

Advances in Stochastic Mortality Modelling

Robust Probabilistic Feature Extraction

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**“Stochastic Period and Cohort Effect State-Space Mortality Models
Incorporating Demographic Factors via Probabilistic Robust Principle
Components”**

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- 2 State Space Stochastic Mortality Models
- 3 State-Space Hybrid Factor Models: Regression Extensions
- 4 Robust Probabilistic Feature Extraction Methods
- 5 Application to Mortality Modelling and Demographic Data
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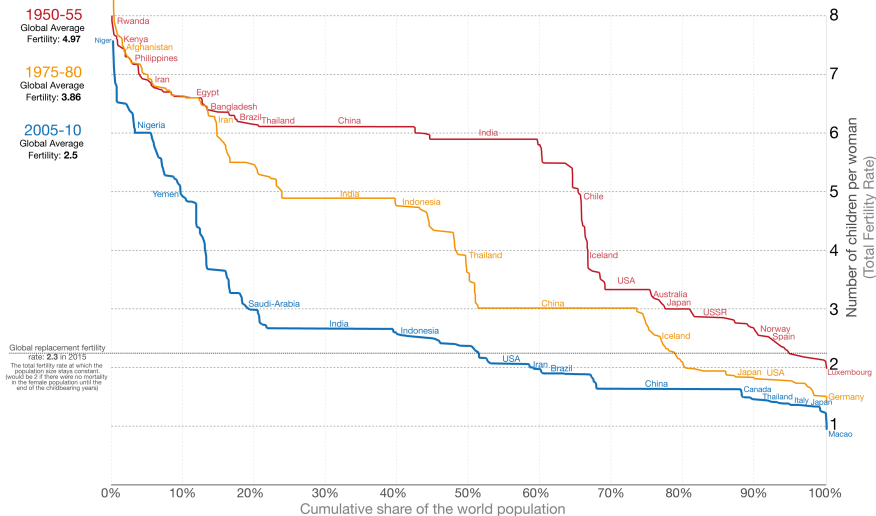
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- Ageing populations are a major challenge for many countries.
 - **Fertility rates are declining while life expectancy is increasing.**
- **longevity risk:** *the adverse financial outcome of people living longer than expected* ⇒ *possibility of outliving their retirement savings.*
 - **long term demographic risk** has significant implications for societies and manifests as a **systematic risk for pension plans and annuity providers.**
- Policymakers rely on **mortality projection** to determine appropriate pension benefits and regulations regarding retirement.

Our World
in Data

World population by level of fertility over time (1950-2010)

On the x-axis you find the cumulative share of the world population. The countries are ordered along the x-axis descending by the total fertility rate of the country.



Data source: United Nations Population Division (2012 revision).

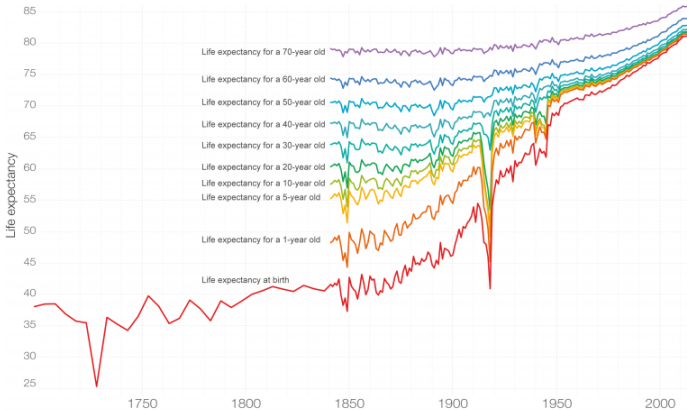
The interactive data visualization is available at OurWorldinData.org. There you find the raw data and more visualizations on this topic.

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Life Expectancy by Age in England and Wales, 1700-2013

Shown is the total life expectancy given that a person reached a certain age.

Our World
in Data



Data source: Life expectancy at birth Clio-Infra. Data on life expectancy at age 1 and older from the Human Mortality Database (www.mortality.org).
The interactive data visualization is available at OurWorldinData.org. There you find the raw data and more visualizations on this topic.

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Enhancing mortality models requires an understanding of common features of mortality behaviour [Cairns, Blake and Dowd, 2008]

- Mortality rates have fallen dramatically at all ages.
- Rate of decrease in mortality has **varied over time and by age group**.
- Absolute decreases have **varied by age group**.
- Aggregate mortality rates have significant **volatility year on year**.

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The uncertainty in future death rates can be divided into two components:

- **Unsystematic mortality risk.** Even if the true mortality rate is known, the number of deaths, will be random.
 - larger population \Rightarrow smaller unsystematic mortality risk (due to pooling of offsetting risks - diversification).
- **Systematic mortality risk.** This is the undiversifiable component of mortality risk that affects all individuals in the same way.
 - **Forecasts of mortality rates in future years are uncertain.**

- [Lee and Carter, 92] proposed a stochastic mortality model (LC) to forecast the trend of age-specific mortality rates.
- Several extensions to Lee-Carter model have been proposed, overview in [Fung et al. 2017].

