

New Directions  
in the Modelling of Longevity Risk

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## Plan

- Motivation
- Genealogy
- New directions in modelling
- Numerical illustrations – single population models
- Remarks on multiple populations

## Motivation

- Application focus:
  - risk measurement and management of longevity risk
  - multiple populations
  - life insurance diversification benefits
  - basis risk in standardised longevity contracts
- industry requires robust models


## Development of New Models

- Many new stochastic mortality models since Lee-Carter
- Are they fit for purpose?
- Are they robust?

# GENEALOGY: 1st GENERATION MODELS

Eilers/Marx  
P-splines

Currie/Richards (M4)  
2-D P-splines  
2002, ...



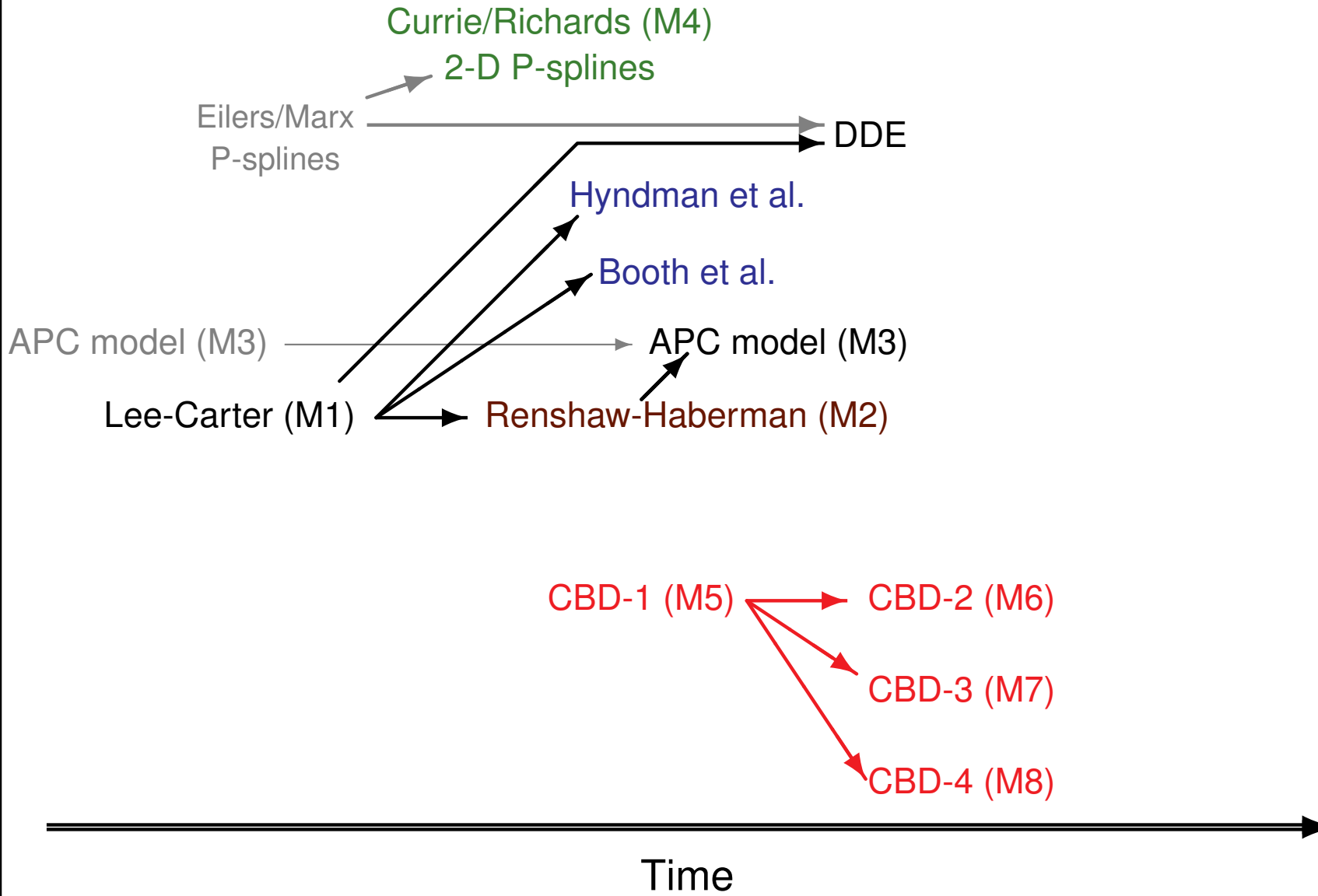
Lee-Carter (M1)  
