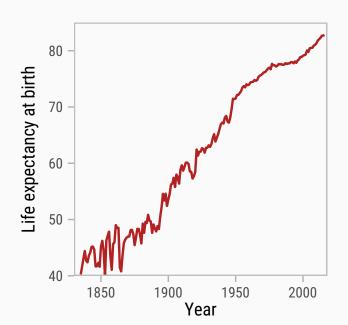
POPULATION DYNAMICS WITH EMPHASIS ON MORTALITY RESEARCH

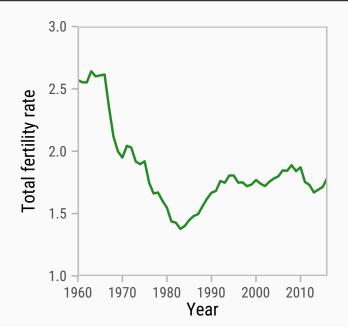
Jesús-Adrián Álvarez

alvarez@sdu.dk

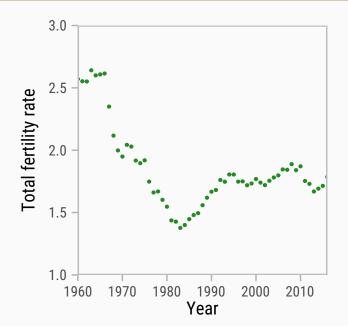
Interdisciplinary Centre on Population Dynamics University of Southern Denmark



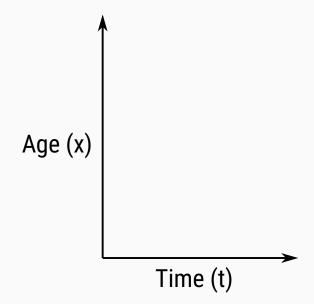
TOTAL FERTILITY RATE, DANISH FEMALES. 1960-2018

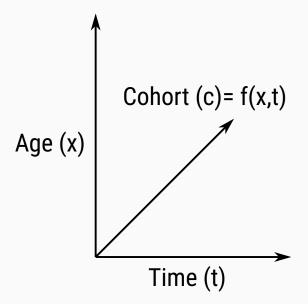


TOTAL FERTILITY RATE, DANISH FEMALES. 1960-2018



Changes over time and age





Mortality

Mortality

· Mortality hazard,

Mortality

- · Mortality hazard,
- · Life expectancy,

Mortality

- · Mortality hazard,
- · Life expectancy,
- · Life table entropy.

Mortality

- · Mortality hazard,
- · Life expectancy,
- · Life table entropy.

Fertility

Mortality

- · Mortality hazard,
- · Life expectancy,
- · Life table entropy.

Fertility

· Fertility rates,

Mortality

- · Mortality hazard,
- · Life expectancy,
- · Life table entropy.

Fertility

- · Fertility rates,
- · Mean age at birth.

Mortality

- · Mortality hazard,
- · Life expectancy,
- Life table entropy.

Fertility

- · Fertility rates,
- · Mean age at birth.

Family demography

• Expected years ever married (Mogi and Canudas-Romo, 2018)

How can I calculate changes over time in...

remaining life expectancy at the 80th percentile?

- · remaining life expectancy at the 80th percentile?
- the age at which remaining life expectancy is 14.5 years (i.e. e(x) = 14.5)

- · remaining life expectancy at the 80th percentile?
- the age at which remaining life expectancy is 14.5 years (i.e. e(x) = 14.5)
- the probability of dying between percentiles 70th and 90th of the death distribution?

- · remaining life expectancy at the 80th percentile?
- the age at which remaining life expectancy is 14.5 years (i.e. e(x) = 14.5)
- the probability of dying between percentiles 70th and 90th of the death distribution?
- the difference in standard deviation between females and males at age 100?