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Survival probability is still consistently underestimated
especially in the last few decades ([IMF, 2012]).

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Survival probability is still consistently underestimated especially in the last few decades ([IMF, 2012]).

This talk considers models aiming to help resolve this issue via

- **Stochastic State-Space Mortality Models** with **Period** and **Cohort** stochastic latent effects (LCC).
- + Extensions to State-Space Hybrid Regression Structures!

(see [Fung et al. 2017] and [Fung et al. 2018])

A state space model has two model components:

- **a stochastic observation equation**; and
- **a stochastic latent Markov state process**.

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Key advantages of state space modelling approach:

- remove awkward identification specifications;
- computational efficiency and numerical robustness;
- accurate in-sample and out-of-sample forecasts;
- optimal statistical efficiency and unbiased estimation;

Period-Cohort effect state-space formulation

Observation equation: log crude death rates, $y_{x,t} = \ln \hat{m}_{x,t}$, follow:

$$\ln \hat{m}_{x,t} = \alpha_x + \beta_x \kappa_t + \beta_x^\gamma \gamma_{t-x} + \varepsilon_{x,t},$$

where $\varepsilon_{x,t}$ is a regression noise term.

- $\alpha = \alpha_{x_1:x_p} := [\alpha_{x_1}, \dots, \alpha_{x_p}]$ represents the *age-profile of the log death rates*
- $\beta = \beta_{x_1:x_p}$ measures the *sensitivity of death rates for different age group* to a change of period effect κ_t .
- $\beta^\gamma = \beta_{x_1:x_p}^\gamma$ measures the *sensitivity of death rates for different age group* to a change of cohort effect γ_{t-x} .

Observation Process: in matrix form.

$$\begin{pmatrix} y_{x_1,t} \\ y_{x_2,t} \\ \vdots \\ y_{x_p,t} \end{pmatrix} = \begin{pmatrix} \alpha_{x_1} \\ \alpha_{x_2} \\ \vdots \\ \alpha_{x_p} \end{pmatrix} + \begin{pmatrix} \beta_{x_1} & \beta_{x_1}^\gamma & 0 & \cdots & 0 \\ \beta_{x_2} & 0 & \beta_{x_2}^\gamma & \cdots & 0 \\ \vdots & \vdots & \vdots & \ddots & \vdots \\ \beta_{x_p} & 0 & 0 & \cdots & \beta_{x_p}^\gamma \end{pmatrix} \begin{pmatrix} \kappa_t \\ \gamma_t^{x_1} \\ \gamma_t^{x_2} \\ \vdots \\ \gamma_t^{x_p} \end{pmatrix} + \begin{pmatrix} \varepsilon_{x_1,t} \\ \varepsilon_{x_2,t} \\ \vdots \\ \varepsilon_{x_p,t} \end{pmatrix}.$$

Here $(\kappa_t, \gamma_t^{x_1}, \dots, \gamma_t^{x_p})^\top$ is the $p + 1$ dimensional latent state vector. $\gamma_t^x := \gamma_{t-x}$

State Equation in matrix form:

$$\begin{pmatrix} \kappa_t \\ \gamma_t^{x_1} \\ \gamma_t^{x_2} \\ \vdots \\ \gamma_t^{x_{p-1}} \\ \gamma_t^{x_p} \end{pmatrix} = \begin{pmatrix} 1 & 0 & 0 & \cdots & 0 & 0 \\ 0 & \lambda_1 & \lambda_2 & \cdots & \lambda_{p-1} & \lambda_p \\ 0 & 1 & 0 & \cdots & 0 & 0 \\ 0 & 0 & 1 & \cdots & 0 & 0 \\ \vdots & \vdots & \vdots & \ddots & \vdots & \vdots \\ 0 & 0 & 0 & \cdots & 1 & 0 \end{pmatrix} \begin{pmatrix} \kappa_{t-1} \\ \gamma_{t-1}^{x_1} \\ \gamma_{t-1}^{x_2} \\ \vdots \\ \gamma_{t-1}^{x_{p-1}} \\ \gamma_{t-1}^{x_p} \end{pmatrix} + \begin{pmatrix} \theta \\ \eta \\ 0 \\ \vdots \\ 0 \\ 0 \end{pmatrix} + \begin{pmatrix} \omega_t^\kappa \\ \omega_t^\gamma \\ 0 \\ \vdots \\ 0 \\ 0 \end{pmatrix}.$$

Period effect κ_t is a random walk with drift process with $\omega_t^\kappa \stackrel{iid}{\sim} N(0, \sigma_\omega^2)$ and Cohort effect $\gamma_t^{x_1}$ is a stationary AR(p) process with $\omega_t^\gamma \stackrel{iid}{\sim} N(0, \sigma_\gamma^2)$

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GOAL: develop *stochastic mortality state-space hybrid factor models*.

