Is there a lost US generation?

# Results from a Bayesian-based period-cohort model of US male and female adult mortality

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#### **Quick summary**

- Advantages of using age and cohort as the primary variables of analysing & forecasting mortality data:
  - Cohort analysis applies to real people, with consequent advantages
  - Age-cohort fits adult mortality data just as well as age period (if not better)
  - Cohort effects appear stable across (adult) life
  - Key: period-related randomness has declined in developed countries,
     creating challenges for using it alone to forecast future mortality variation
- Disadvantages: estimation difficulties for recent cohorts
- Solution: use Bayesian maximum a posteriori estimation to jointly estimate cohort mortality parameters given a prior about how they change over time (that is, slowly)
- Analysis: Apply approach to a variant of the Cairns-Blake-Dowd (2006) model, estimate it on US M & F data
- Findings: Approach seems to produce sensible estimates & reasonable ranges for future cohort adult life expectancy
- Evidence of a significant reduction aggregate mortality improvement for those born in the 1950's, but not as bad as LC The University of Georgia TERRY COLLEGE OF BUSINESS

#### Some history

- First demographers looked at the effect of <u>age only</u> on mortality (e.g. Ulpian, ~150CE, Halley, 1693)
- Next, added period: age-period combinations are the data produced by mortality investigations (Lee & Carter, 1992; Cairns, Blake & Dowd, 2006)
- Next, looked at cohort data in the context of existing age-period models (Willetts, 2003, Haberman & Renshaw, 2006 etc)

#### BUT:

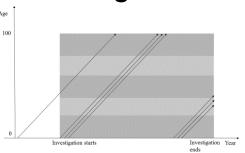
- Age, period & cohort are collinear
- Including all three effects difficult and contentious (across the social sciences, Yang et al (2008) w/ Luo (2013), Chauvel & Leist (2016) etc)
- Uncertainty in period-related fluctuations in mortality rates appears to be dying away (in rich countries, at least)
- Real people have fixed cohorts

#### SO:

– What about using age and cohort as the first two variables, with period as an afterthought?
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# An ad-hoc investigation on US data (I)

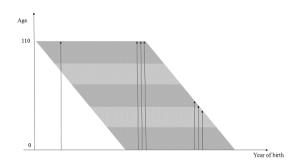
#### Age-period regressions



$$\log(q_{x,t}) = \alpha + \beta' I_x + \gamma' I_t + \varepsilon_{x,t}$$

Subsample	Males					
	Adj-R	Root mean				
	squared	square error				
Age 0-9	0.9961	0.08454				
Age 10-19	0.9410	0.18702				
Age 20-29	0.9660	0.05286				
Age 30-39	0.9856	0.04293				
Age 40-49	0.9897	0.04083				
Age 50-59	0.9932	0.03237				
Age 60-69	0.9948	0.02654				
Age 70-79	0.9944	0.02574				
Age 80-89	0.9853	0.03558				
Age 90-99	0.9557	0.03815				

#### Age-cohort regressions

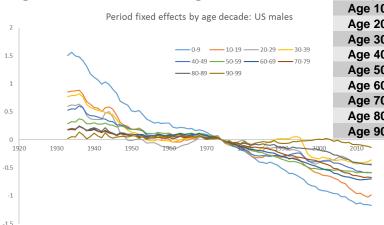


$$\log(q_{x,c}) = \alpha + \beta' I_x + \xi' I_c + \varepsilon_{x,c}$$

Subsample	Males						
	Adj-R	Root mean					
	squared	square error					
Age 0-9	0.9945	0.10044					
Age 10-19	0.9385	0.19097					
Age 20-29	0.9176	0.08231					
Age 30-39	0.9565	0.07461					
Age 40-49	0.9831	0.05227					
Age 50-59	0.9952*	0.02728*					
Age 60-69	0.9952*	0.02544*					
Age 70-79	0.9933	0.02837					
Age 80-89	0.9898*	0.02962*					
Age 90-99	0.9487	0.04103					