- · remaining life expectancy at the 80th percentile?
- the age at which remaining life expectancy is 14.5 years (i.e. e(x) = 14.5)
- the probability of dying between percentiles 70th and 90th of the death distribution?
- the difference in standard deviation between females and males at age 100?
- · the modal age at death?

The struggle

We develop neat and elegant equations in continuous mathematics but....

We develop neat and **elegant equations** in continuous mathematics but.... when we deal with **real data**, everything gets **messy**.

We develop neat and **elegant equations** in continuous mathematics but.... when we deal with **real data**, everything gets **messy**.

In particular, when data is scarce (e.g. populations are small)

We develop neat and **elegant equations** in continuous mathematics but.... when we deal with **real data**, everything gets **messy**.

In particular, when data is scarce (e.g. populations are small)

Advantages of working in continuous time:

We develop neat and **elegant equations** in continuous mathematics but.... when we deal with **real data**, everything gets **messy**.

In particular, when data is scarce (e.g. populations are small)

Advantages of working in continuous time:

· We can use calculus,

We develop neat and **elegant equations** in continuous mathematics but.... when we deal with **real data**, everything gets **messy**.

In particular, when data is scarce (e.g. populations are small)

Advantages of working in continuous time:

- · We can use calculus,
- More intuitive when we think about the dynamics and processes of populations.

We develop neat and **elegant equations** in continuous mathematics but.... when we deal with **real data**, everything gets **messy**.

In particular, when data is scarce (e.g. populations are small)

Advantages of working in continuous time:

- · We can use calculus,
- More intuitive when we think about the dynamics and processes of populations.

Limitations:

Bad approximations from discrete to continuous measures

Everything we do (or most of it) is based on life tables.

Life tables are powerful but imply many assumptions behind:

Everything we do (or most of it) is based on life tables.

Life tables are powerful but imply many assumptions behind:

• How deaths are distributed between age groups (i.e. ax = 0.5)

Everything we do (or most of it) is based on life tables.

Life tables are powerful but imply many assumptions behind:

- How deaths are distributed between age groups (i.e. ax = 0.5)
- The open age interval (i.e. age 110+)

Everything we do (or most of it) is based on life tables.

Life tables are powerful but imply many assumptions behind:

- How deaths are distributed between age groups (i.e. ax = 0.5)
- The open age interval (i.e. age 110+)

Continuous representation

$$e(0) = \int_0^\infty \ell(x) dx$$

Everything we do (or most of it) is based on life tables.

Life tables are powerful but imply many assumptions behind:

- How deaths are distributed between age groups (i.e. ax = 0.5)
- The open age interval (i.e. age 110+)

Continuous representation

$$e(0) = \int_0^\infty \ell(x) dx$$

Discrete approach

$$e(0) = L_0 + L_1 + L_2 + ...L_{100+}$$

How to calculate changes over time in demographic measures?

How to calculate changes over time in demographic measures?

How to apply neat continuous formulas to messy discrete data?

Data

1. Individual data

1. Individual data

 Date of birth, date of death, date of migration, individual characteristics, etc.

1. Individual data

- Date of birth, date of death, date of migration, individual characteristics, etc.
- Sources: Statistics Denmark, Statistics Netherlands, International Database on Longevity, Surveys, etc.

1. Individual data

- Date of birth, date of death, date of migration, individual characteristics, etc.
- Sources: Statistics Denmark, Statistics Netherlands, International Database on Longevity, Surveys, etc.

2. Aggregated data

1. Individual data

- Date of birth, date of death, date of migration, individual characteristics, etc.
- Sources: Statistics Denmark, Statistics Netherlands, International Database on Longevity, Surveys, etc.

2. Aggregated data

· Death counts and exposures by age-group and year,

1. Individual data

- Date of birth, date of death, date of migration, individual characteristics, etc.
- Sources: Statistics Denmark, Statistics Netherlands, International Database on Longevity, Surveys, etc.