- **Hybrid** := Stochastic Latent Factors + Observable Covariate Features
- **observable features extracted** from demographic data
- Feature extraction should aim for dimension reduction
⇒ model parsimony.

[Toczydlowska and Peters, 2017] address important aspects of feature extraction:

- 1 **missing data** in time-series and panel (matrix) structured real demographic data;
- 2 **noisy observations and outliers** (in real data);

Two fundamental approaches to develop Hybrid Factor Models:

- 1 time varying factor with **static loading coefficient** (*classical distributed lag regressions such as ARDL models*);
 - 2 static factor with **time varying stochastic loading coefficients**. (*state space models e.g. dynamic Nelson-Siegel yield curves*).
- Option 2: suitable for **high dimensional** data, **time series** / **panel structured** but represented by relatively “**short time series**” lengths.
 - \Rightarrow *particularly prevalent in demographic studies!*

Consider the State-Space Hybrid Period-Cohort-Demographic Model

$$\begin{aligned} \mathbf{y}_t &= \boldsymbol{\alpha} + \tilde{\mathbf{B}}_t \tilde{\boldsymbol{\varphi}}_t + \boldsymbol{\varepsilon}_t, \quad \boldsymbol{\varepsilon}_t \stackrel{iid}{\sim} \mathcal{N}(\mathbf{0}, \sigma_\varepsilon^2 \mathbb{I}_p), \\ \tilde{\boldsymbol{\varphi}}_t &= \tilde{\Lambda} \tilde{\boldsymbol{\varphi}}_{t-1} + \tilde{\Theta} + \tilde{\boldsymbol{\omega}}_t, \quad \tilde{\boldsymbol{\omega}}_t \stackrel{iid}{\sim} \mathcal{N}(\mathbf{0}, \tilde{\Upsilon}) \end{aligned}$$

where $\tilde{\boldsymbol{\varphi}}_t = (\boldsymbol{\varphi}_t, \boldsymbol{\varrho}_t)$ is a $(p + pk + 1) \times 1$ latent process vector of $\boldsymbol{\varphi}_t$ **stochastic mortality factors** (period-cohort) and $\boldsymbol{\varrho}_t$ **dynamic factor loadings**, with

$$\tilde{\Theta} = \begin{pmatrix} \Theta_{(p+1) \times 1} \\ \Psi_{pk \times 1} \end{pmatrix}_{(p+pk+1) \times 1}$$

a vector of drift parameters for state equations.

Consider three models:

Case 1: Factors in
Observation
Equation Only;

Case 2: Factors in
Period Effect State
Equation Only;

Case 3: Factors in
Cohort Effect State
Equation Only.

$$\tilde{\mathbf{B}}_{t \, p \times (\rho + \rho k + 1)} = \begin{cases} \left(\begin{array}{c|c} \mathbf{B}_{\rho \times (\rho + 1)} & \tilde{\mathbf{F}}_t \end{array} \right) & \text{for Case 1,} \\ \left(\begin{array}{c|c} \mathbf{B}_{\rho \times (\rho + 1)} & \mathbf{0}_{\rho \times \rho k} \end{array} \right) & \text{otherwise,} \end{cases}$$

$$\tilde{\mathbf{\Lambda}}_{(\rho + \rho k + 1) \times (\rho + \rho k + 1)} = \begin{cases} \left(\begin{array}{c|c} \mathbf{\Lambda}_{(\rho + 1) \times (\rho + 1)} & \mathbf{0}_{(\rho + 1) \times \rho k} \\ \mathbf{0}_{\rho k \times (\rho + 1)} & \mathbf{\Omega}_{\rho k \times \rho k} \end{array} \right) & \text{for Case 1,} \\ \left(\begin{array}{c|c} \mathbf{\Lambda}_{(\rho + 1) \times (\rho + 1)} & \tilde{\mathbf{f}}_t^T \\ \mathbf{0}_{\rho k \times (\rho + 1)} & \mathbf{\Omega}_{\rho k \times \rho k} \end{array} \right) & \text{for Case 2,} \\ \left(\begin{array}{c|c} \mathbf{\Lambda}_{(\rho + 1) \times (\rho + 1)} & \mathbf{0}_{1 \times \rho k} \\ \mathbf{0}_{\rho k \times (\rho + 1)} & \tilde{\mathbf{F}}_t \end{array} \right) & \text{for Case 3.} \end{cases}$$

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- Data \mathbf{Y}_t is observed (or partially observed) over periods $t \in \{1, \dots, n\}$ and will be reduced to factors $\tilde{\mathbf{F}}_t$

Example: d countries demographic data and p denotes the number of different demographic attributes observed
 \Rightarrow then $p \times d$ matrix of data in year t is \mathbf{Y}_t .