# An ad-hoc investigation on US data (II)

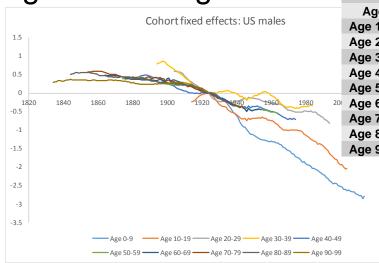
#### Age-period regressions



					Subsa	mple				
	Age									
	0-9	10-19	20-29	30-39	40-49	50-59	60-69	70-79	80-89	90-99
Age 0-9	1.000									
Age 10-19	0.656	1.000								
Age 20-29	0.554	0.644	1.000							
Age 30-39	0.537	0.629	0.826	1.000						
Age 40-49	0.529	0.588	0.704	0.770	1.000					
Age 50-59	0.454	0.481	0.646	0.693	0.752	1.000				
Age 60-69	0.348	0.461	0.474	0.539	0.660	0.760	1.000			
Age 70-79	0.288	0.408	0.351	0.440	0.612	0.697	0.906	1.000		
Age 80-89	0.227	0.309	0.267	0.349	0.532	0.606	0.834	0.922	1.000	
Age 90-99	0.242	0.308	0.243	0.310	0.499	0.598	0.787	0.877	0.958	1.000
2000										

$$corr(\gamma_t^i - \gamma_{t-1}^i, \gamma_t^j - \gamma_{t-1}^j) \quad [H_0: LC = 1]$$

#### Age-cohort regressions



		Age 0-9	Age	Age	Age	Age	Age	Age	Age	Age	Age
		Age 0-9	10-19	20-29	30-39	40-49	50-59	60-69	70-79	80-89	90-99
	Age 0-9	1.000									
	Age 10-19	0.122	1.000								
	Age 20-29	-0.174	0.155	1.000							
	Age 30-39	0.115	0.147	0.393	1.000						
	Age 40-49	0.169	0.530	0.338	0.345	1.000					
	Age 50-59	0.181	0.354	0.397	0.357	0.574	1.000				
201	Age 60-69	0.092	0.391	0.460	0.389	0.508	0.684	1.000			
	Age 70-79	-0.269	0.019	0.193	0.083	0.384	0.536	0.703	1.000		
	Age 80-89	-0.379	0.164	0.093	-0.226	0.107	0.539	0.498	0.513	1.000	
	Age 90-99	-	0.278	0.268	-0.014	-0.293	0.178	0.271	0.281	0.449	1.000
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Subsample

$$corr(\xi_t^i - \xi_{t-1}^i, \xi_t^j - \xi_{t-1}^j) \quad [H_0 : LC^C = 1]$$

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#### Is variance in period innovations declining?

- Evidence that the variance of random innovations in period effects are declining comes from three sources
  - Ad-hoc investigation. Variance ratio tests of the equality of the variance of differenced period coefficients in first and second halves of sample rejected in all subsamples of the data except one (variance of differenced cohort effects, if anything, may be increasing)

		Cohort	effects	Period effects				
	Statistic	DF1	DF2	p-value	Statistic	DF1	DF2	p-value
Age 0-9	-	-	-	-	1.8086	39	39	0.034
Age 10-19	0.1290	8	79	0.002	2.2458	39	39	0.007
Age 20-29	0.6875	18	69	0.188	2.8957	39	39	0.001
Age 30-39	0.9978	28	59	0.513	1.7137	39	39	0.048
Age 40-49	0.3621	38	49	0.001	1.3869	39	39	0.156
Age 50-59	0.3962	48	39	0.001	2.8624	39	39	0.001
Age 60-69	0.7900	58	29	0.220	2.4701	39	39	0.003
Age 70-79	0.9873	68	19	0.459	2.7500	39	39	0.001
Age 80-89	0.7194	78	9	0.207	2.1874	39	39	0.008
Age 90-99	-	-	-	-	2.0646	39	39	0.013

- 2. Estimates of Lee-Carter model clearly show the decline in variation in time parameter (e.g. McCarthy & Miles 2014, \*\*\* 2017)
- 3. McCarthy and Wang (2017), presented in this conference, fit this model to ~30 countries, variance ratio tests within countries and  $\chi^2$  tests across them rejected
- Conclusion: cannot rely on variance in period effects to generate *all* mortality uncertainty

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#### So what to do?

- Two main options:
  - Cohort version of Lee-Carter (1992):

$$\log(m_{x,c}) = \alpha_x + \beta_x k_c + \delta' I_{c+x} + \varepsilon_{x,c}$$

- (we're working on it)
- Cohort version of Cairns-Blake-Dowd (2006):

$$\log(m_{x,c}) = \alpha' I_c + \beta' I_c \frac{(x-k_1)}{k_2} + \gamma' I_c (\frac{x-k_1}{k_2})^2 + \delta' I_{c+x} + \varepsilon_{x,c}$$

- (this paper)
- Difficulty:
  - How to fit model to recent cohorts where not much data is available?
  - We use Bayesian maximum a posteriori estimation (de Groot, 1970)



#### **Model description**

Modified version of CBD (2006)

$$\log(m_{x,c}) = \alpha' I_c + \beta' I_c \frac{(x-k_1)}{k_2} + \gamma' I_c (\frac{x-k_1}{k_2})^2 + \delta' I_{c+x} + \varepsilon_{x,c}$$

