
MULTI-POPULATION MORTALITY MODELLING

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Joint work with

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Plan

- Motivation and challenges
- Danish males data
 - 10 sub-populations grouped by wealth
- An extended CBD multi-population model
- Bayesian implementation and results

Motivation for stochastic mortality modelling

- Life expectancy is increasing/mortality falling
⇒ potential impact on
 - pension plan finances; costs ↗
 - life insurance premiums and reserves
- Past patterns ⇒ future improvements uncertain
- Need good stochastic models for
 - central forecasts
 - assessment of uncertainty around central trend
 - development of risk management strategies

Motivation for multi-population modelling

A: Risk assessment

- Multi-country (e.g. consistent demographic projections)
- Males/Females (e.g. consistent demographic projections)
- Socio-economic subgroups (e.g. blue or white collar)
- Smokers/Non-smokers
- Annuities/Life insurance
- Limited data \Rightarrow learn from other populations

Motivation for two-population modelling

B: Risk management for pension plans and insurers

- Retain systematic mortality risk; versus:
- ‘Over-the-counter’ deals (e.g. longevity swap)
 - own experience \Rightarrow 100% risk reduction
 - potentially expensive
- Standardised mortality-linked securities
 - linked to national mortality index
 - $< 100\%$ risk reduction
 - less expensive
 - potential secondary market

Two or more populations

- Linked in some way
- But not identical
- Desire for consistent forecasts
 - distributions
 - individual future scenarios

Key hypothesis

- $m^{(k)}(t, x)$ = pop. k death rate in year t at age x

- Hypothesis (e.g. Li and Lee, 2005):

For each age x , and for any two populations j and k

$$\frac{m^{(j)}(t, x)}{m^{(k)}(t, x)} \text{ does not diverge over time}$$

- Hypothesis \Rightarrow Consequences depend on your choice of stochastic mortality model

Challenges

- Data availability
- Data quality and depth
- Model complexity
 - single population models can be complex
 - 2-population versions are more complex
 - multi-pop
- Multi-population modelling requires
 - (fairly) simple single-population models
 - simple dependencies between populations

A New Case Study and a New Model

- Sub-populations differ from national population
 - socio-economic factors
 - geographical variation
 - other factors
- Denmark
 - High quality data on ALL residents
 - 1981-2005 available
 - Can subdivide population using covariates on the database

Danish Data

- Key covariates
 - Net assets
 - Net income

Problem

- High income \Rightarrow “wealthy” *and healthy* BUT
- Low income $\not\Rightarrow$ not wealthy, poor health
- High assets \Rightarrow “wealthy” *and healthy* BUT
- Low assets $\not\Rightarrow$ not wealthy, poor health

Solution: use a combination

- Wealth, $W = \text{assets} + K \times \text{income}$
- $K = 15$ seems to work well *statistically* as a predictor
- Low wealth, W , predicts poor mortality

Subdividing Data

- Males resident in Denmark for the previous 12 months
- Divide population in year t
 - into 10 equal sized Groups (approx)
 - using wealth in year $t - 1$
- Individuals can change groups up to age 67
- Group is locked down at age 67

