

Lecture 5:

Coherent Mortality Forecasting: a short introduction

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Advances in Mortality Forecasting

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Background

- Individual forecasts likely imply increasing divergence in life expectancy, against the observed global convergence in mortality levels (Wilson 2001)
- In the recent past, there has been a growing body of research on so-called coherent forecasting
- What does it mean? Simultaneous forecast of:
 - males and females from the same population
 - a group of countries
 - more cause-of-death
 - small sub-population belonging to the same country
- Let's have a look at what has been done

“Playing” on the Lee-Carter idea

- Li, N. and R. D. Lee (2005). Coherent mortality forecasts for a group of populations: An extension of the Lee-Carter method. *Demography* **42**, 575-594.
- For a given age x , year t and population i :

$$\ln(m_{x,t,i}) = a_{x,i} + B_x K_t + b_{x,i} k_{t,i}$$

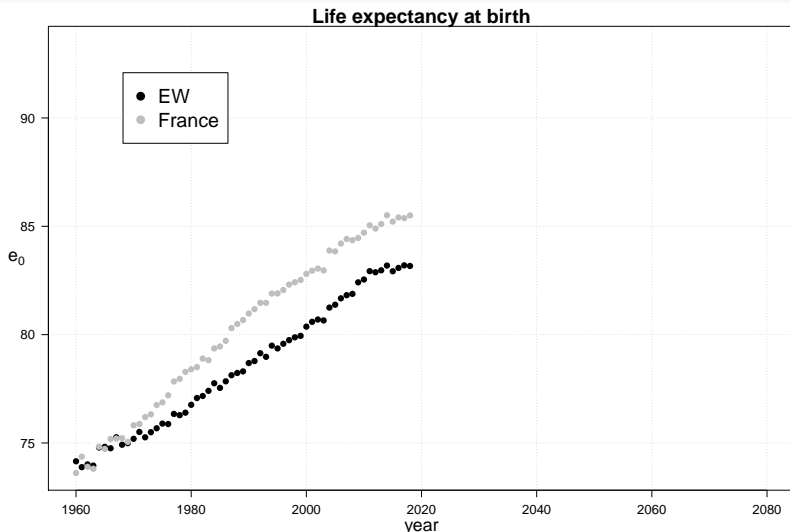
where

- $a_{x,i}$: population-specific average log-mortality
- K_t : time index for the whole group of populations
- B_x : sensitivity of the log-mortality at age x to variations in the time index for the whole group of populations
- $b_{x,i}$ and $k_{t,i}$ population-specific parameters-vectors
- In the original paper, they estimate the model using SVD

Li-Lee: estimation & forecast

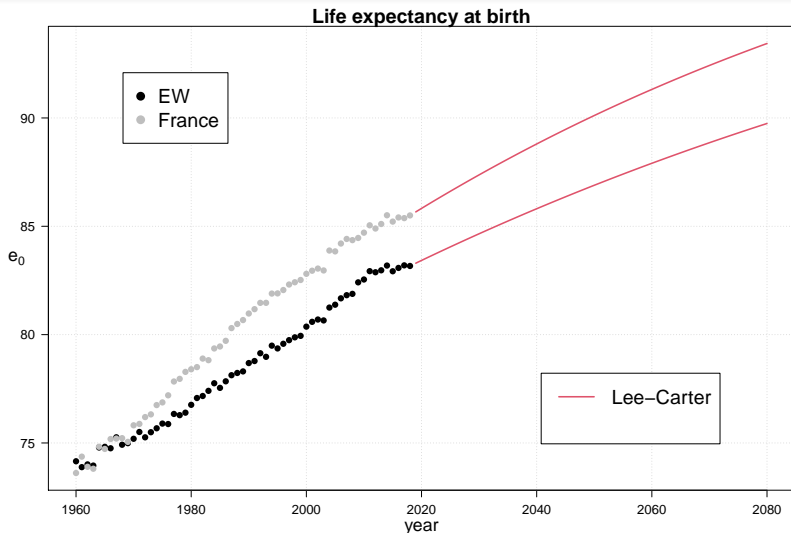
- ① Estimate the common factor $B_x K_t$ by applying LC model to the whole group, adjusting K_t to fit the group's average life expectancy
- ② Estimate the population-specific $a_{x,i}$ as the average age-specific mortality in group i over the observed time period
- ③ Estimate specific factor $b_{x,i} k_{t,i}$ by employing SVD on the matrix $\ln(m_{x,t,i}) - a_{x,i} - B_x K_t$ (no need of second step adjustment to $k_{t,i}$)
- ④ Forecasting:
 - K_t : random walk with drift (as in the LC model)
 - $k_{t,i}$: AR(1) model (or RW) **without** drift (to ensure convergent forecasts)