2. Aggregated data

- · Death counts and exposures by age-group and year,
- · Life tables,

1. Individual data

- Date of birth, date of death, date of migration, individual characteristics, etc.
- Sources: Statistics Denmark, Statistics Netherlands, International Database on Longevity, Surveys, etc.

2. Aggregated data

- · Death counts and exposures by age-group and year,
- · Life tables,
- · Sources: HMD, UN, WHO, LAMBdA (Latin America), etc.

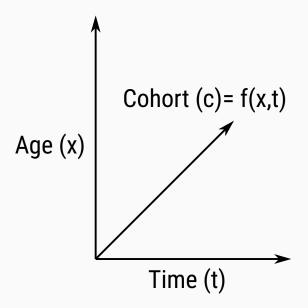
change?

How do we measure continuous

How do we measure continuous change?

$$\frac{df(x)}{dx} = \lim_{h \to 0} \frac{f(x+h) - f(x)}{h}$$

In what direction should we derive?



IN WHAT DIRECTION SHOULD WE DERIVE?

Rate of ageing (in the age direction)

$$b(x,t) = \frac{\frac{\partial \mu(x,t)}{\partial x}}{\frac{\partial \mu(x,t)}{\mu(x,t)}}$$

IN WHAT DIRECTION SHOULD WE DERIVE?

Rate of ageing (in the age direction)

$$b(x,t) = \frac{\frac{\partial \mu(x,t)}{\partial x}}{\frac{\partial \mu(x,t)}{\partial x}}$$

Rate of mortality improvement (in the time direction)

$$\rho(x,t) = -\frac{\frac{\partial \mu(x,t)}{\partial t}}{\mu(x,t)}$$

IN WHAT DIRECTION SHOULD WE DERIVE?

Rate of ageing (in the age direction)

$$b(x,t) = \frac{\frac{\partial \mu(x,t)}{\partial x}}{\frac{\partial \mu(x,t)}{\partial x}}$$

Rate of mortality improvement (in the time direction)

$$\rho(\mathbf{x},t) = -\frac{\frac{\partial \mu(\mathbf{x},t)}{\partial t}}{\frac{\partial \mu(\mathbf{x},t)}{\partial t}}$$

Rate of ageing (in the cohort direction)

$$\beta(\mathbf{x},t) = \frac{\frac{\partial \mu(\mathbf{x},t)}{\partial t \partial \mathbf{x}}}{\frac{\mu(\mathbf{x},t)}{\mu(\mathbf{x},t)}}$$

$$\beta(\mathbf{x},t) = b(\mathbf{x},t) - \rho(\mathbf{x},t)$$

Approximating derivatives

$$f'(x,t) = \frac{\partial f(x,t)}{\partial t} \cong \frac{f(x,t) + f(x,t+h)}{h}$$

$$f'(x,t) = \frac{\partial f(x,t)}{\partial t} \cong \frac{f(x,t) + f(x,t+h)}{h}$$

and

$$f'(x,t) = \frac{\partial f(x,t)}{\partial t} \cong \frac{f(x,t) + f(x,t+h)}{h}$$

and

$$\frac{\frac{\partial f(x,t)}{\partial t}}{f(x,t)} \cong \frac{\ln\left(\frac{f(x,t+h)}{f(x,t)}\right)}{h}$$

$$f'(x,t) = \frac{\partial f(x,t)}{\partial t} \cong \frac{f(x,t) + f(x,t+h)}{h}$$

and

$$\frac{\frac{\partial f(x,t)}{\partial t}}{f(x,t)} \cong \frac{\ln\left(\frac{f(x,t+h)}{f(x,t)}\right)}{h}$$

The approximation depends on the size of h.