1992

CBD-1 (M5)  
2006

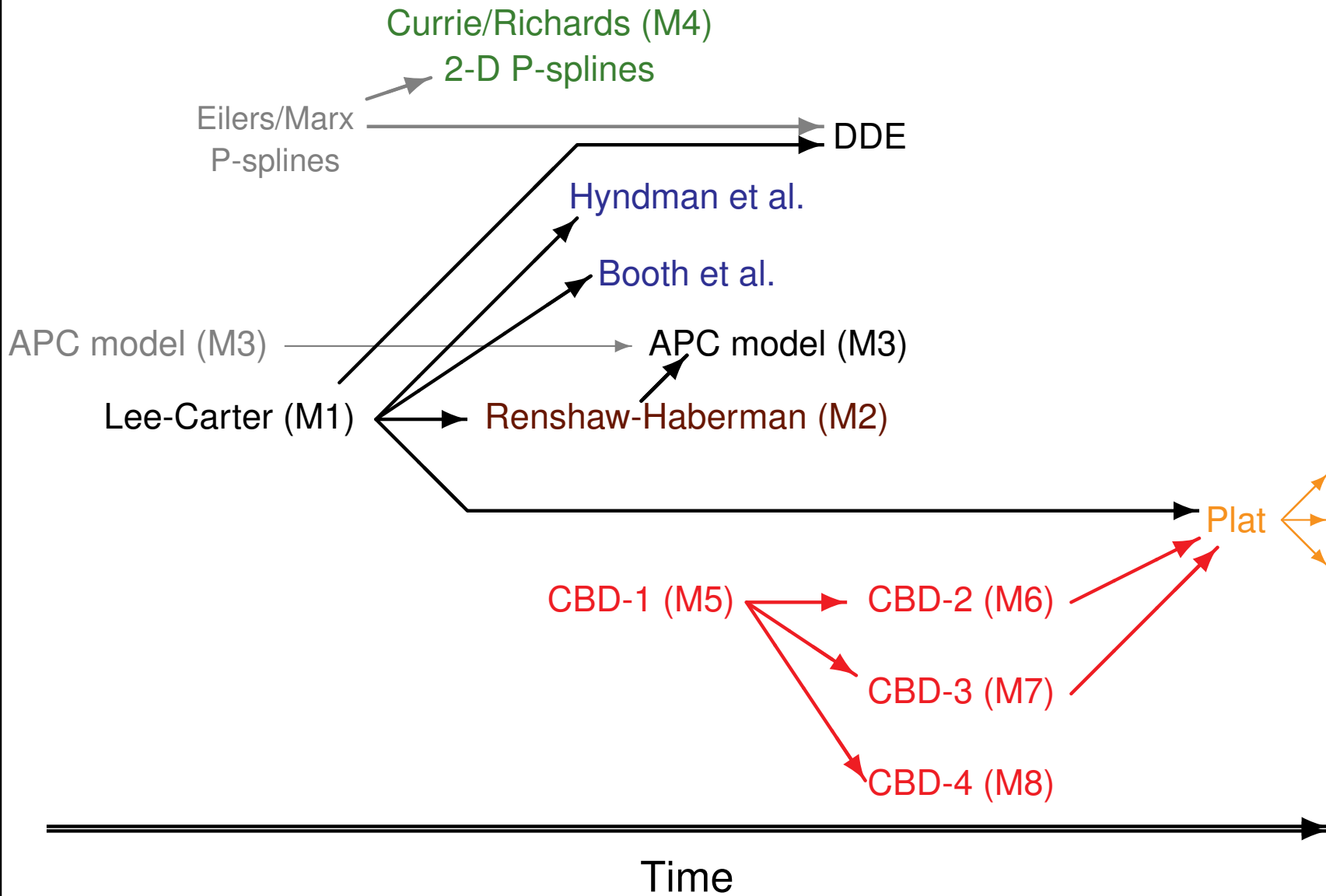


Time

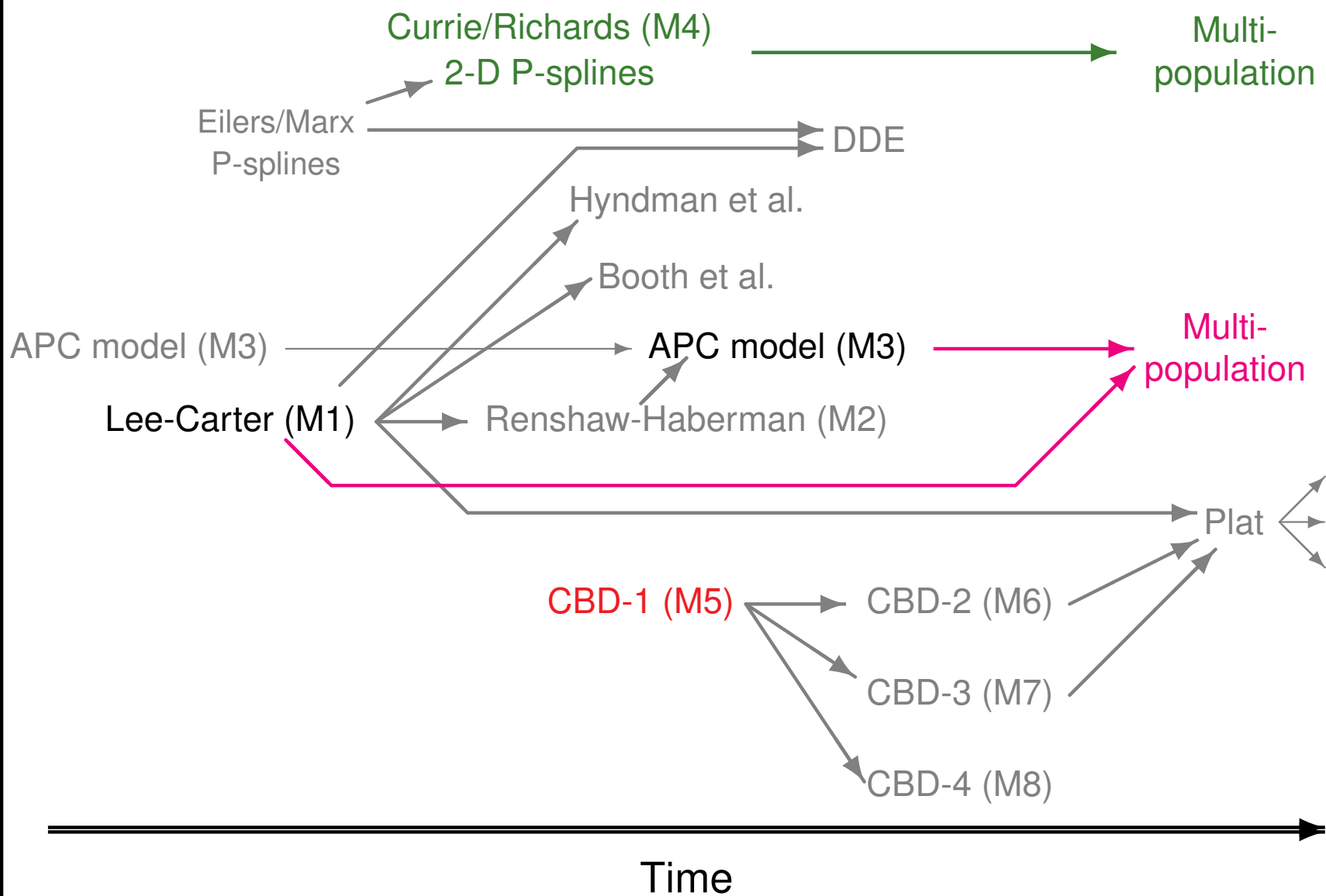
# Improvements + more complexity



# More improvements + even more complexity



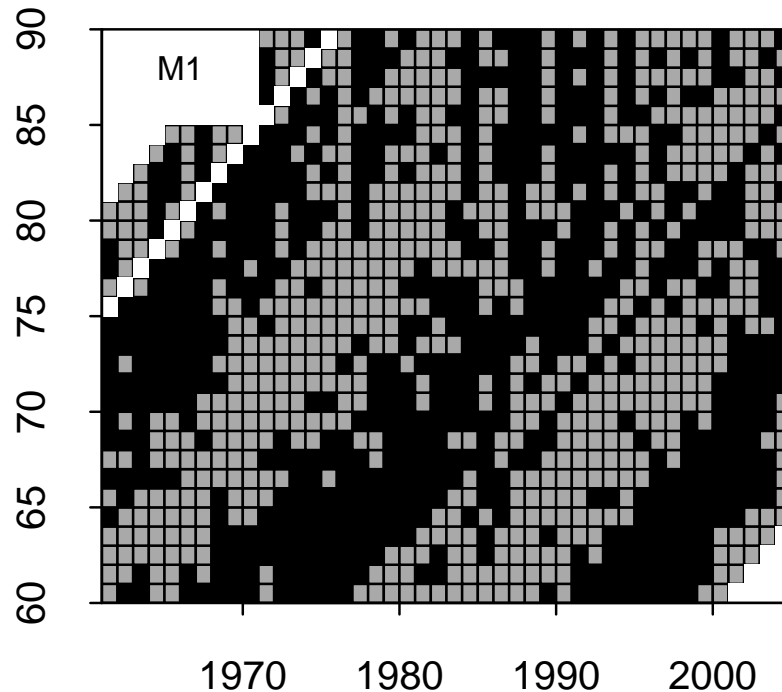
# Multiple population modelling



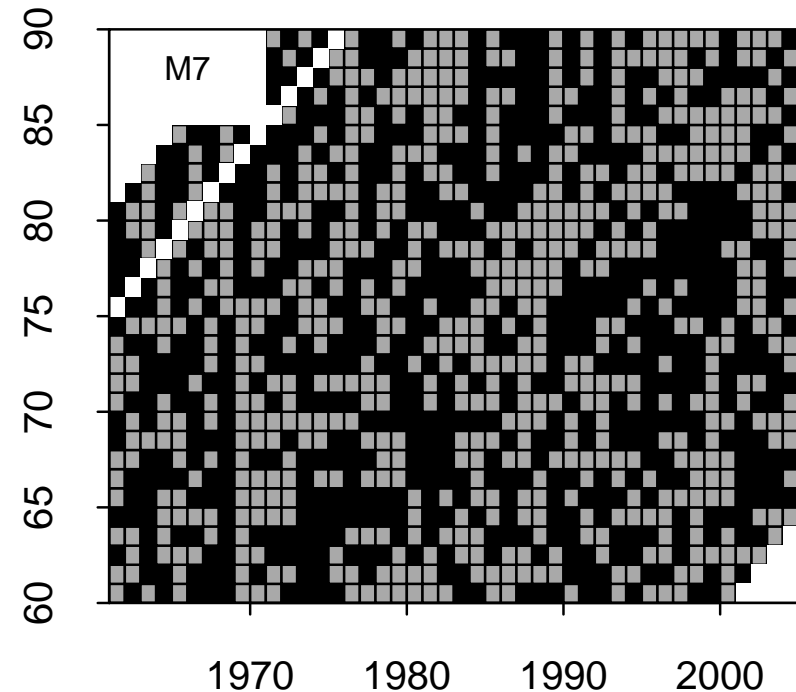


# Why do we need complexity?

Lee-Carter Model



CBD Model + Cohort Effect



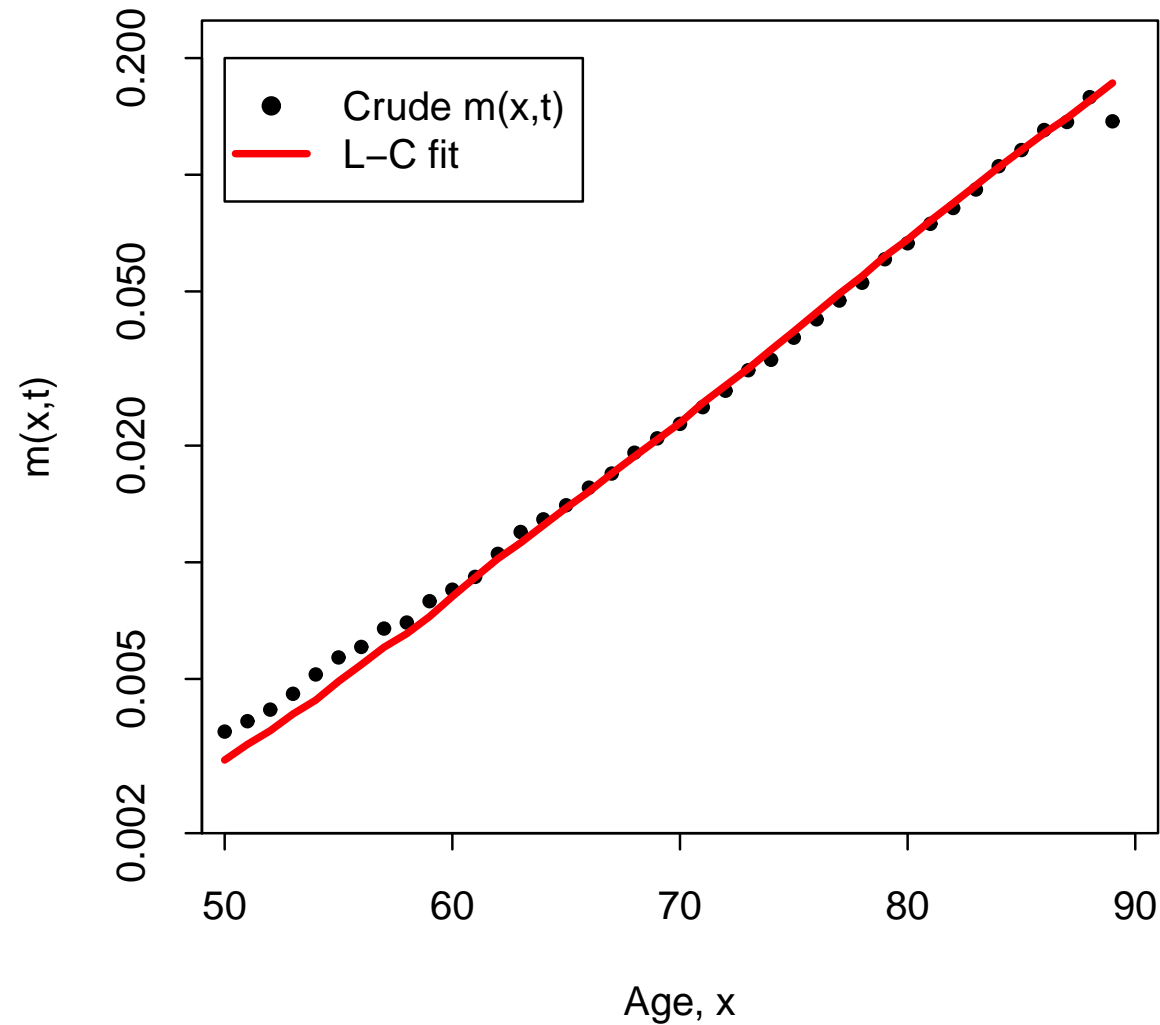
Black  $\Rightarrow$  model *over*-estimates  $m(x, t)$  death rate

Gray  $\Rightarrow$  model *under*-estimates  $m(x, t)$  death rate

LC: non-random clusters + errors are too big

# Need for complexity: accurate base table for forecasts

EW males 1971–2008: Lee–Carter Fit  
 $m(x,2008)$  (log scale)



## Issues on complexity

- More complex  $\Rightarrow$  More random processes