- $-\alpha$ ,  $\beta$ ,  $\gamma$  constant for each *cohort* (this is the identifying assumption aka Fienberg and Mason (1978))
- Model log of central rate of mortality (estimated log hazard rate) to enable easy conversion to survival distribution
- Add a quadratic term (only needed for US females)
- Add period random effects to capture non-linear period-related mortality shocks  $\delta$ :  $\delta_{c+x} \sim N(0, \sigma_{\delta}^2)$  (remember: (c+x=t); linear changes in mortality captured by cohort parameters)
- Make an appropriate assumption about error term, theory suggests for perfect model:

$$\varepsilon_{x,c} \sim N(0,\omega_{x,c}), \quad \omega_{x,c} = \frac{1}{m_{x,c}^* E_{x,c}} \cong \frac{1}{D_{x,c}}$$

for our imperfect model we assume:

$$\varepsilon_{x,c} \sim N(0, \sigma_{\varepsilon}^2 \omega_{x,c})$$



# Bayesian maximum a posteriori estimation

- Step 1
  - Estimate a prior distribution for the parameters  $\alpha$ ,  $\beta$  [,  $\gamma$ ]
- Step 2
  - Calculate the posterior distribution of these parameters, conditional on the data and the chosen prior
- Step 3
  - For point estimates, choose the mode of the posterior distribution
  - For actual posterior distribution, can do Metropolis-Hastings, we use an approximation due to Laplace
- Step 4
  - Point estimates for calculating fitted values, estimating random effects  $\delta$ , measuring goodness-of-fit
  - Post. dstbn & VAR for forecasting

#### Step 1: choosing the prior

• Assume that first differences of  $\theta_c = (\alpha, \beta, \gamma)_c$  follow a first-order VAR, with exogenous variables (also differenced), x to capture known causes of mortality variation (e.g. smoking)

$$(\Delta \theta)_c - \hat{\mu} - \hat{\Delta}_1 (\Delta \theta)_{c-1} - \hat{\kappa}_0 (\Delta x)_c = v_c, \quad v_c \sim N(0, \Sigma) \text{ iid}$$

Hence, the prior distribution is:

$$p_{1}(\theta \mid \xi \equiv \hat{\mu}, \hat{\Delta}_{1}, \hat{\Sigma}, \hat{\kappa}) = \prod_{all \ c} \phi((\Delta \theta)_{c} - \hat{\mu} - \hat{\Delta}_{1}(\Delta \theta)_{c-1} - \hat{\kappa}_{0}(\Delta x)_{c}, 0, \hat{\Sigma})$$

- Values of fitted parameters estimated off VAR fitted to cohort-by-cohort estimates of  $\theta_c = (\alpha, \beta, \gamma)_c$  obtained from regressions on reasonably complete cohorts, fitted using standard ML
- Prior then reflects how we believe the value of  $\theta_c = (\alpha, \beta, \gamma)_c$  changes from cohort to cohort



# Step 2: obtaining the posterior

Bayes' Theorem gives the posterior distribution as

$$p_{2}(\theta,(\sigma_{\delta}^{2},\sigma_{\varepsilon}^{2})|\xi,\{m_{x,c},\omega_{x,c}\}) = \frac{\ell(\{m_{x,c},\omega_{x,c}\}|\theta,(\sigma_{\delta}^{2},\sigma_{\varepsilon}^{2}))p_{1}(\theta|\xi)}{\int \ell(\{m_{x,c},\omega_{x,c}\}|\theta,(\sigma_{\delta}^{2},\sigma_{\varepsilon}^{2}))p_{1}(\theta|\xi)d\theta}$$

or:

$$\begin{split} \ell p_2(\theta, (\sigma_\delta^2, \sigma_\varepsilon^2) \,|\, \xi, \{m_{x,c}, \omega_{x,c}\}) &= K + \underbrace{\ell \ell(\{m_{x,c}, \omega_{x,c}\} \,|\, \theta, (\sigma_\delta^2, \sigma_\varepsilon^2))}_{\text{Fisherian log-likelihood of data } \{m_{x,c}, \omega_{x,c}\}} \\ &+ \underbrace{\ell p_1(\theta \,|\, \xi)}_{\text{logged prior distribution}} \end{split}$$

(Hence, BMAP estimator is a penalized ML estimator)

#### Step 3: selecting estimates

For point estimates choose the mode of the posterior dstbn:

$$(\hat{\theta}, (\hat{\sigma}_{\delta}^{2}, \hat{\sigma}_{\varepsilon}^{2})) = \underset{\theta, (\sigma_{\delta}^{2}, \sigma_{\varepsilon}^{2})}{\arg \max} \ell p_{2}(\theta, (\sigma_{\delta}^{2}, \sigma_{\varepsilon}^{2}) | \xi, \{m_{x,c}, \omega_{x,c}\})$$