Approximating integrals

$$\int_a^b f(x,t)dx$$

$$\int_a^b f(x,t)dx$$

Trapezoidal rule

$$\int_a^b f(x,t)dx$$

Trapezoidal rule

$$\int_a^b f(x,t)dx \cong (b-a)\frac{f(a)+f(b)}{2}$$

$$\int_a^b f(x,t)dx$$

Trapezoidal rule

$$\int_a^b f(x,t)dx \cong (b-a)\frac{f(a)+f(b)}{2}$$

Simpson's rule

$$\int_a^b f(x,t)dx$$

Trapezoidal rule

$$\int_a^b f(x,t)dx \cong (b-a)\frac{f(a)+f(b)}{2}$$

Simpson's rule

$$\int_a^b f(x,t)dx \cong \frac{b-a}{6}[f(a)+4f(\frac{a+b}{2})+f(b)]$$

$$\int_a^b f(x,t)dx$$

Trapezoidal rule

$$\int_a^b f(x,t)dx \cong (b-a)\frac{f(a)+f(b)}{2}$$

Simpson's rule

$$\int_a^b f(x,t)dx \cong \frac{b-a}{6}[f(a)+4f(\frac{a+b}{2})+f(b)]$$

Simpson's 3/8 rule

$$\int_a^b f(x,t)dx$$

Trapezoidal rule

$$\int_a^b f(x,t)dx \cong (b-a)\frac{f(a)+f(b)}{2}$$

Simpson's rule

$$\int_{a}^{b} f(x,t) dx \cong \frac{b-a}{6} [f(a) + 4f(\frac{a+b}{2}) + f(b)]$$

Simpson's 3/8 rule

$$\int_{a}^{b} f(x,t)dx \cong \frac{b-a}{8} [f(a) + 3f(\frac{2a+b}{3}) + 3f(\frac{a+2b}{3}) + f(b)]$$

Producing continuous estimates

1. Approximations based on life table functions

- 1. Approximations based on life table functions
 - Appendix of every new method in continuous time

Some approaches to produce continuous estimates

- 1. Approximations based on life table functions
 - Appendix of every new method in continuous time
 - Wilmoth and Horuchi (2009),

Some approaches to produce continuous estimates

- 1. Approximations based on life table functions
 - Appendix of every new method in continuous time
 - · Wilmoth and Horuchi (2009),
 - · Beltran-Sanchez and Fernandez (2015),

- 1. Approximations based on life table functions
 - Appendix of every new method in continuous time
 - · Wilmoth and Horuchi (2009),
 - · Beltran-Sanchez and Fernandez (2015),
 - Vaupel and Canudas-Romo (2003).

- 1. Approximations based on life table functions
 - · Appendix of every new method in continuous time
 - · Wilmoth and Horuchi (2009),
 - Beltran-Sanchez and Fernandez (2015),
 - Vaupel and Canudas-Romo (2003).
- 2. Rolling window

- 1. Approximations based on life table functions
 - · Appendix of every new method in continuous time
 - · Wilmoth and Horuchi (2009),
 - Beltran-Sanchez and Fernandez (2015),
 - Vaupel and Canudas-Romo (2003).
- 2. Rolling window
- Simulation of lifespans using exponential distribution with piecewise constant rate (Willikens, 2009)

- 1. Approximations based on life table functions
 - · Appendix of every new method in continuous time
 - · Wilmoth and Horuchi (2009),
 - Beltran-Sanchez and Fernandez (2015),
 - Vaupel and Canudas-Romo (2003).
- 2. Rolling window
- Simulation of lifespans using exponential distribution with piecewise constant rate (Willikens, 2009)
- 4. Smoothing

- 1. Approximations based on life table functions
 - · Appendix of every new method in continuous time
 - · Wilmoth and Horuchi (2009),
 - Beltran-Sanchez and Fernandez (2015),
 - Vaupel and Canudas-Romo (2003).
- 2. Rolling window
- Simulation of lifespans using exponential distribution with piecewise constant rate (Willikens, 2009)
- 4. Smoothing
 - Parametric
 - Gompertz

