



Actuarial Research Centre

Institute and Faculty
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Actuarial Research Centre (ARC)

PhD studentship output

The Actuarial Research Centre (ARC) is the Institute and Faculty of Actuaries' network of actuarial researchers around the world. The ARC seeks to deliver research programmes that bridge academic rigour with practitioner needs by working collaboratively with academics, industry and other actuarial bodies.

The ARC supports actuarial researchers around the world in the delivery of cutting-edge research programmes that aim to address some of the significant challenges in actuarial science.

Multi-population mortality models

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joint work with

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Actuarial
Research Centre

Motivation and objectives

- Multi-population mortality models at older ages
- Research is pretty limited in comparison to the single-population mortality models
- Comparison
 - Fitting solution for the models
 - Identifiability problems correction
 - Joint mortality scenario forecast
- Rank the models value

The models

Model 0 $\log[m_{(x,t,i)}] = \alpha_{(x,i)} + B_{(x)}K_{(t)} + \beta_{(x,i)}\kappa_{(t,i)} + \varepsilon_{(x,t,i)}$

The Li and Lee model (2005)

$m_{(x,t,i)}$	Crude mortality rates (deaths over exposure)	$x \times t \times i$
$\alpha_{(x,i)}$	Determines the general mortality shape	$x \times i$
$K_{(t)}$	Explains the evolution of the global mortality over time	$t \times 1$
$B_{(x)}$	Shows which rates change more rapidly in response to $K_{(t)}$	$x \times 1$
$\kappa_{(t,i)}$	Explains the country specific evolution of the mortality over time	$t \times i$
$\beta_{(x,i)}$	Shows which rates change more rapidly in response to $\kappa_{(t,i)}$	$x \times i$

The models

Model 0 $\log[m_{(x,t,i)}] = \alpha_{(x,i)} + B_{(x)}K_{(t)} + \beta_{(x,i)}\kappa_{(t,i)} + \varepsilon_{(x,t,i)}$

Model 1 $\log[m_{(x,t,i)}] = \alpha_{(x,i)} + B_{(x)}K_{(t)} + \beta_{(x)}\kappa_{(t,i)} + \varepsilon_{(x,t,i)}$

↳ Common parameter: $\beta_{(x,i)} \rightsquigarrow \beta_{(x)}$

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The models

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Model 1 $\log[m_{(x,t,i)}] = \alpha_{(x,i)} + B_{(x)}K_{(t)} + \beta_{(x)}\kappa_{(t,i)} + \varepsilon_{(x,t,i)}$

↳ Common parameter: $\beta_{(x,i)} \rightsquigarrow \beta_{(x)}$

Model 2 $\log[m_{(x,t,i)}] = \alpha_{(x,i)} + B_{(x)}K_{(t)} + B_{(x)}\kappa_{(t,i)} + \varepsilon_{(x,t,i)}$

↳ Common parameter: $\beta_{(x,i)} \rightsquigarrow B_{(x)}$

$m_{(x,t,i)}$	Crude mortality rates (deaths over exposure)	$x \times t \times i$
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The models

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Model 1 $\log[m_{(x,t,i)}] = \alpha_{(x,i)} + B_{(x)}K_{(t)} + \beta_{(x)}\kappa_{(t,i)} + \varepsilon_{(x,t,i)}$

↳ Common parameter: $\beta_{(x,i)} \rightsquigarrow \beta_{(x)}$

Model 2 $\log[m_{(x,t,i)}] = \alpha_{(x,i)} + B_{(x)}K_{(t)} + B_{(x)}\kappa_{(t,i)} + \varepsilon_{(x,t,i)}$

The Kleinow model (2015)

↳ Common parameter: $\beta_{(x,i)} \rightsquigarrow B_{(x)}$

Model 3 $\log[m_{(x,t,i)}] = \alpha_{(x,i)} + \beta_{(x)}^1 \kappa_{(t,i)}^1 + \beta_{(x)}^2 \kappa_{(t,i)}^2 + \varepsilon_{(x,t,i)}$