 To approximate the posterior distribution itself, use a result from Laplace:

$$p_{2}(\theta) \approx K(\hat{\theta}) \exp \left[ \frac{1}{2} (\theta - \hat{\theta})' \ell p_{2,\theta\theta}(\hat{\theta}) (\theta - \hat{\theta}) \right]$$

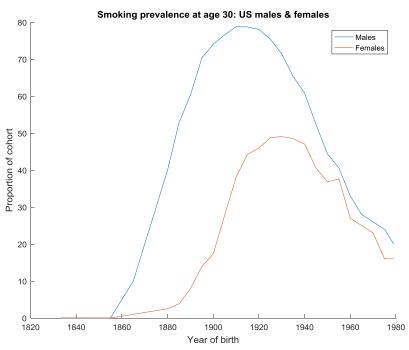
[Note: posterior is approximately MVN (property shared by all modal estimators, including ML)]

(We are also using the Metropolis-Hastings algorithm to test accuracy for marginals)

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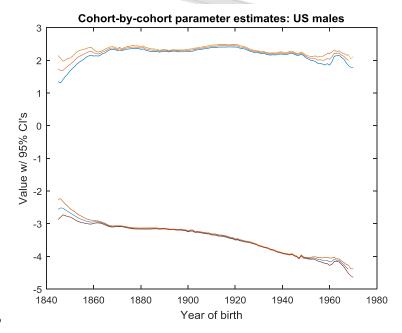
#### **Application**

- US male and female mortality data, 1933 2014, from <u>www.mortality.org</u>, focus on adult mortality (age 35-100)
- For exogenous variable, use cohort smoking prevalence data collected from Forey et al (1997), updated by our own estimates from US survey data (NHIS)
  - Use proportion of the cohort that smoked (cigarettes) at age 30 (usu. ~peak smoking)
  - Pattern similar across cohorts
  - Peaked for men born ~1910
    - (Later for women)
  - Rapid decline
    - (Slower for women)
  - M & F proportion now similar



# Step 1: choosing the prior

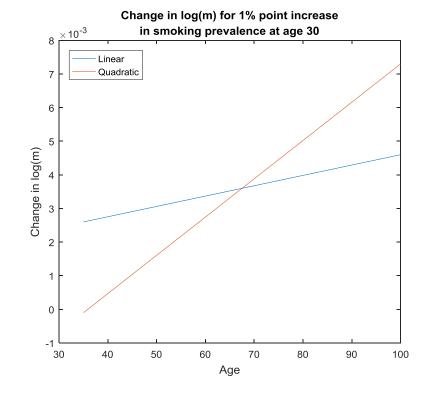
- Use standard ML to fit cohort-bycohort estimates to α, β [, γ]
- Results shown for US males (earliest and latest cohorts excluded)
- Cohorts born after 1870 and before 1942 are reasonably precise
- Fit a VAR to first differences of these estimates only
- Include first differences of exogenous variables, but estimate smoking effect using a SU-VAR on males and female cohorts jointly