Li-Lee: an example



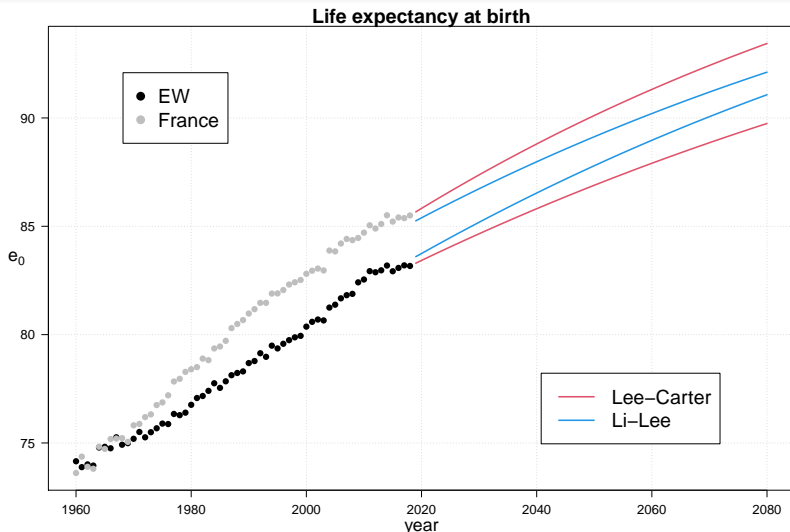
Females aged 0–105+ in England & Wales and France, fitting period 1960–2018, forecast 2019–2080. Source: HMD (2021)

Li-Lee: an example



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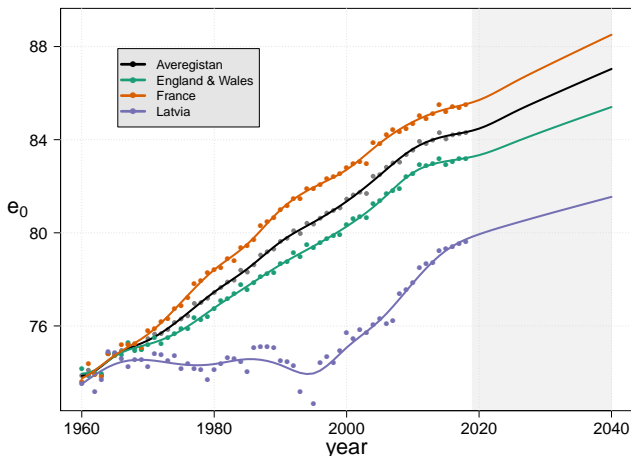
Coherent *CP*-splines

The idea:

given forecast values for a mean of populations, constrain mortality differences between each population and the forecast mean to lay within (a range of) observed past differences

- Let's take two examples:
 - France, England & Wales and *Latvia*, females, 1960-2018, ages 0-105
 - Australia and its 8 territories, both sexes, 1971-2016, ages 0-100

CP-splines on “Averegistan” and each component

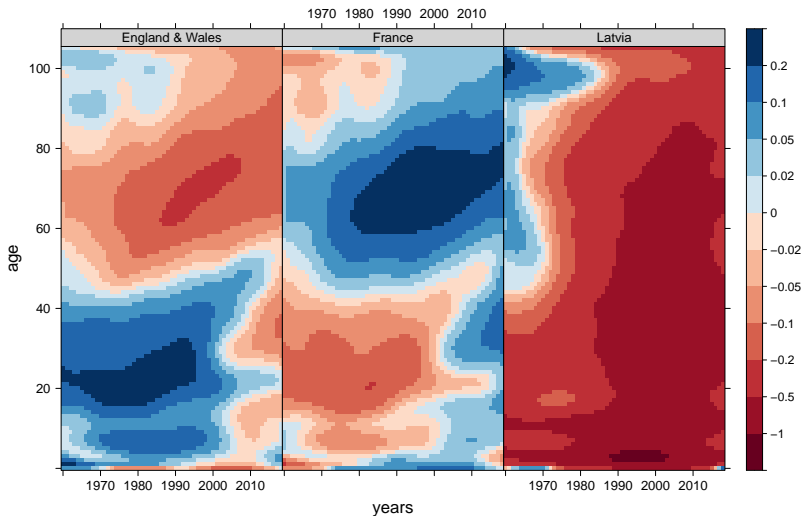


Actual, estimated and forecast life expectancy by *CP*-splines.
“Averegistan”, France, England & Wales and Latvia



