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Fast Estimation of the Renshaw-Haberman Model and Its Variants

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Motivat	tion			

- Three main components in mortality modeling: Age, Time, **Cohort**.
 - Importance of cohort effects (particularly for UK data).
- Challenges of stochastic mortality models involving cohort effects.
 - Slow model fitting process.
 - Limited types of model fitting approaches (only likelihood-based).
- Our main objective Develop a fast **non-likelihood-based** estimation approach.

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Outline				

- Review of the Lee-Carter model and Renshaw-Haberman model (generalized APC model).
- 2 Main proposed methodology.
- Oumerical studies.
- Onclusion and further research.

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 Review:
 The Lee-Carter Model

The Lee-Carter model (Lee and Carter, 1992):

$$y_{x,t} := \log(m_{x,t}) = a_x + b_x k_t + \varepsilon_{x,t}, \quad (1)$$

with $\sum_{x} b_{x} = 1$ and $\sum_{t} k_{t} = 0$,

- $m_{x,t}$: Central mortality rate for age x and time t.
- a_x : Average log mortality rate for age x.
- b_x : Age effect (sensitivity) for age x.
- *k_t*: Time trend.

It involves two stage:

- Estimating a_x , b_x and k_t ;
- **2** Time series modeling on $\{k_t\}$ and forecasting.

Two Parameter Estimation Methods for the Lee-Carter Model

SVD, non-likelihood-based (Lee and Carter, 1992):

$$\min_{(a,b,k)} \sum_{x,t} (y_{x,t} - (a_x + b_x k_t))^2.$$
 (2)

The solution is obtained via PCA:

$$\hat{\boldsymbol{a}} = \bar{\boldsymbol{y}}, \quad \hat{\boldsymbol{b}} = \frac{\boldsymbol{u}}{\mathbf{1}^T \boldsymbol{u}}, \quad \hat{\boldsymbol{k}} = (\mathbf{1}^T \boldsymbol{u}) \cdot (\boldsymbol{Y} - \bar{\boldsymbol{Y}})^T \boldsymbol{u}.$$
 (3)

Poisson regression, likelihood-based (Brouhns et al., 2002):

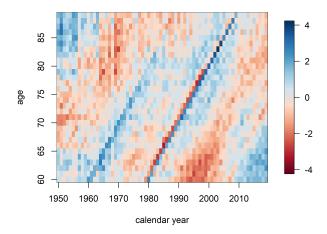
$$\max_{(\boldsymbol{a},\boldsymbol{b},\boldsymbol{k})}\sum_{x,t}\left(D_{x,t}(\boldsymbol{a}_x+\boldsymbol{b}_x\boldsymbol{k}_t)-N_{x,t}\boldsymbol{e}^{\boldsymbol{a}_x+\boldsymbol{b}_x\boldsymbol{k}_t}\right).$$
 (4)

The MLE is obtained via the iterative Newton-Raphson.

Comparison of the two methods: Interpretability? Flexibility? Computation?



Residual heatmap of England and Wales data:



Cohort effects (diagonal) are significant!

The Renshaw-Haberman model or generalized APC model (Renshaw and Haberman, 2006):

$$y_{x,t} := \log(m_{x,t}) = a_x + b_x k_t + c_x \gamma_{t-x},$$
 (5)

with
$$\sum_{x} b_x = \sum_{x} c_x = 1$$
 and $\sum_{t} k_t = \sum_{t-x} \gamma_{t-x} = 0$.

- c_x : Age effect (sensitivity) with respect to the cohort effects.
- γ_{t-x} : Cohort effects.

Simplified variants of the model can be obtained by setting $c_x = 1/p$ or $b_x = c_x = 1/p$, where p is the number of ages.

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Parameter Estimation and Slow Convergence

The parameter estimation is likelihood-based Poisson GLM framework (via iterative Newton-Raphson):

$$\max_{(\boldsymbol{a},\boldsymbol{b},\boldsymbol{k},\boldsymbol{c},\gamma)}\sum_{x,t}\left(D_{x,t}(\boldsymbol{a}_{x}+\boldsymbol{b}_{x}\boldsymbol{k}_{t}+\boldsymbol{c}_{x}\gamma_{t-x})-N_{x,t}\boldsymbol{e}^{\boldsymbol{a}_{x}+\boldsymbol{b}_{x}\boldsymbol{k}_{t}+\boldsymbol{c}_{x}\gamma_{t-x}}\right).$$
(6)

- Critical issue: Slow convergence rate (Cairns et al., 2009, 2011; Haberman and Renshaw, 2009, 2011).
- It hinders the use Monte-Carlo or bootstrap to examine parameter/model uncertainty.
- Some literature has worked on improving the convergence: Hunt and Villegas (2015) and Currie (2016).



To the best of our knowledge, all the existing estimation methods are likelihood-based.

Our objectives:

- Propose a non-likelihood-based (PCA-based) estimating method for the Renshaw-Haberman model and its variants.
- 2 Accelerate the parameter estimation process.



The non-likelihood-based approach to fit the Lee-Carter model is via SVD or PCA:

$$\min_{(a,b,k)} \sum_{x,t} (y_{x,t} - (a_x + b_x k_t))^2.$$
 (7)

Similarly, we aim at **directly minimizing the** L^2 **error** of the estimated log mortality rates under the RH framework:

$$\min_{(\boldsymbol{a},\boldsymbol{b},\boldsymbol{k},\boldsymbol{c},\boldsymbol{\gamma})}\sum_{x,t}\left(y_{x,t}-\left(a_{x}+b_{x}k_{t}+c_{x}\gamma_{t-x}\right)\right)^{2}.$$
 (8)

No close-form solution exists, due to the non-orthogonality.