- **We do not wish to utilise the raw demographic data**
 $\tilde{\mathbf{F}}_t \neq \mathbf{Y}_t$:
in general it will produce a model with too many parameters
- [Toczydlowska and Peters, 2017] considered stochastic projection methods of dimensionality reduction
 \Rightarrow **Probabilistic Principal Component Analysis (PPCA)** and **Robust** extensions.

PCA by means of Factor Analysis: with n realisations of the $(p \times d)$ -dimensional observed demographic data, vectorized into columns \mathbf{Y} .

Consider linear decompositions:

$$\mathbf{Y}_{n \times pd} = \mathbf{X}_{n \times pd} \mathbf{W}_{pd \times pd}^T + \epsilon_{n \times pd}.$$

Factor analysis assumes diagonal covariance for ϵ_t .

Stochastic Factor PCA: differs from deterministic PCA as components \mathbf{x}_t and **factor loading matrix** \mathbf{W} account for correlation between elements of \mathbf{y}_t and only part of the variation:

$$\mathbb{E} \mathbf{y}_t^T \mathbf{y}_t = \mathbb{E} \left[\left(\mathbf{x}_t \mathbf{W}^T + \epsilon_t \right)^T \left(\mathbf{x}_t \mathbf{W}^T + \epsilon_t \right) \right] = \mathbf{W} \mathbf{\Lambda} \mathbf{W}^T + \mathbf{\Psi}.$$

In standard PCA \mathbf{x}_t and \mathbf{W} account for the entire covariance.

Show \mathbf{x}_t and \mathbf{W} account for correlation!

Example: assume $\mathbf{x}_t \sim \mathcal{N}(\mathbf{0}, \mathbb{I}_d)$ and $\epsilon_t \sim \mathcal{N}(\mathbf{0}, \Psi)$ to obtain,

$$\mathbf{y}_t | \mathbf{x}_t, \mathbf{W}, \Psi \sim \mathcal{N}(\mathbf{x}_t \mathbf{W}^T, \Psi),$$

$$\pi(\mathbf{y}_t | \mathbf{W}, \Psi) = \int_{\mathbb{R}^d} \pi(\mathbf{y}_t, \mathbf{x}_t | \mathbf{W}, \Psi) d\mathbf{x}_t = (2\pi)^{-\frac{d}{2}} |\mathbf{C}|^{-1} \exp\left\{-\frac{1}{2} \mathbf{y}_t \mathbf{C}^{-1} \mathbf{y}_t^T\right\}$$

for $\mathbf{C} = \mathbf{W}\mathbf{W}^T + \Psi$ where $|\mathbf{C}|$ denotes the determinant of the matrix.

- Notice that since Ψ is diagonal, the correlation structure between components \mathbf{y}_t is specified by the matrix \mathbf{W} .

Probabilistic Feature Extraction

Show \mathbf{x}_t and \mathbf{W} account for correlation cont.

Eigen decomposition of covariance $\mathbf{C} = \mathbf{U}_{d \times d} \mathbf{L}_{d \times d} \mathbf{U}^T$, for diagonal \mathbf{L} and orthonormal \mathbf{U} , gives

$$\mathbf{0} = (\mathbf{C} - \mathbf{L})\mathbf{U} = (\mathbf{W}^T \mathbf{W} + \sigma^2 \mathbb{I}_d - \mathbf{L}) \mathbf{U} = (\mathbf{W} \mathbf{W}^T - (\mathbf{L} - \sigma^2 \mathbb{I}_d)) \mathbf{U}.$$

- Thus, the matrix $\mathbf{\Lambda} = (\mathbf{L} - \sigma^2 \mathbb{I}_d)$ and \mathbf{U} are matrices of eigenvalues and corresponding eigenvectors of $\mathbf{W} \mathbf{W}^T$.
- Since $\lambda_j = l_j - \sigma^2 \geq 0$, the scalar σ^2 can be chosen as the smallest diagonal element of $\mathbf{\Lambda}$.
- **Factor loadings are given by $\mathbf{U} \mathbf{\Lambda}^{\frac{1}{2}}$.**

Assuming the error term ϵ_t is homogeneous s.t. $\mathbf{\Psi} = \sigma^2 \mathbb{I}_d$, then estimating \mathbf{W} via PCA given $\mathbf{C} = \mathbf{W} \mathbf{W}^T + \sigma^2 \mathbb{I}_d$ is identifiable.

Probabilistic Feature Extraction

Feature Extraction via EM Algorithm Estimation!

Goal is to estimate:

- projection matrix \mathbf{W} ,
- vector $\boldsymbol{\mu}$ and
- scalar σ^2

given marginal distribution of \mathbf{Y}_t

$$\mathbf{Y}_t | \Psi \sim \mathcal{N} \left(\boldsymbol{\mu}, \mathbf{W}\mathbf{W}^T + \sigma^2 \mathbb{I}_d \right)$$

for the vector of static parameters $\Psi = [\mathbf{W}, \boldsymbol{\mu}, \sigma^2]$ of the model.