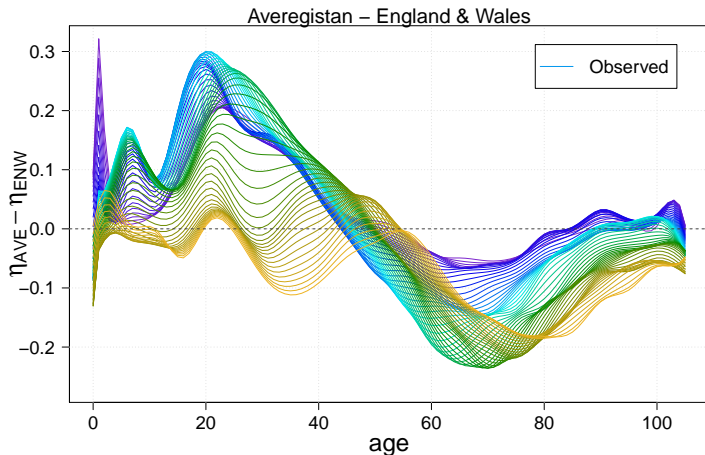
Differences in log-mortality: only England & Wales

Differences in log-mortality: all 3 populations



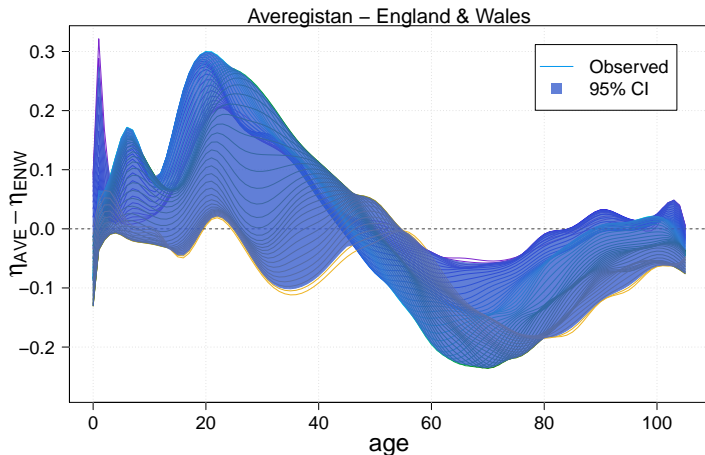
Observed differences in smooth log-mortality: “Averegistan” - each population

Differences in future log-mortality with *CP*-splines



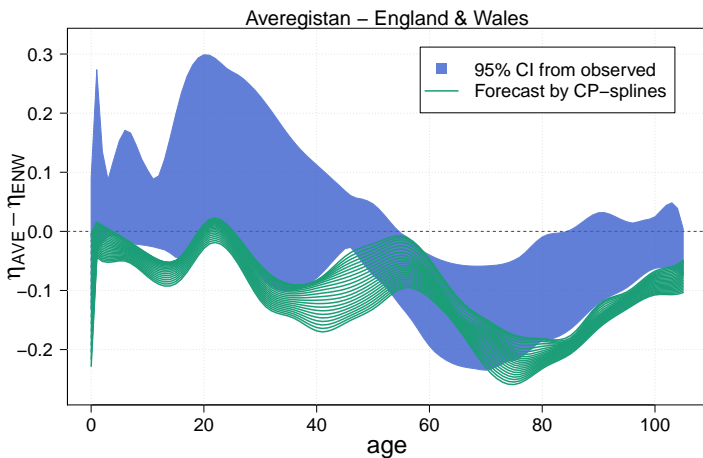
Differences in estimated smooth log-mortality

