



A Comprehensive Analysis of the Patterns of Worldwide Mortality Evolution

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Motivation

- For example, life expectancy increases in most countries all over the world.
- But changes in life expectancy (and other typically used statistics) are only a consequence of the underlying change of the age distribution of deaths.



- How does the shape of the deaths curve change over time and how can we classify that?
- Are there different evolutions for different countries? And are there any common patterns and/or exceptions?

Agenda

Motivation

Data: What do we look at?

Toolkit: How do we compress the data?

The Classification Framework of Börger et al. Detection of Trends

Results

Summary and Outlook



Data: What do we look at?

The Human Mortality Database (HMD) provides mortality data for

- 42 different populations and
- for females and males all over the world.
- For 34 populations the data history is longer than 40 years.
- We calculate **death curves** for each
 - population,
 - sex,
 - starting age, and
 - calendar year, where we have data.

We analyze the changes of these deaths curves over time and compare them to each other.



The Classification Framework (1)

- There exists a variety of literature on the question how the age distribution of deaths changes over time.
- We have identified some shortcomings there (for details, see Börger et al. 2016):
 - Often the established scenarios were defined imprecisely, e.g. compression, extension, rectangularization, ...
 - Some of these scenarios were supposed to be **mutually exclusive**, which is demonstrably false.
 - Several often used statistics are insufficient or even misleading.
 - Often effects caused by the choice of a certain age range under observation were not considered.
- We recently introduced a **unique classification system** for mortality evolution patterns , which...
 - ... rules out the shortcomings which we mentioned above,
 - ... uniquely classifies any kind of change in the age distribution of deaths,
 - ... is based on 4 statistics.



The Classification Framework (2)



Detection of Trends – Examples: M and DoI for Swedish females, starting age 0



- When we compute these statistics on real data, we can get such plots.
- We can observe three things:
 - We must find a formalization of the statements "increasing", "neutral", and "decreasing".
 - We have to find a method, with which we can explicitly identify changes in the trend.
 - We observe heteroscedasticity.



Detection of Trends - Examples: M and DoI for Swedish females, starting age 0

- Muggeo (2003) proposed a method, with which historical data can be fitted with a continuous and piecewise linear curve.
 - We minimize the Akaike Information Criterion (AIC) to find the best trade-off between goodness of fit and number trend changes. The AIC has the advantage to be time invariant.
- This method was modified by Börger et al. (2015), such that it considers heteroscedasticity in the input data.
- With a test for significance we determine the trend of the statistics in this intervals and can classify it in the three categories "increasing", "neutral", and "decreasing".



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Females, starting age 0, M



left-shift
neutral
right-shift

- Right shift seems to be an almost global trend for the last decades.
 - But some East European countries experience left shift or at least no shift after 1970 and around 1990.



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Females, starting age 0, UB



contraction
neutral
extension

- No global trend concerning extension / contraction.
- For example, in the second half of the 20th century, we observe...
 - ... extension in most Central European countries,
 - ... no change in Australia and New Zealand, but also for Luxembourg or Iceland,
 - European countries for some years.



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Females, starting age 0, DoI



decompression
neutral
compression

There seems to be compression for most countries in the last decades.

Exception: Some East European countries around 1990 and single "outliers".



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Females, starting age 0, d(M)



diffusion
neutral
concentration

- Concentration also seems to be a global effect in the last two decades, but there are some exceptions:
 - Almost all East European countries experience diffusion around 1990.
 - The North European countries have a relative long period of diffusion or neutral starting in the 1950s / 60s. Exception: Norway
 - There was diffusion in Japan in the 1990s and in the USA in the late 2000s.



Conclusion

Is clustering a reasonable idea?

- We can detect "vertical patterns".
 - There are common trend changes for some countries, e.g.
 - East European countries around 1970 and 1990, or
 - Central European countries in the 1960s / 70s.
- But there are some countries which do not experience the same trend changes as their neighboring countries. For example:
 - The mortality evolution for females in Belgium shows different trend breaks, than that for females in the Netherlands or Luxembourg.
 - Norway does not experience diffusion in the 1950s/60s.

Thus, clustering is a reasonable idea but it must be handled with care.





Summary and Outlook (1)

What have we seen?

- We introduced a framework, which we use for the classification of changes in mortality evolution patterns.
- Calculation of these four statistics give us results partly with rather strong variance.
- With help of the method of Muggeo (2003) and Börger et al. (2015) and a statistical test, we can formalize the terms "increasing", "neutral", and "decreasing" in mathematical terms.
- We plot these trends for every population over the time axis. With these plots we can compare mortality evolutions of every country to each other.
- For females with the starting age 0, we analyzed these results and have seen...
 - ... that there are "vertical patterns", which show common trend changes between countries,
 - ... that there are some "outliers" within a cluster of neighboring countries.

Summary and Outlook (2)

Outlook: What are the next steps?

- We have to do **further research**:
 - Analysis of more populations, especially men
 - Analysis of different starting ages, e.g. starting age 60.
- These results could for example help in the **application of multi-population models**.
 - Is it reasonable to model the mortality of small populations (e.g. Luxembourg) depending on the mortality of bigger populations (e.g. West Germany)?

Thank you for your attention!

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Literature

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