- 1. Approximations based on life table functions
 - · Appendix of every new method in continuous time
 - · Wilmoth and Horuchi (2009),
 - Beltran-Sanchez and Fernandez (2015),
 - · Vaupel and Canudas-Romo (2003).
- 2. Rolling window
- Simulation of lifespans using exponential distribution with piecewise constant rate (Willikens, 2009)
- 4. Smoothing
 - Parametric
 - · Gompertz
 - · Kannisto (HMD)

- 1. Approximations based on life table functions
 - · Appendix of every new method in continuous time
 - · Wilmoth and Horuchi (2009),
 - Beltran-Sanchez and Fernandez (2015),
 - · Vaupel and Canudas-Romo (2003).
- 2. Rolling window
- Simulation of lifespans using exponential distribution with piecewise constant rate (Willikens, 2009)
- 4. Smoothing
 - · Parametric
 - · Gompertz
 - · Kannisto (HMD)
 - · Gamma-Gompertz

- 1. Approximations based on life table functions
 - · Appendix of every new method in continuous time
 - · Wilmoth and Horuchi (2009),
 - Beltran-Sanchez and Fernandez (2015),
 - · Vaupel and Canudas-Romo (2003).
- 2. Rolling window
- Simulation of lifespans using exponential distribution with piecewise constant rate (Willikens, 2009)
- 4. Smoothing
 - Parametric
 - · Gompertz
 - · Kannisto (HMD)
 - · Gamma-Gompertz
- Non-Parametric
 - Splines, P-Splines (Camarda, 2012)

- 1. Approximations based on life table functions
 - · Appendix of every new method in continuous time
 - · Wilmoth and Horuchi (2009),
 - Beltran-Sanchez and Fernandez (2015),
 - · Vaupel and Canudas-Romo (2003).
- 2. Rolling window
- Simulation of lifespans using exponential distribution with piecewise constant rate (Willikens, 2009)
- 4. Smoothing
 - · Parametric
 - Gompertz
 - · Kannisto (HMD)
 - · Gamma-Gompertz
- 5. Non-Parametric
 - · Splines, P-Splines (Camarda, 2012)
 - · Ungrouping (Rizzi, 2015)

Discrete approximation from life tables

VAUPEL AND CANUDAS-ROMO (2003) APPROXIMATION

$$\rho(x,t+h/2) \cong \frac{\ln\left(\frac{\mu(x,t+h)}{\mu(x,t)}\right)}{h}$$

and

$$\mu(x,t+h/2)\cong\frac{M(x,t+h)+M(x,t)}{2}$$

VAUPEL AND CANUDAS-ROMO (2003) APPROXIMATION

$$\rho(x,t+h/2) \cong \frac{\ln\left(\frac{\mu(x,t+h)}{\mu(x,t)}\right)}{h}$$

and

$$\mu(x,t+h/2)\cong\frac{M(x,t+h)+M(x,t)}{2}$$

Advantages

- Easy to estimate from life tables,
- It can be applied to any function (changes over time and age).

VAUPEL AND CANUDAS-ROMO (2003) APPROXIMATION

$$\rho(x,t+h/2) \cong \frac{\ln\left(\frac{\mu(x,t+h)}{\mu(x,t)}\right)}{h}$$

and

$$\mu(x,t+h/2)\cong\frac{M(x,t+h)+M(x,t)}{2}$$

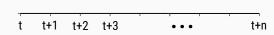
Advantages

- Easy to estimate from life tables,
- It can be applied to any function (changes over time and age).

Disadvantages

Not very precise.







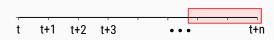














Advantages

- In R, it can be computed easily with the ${\tt zoo}$ package.



Advantages

- In R, it can be computed easily with the ${\tt zoo}$ package.
- · It can be applied to any function.



Advantages

- In R, it can be computed easily with the zoo package.
- It can be applied to any function.

Disadvantages



Advantages

- In R, it can be computed easily with the zoo package.
- It can be applied to any function.