Country specific parameter: $K_{(t)} \rightsquigarrow \kappa_{(t,i)}^1$ ↯

↳ Common parameter: $\beta_{(x,i)} \rightsquigarrow \beta_{(x)}^2$

$m_{(x,t,i)}$	Crude mortality rates (deaths over exposure)	$x \times t \times i$
$\alpha_{(x,i)}$	Determines the general mortality shape	$x \times i$
$\kappa_{(t,i)}^1$	Explains the evolution of the global mortality over time	$t \times 1$
$\beta_{(x)}^1$	Shows which rates change more rapidly in response to $\kappa_{(t,i)}^1$	$x \times 1$
$\kappa_{(t,i)}^2$	Explains the country specific evolution of the mortality over time	$t \times i$
$\beta_{(x)}^2$	Shows which rates change more rapidly in response to $\kappa_{(t,i)}^2$	$x \times 1$

Identifiability problems

■ The models are undetermined

■ Parametrisation I

Model 0

$$\log[m_{(x,t,i)}] = \tilde{\alpha}_{(x,i)} + \tilde{B}_{(x)} \tilde{K}_{(t)} + \tilde{\beta}_{(x,i)} \tilde{\kappa}_{(t,i)} + \varepsilon_{(x,t,i)}$$

$$\begin{aligned} \tilde{\alpha}_{(x,i)} &= \alpha_{(x,i)} + aB_{(x)} + c\beta_{(x,i)} & \tilde{B}_{(x)} &= \frac{B_{(x)}}{b} \\ \tilde{\beta}_{(x,i)} &= \frac{\beta_{(x,i)}}{d} & \tilde{K}_{(t)} &= b(K_{(t)} - a) \\ \tilde{\kappa}_{(t,i)} &= d(\kappa_{(t,i)} - c) \end{aligned}$$

■ Parametrisation II

Model 2

$$\log[m_{(x,t,i)}] = \alpha_{(x,i)} + B_{(x)} \left[\underbrace{K_{(t)} + C_{(t)}}_{\tilde{K}_{(t)}} + \underbrace{\kappa_{(t,i)} - C_{(t)}}_{\tilde{\kappa}_{(t,i)}} \right] + \varepsilon_{(x,t,i)}$$

Parameter constraints

Model 0	Model 1	Model 2	Model 3
<i>Common constraints</i>			
$\sum B(x) = 1$	$\sum B(x) = 1$	$\sum B(x) = 1$	$\sum \beta_{(x)}^1 = 1$
$\sum_t K(t) = 0$	$\sum_t K(t) = 0$	$\sum_t K(t) = 0$	$\sum_x \beta_{(x)}^2 = 1$
-	$\sum_x \beta_{(x)} = 1$	-	-
<i>Country specific constraints</i>			
For each i :			
$\sum \beta_{(x,i)} = 1$	-	-	$\sum_t \kappa_{(t,i)}^1 = 0$
$\sum_t \kappa_{(t,i)} = 0$	$\sum_t \kappa_{(t,i)} = 0$	$\sum_t \kappa_{(t,i)} = 0$	$\sum_t \kappa_{(t,i)}^2 = 0$
<i>Time specific constraints</i>			
For each t :			
-	$\sum_i \kappa_{(t,i)} = 0$	$\sum_i \kappa_{(t,i)} = 0$	$\sum_i \kappa_{(t,i)}^2 = 0$
-	Quasi identifiability constraint	True identifiability constraint	Quasi identifiability constraint

Parameters estimation

■ The used countries data

<i>i</i>	Country	Exposure to risk at the age of 60 in 2010	
		Male	Female
1	Austria	47023.17	49526.5
2	Belgium	65344.67	66434.33
3	Czech Republic	71575.69	71575.69
4	Denmark	34420.17	35132.33
5	Sweden	59759.83	59742.67
6	Switzerland	46527.67	47078.67

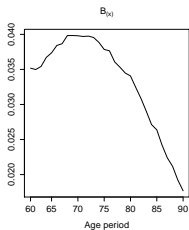
■ Single step maximum likelihood estimation

$$l = \sum_{x,t,i} \left[D_{(x,t,i)} \log(m_{(x,t,i)}) - E_{(x,t,i)} m_{(x,t,i)} \right] + \mathbf{C}_{(x,t,i)}$$