- More random processes  $\Rightarrow$   
MUCH more difficult to model **multiple populations**
- Excessive complexity  $\Rightarrow$   
**potentially less robust forecasts**

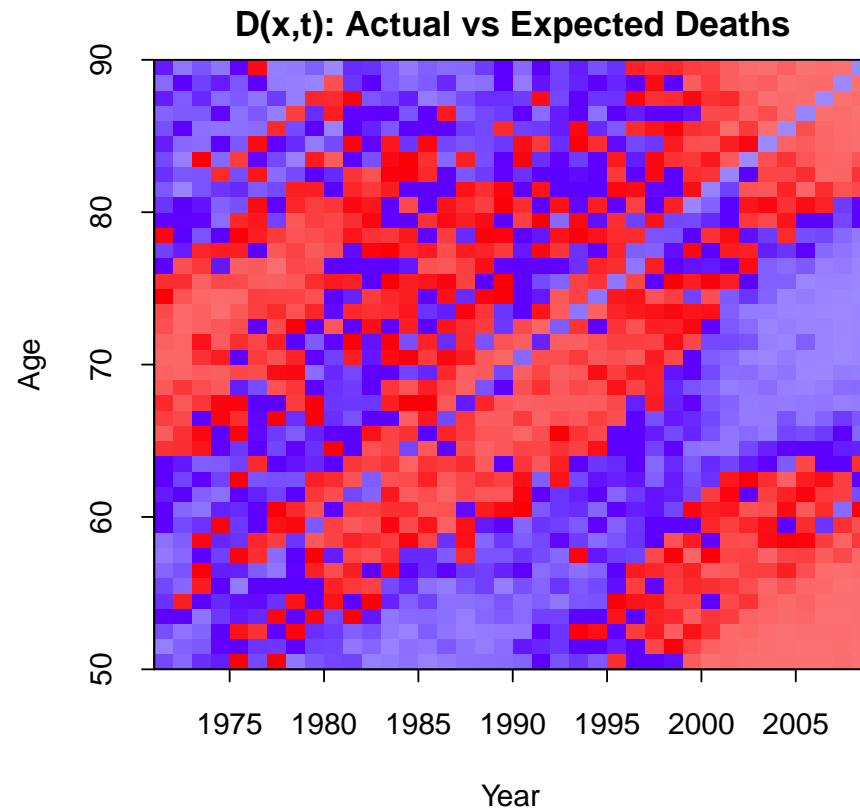
## A Possible Way Forward

### Single-population models

- Focus on **small number of key drivers**  
⇒ much easier to extend to multi-populations
- Focus on greater robustness of forecasts

## Case Study: CBD/Plat Revisited

$$\log m(x, t) = \beta(x) + \kappa_1(t) + \kappa_2(t)(x - \bar{x})$$



Red  $\Rightarrow$  actual deaths  $>$  expected deaths

## CBD/Plat Revisited: Key Idea: Possible responses

$$\log m(x, t) = \beta(x) + \kappa_1(t) + \kappa_2(t)(x - \bar{x})$$

Add:

- Cohort effect,  $\gamma(t - x)$
- Extra age-period effects
- Do something new .....