#### Step 1: choosing the prior

Quadratic   Linear   Females   Males   Males   D.Alpha:   D.Alpha.L1   0.2779   0.3863   (0.0670)   (0.0763)   Change in smoking prevalence   (0.0007)   (0.0007)   (0.0007)   (0.0007)   (0.0026)   (0.0026)   (0.0023)   RMSE   0.0196   0.0169   R-squared   -0.2632   -0.1593		VAR estim	ates	
D.Alpha:         D.Alpha.L1         0.2779         0.3863           Change in smoking prevalence         0.0036         0.0036           (0.0007)         (0.0007)         (0.0007)           Constant         -0.0127         -0.0095           (0.0026)         (0.0023)           RMSE         0.0196         0.0169           R-squared         -0.2632         -0.1593           D.Beta:         D.Beta.L1         0.2713         0.3398           (0.0811)         (0.0870)           Change in smoking prevalence         (0.0011)         (0.00870)           Constant         -0.0016         -0.0024           (0.0021)         (0.0020)           RMSE         0.0169         0.0163           R-squared         0.2736         0.1161           D.Gamma:         D.Gamma.L1         0.2870           (0.0626)         Change in smoking prevalence         0.0000		77		Linear
Change in smoking prevalence (0.0007) (0.0007)  Constant -0.0127 -0.0095 (0.0026) (0.0023)  RMSE 0.0196 0.0169  R-squared -0.2632 -0.1593  D.Beta: D.Beta.L1 0.2713 0.3398 (0.0811) (0.0870)  Change in smoking prevalence (0.0011) (0.0010)  Constant -0.0016 -0.0024 (0.0021) (0.0020)  RMSE 0.0169 0.0163  R-squared 0.2736 0.1161  D.Gamma: D.Gamma.L1 0.2870 (0.0626)  Change in smoking prevalence (0.0020)			Females	Males
Change in smoking prevalence (0.0007) (0.0007)  Constant -0.0127 -0.0095 (0.0023)  RMSE 0.0196 0.0169  R-squared -0.2632 -0.1593  D.Beta: D.Beta.L1 0.2713 0.3398 (0.0811) (0.0870)  Change in smoking prevalence (0.0011) (0.0010)  Constant -0.0016 -0.0024 (0.0021) (0.0020)  RMSE 0.0169 0.0163  R-squared 0.2736 0.1161  D.Gamma: D.Gamma.L1 0.2870 (0.0626)  Change in smoking prevalence (0.0020)	D.Alpha:	D.Alpha.L1	0.2779	0.3863
Smoking prevalence			(0.0670)	(0.0763)
Constant -0.0127 -0.0095		smoking		
RMSE 0.0196 0.0169 R-squared -0.2632 -0.1593  D.Beta: D.Beta.L1 0.2713 0.3398				, ,
RMSE 0.0196 0.0169 R-squared -0.2632 -0.1593  D.Beta: D.Beta.L1 0.2713 0.3398 (0.0811) (0.0870) Change in smoking 0.0037 0.0010 prevalence (0.0011) (0.0010) Constant -0.0016 -0.0024 (0.0021) (0.0020) RMSE 0.0169 0.0163 R-squared 0.2736 0.1161  D.Gamma: D.Gamma.L1 0.2870 (0.0626) Change in smoking prevalence (0.0020)		Constant		
D.Beta: D.Beta.L1 0.2713 0.3398 (0.0811) (0.0870)  Change in smoking 0.0037 0.0010 prevalence (0.0011) (0.0010)  Constant -0.0016 -0.0024 (0.0021) (0.0020)  RMSE 0.0169 0.0163 R-squared 0.2736 0.1161  D.Gamma: D.Gamma.L1 0.2870 (0.0626)  Change in smoking prevalence (0.0020)			,	, ,
D.Beta: D.Beta.L1 0.2713 0.3398 (0.0811) (0.0870)  Change in smoking 0.0037 0.0010 prevalence (0.0011) (0.0010)  Constant -0.0016 -0.0024 (0.0021) (0.0020)  RMSE 0.0169 0.0163 R-squared 0.2736 0.1161  D.Gamma: D.Gamma.L1 0.2870 (0.0626)  Change in smoking prevalence (0.0020)				
Change in smoking prevalence (0.0811) (0.0870)  Change in smoking prevalence (0.0011) (0.0010)  Constant -0.0016 -0.0024 (0.0021) (0.0020)  RMSE 0.0169 0.0163  R-squared 0.2736 0.1161  D.Gamma: D.Gamma.L1 0.2870 (0.0626)  Change in smoking prevalence (0.0020)		R-squared	-0.2632	-0.1593
Change in smoking prevalence (0.0811) (0.0870)  Change in smoking prevalence (0.0011) (0.0010)  Constant -0.0016 -0.0024 (0.0021) (0.0020)  RMSE 0.0169 0.0163  R-squared 0.2736 0.1161  D.Gamma: D.Gamma.L1 0.2870 (0.0626)  Change in smoking prevalence (0.0020)				
Change in smoking prevalence (0.0811) (0.0870)  Change in smoking prevalence (0.0011) (0.0010)  Constant -0.0016 -0.0024 (0.0021) (0.0020)  RMSE 0.0169 0.0163  R-squared 0.2736 0.1161  D.Gamma: D.Gamma.L1 0.2870 (0.0626)  Change in smoking prevalence (0.0020)				
Change in smoking prevalence (0.0011) (0.0010)  Constant -0.0016 -0.0024 (0.0021) (0.0020)  RMSE 0.0169 0.0163  R-squared 0.2736 0.1161  D.Gamma: D.Gamma.L1 0.2870 (0.0626)  Change in smoking prevalence (0.0020)	D.Beta:	D.Beta.L1	0	
Smoking prevalence   0.0037   0.0010     (0.0011)   (0.0010)     (0.0011)   (0.0024   (0.0021)   (0.0020)     (0.0021)   (0.0020)     (0.0026)     (0.0026)     (0.0626)     (0.0626)     (0.0626)     (0.0020)       (0.0020)     (0.0020)     (0.0020)     (0.0020)     (0.0020)       (0.0020)     (0.0020)       (0.0020)       (0.0020)       (0.0020)			(0.0811)	(0.0870)
Constant -0.0016 -0.0024 (0.0021) (0.0020)  RMSE 0.0169 0.0163  R-squared 0.2736 0.1161  D.Gamma: D.Gamma.L1 0.2870 (0.0626)  Change in smoking prevalence (0.0020)		smoking	0.0037	0.0010
(0.0021) (0.0020)   RMSE		·	(0.0011)	(0.0010)
RMSE 0.0169 0.0163 R-squared 0.2736 0.1161  D.Gamma: D.Gamma.L1 0.2870 (0.0626) Change in smoking 0.0000 prevalence (0.0020)		Constant	-0.0016	-0.0024
R-squared 0.2736 0.1161  D.Gamma: D.Gamma.L1 0.2870 (0.0626)  Change in smoking prevalence (0.0020)			(0.0021)	(0.0020)
D.Gamma: D.Gamma.L1 0.2870 (0.0626)  Change in smoking prevalence (0.0020)		RMSE	0.0169	0.0163
(0.0626)  Change in smoking 0.0000 prevalence (0.0020)		R-squared	0.2736	0.1161
(0.0626)  Change in smoking 0.0000 prevalence (0.0020)				
(0.0626)  Change in smoking 0.0000 prevalence (0.0020)				
Change in smoking 0.0000 prevalence (0.0020)	D.Gamma:	D.Gamma.L1		
smoking 0.0000 prevalence (0.0020)			(0.0626)	
,		smoking	0.0000	
		·	(0.0020)	
Constant -0.0038		Constant	-0.0038	
(0.0043)				
RMSE 0.0354		RMSE	0.0354	
R-squared 0.0636		R-squared	0.0636	