Disadvantages

• The greater the window, the more information we lose.



Advantages

- In R, it can be computed easily with the zoo package.
- · It can be applied to any function.

Disadvantages

- · The greater the window, the more information we lose.
- It requires many years/ages to provide meaningful results.

Simulating lifespans using exponential distribution with

piecewise constant rate

• We start with a vector of death rates M(x, t),

- We start with a vector of death rates M(x, t),
- Simulate a great number of lifespans (>100,000) using M(x, t) as the vector of rates in the function $\mathtt{rpexp}()$ of the package \mathtt{msm} .

$$s(x,t) = e^{-nM(x,t)}$$
 and $\mu(x,t) = \frac{f(x,t)}{s(x,t)}$

- We start with a vector of death rates M(x, t),
- Simulate a great number of lifespans (>100,000) using M(x, t) as the vector of rates in the function $\mathtt{rpexp}()$ of the package \mathtt{msm} .

$$s(x,t) = e^{-nM(x,t)}$$
 and $\mu(x,t) = \frac{f(x,t)}{s(x,t)}$

Advantages

 Can calculate hazard, density, survival function, life expectancy, entropy, quantiles and associated confidence intervals in a continuous setting.

- We start with a vector of death rates M(x, t),
- Simulate a great number of lifespans (>100,000) using M(x, t) as the vector of rates in the function rpexp() of the package msm.

$$s(x,t) = e^{-nM(x,t)}$$
 and $\mu(x,t) = \frac{f(x,t)}{s(x,t)}$

Advantages

- Can calculate hazard, density, survival function, life expectancy, entropy, quantiles and associated confidence intervals in a continuous setting.
- It is equivalent to calculating life tables in a continuous setting.

- We start with a vector of death rates M(x, t),
- Simulate a great number of lifespans (>100,000) using M(x, t) as the vector of rates in the function $\mathtt{rpexp}()$ of the package \mathtt{msm} .

$$s(x,t) = e^{-nM(x,t)}$$
 and $\mu(x,t) = \frac{f(x,t)}{s(x,t)}$

Advantages

- Can calculate hazard, density, survival function, life expectancy, entropy, quantiles and associated confidence intervals in a continuous setting.
- It is equivalent to calculating life tables in a continuous setting.
- Can derive and integrate any function (with trapz() from pracma).

- We start with a vector of death rates M(x, t),
- Simulate a great number of lifespans (>100,000) using M(x, t) as the vector of rates in the function rpexp() of the package msm.

$$s(x,t) = e^{-nM(x,t)}$$
 and $\mu(x,t) = \frac{f(x,t)}{s(x,t)}$

Advantages

- Can calculate hazard, density, survival function, life expectancy, entropy, quantiles and associated confidence intervals in a continuous setting.
- It is equivalent to calculating life tables in a continuous setting.
- Can derive and integrate any function (with trapz() from pracma).

Disadvantages

It only helps when you are looking at changes over age

- We start with a vector of death rates M(x, t),
- Simulate a great number of lifespans (>100,000) using M(x, t) as the vector of rates in the function $\mathtt{rpexp}()$ of the package \mathtt{msm} .

$$s(x,t) = e^{-nM(x,t)}$$
 and $\mu(x,t) = \frac{f(x,t)}{s(x,t)}$

Advantages

- Can calculate hazard, density, survival function, life expectancy, entropy, quantiles and associated confidence intervals in a continuous setting.
- It is equivalent to calculating life tables in a continuous setting.
- Can derive and integrate any function (with trapz() from pracma).

Disadvantages

- · It only helps when you are looking at changes over age
- It requires a lot of RAM (and time).

Smoothing

Poisson counts with P-splines (Camarda, 2012)

Age and year specific death counts are assumed to follow a Poisson distribution. P-splines and GLM are employed.

Poisson counts with P-splines (Camarda, 2012)

Age and year specific death counts are assumed to follow a Poisson distribution. P-splines and GLM are employed.