## Key Idea: CBD/Plat Revisited

Underlying  $\log m(x, t) =$

- $\beta(x) + \kappa_1(t) + \kappa_2(t)(x - \bar{x})$ : **two key drivers**

PLUS

$R(x, t)$  *Residuals*

- Assume: vector  $R(t) \rightarrow R(t + 1)$  mean reverting process

$\Rightarrow$  long term risk depends on **two key drivers**

## Specific Model

$$\log m(x, t) = \beta(x) + \kappa_1(t) + \kappa_2(t)(x - \bar{x}) + R(x, t)$$

- $(\kappa_1(t), \kappa_2(t))$ : bivariate random walk
- $R(t) = (n_x \times 1 \text{ vector})$  VAR(2), reverting to 0

$$R(t) = AR(t - 1) + BR(t - 2) + Z(t)$$

- $Z(x, t) \sim \text{i.i.d. } N(0, \sigma_Z^2)$
- $A = A_1 + A_2$  and  $B = -A_2A_1$



## VAR matrices $A_1$ and $A_2$

$$A_i = \begin{pmatrix} a_i & 0 & 0 & \dots & & & & & \\ c_i & d_i & 0 & 0 & \dots & & & & \\ d_i/2 & c_i & d_i/2 & 0 & 0 & \dots & & & \\ 0 & d_i/2 & c_i & d_i/2 & 0 & 0 & \dots & & \\ 0 & 0 & d_i/2 & c_i & d_i/2 & 0 & 0 & \dots & \\ \vdots & & \ddots & \ddots & \ddots & \ddots & \ddots & \ddots & \\ & & & & & & & & \end{pmatrix}$$

$a_i$  = AR terms for new members;

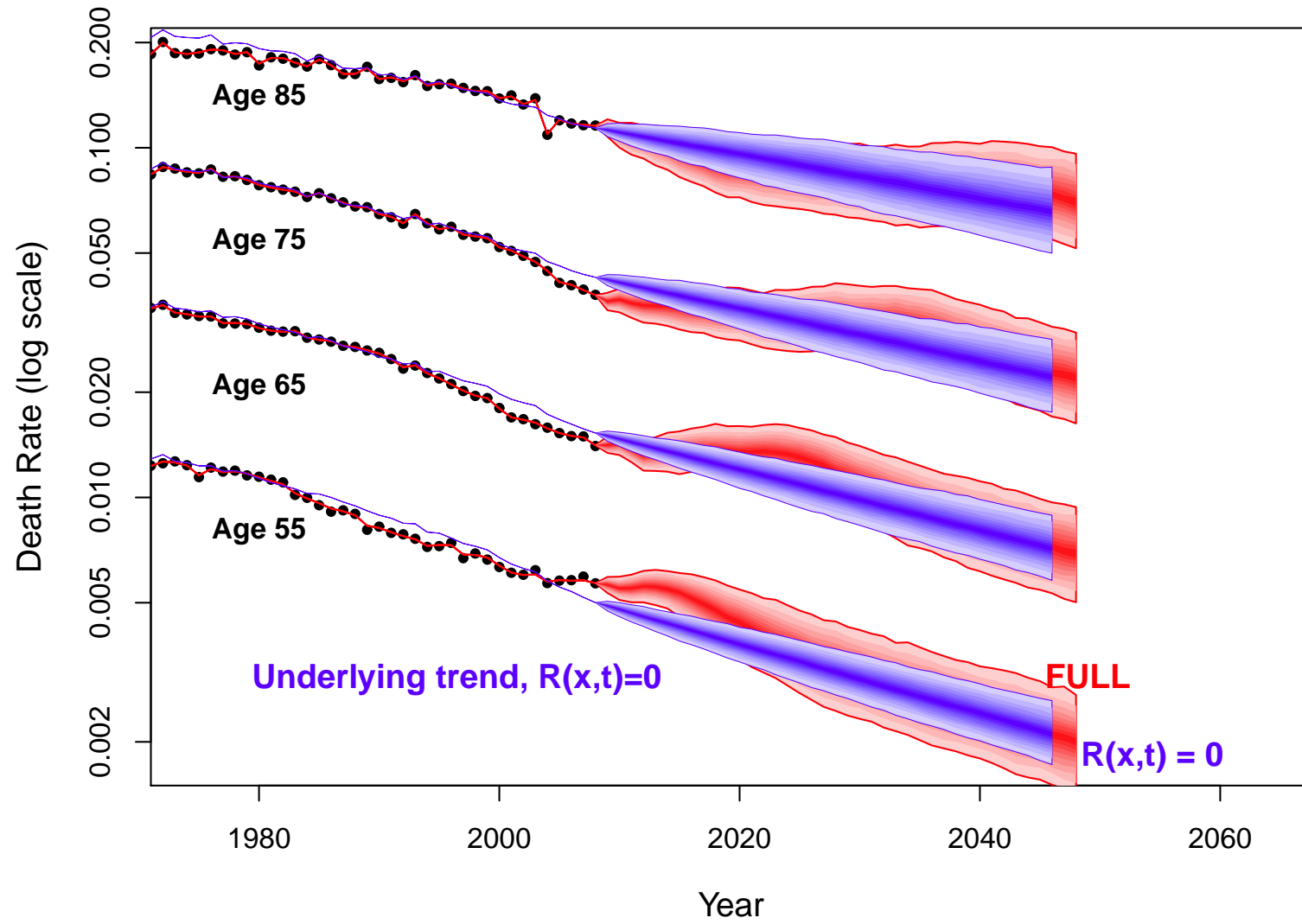
$c_i$  = cohort persistence;

$d_i$  = diffusion coeff.