Differences in future log-mortality with *CP*-splines



Differences with the 95% CI in estimated log-mortality

Differences in future log-mortality with *CP*-splines



95% CI of estimated & forecast (*CP*-splines) differences in log-mortality

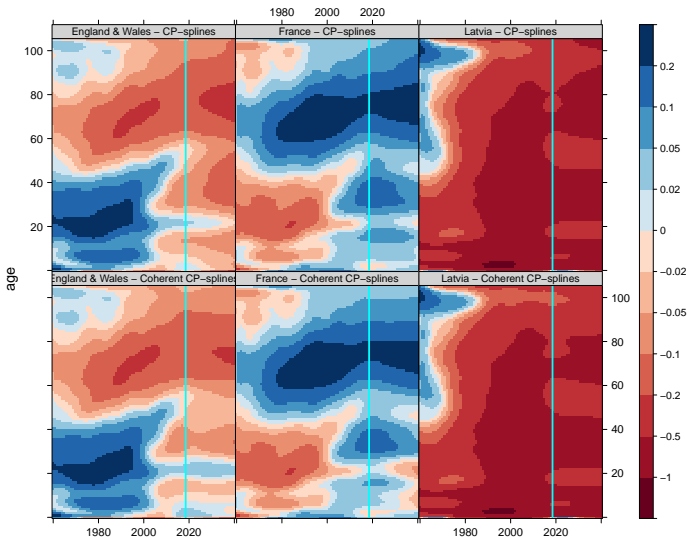
Constraining differences: Coherent CP -splines

- Incorporate constraints on future differences as CP -splines do with relative derivatives
- Asymmetric penalty can be adapted
- Coherence with respect to known overall mortality preserved
- Each (sub-)population can be treated independently
- We achieve age-specific coherence in future years simultaneously
- Limits:
 - Knowledge about the future mortality in the “Averegistan” is necessary
 - Range from past differences are kept in the future

Asymmetric penalty in action on differences

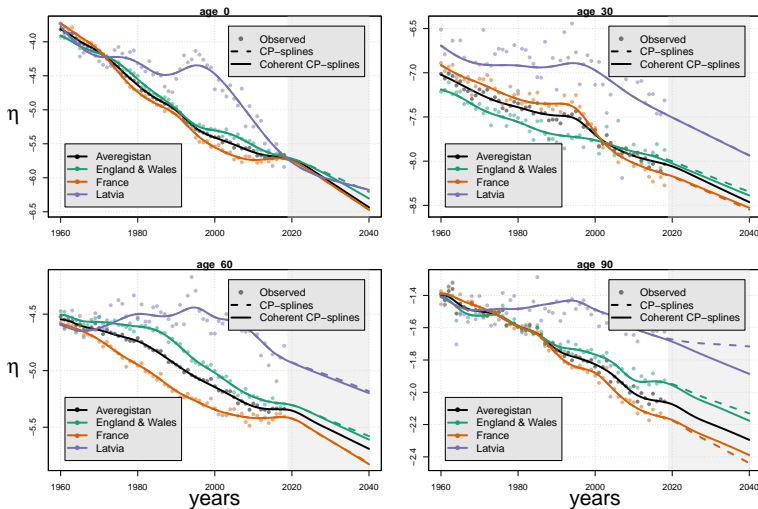
95% CI of estimated & forecast (Coherent *CP*-splines)
differences in log-mortality

Forecast differences: Averegistan - each population



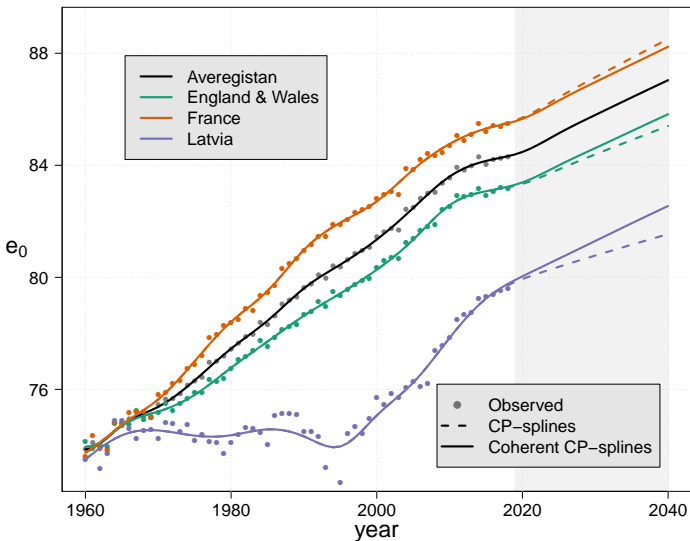
Estimated and forecast differences in log-mortality