- Own first-order lags significant
- Steady-state: decline in alpha, beta and gamma
- Increase in smoking prevalence increases cohort mortality at all ages, but more at older ages



# Step 2: obtaining the posterior

• Given the estimated parameters of the VAR,  $\xi$ , the logged posterior pdf is easily calculated for *any* choice of parameters  $\{\theta, (\sigma_{\delta}^2, \sigma_{\varepsilon}^2)\}$ , from Bayes' Theorem:

$$\ell p_{2}(\theta, (\sigma_{\delta}^{2}, \sigma_{\varepsilon}^{2}) \mid \xi, \{m_{x,c}, \omega_{x,c}\}) = K + \ell \ell (\{m_{x,c}, \omega_{x,c}\} \mid \theta, (\sigma_{\delta}^{2}, \sigma_{\varepsilon}^{2}))$$
Fisherian log-likelihood of data  $\{m_{x,c}, \omega_{x,c}\}$ 

$$+ \ell p_{1}(\theta \mid \xi)$$
logged prior distribution

 The log likelihood function of the data is penalized by the addition of the prior information

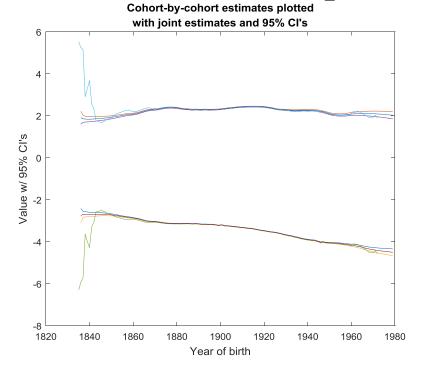
#### Step 3: obtaining the estimates

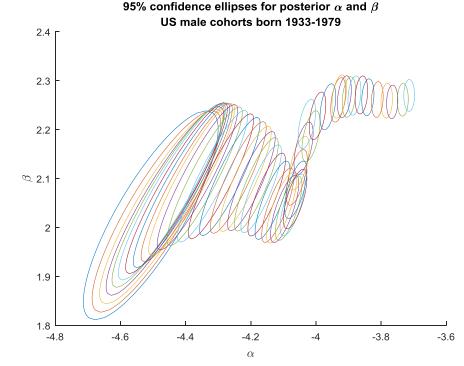
• Use numerical software to obtain point estimates for  $\{\theta, (\sigma_{\delta}^2, \sigma_{\varepsilon}^2)\}$  by solving:

$$(\hat{\theta}, (\hat{\sigma}_{\delta}^{2}, \hat{\sigma}_{\varepsilon}^{2})) = \underset{\theta, (\sigma_{\delta}^{2}, \sigma_{\varepsilon}^{2})}{\arg \max} \ell p_{2}(\theta, (\sigma_{\delta}^{2}, \sigma_{\varepsilon}^{2}) | \xi, \{m_{x,c}, \omega_{x,c}\})$$