## Further details

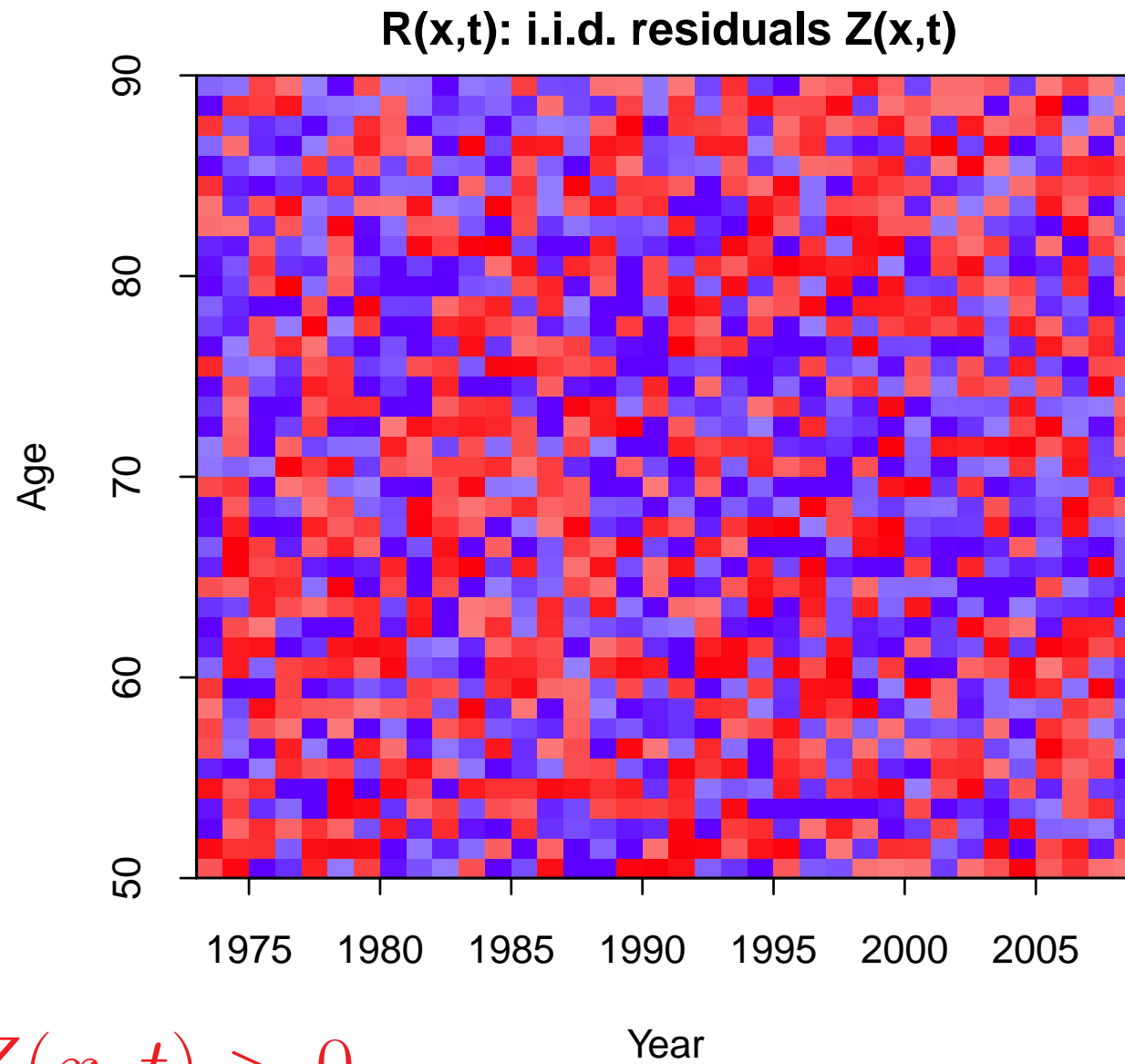
- Deaths:  $D(x, t) \sim \text{Poisson}(m(x, t)E(x, t))$
- Bayesian approach:  
posterior density = likelihood  $\times$  prior
- Upcoming results: mode of posterior density
- Further work: Bayesian parameter uncertainty