Outcomes for both approaches



Log-mortality over years for selected ages

Outcomes for both approaches



Life expectancy at birth

Coherent CP -splines. Australian territories e_0

Females	CP -splines	Coherent CP -splines	Δ
Australia	89.69	-	-
Australian Capital Territory	90.05	89.91	-0.14
New South Wales	89.75	89.64	-0.11
Northern Territory	86.57	86.11	-0.46
Queensland	90.18	89.90	-0.27
South Australia	89.49	89.73	0.24
Tasmania	87.64	88.32	0.68
Victoria	90.28	89.92	-0.36
Western Australia	90.20	90.08	-0.12

Forecast female life expectancy in 2050 by (coherent) CP -splines.
Australia and its territories

Coherent CP -splines. Australian territories e_0

Males	CP -splines	Coherent CP -splines	Δ
Australia	87.04	-	-
Australian Capital Territory	88.20	87.92	-0.29
New South Wales	87.47	87.08	-0.40
Northern Territory	81.78	83.13	1.35
Queensland	86.75	86.79	0.04
South Australia	86.88	86.93	0.05
Tasmania	84.94	85.78	0.85
Victoria	87.97	87.47	-0.51
Western Australia	87.49	87.33	-0.16

Forecast male life expectancy in 2050 by (coherent) CP -splines.
Australia and its territories

A Functional Data approach

- Hyndman R. et al. (2013). Coherent Mortality Forecasting: The Product-Ratio Method With Functional Time Series Models. *Demography*, **50**, 261-283.
 - for each t, i , smoothing $m_{x,t,i}$ over x with weighted penalised regression splines with monotonic increase after 65+
 - compute

$$\text{product}_{x,t} = \left[\prod_{i=1}^I m_{x,t,i} \right]^{1/I} \quad \text{ratio}_{x,t,i} = m_{x,t,i} / \text{product}_{x,t}$$

- Apply functional principal component analysis:

$$\begin{aligned} \ln(\text{product}_{x,t}) &= a_x^{\text{product}} + \sum_{k=1}^K \beta_{t,k} \phi_{x,k} \\ \ln(\text{ratio}_{x,t,i}) &= a_{x,i}^{\text{ratio}} + \sum_{l=1}^L \delta_{x,l,i} \psi_{x,l,i} \end{aligned}$$

with K and L up to 6

A Compositional Data approach

- Bergeron-Boucher, M.-P. et al. (2017). Coherent forecasts of mortality with compositional data analysis. *Demographic Research* **37**, 527-566.
- They consider $d_{x,t,i}$ (deaths from life-table, i.e. density) instead of rates as compositional data: vectors of components which are strictly positive, carry only relative information, and always sum to a constant
- use the *clr* transformation defined as the logarithm of the composition divided by its geometric mean:

$$clr(d_{t,x}) = \ln \left(\frac{d_{x,t,i}}{g_t} \right)$$

- They translate Li-Lee model in the new simplex:

$$clr(d_{x,t,i} \ominus a_{x,i} \ominus C[e^{K_t B_x}]) = k_{t,i} b_{x,i}$$

A Bayesian Hierarchical approach

- Ševčíková, H. et al. (2016). Age-Specific Mortality and Fertility Rates for Probabilistic Population Projections. In R. Schoen (Ed.): Demographic methods and population analysis. pp. 285-310. Springer.
- They take life expectancy for all the world and forecast it by Bayesian hierarchical models
- For a given country, they convert overall levels
 - for ages above 100, they fit (on 80-99) and the extrapolate:

$$m_x = \frac{ae^{bx}}{1 + ae^{bx}}$$

where both sexes share common b

- for ages 0-99, they fit a variant of a Lee-Carter where both β_x and κ_t is common between sexes
- They estimate and forecast by 5-years age/time groups

Final slide

- Concluding remarks:

- (Good) mortality forecasting is a (laborious) process
- Details matter, be prepared to spend time on them
- Fortunately, software is getting better
- After some experience and this course, you ought master some mortality forecasting models
- Believe us: it is a lot of fun!
- And finally remember:
"It is very difficult to predict – especially the future"

Niels Bohr (maybe)

- Acknowledgments:

- Heiner Maier for organizing the logistics behind the course
- Emilio Zagheni and Mikko Myrskylä for welcoming this course as part of the PHDS and IDEM programs
- All of you for attending the course
- **Final hope:** to meet all in person sooner than later!