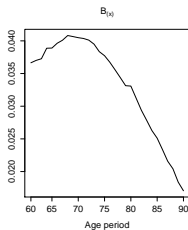
■ Optimizing l with the Newton-Raphson method

Common parameters plots

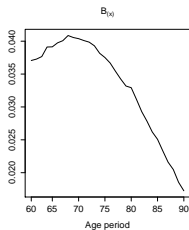
Model 0



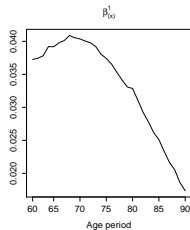
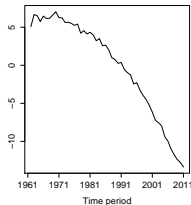
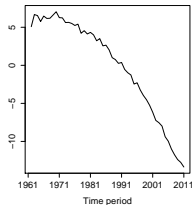
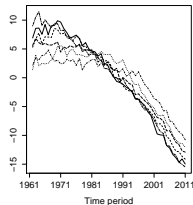
Model 1



Model 2

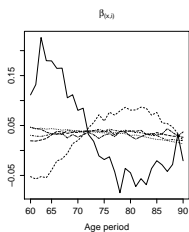


Model 3

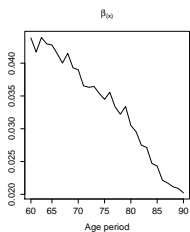
 $K_{(t)}$  $K_{(t)}$  $K_{(t)}$  $K_{(t,\beta)}^1$ 

Country specific parameters plots

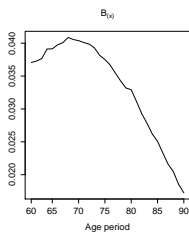
Model 0



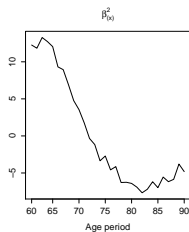
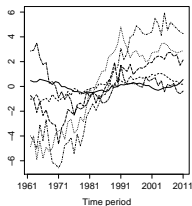
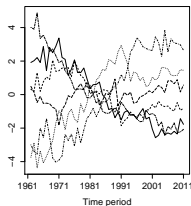
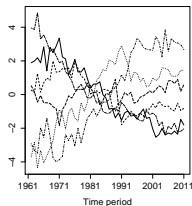
Model 1



Model 2



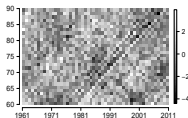
Model 3

 $\kappa_{(t,i)}$  $\kappa_{(t)}$  $\kappa_{(t,i)}$  $\kappa_{(t,i)}^2$

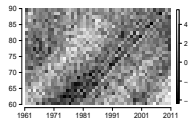
Time period

Standardized residuals plots

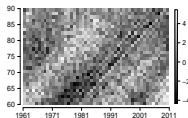
Model 0

**Austria**

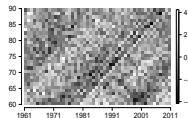
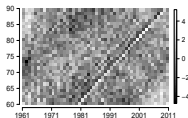
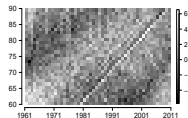
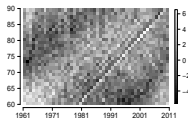
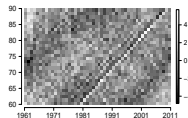
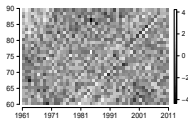
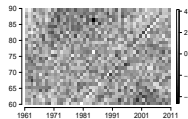
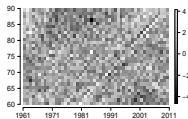
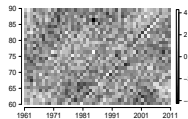
Model 1

**Austria**

Model 2

**Austria**

Model 3

**Austria****Belgium****Belgium****Belgium****Belgium****Sweden****Sweden****Sweden****Sweden**

Comparison of the models

■ Information criterion - ranking the models

$$\text{BIC value} = -2[\log -\text{likelihood value}] + \log(N)k^{\text{effective}}$$