Obtain confidence intervals & ellipses using:

$$p_2(\theta) \approx K(\hat{\theta}) \exp\left[\frac{1}{2}(\theta - \hat{\theta})' \ell p_{2,\theta\theta}(\hat{\theta})(\theta - \hat{\theta})\right]$$

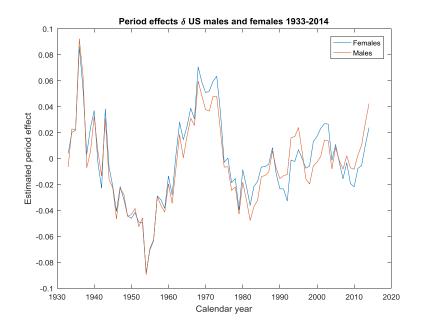




#### Step 4A: goodness of fit

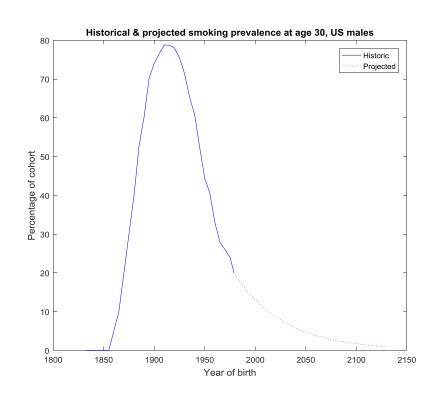
- Model fits historical data extremely well
- Using adjusted R-squared as a model selection criterion suggests linear model for males & quadratic model for females, as does a check of the residuals (not shown)
- Estimated period effects highly correlated across M & F

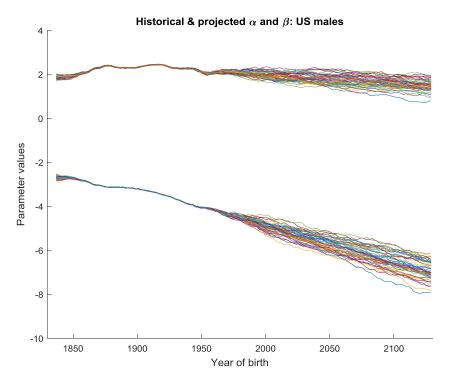
	Males						
Alpha (147 cohorts)	X	Х	Х	X			
Beta (147 cohorts)	X	X	X	X			
Gamma (147 cohorts)	X	Χ	-	-			
Random period effects	X	-	X	-			
log(sigma²(epsilon))	0.4947	0.5977	0.5219	0.6663			
	(0.0199)	(0.0177)	(0.0197)	(0.0188)			
log(sigma²(delta))	-3.7114	-	-3.5768	-			
	(0.1957)		(0.1906)				
R-squared	0.9989	0.9984	0.9989	0.9985			
Adjusted R-squared	0.9988	0.9983	0.9989	0.9984			
N	5412	5412	5412	5412			
	Females						
Alpha (147 cohorts)	X	X	X	X			
Beta (147 cohorts)	X	X	X	X			
Gamma (147 cohorts)	X	X	-	-			
Random period effects	X	-	X	-			
log(sigma²(epsilon))	0.4500	0.5881	0.6137	1.3642			
	(0.0197)	(0.0174)	(0.0196)	(0.0378)			
log(sigma²(delta))	-3.5289	-	-1.8582	-			
	(0.1920)		(0.1759)				
R-squared	0.9993	0.9989	0.9918	0.9932			
Adjusted R-squared	0.9992	0.9988	0.9913	0.9928			
N	5412	5412	5412	5412			





- Project what will happen to smoking rates (3% p.b.y. decline seems to fit well)
- Take random draws from posterior distribution of fitted parameters to get starting point
- Use fitted VAR to project these forward (including shocks & projected changes in smoking behaviour)



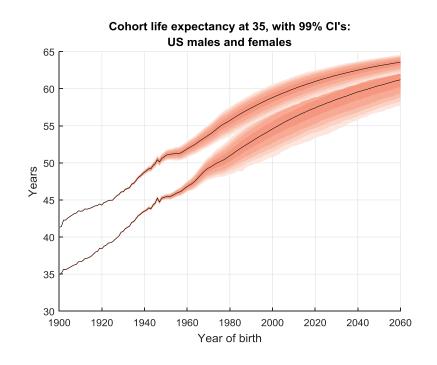


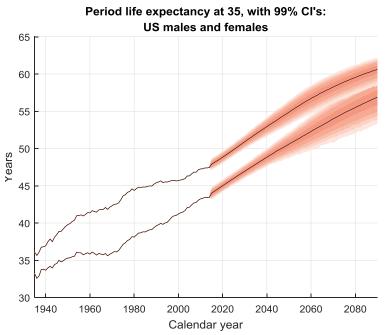
• Use projected values of  $\alpha$ ,  $\beta$ ,  $\gamma$ , fitted values of  $\{\log(\sigma_{\delta}^2), \log(\sigma_{\varepsilon}^2)\}$  to project future mortality hazard rates using:

$$\log(m_{x,c}) = \alpha' I_c + \beta' I_c \frac{(x-k_1)}{k_2} + \gamma' I_c (\frac{x-k_1}{k_2})^2 + \delta' I_{c+x} + \varepsilon_{x,c}$$