The EM algorithm uses logarithm of the the complete data likelihood, i.e. the joint distribution of $\mathbf{Y}_{1:N}, \mathbf{X}_{1:N} | \Psi$ given by

$$\begin{aligned} \pi_{\mathbf{Y}_{1:N}, \mathbf{X}_{1:N} | \Psi}(\mathbf{y}_{1:N}, \mathbf{x}_{1:N}) &= \prod_{t=1}^N \pi_{\mathbf{Y}_t | \mathbf{X}_t, \Psi}(\mathbf{y}_t) \pi_{\mathbf{X}_t | \Psi}(\mathbf{x}_t) \\ &= (2\pi)^{-N \frac{d+k}{2}} (\sigma^2)^{-N \frac{d}{2}} \exp \left\{ -\frac{1}{2\sigma^2} \sum_{t=1}^N (\mathbf{y}_t - \boldsymbol{\mu} - \mathbf{x}_t \mathbf{W}^T) (\mathbf{y}_t - \boldsymbol{\mu} - \mathbf{x}_t \mathbf{W}^T)^T - \frac{1}{2} \sum_{t=1}^N \mathbf{x}_t \mathbf{x}_t^T \right\}. \end{aligned}$$

Feature Extraction via EM Algorithm Estimation!

1. **Expectation step:** Expectation of the loglikelihood function of the joint distribution of $\mathbf{Y}_{1:N}, \mathbf{X}_{1:N} | \Psi$ for a fixed vector of static parameters Ψ^* with respect to the conditional distribution $\mathbf{X}_{1:N} | \mathbf{Y}_{1:N}, \Psi$

$$Q(\Psi, \Psi^*) = \mathbb{E}_{\mathbf{X}_{1:N} | \mathbf{Y}_{1:N}, \Psi} \left[\log \pi_{\mathbf{Y}_{1:N}, \mathbf{X}_{1:N} | \Psi^*}(\mathbf{y}_{1:N}, \mathbf{x}_{1:N}) \right]$$

2. **Maximisation step:** Finding $\mathbf{W}^*, \boldsymbol{\mu}^*$ and σ^{*2} that maximize $Q(\Psi | \Psi^*)$

$$\left(\mathbf{W}^*, \boldsymbol{\mu}^*, \sigma^{*2} \right) = \underset{\mathbf{W}^* \in \mathbb{R}^{d \times k}, \boldsymbol{\mu}^* \in \mathbb{R}^d, \sigma^{*2} > 0}{\operatorname{argmax}} Q(\Psi, \Psi^*)$$

Theorem

The E-step of the EM algorithm for Gaussian Probabilistic Principal Component Analysis given N realisations of the observation vector \mathbf{Y}_t denoted by $\mathbf{y}_{1:N} = \{\mathbf{y}_1, \dots, \mathbf{y}_N\}$ is obtained in the closed form as follows

$$\begin{aligned} Q(\Psi, \Psi^*) &= \mathbb{E}_{\mathbf{x}_{1:N} | \mathbf{y}_{1:N}, \Psi} \left[\log \pi_{\mathbf{y}_{1:N}, \mathbf{x}_{1:N} | \Psi^*}(\mathbf{y}_{1:N}, \mathbf{x}_{1:N}) \right] \\ &= -\frac{N(d+k)}{2} \log 2\pi - \frac{Nd}{2} \log \sigma^{*2} - \frac{1}{2} \sum_{t=1}^N \left\{ \frac{1}{\sigma^{*2}} \text{Tr} \{ \mathbf{y}_t^T \mathbf{y}_t \} \right. \\ &\quad - \frac{2}{\sigma^{*2}} \mathbf{y}_t \boldsymbol{\mu}^{*T} + \frac{1}{\sigma^{*2}} \boldsymbol{\mu}^* \boldsymbol{\mu}^{*T} - \frac{2}{\sigma^{*2}} \text{Tr} \left\{ \mathbf{W}^* \mathbb{E}_{\mathbf{x}_t | \mathbf{y}_t, \Psi} [\mathbf{X}_t^T] \mathbf{y}_t \right\} \\ &\quad \left. + \frac{2}{\sigma^{*2}} \mathbb{E}_{\mathbf{x}_t | \mathbf{y}_t, \Psi} [\mathbf{X}_t] \mathbf{W}^{*T} \boldsymbol{\mu}^{*T} + \text{Tr} \left\{ \left(\frac{1}{\sigma^{*2}} \mathbf{W}^{*T} \mathbf{W}^* + \mathbb{I}_k \right) \mathbb{E}_{\mathbf{x}_t | \mathbf{y}_t, \Psi} [\mathbf{X}_t^T \mathbf{X}_t] \right\} \right\} \end{aligned}$$

see details of expectations and proof in [Toczydlowska and Peters, 2017].

