquartile range	IQR	$\hat{x}_3 - \hat{x}_1$
European Doctoral School of Demography		Day 1 14

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)

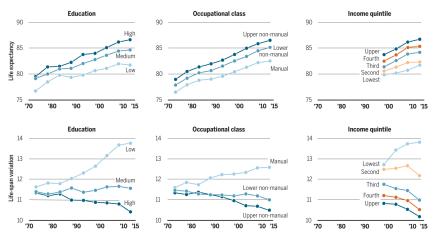
	e^{\dagger}	G	T	MLD	S	V	IQR
e^{\dagger}	1.000						
G	.978	1.000					
T	.947	.991	1.000				
MLD	.965	.991	.992	1.000			
S	.981	.933	.893	.930	1.000		
V	.987	.945	.911	.944	.996	1.000	
IQR	.967	.966	.948	.956	.920	.944	1.000
	e_{10}^{\dagger}	G_{10}	T_{10}	MLD_{10}	S_{10}	V_{10}	IQR_{10}
e_{10}^{\dagger}	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

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- ▶ 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 $\mathbf{x} = [x_1, x_2, \dots, x_n]$.

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- ▶ We have observations at two time points and x is a differentiable vector function of t between t_1 and t_2 .

$$y(t) = f(x(t)) = f(x_1(t), x_2(t), \dots, x_n(t))$$
 (4)

By the fundamental theorem of calculus

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

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 (5)

Applying the chain rule for partial derivatives of a composite function

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

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

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

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 (7)

(Simplifying notation)

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

$$y_2 - y_1 = \sum_{i=1}^n c_i \tag{9}$$

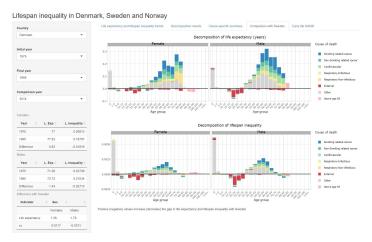
 c_i is the total change in y produced by changes in the i-th covariate, x_i .

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.

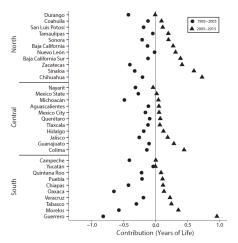
Exercise 1: by age



Extending to causes of death



https://jmaburto.shinyapps.io/DK_App/



Note This figure shows how homiddes contributed to changes in life span inequality (i.e., panel b of Figure 1) in 2 periods; positive values suggest increases in years of life lost because of homiddes and regative values correspond to reductions in life years (so because of death.) This figure shows each of the 32 Mexican states grouped in broad regions; north, central, and south. Within each region, states are ordered according to the maintude of the inmact of homiddes to life span inequality at a set is vers in the period 2005 to 2015.

FIGURE 2—Contribution of Homicide Mortality to Changes in Male Life Span Inequality by State and Period: Mexico, 1995–2005 and 2005–2015

Exercise 2: by cause of death

Outline
Cause-deleted life tables
Lifespan variation measures
Linear integral decomposition

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.