## England and Wales, Males 1971–2008

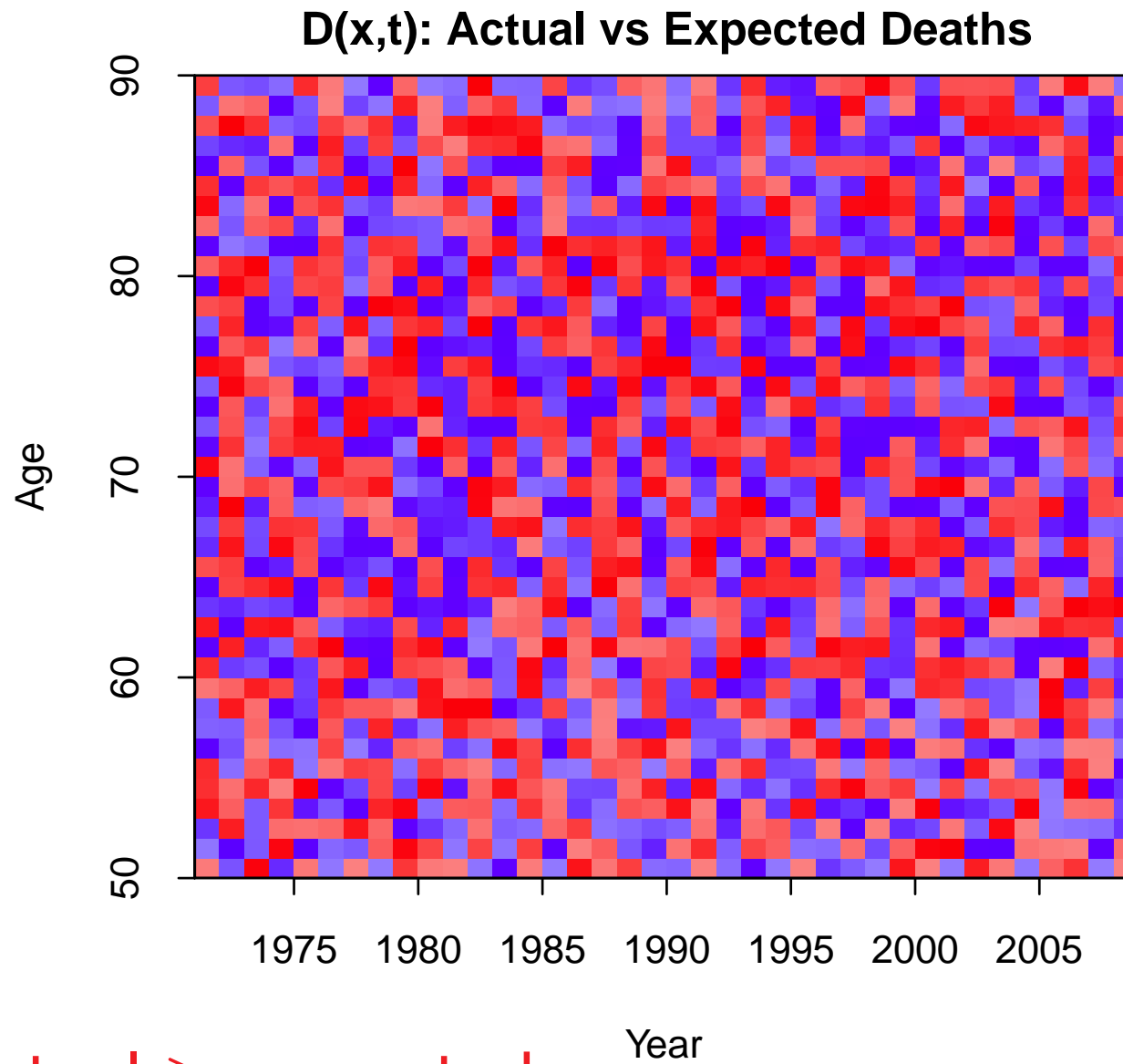


# Cohort-type effects





Red  $\Rightarrow Z(x,t) > 0$



Red  $\Rightarrow$  actual  $>$  expected

## Comparison with related models

$$\log m(x, t) = \beta(x) + \kappa_1(t) + \kappa_2(t)(x - \bar{x})$$

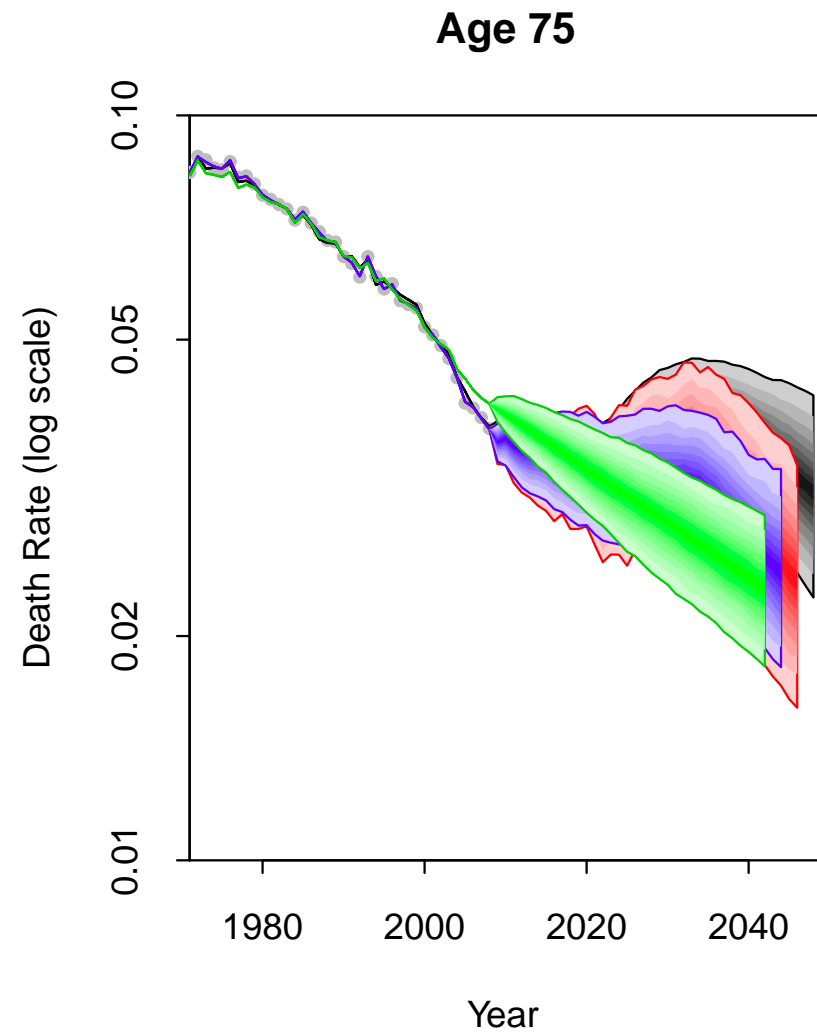
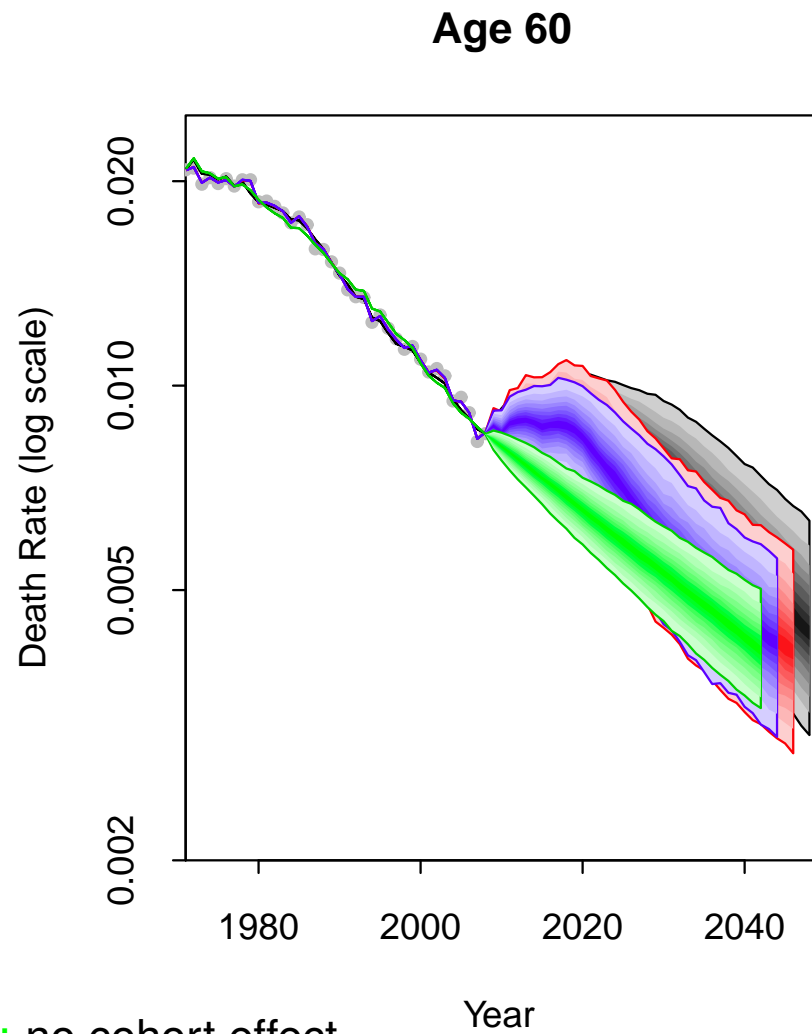
$$\log m(x, t) = \beta(x) + \kappa_1(t) + \kappa_2(t)(x - \bar{x}) + \gamma(t - x)$$