	N	k	$k^{\text{effective}}$	log-likelihood	BIC	rank
Model 0	9000	740	726	-46716.43	86822.64	(2)
Model 1	9000	590	531	-48396.82	91958.89	(3)
Model 2	9000	560	502	-48477.21	92383.72	(4)
Model 3	9000	840	776	-46369.96	85674.45	(1)

■ Estimation time - iterations count

	Model 0	Model 1	Model 2	Model 3
Number of iterations	200+	10+	5+	40+

Forecasting joint mortality rates

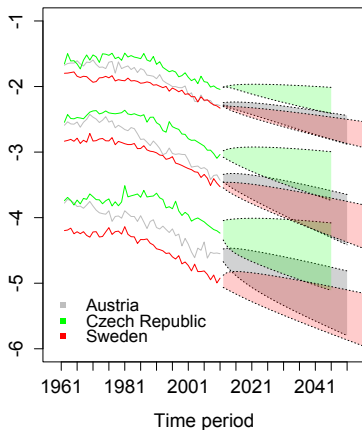
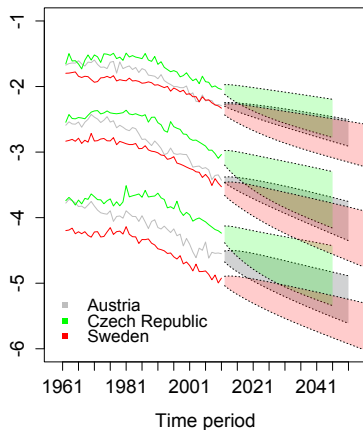
- Choosing time series process
 - Dependent on the parameters shape
 - Constraints over the parameters
 - In sample residuals correlation ratios
- Common parameters

Model 0	Model 1	Model 2	Model 3
Random walk with drift	Random walk with drift	Random walk with drift	Multi variate random walk with common drift
$K_{(t)} = d + K_{(t-1)} + \sigma Z_{(t)}$	$K_{(t)} = d + K_{(t-1)} + \sigma Z_{(t)}$	$K_{(t)} = d + K_{(t-1)} + \sigma Z_{(t)}$	$\kappa_{(t,i)}^1 = d_c + \kappa_{(t-1,i)}^1 + CZ_{(t)}$

- Country specific parameters

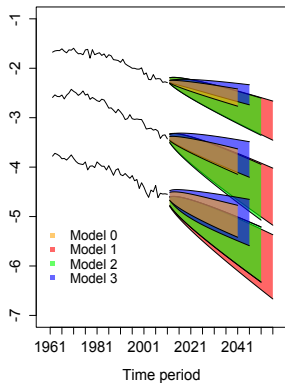
Model 0	Model 1	Model 2	Model 3
First order vector autoregression process	Multi variate random walk with drift	Multi variate random walk with drift	First order vector autoregression process (Φ is diagonal matrix)
$\kappa_{(t,i)} = \Phi \kappa_{(t-1,i)} + CZ_{(t)}$	$\kappa_{(t,i)} = d + \kappa_{(t-1,i)} + CZ_{(t)}$	$\kappa_{(t,i)} = d + \kappa_{(t-1,i)} + CZ_{(t)}$	$\kappa_{(t,i)}^2 = \Phi \kappa_{(t-1,i)}^2 + CZ_{(t)}$

Forecasting results

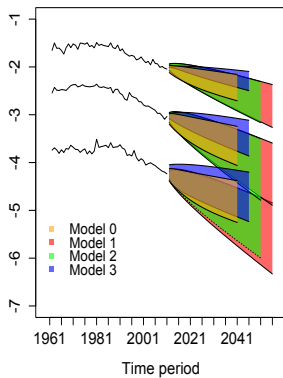
Model 3**Model 0**

Forecasting results

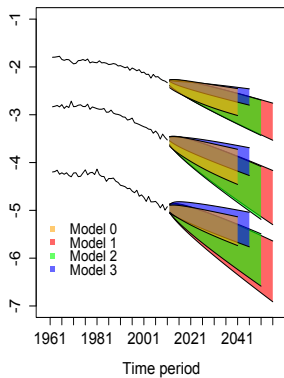
Austria



Czech Republic



Sweden



Conclusion

- Comparison of multi-population models
- Reduced parameter number
- Models that are estimated faster
- Generation of joint scenarios for future mortality rates