Theorem

The maximizers of the function $Q(\Psi, \Psi^*)$ are given by

$$\mu^* = \bar{\mu}(\mathbf{y}_{1:N}; \Psi) \left(\mathbb{I}_d - \mathbf{W}\mathbf{M}^{-1}\mathbf{W}^{*T} \right) + \mu\mathbf{W}\mathbf{M}^{-1}\mathbf{W}^{*T}$$

$$\mathbf{W}^* = \bar{\mathbf{C}}_{\mu, \mu^*}(\mathbf{y}_{1:N}; \Psi, \Psi^*)\mathbf{W}\mathbf{M}^{-1} \left(\sigma^2\mathbf{M}^{-1} + \mathbf{M}^{-1}\mathbf{W}^T\bar{\mathbf{C}}_{\mu}(\mathbf{y}_{1:N}; \Psi)\mathbf{W}\mathbf{M}^{-1} \right)^{-1}$$

$$\sigma^{*2} = \frac{1}{d} \text{Tr} \left\{ \bar{\mathbf{C}}_{\mu^*}(\mathbf{y}_{1:N}; \Psi, \Psi^*) - 2\mathbf{W}^*\mathbf{M}^{-1}\mathbf{W}^T\bar{\mathbf{C}}_{\mu, \mu^*}(\mathbf{y}_{1:N}; \Psi, \Psi^*) \right. \\ \left. + \mathbf{W}^* \left(\sigma^2\mathbf{M}^{-1} + \mathbf{M}^{-1}\mathbf{W}^T\bar{\mathbf{C}}_{\mu}(\mathbf{y}_{1:N}; \Psi)\mathbf{W}\mathbf{M}^{-1} \right) \mathbf{W}^{*T} \right\}$$

see details of components and proof in [\[Toczydlowska and Peters, 2017\]](#).

Probabilistic PCA with Missing Data:

Until now, we assumed the data did not contain any missing observations!

- Real demographic time series data have **numerous types of missingness**.
- \Rightarrow *missingness is an important aspect to address in the feature extraction!*

[Toczydlowska and Peters, (2017), (2018)] *address different components of **PPCA in missing data** estimation settings via **robust** versions of Expectation-Maximisation.*

- Distributional Extensions: **Student-t, Skewed and Grouped Student-t cases.**

Probabilistic Feature Extraction

Feature Extraction via EM Algorithm with **MISSING DATA!**

Define the indicator random variable \mathbf{R}_t which decides which entries of \mathbf{Y}_t are missing denoting them by 1, otherwise 0.

- Each observation consists of the pair $[\mathbf{Y}_t^o, \mathbf{R}_t]$ with distribution parameterized according to parameters $[\Psi, \Theta]$ respectively.

Likelihood is given by conditional probability $\mathbf{Y}_t^o, \mathbf{R}_t | \Psi, \Theta$:

$$\begin{aligned}\pi_{\mathbf{Y}_t^o, \mathbf{R}_t | \Psi, \Theta}(\mathbf{y}_t^o, \mathbf{r}_t) &= \int \pi_{\mathbf{Y}_t^o, \mathbf{Y}_t^m, \mathbf{R}_t | \Psi, \Theta}(\mathbf{y}_t^o, \mathbf{y}_t^m, \mathbf{r}_t) d\mathbf{y}_t^m \\ &= \int \pi_{\mathbf{R}_t | \mathbf{Y}_t, \Psi, \Theta}(\mathbf{r}_t) \pi_{\mathbf{Y}_t | \Psi, \Theta}(\mathbf{y}_t) d\mathbf{y}_t^m\end{aligned}$$

We assume for simplicity a pattern of missing data according to **MAR** - missing at random

- The assumptions imposes the indicator variable \mathbf{R}_t to be independent of the value of missing data.

Then the vector \mathbf{Y}_t which is MAR satisfies

$$\pi_{\mathbf{R}_t|\mathbf{Y}_t,\Psi}(\mathbf{r}_t) = \pi_{\mathbf{R}_t|\mathbf{Y}_t^o,\Psi}(\mathbf{r}_t)$$

resulting in

$$\begin{aligned}\pi_{\mathbf{Y}_t^o,\mathbf{R}_t|\Psi,\Theta}(\mathbf{y}_t^o) &= \pi_{\mathbf{R}_t|\mathbf{Y}_t^o,\Theta}(\mathbf{r}_t) \int \pi_{\mathbf{Y}_t|\Psi}(\mathbf{y}_t) d\mathbf{y}_t^m \\ &= \pi_{\mathbf{R}_t|\mathbf{Y}_t^o,\Theta}(\mathbf{r}_t) \pi_{\mathbf{Y}_t^o|\Psi}(\mathbf{y}_t^o).\end{aligned}$$

⇒ Under the MAR assumption, the estimation of Ψ via maximum likelihood of the joint distribution $\mathbf{Y}_t^o, \mathbf{R}_t|\Psi, \Theta$ is equivalent to the maximisation of the likelihood of the marginal distribution $\mathbf{Y}_t^o|\Psi$.