1. Fit model.

Age and year specific death counts are assumed to follow a Poisson distribution. P-splines and GLM are employed.

- 1. Fit model.
- 2. Predict log-hazard for very small intervals x = 0, 0.01, 0.02... and t = 0, 0.01, 0.02...

Age and year specific death counts are assumed to follow a Poisson distribution. P-splines and GLM are employed.

- 1. Fit model.
- 2. Predict log-hazard for very small intervals x = 0, 0.01, 0.02... and t = 0, 0.01, 0.02...

Advantages

• In R, MortalitySmooth package.

Age and year specific death counts are assumed to follow a Poisson distribution. P-splines and GLM are employed.

- 1. Fit model.
- 2. Predict log-hazard for very small intervals x = 0, 0.01, 0.02... and t = 0, 0.01, 0.02...

- In R, MortalitySmooth package.
- It can smooth in two dimensions (age and time).

Age and year specific death counts are assumed to follow a Poisson distribution. P-splines and GLM are employed.

- 1. Fit model.
- 2. Predict log-hazard for very small intervals x = 0, 0.01, 0.02... and t = 0, 0.01, 0.02...

- In R, MortalitySmooth package.
- It can smooth in two dimensions (age and time).
- Can derive and integrate any function (with trapz() from pracma).

Age and year specific death counts are assumed to follow a Poisson distribution. P-splines and GLM are employed.

- 1. Fit model.
- 2. Predict log-hazard for very small intervals x = 0, 0.01, 0.02... and t = 0, 0.01, 0.02...

- In R, MortalitySmooth package.
- It can smooth in two dimensions (age and time).
- Can derive and integrate any function (with trapz() from pracma).
- · It can produce confidence intervals.

Age and year specific death counts are assumed to follow a Poisson distribution. P-splines and GLM are employed.

- 1. Fit model.
- 2. Predict log-hazard for very small intervals x = 0, 0.01, 0.02... and t = 0, 0.01, 0.02...

Advantages

- In R, MortalitySmooth package.
- It can smooth in two dimensions (age and time).
- Can derive and integrate any function (with trapz() from pracma).
- · It can produce confidence intervals.

Disadvantages

Age and year specific death counts are assumed to follow a Poisson distribution. P-splines and GLM are employed.

- 1. Fit model.
- 2. Predict log-hazard for very small intervals x = 0, 0.01, 0.02... and t = 0, 0.01, 0.02...

Advantages

- In R, MortalitySmooth package.
- It can smooth in two dimensions (age and time).
- Can derive and integrate any function (with trapz() from pracma).
- It can produce confidence intervals.

Disadvantages

It is only designed for mortality data.

Age and year specific death counts are assumed to follow a Poisson distribution. P-splines and GLM are employed.

- 1. Fit model.
- 2. Predict log-hazard for very small intervals x = 0, 0.01, 0.02... and t = 0, 0.01, 0.02...

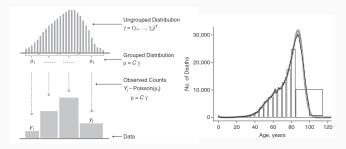
Advantages

- In R, MortalitySmooth package.
- It can smooth in two dimensions (age and time).
- Can derive and integrate any function (with trapz() from pracma).
- It can produce confidence intervals.

Disadvantages

It is only designed for mortality data.

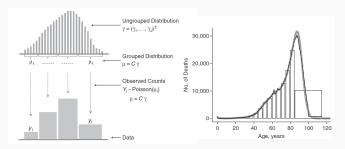
Ungroup age groups and smooth death counts:



Advantages

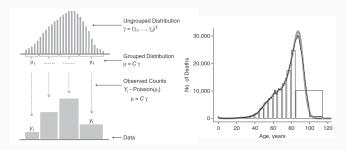
• In R, ungroup package.