$$\log m(x, t) = \beta(x) + \kappa_1(t) + \kappa_2(t)(x - \bar{x}) + R(x, t)$$

$$R(t) = AR(t - 1) + BR(t - 2) + Z(t) \text{ (A, B as specified earlier)}$$

$$\log m(x, t) = \beta(x) + \kappa_1(t) + \kappa_2(t)(x - \bar{x}) + R(x, t)$$

$$R(t) = AR(t - 1) + BR(t - 2) + Z(t) \text{ (simplified A, B)}$$



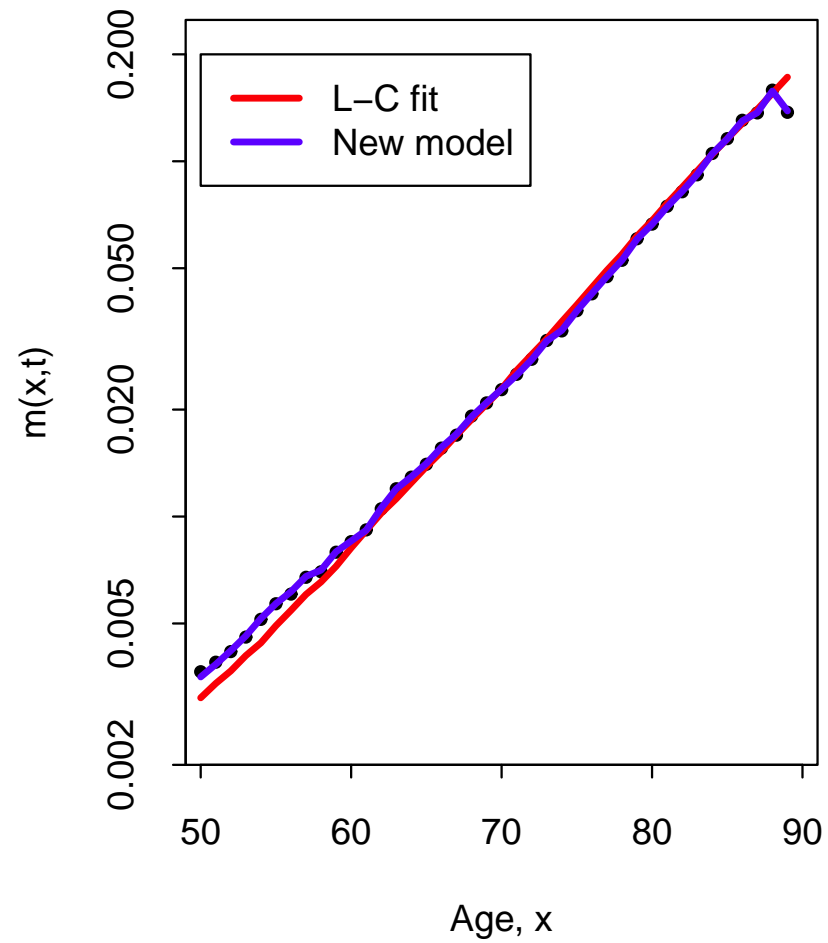
Green: no cohort effect

Red, Blue, Gray: similar trends but also differences

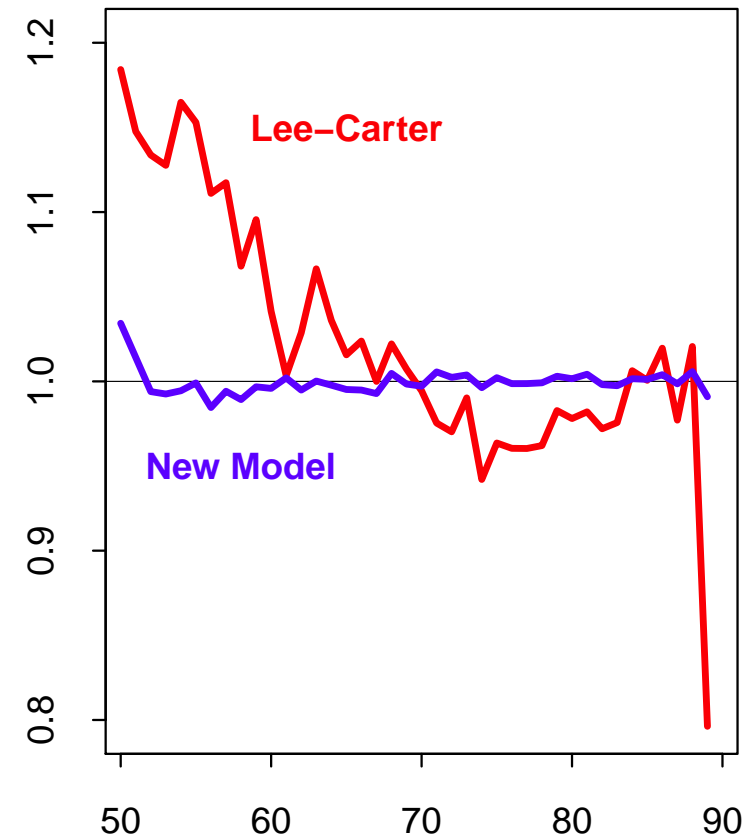


# Base Table Accuracy

EW males 1971–2008: two models  
 $m(x, 2008)$  (log scale)