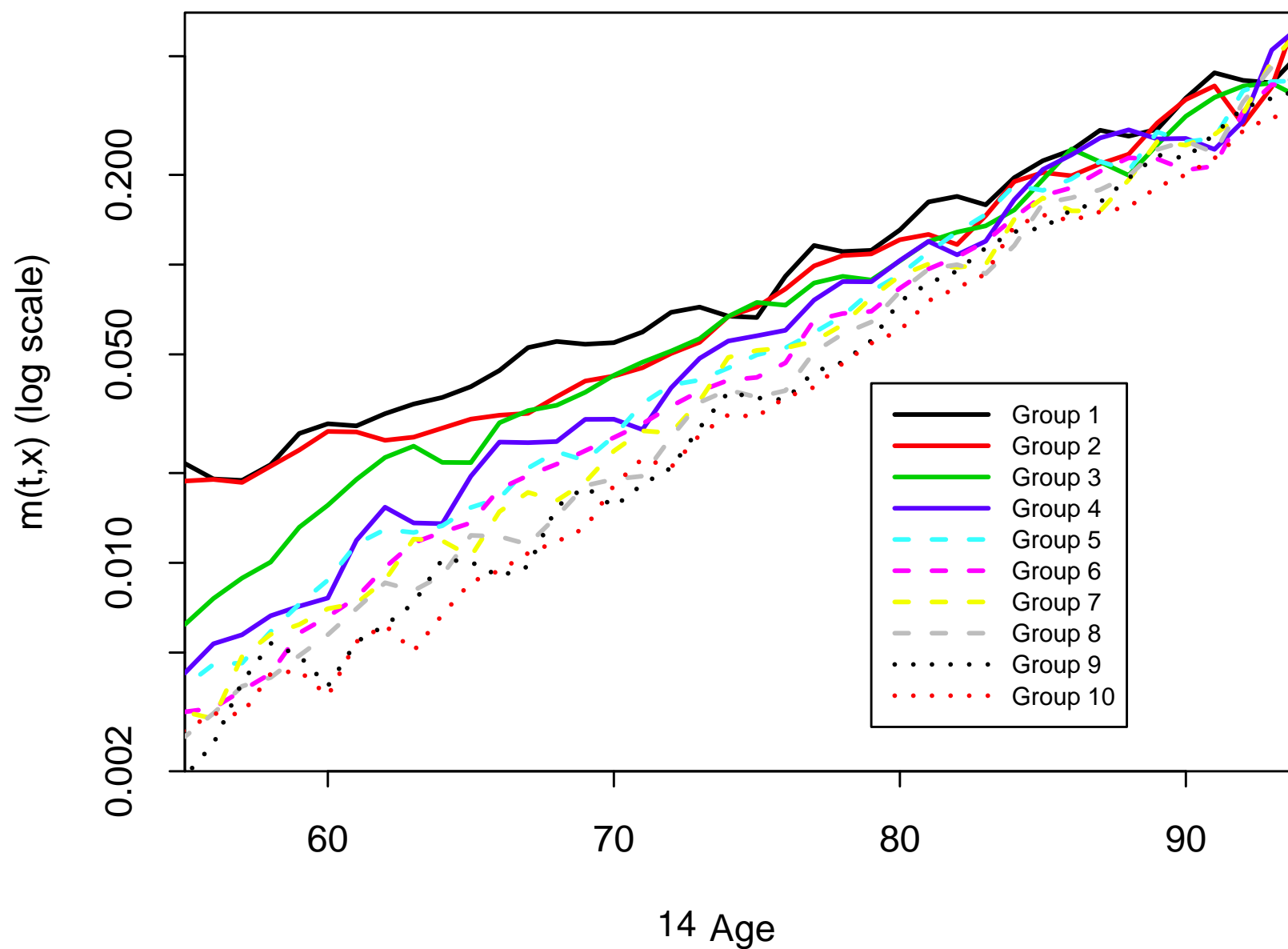
(better than not locking down at age 67)

Subdivided Data

- Exposures $E^{(i)}(t, x)$ for groups $i = 1, \dots, 10$
range from over 4000 down to 20
- Deaths $D^{(i)}(t, x)$
range from 150 down to 6
- Crude death rates $\hat{m}^{(i)}(t, x) = D^{(i)}(t, x) / E^{(i)}(t, x)$
- Small groups \Rightarrow Poisson risk is important