Ungroup age groups and smooth death counts:



- In R, ungroup package.
- It can smooth in two dimensions (age and time).

Ungroup age groups and smooth death counts:

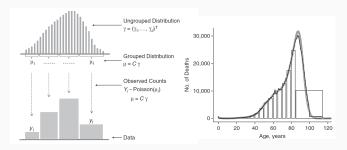


Advantages

- In R, ungroup package.
- It can smooth in two dimensions (age and time).

Disadvantages

Ungroup age groups and smooth death counts:



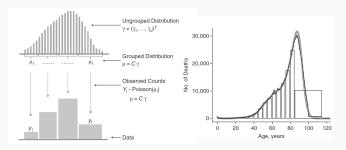
Advantages

- In R, ungroup package.
- It can smooth in two dimensions (age and time).

Disadvantages

· It is only designed for mortality data.

Ungroup age groups and smooth death counts:



Advantages

- In R, ungroup package.
- It can smooth in two dimensions (age and time).

Disadvantages

- · It is only designed for mortality data.
- · Not able to produce confidence intervals (so far).

Does Demography need differential equations?

by Thomas K. Burch

Why has demography made relatively little use of differential equations?

Why has demography made relatively little use of differential equations?

"Demography is or aspires to be an autonomous science, not just a branch of applied statistics".

Why has demography made relatively little use of differential equations?

"Demography is or aspires to be an autonomous science, not just a branch of applied statistics".

"The average demographer has little competence in the use of differential equations."

Why has demography made relatively little use of differential equations?

"Demography is or aspires to be an autonomous science, not just a branch of applied statistics".

"The average demographer has little competence in the use of differential equations."

"That level of mathematics has not been required for entrance into, or successful completion of, most graduate programs."

Why has demography made relatively little use of differential equations?

"Demography is or aspires to be an autonomous science, not just a branch of applied statistics".

"The average demographer has little competence in the use of differential equations."

"That level of mathematics has not been required for entrance into, or successful completion of, most graduate programs."

"Demographers generally were not schooled in differential equations, so we did not try to use them, and avoided topics that required their use even at the most elementary level."

"Demography is generally thought to be rich in data and technique, and poor in theory."

"Demography is generally thought to be rich in data and technique, and poor in theory."

"I think Demography might be an even stronger discipline if it had assimilated the regular use of differential equations in general, and systems dynamics software in particular."

"Demography is generally thought to be rich in data and technique, and poor in theory."

"I think Demography might be an even stronger discipline if it had assimilated the regular use of differential equations in general, and systems dynamics software in particular."

"...their use would encourage us to think more about dynamics and process, and not just cross-sectional relationships and equilibria. They could help us think better about complex social and demographic systems containing non-linear relationships and feedbacks."

"Demography is generally thought to be rich in data and technique, and poor in theory."

"I think Demography might be an even stronger discipline if it had assimilated the regular use of differential equations in general, and systems dynamics software in particular."

"...their use would encourage us to think more about dynamics and process, and not just cross-sectional relationships and equilibria. They could help us think better about complex social and demographic systems containing non-linear relationships and feedbacks."

"Our body of theory could be richer still if we were to take advantage of both classic (differential equations) and contemporary (software) tools for the statement and manipulation of theoretical ideas about demographic processes".

"Demography is generally thought to be rich in data and technique, and poor in theory."

"I think Demography might be an even stronger discipline if it had assimilated the regular use of differential equations in general, and systems dynamics software in particular."

"...their use would encourage us to think more about dynamics and process, and not just cross-sectional relationships and equilibria. They could help us think better about complex social and demographic systems containing non-linear relationships and feedbacks."

"Our body of theory could be richer still if we were to take advantage of both classic (differential equations) and contemporary (software) tools for the statement and manipulation of theoretical ideas about demographic processes".

Burch, Thomas K. Model-based demography: Essays on integrating data, technique and theory. Springer Nature, 2018.

Questions?