- Use these to estimate future life expectancy (or other quantities of interest, e.g. pension fund, annuity, SS liabilities)
- Two options
  - Do lots of Monte-Carlo runs to obtain Cl's for variables of interest
  - Use (known) distribution of  $\alpha$ ,  $\beta$ ,  $\gamma$  & known VAR to obtain computationally quicker but approximate theoretical distributions using survival distribution theory
    - Generate distributions of 'pseudo-parameters', use these & known Bayesian results to generate approximate posterior

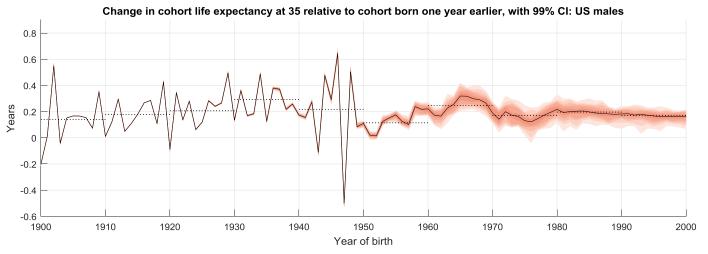
- Use forecasts of mortality rates to generate
  - Period life expectancy at 35 and 65, by calendar year
  - Cohort life expectancy at 35 and 65, by year of birth

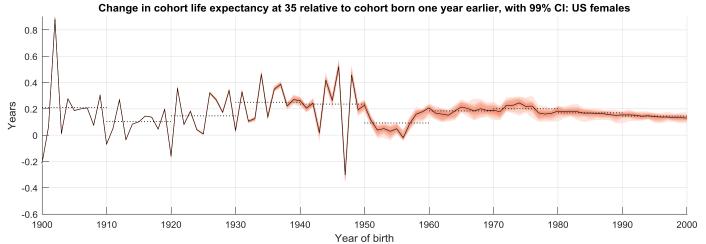






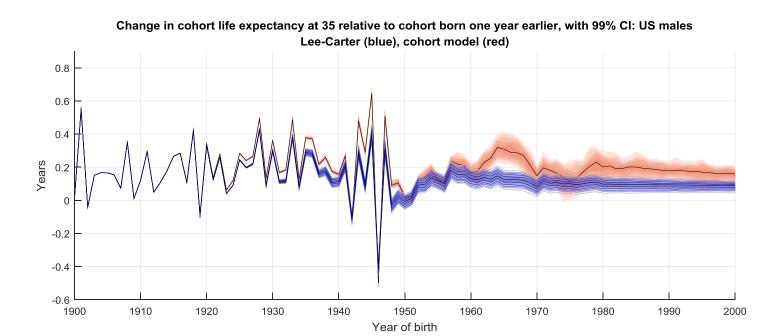
 Decadal cohort born 1950-1960 has lowest average rate of improvement in life expectancy at 35 of any such cohort in our dataset







- Fitted Lee-Carter model with parameter uncertainty to same data using ML (shown for males)
- Cohort-based model predicts:
  - Greater increases in life expectancy of cohorts starting in 1925 (largely due to projected decline in smoking rates)
  - Greater improvement in LE of those born in 1960-1970 (cohort effect)
  - Models converge from ~2040 (not shown)



#### Conclusion

- Proposed use of Bayesian maximum a posteriori estimation to jointly estimate cohort mortality parameters given a prior about how they change over time (that is, slowly)
- Applied approach to a variant of the Cairns-Blake-Dowd (2006) model, with period effects, estimated it on US M & F data
- Findings: Approach seems to produce sensible estimates & reasonable ranges for future cohort adult life expectancy
- Evidence of a significant reduction aggregate mortality improvement for those born in the 1950's
  - But projected mortality improvements appear to be better than the LC model would predict
  - Largely due to projected effect of reduced smoking rates on cohort mortality
  - (Models converge in differences after around 2040)
- Future work
  - Use this model to investigate international mortality patterns (w/ Wang)
  - Estimate cohort-based LC using BMAP