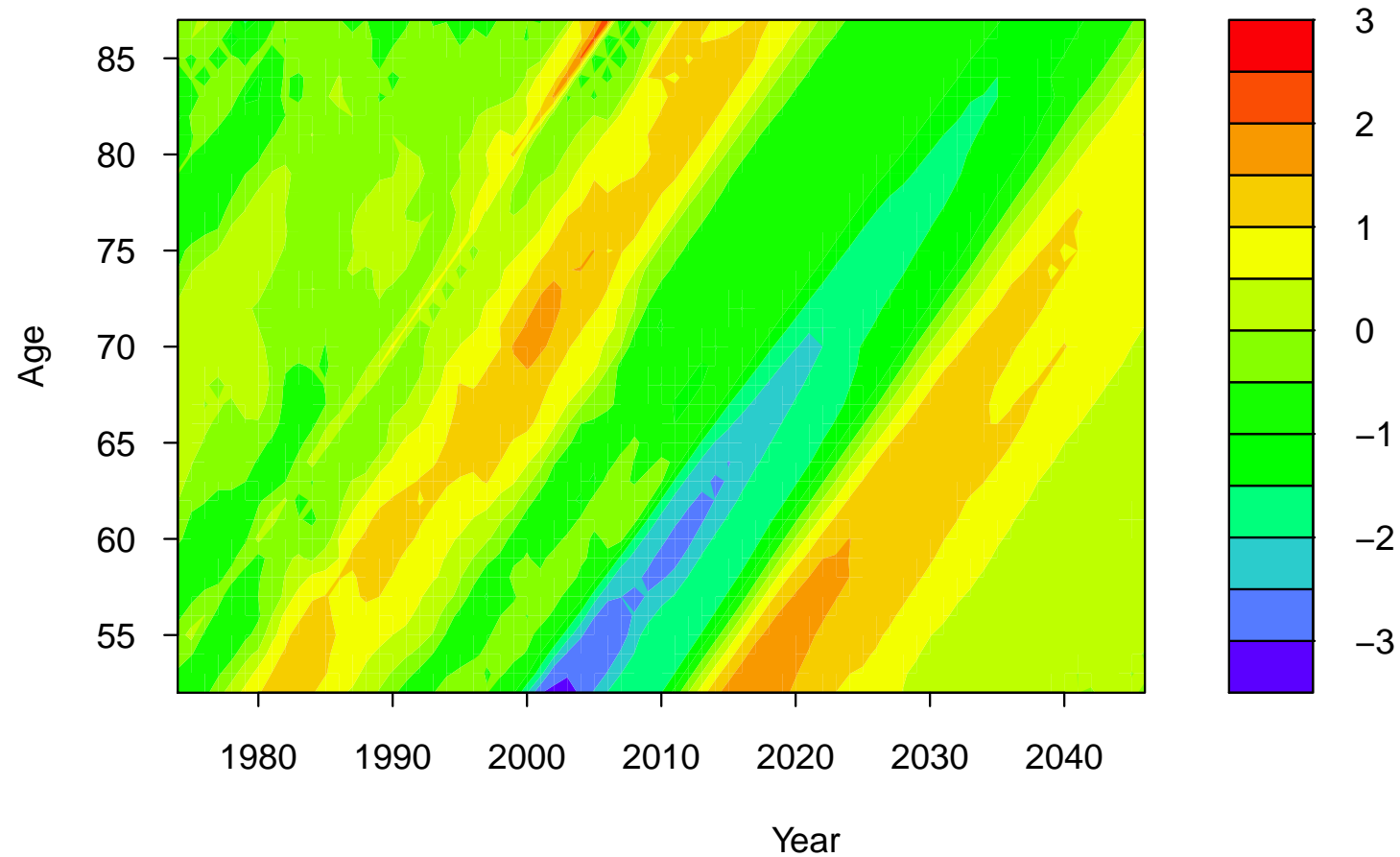


$m_c(x, t) / \hat{m}(x, t)$

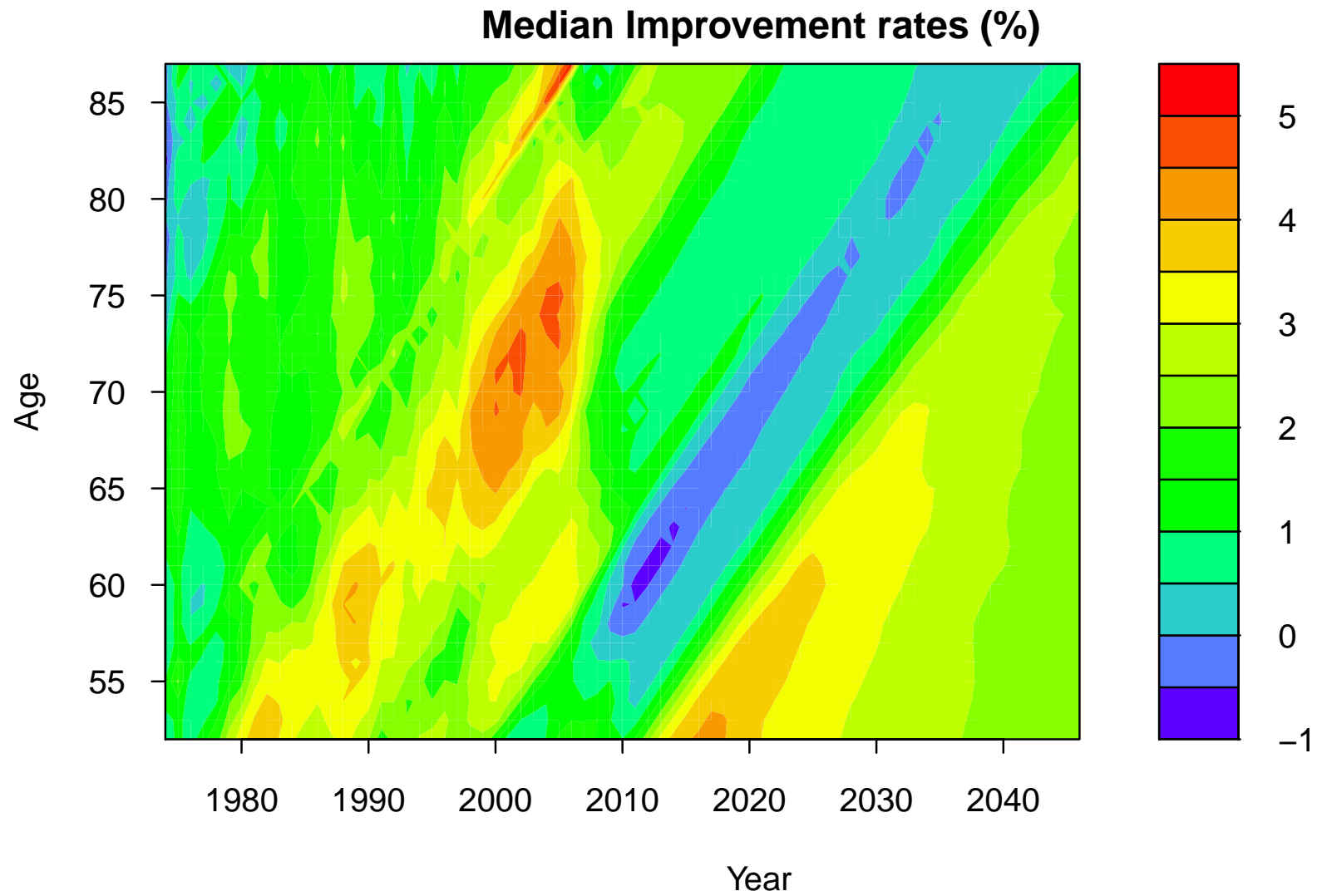


# Contribution of $R(x, t)$ to improvement rates

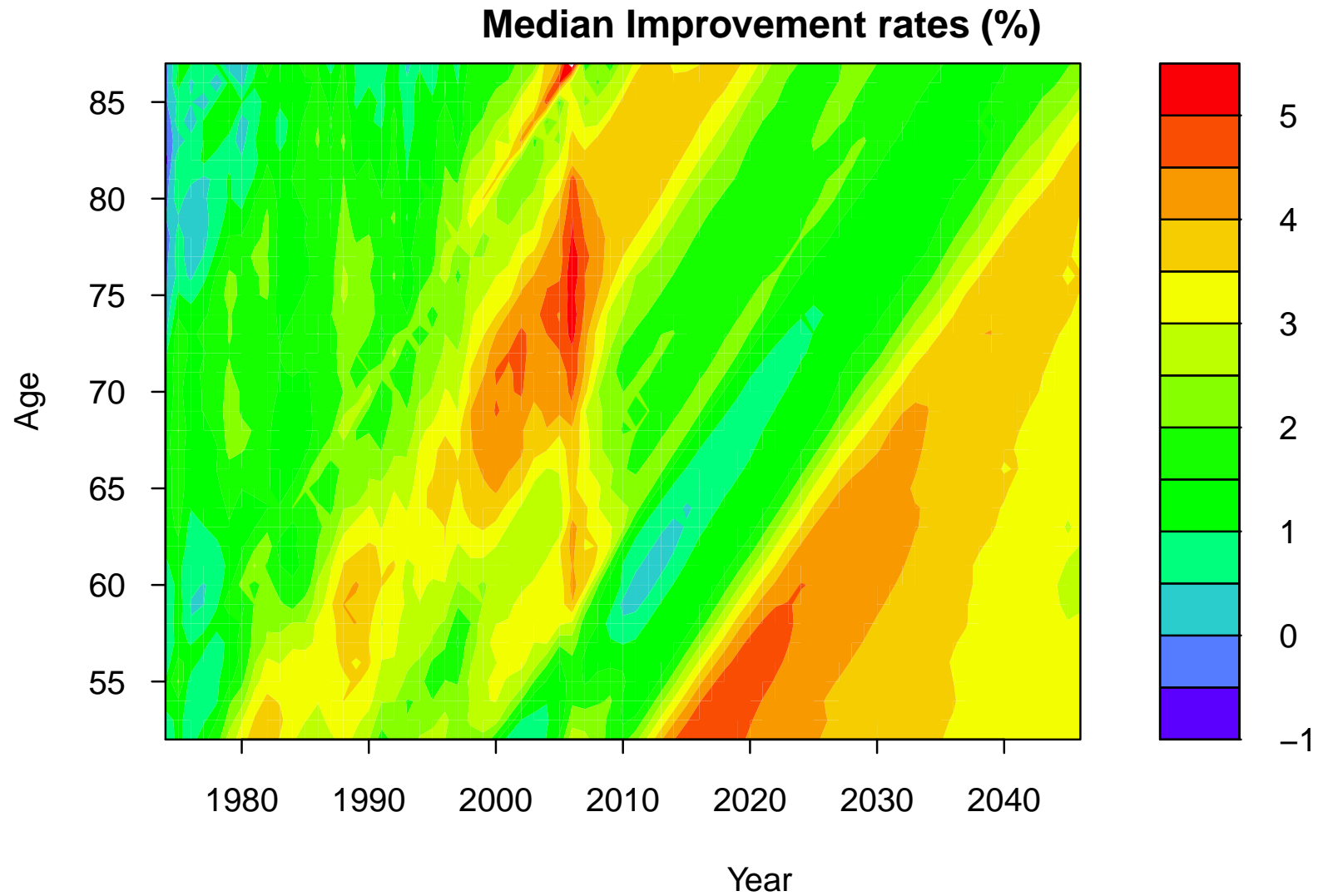
Contribution of  $R(x,t)$  to  
Median Improvement rates (%)



# Median total improvement rates



# Median total improvement rates: Adjusted $\kappa(t)$



## Multiple populations: some thoughts

- Aim for a parsimonious structure
- Items to deal with:
  - Population,  $P$ , specific  $\kappa_i^{(P)}(t)$
  - Population,  $P$ , specific  $R^{(P)}(x, t)$

## Multiple populations: possible structures

### Mortality – version 1:

- Population,  $P$ , specific  $\kappa_i^{(P)}(t)$  correlated
- $R^{(P)}(x, t)$ : assume independent

### Mortality – version 2:

- All populations have the same  $\kappa_i(t)$
- $R^{(P)}(x, t)$ : assume independent
- Greater role for  $R^{(P)}(x, t)$  as country specific effect

## Conclusions

- New model
  - focus on a small number of core period effects
  - adds alternative  $R(x, t)$  to popular cohort effects,  $\gamma(t - x)$
- Model risk more evident in the mean reverting  $R(x, t)$
- But general framework should prove to be more robust:  
long term underlying trends ( $\kappa(t)$ ) are reasonably consistent

## Further work

- Bayesian parameter uncertainty
- Multiple populations: focus on underlying  $\kappa(t)$ 
  - ⇒ less complexity



# Questions

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