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Demographic data that we extract “Observable” covariate regression Features from:

- Data from Human Mortality Database (<http://www.mortality.org>).
- We use four different data sets:
 - Birth counts;
 - Death counts;
 - Life tables: Life Expectancy at Birth and Death Rates.
- The time series vary in terms of **data structure, the number of available observations and the missingness attributes** of the records.

TYPES OF DATA:

- **One dimensional time series data per country per gender**
(31 countries, M and F, gives 124 time series):
 - Birth counts and
 - Life expectancy at Birth.
- **Multivariate cross sectional time series data per country & gender:** age specific data for Death counts and Death Rates.
- **A single observation per country in time t describes:**
 - number of deaths of people with ages from 0 to 110+ (Death counts) or;
 - number of deaths for ages from 0 to 110+ scaled to the size of that population, per unit of time (Death Rates).

Model estimation performed by Forward-Backward Kalman Filter within Rao-Blackwellised Adaptive Gibbs Sampler (MCMC).

The state space models we considered in our studies were of type:

- 1 [LCC:] Lee-Carter model with the stochastic period + cohort effect.
- 2 [DFM-PC:] demographic factor model versions of Lee-Carter (Period-Cohort).

The factors are obtained by performing **Probabilistic Principle Component Analysis PPCA** jointly on the set of data for all countries listed, excluding:

United Kingdom (response variable)

LC State Space Model - only a Period Effect κ_t included.

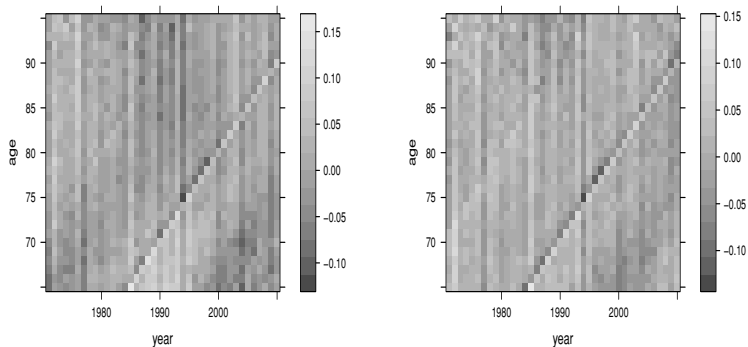


Figure: In sample analysis residuals (left Female, right Male).

LCC State Space Model - with Period + Cohort Effects κ_t, γ_{t-x} included.

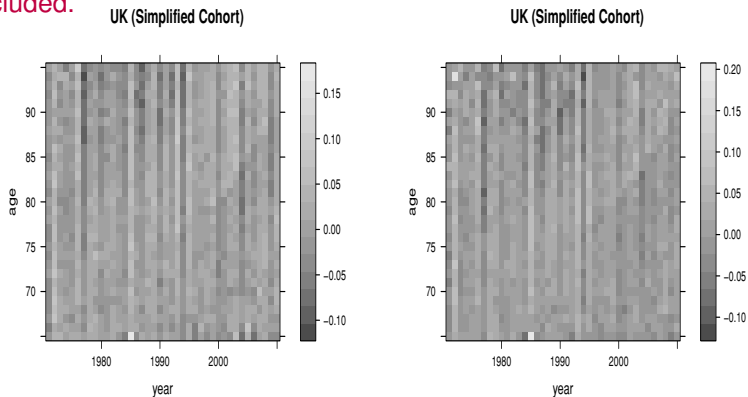


Figure: In sample analysis residuals (left Female, right Male).

- [DFM-PC-B:] the mean of first principal component of Birth counts as a static parameter, age specific element of ϱ_t ;
- [DFM-PC-D-r/s:] the first principal component of Death counts (which is age and country specific) as an exogenous factor, one element of ϱ_t corresponds to a country specific subvector of the component.;
- [DFM-PC-Mx-r/s:] the first principal component of Death Rates (which is age and country specific) as an exogenous factor, one element of ϱ_t corresponds to a country specific subvector of the component.

r/s - is robust vs standard

- **Out-of-Sample Study:** Model calibration period is 1922 – 2002
⇒ forecast performance analysis for 2003 – 2013

Model	MSE	DIC	MSEP _{MCMC}	MSEP _{Kalman}
LCC	0.0097	-3627	0.1778	0.1774
DFM-PC-B	0.0072	-6500	0.0057	0.0062
DFM-PC-D-r	0.0182	-6380	0.0177	0.0251
DFM-PC-D-s	0.0065	-5996	0.0185	0.0156
DFM-PC-Mx-r	0.0081	-8225	0.0111	0.0129
DFM-PC-Mx-s	0.0174	-3951	0.0692	0.0285

- **The results confirm that adding demographic features, as additional explanatory variables to the LCC model, improves both in-sample fit out-of-sample fit and therefore the predictability of log death rates.**