- The minimization of (8) is via an iterative scheme:
 - **1** Set initial values of $\theta := (a, b, k, c, \gamma)$.
 - 2 Fixing **b**, **k**, **c** and γ , update **a**:

$$\min_{\boldsymbol{a}} \sum_{x,t} [\underbrace{(y_{x,t} - b_x k_t - c_x \gamma_{t-x})}_{\text{given}} - a_x]^2. \quad \text{(Easy)} \qquad (9)$$

3) Fixing \pmb{a} , \pmb{c} and $\pmb{\gamma}$, update \pmb{b} and \pmb{k} :

$$\min_{(\boldsymbol{b},\boldsymbol{k})} \sum_{x,t} [\underbrace{(y_{x,t} - a_x - c_x \gamma_{t-x})}_{\text{given}} - b_x k_t]^2. \quad \text{(Easy)}$$
(10)

• Fixing $\boldsymbol{a}_{\!\scriptscriptstyle X}$, \boldsymbol{b} and \boldsymbol{k} , update \boldsymbol{c} and $\boldsymbol{\gamma}$:

$$\min_{(\boldsymbol{c},\boldsymbol{\gamma})} \sum_{x,t} \underbrace{[(\boldsymbol{y}_{x,t} - \boldsymbol{a}_x - \boldsymbol{b}_x \boldsymbol{k}_t)]}_{\text{given}} - c_x \gamma_{t-x}]^2. \quad \text{(Difficult)} \quad (11)$$

If the objective function has not converged, go back to Step 2.



Step 4 is much more challenging and has no explicit solution:

$$\min_{(\mathbf{c},\gamma)}\sum_{x,t}\left[\underbrace{(y_{x,t}-a_x-b_xk_t)}_{\text{given}}-c_x\gamma_{t-x}\right]^2.$$

But it turns out that Step 4 is equivalent to solving a **PCA** with missing values.

Cohort-effect Estimation: Why PCA with Missing Values?

Denote the age range and time range by (x_1, \dots, x_p) and (t_1, \dots, t_n) . Letting $z_{x,t} := y_{x,t} - a_x - b_x k_t$, we can transform the age-time matrix to the new age-cohort matrix¹

¹This is guaranteed by the one-to-one transformation between (x, t) and (x, t - x). Also, \times represents missing values.



Values - Formulation and Algorithms

• Letting s = t - x, the PCA with missing values problem can be formulated as:

$$\min_{(\boldsymbol{a},\boldsymbol{c},\boldsymbol{\gamma})} \sum_{(x,t-x)\in\mathcal{O}} [z_{x,t-x} - c_x \gamma_{t-x}]^2,$$
(12)

where $\ensuremath{\mathcal{O}}$ is the set of the indices of the observed values.

- Different approaches exist to solve PCA with missing values:
 - Small-scale: Iterative SVD (focus of this project);
 - 2 Large-Scale and sparse: matrix completion with nuclear norm regularization (Mazumder et al., 2010), etc.

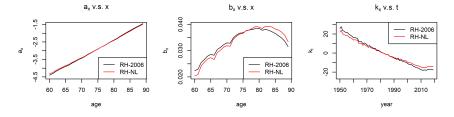
Iterative SVD algorithm (also known as hard-imputation):

- Initialization: Impute the missing values and initialize the approximate complete matrix Z_c .
- Approximate PCA: Implement PCA via SVD to the approximate complete matrix Z_c.
- Imputation: Update the missing values by using the corresponding PCA reconstructions in Step 2.
- Repeat Steps 2 and 3 until the convergence criterion is satisfied.

We proved the convergence of the iterative SVD algorithm within our context.

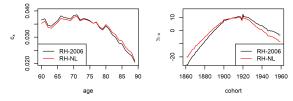
- In this talk, we compare two methods:
 - *RH-2006*: Standard iterative Newton-Raphson method (Renshaw and Haberman, 2006).
 - *RH-NL*: Proposed non-likelihood-based method.
- Data set: England & Wales (EW) and US male mortality rates, from 1950 to 2019, age from 60-89.
- We compare the fitted L^2 errors, log-likelihoods and computation time.

Plots: RH Model, EW Data



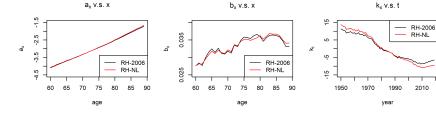


 γ_{t-x} v.s. t-x



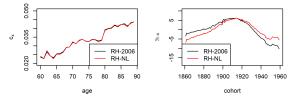
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Plots: RH Model, US Data





 γ_{t-x} v.s. t-x



RH Model	Data	RH-2006
 L ² error	EW 0.5	0.578
L^{-} error		0 470

L^2 error	EW	0.578	0.565
Leno	US	0.472	0.465
Log-likelihood	EW	-12843	-12890
Log-likelihood	US	-17736	-17828
Time (coc)	EW	336.68	38.72
Time (sec)	US	228.72	10.44

RH-NI

We have chosen the same tolerance levels (relative changes of the objective function) to make the comparison fair.

Conclusion and Future Research

- We proposed a non-likelihood-based estimating method for the Renshaw-Haberman model
- The proposed method is significantly faster than the traditional likelihood-based method, while producing satisfactory estimation results.
- Possible future research:
 - Further accelerate the algorithm by incorporating additional identification constraints.
 - Extension to multi-component structure, with proper regularization.

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