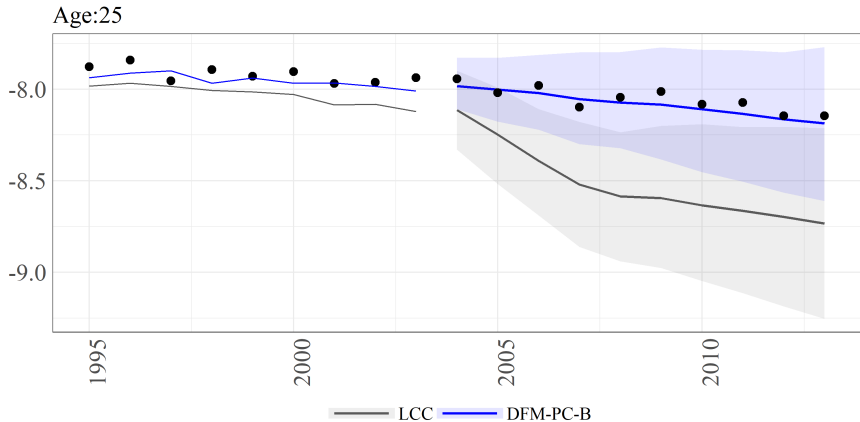


Figure: 10-year out-of-sample forecasted log death (y axis) rates by age with corresponding prediction intervals.

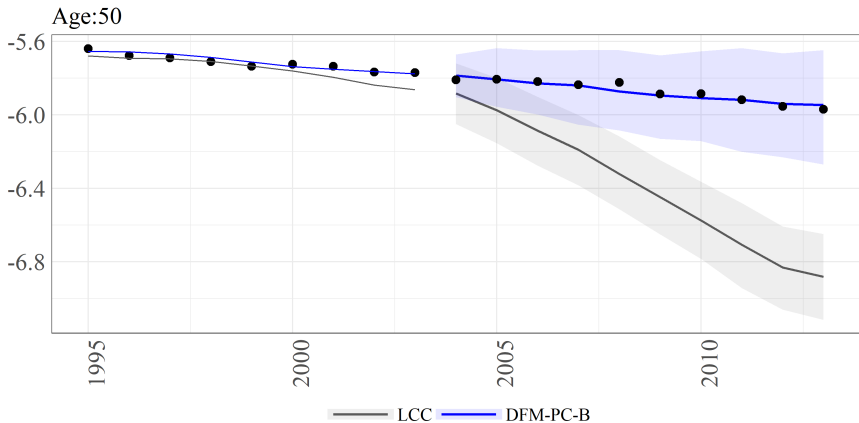


Figure: 10-year out-of-sample forecasted log death (y axis) rates by age with corresponding prediction intervals.

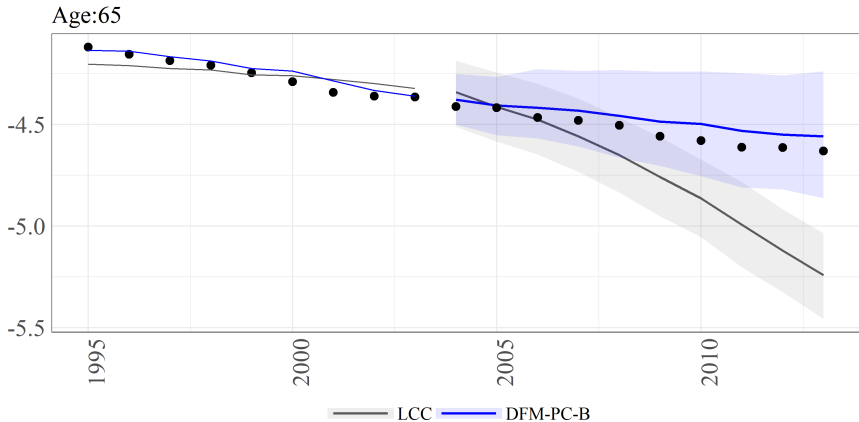


Figure: 10-year out-of-sample forecasted log death (y axis) rates by age with corresponding prediction intervals.

- We explored how to construct a state space formulation of the stochastic mortality models for Period and Cohort factors
- We explored how to extend to Hybrid Multi-Factor Stochastic State-Space Mortality models with Period-Cohort factors as well as demographic regressors.
- We briefly learnt about feature/covariate extraction methods to extract the demographic factors used in the extended HMF Stochastic State-Space Mortality models.
- Standard Lee-Carter Period-Cohort model consistently under estimates forecast log-death rates
- Extended models proposed improve significantly the forecast performance of log-death rates.

- 1 Mortality Modelling Context
- 2 State Space Stochastic Mortality Models
- 3 State-Space Hybrid Factor Models: Regression Extensions
- 4 Robust Probabilistic Feature Extraction Methods
- 5 Application to Mortality Modelling and Demographic Data
- 6 Appendix

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