Crude death rates 2005

Males Crude $m(t,x)$; 2005



Modelling the death rates, $m_k(t, x)$

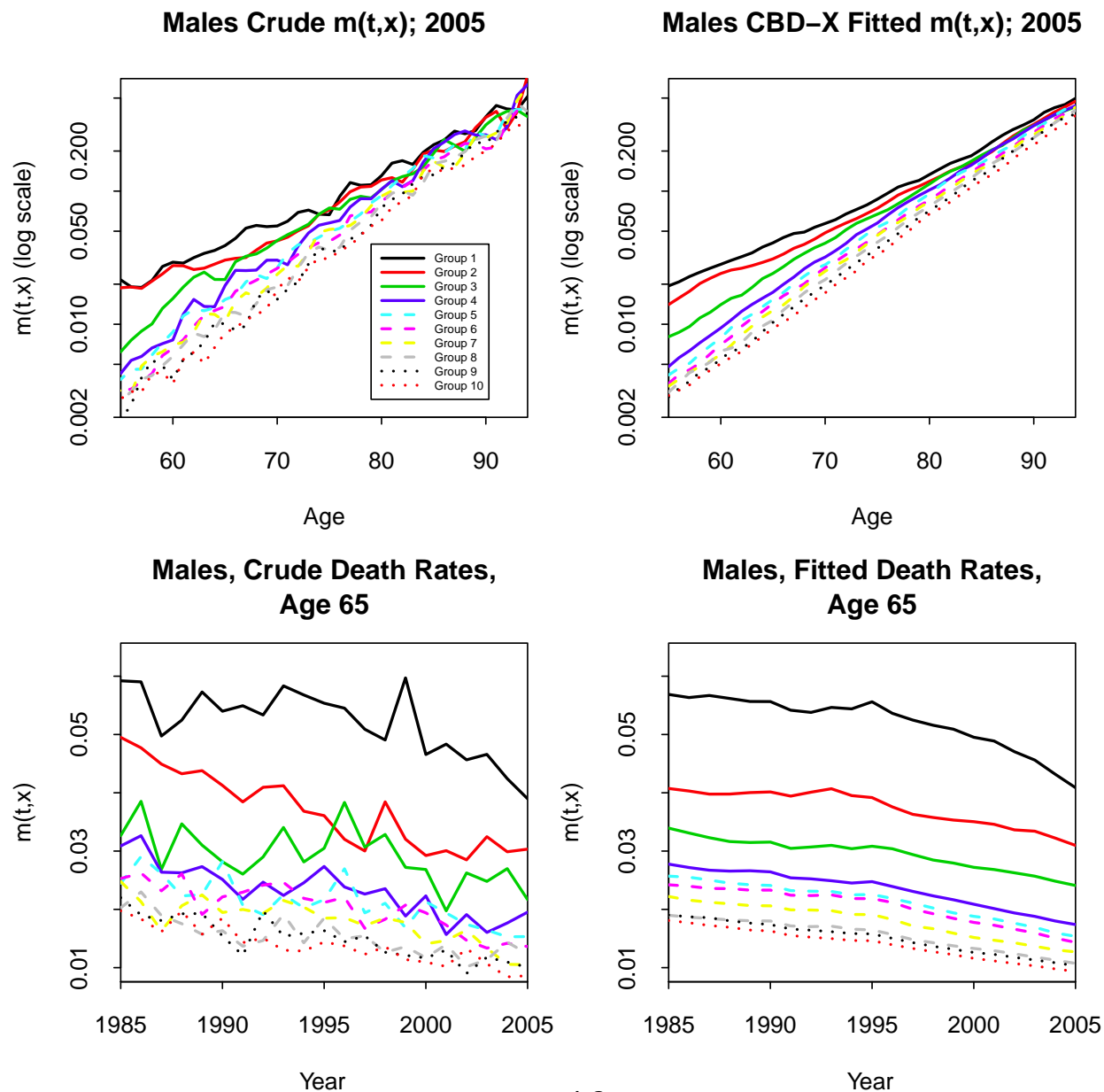
Population k , year t , age x

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

(Extended CBD with a non-parametric base table, $\beta^{(k)}(x)$)

- 10 groups, $k = 1, \dots, 10$ (low to high wealth)
- 21 years, $t = 1985, \dots, 2005$
- 40 ages, $x = 55, \dots, 94$

Model-Inferred Underlying Death Rates 2005



Modelling the death rates, $m_k(t, x)$

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

- Model fits the 10 groups well without a cohort effect
- Non-parametric $\beta^{(k)}(x)$ is essential to preserve group rankings
 - Rankings are evident in crude data
 - “*Biological reasonableness*”: wealthier \Rightarrow healthier

Bayesian modelling

- Combines
 - conditional Poisson likelihood
 - time series likelihood for the $\kappa_j^{(k)}(t)$
 - (uninformative) prior distributions for process parameters
- Output *posterior distribution* for
 - $\beta^{(k)}(x)$, $\kappa_1^{(k)}(t)$, $\kappa_2^{(k)}(t)$ latent state variables
 - time series process parameters

Time series modelling

- $t \rightarrow t + 1$: Allow for correlation
 - between $\kappa_1^{(k)}(t + 1)$ and $\kappa_2^{(k)}(t + 1)$
 - between groups $k = 1, \dots, 10$
- Biological reasonableness \Rightarrow key hypothesis
groups should not diverge
- Sufficient that we have mean reversion in
 $\kappa_1^{(j)}(t) - \kappa_1^{(k)}(t)$ and $\kappa_2^{(j)}(t) - \kappa_2^{(k)}(t)$

A specific model

$$\begin{aligned}\kappa_1^{(i)}(t) &= \kappa_1^{(i)}(t-1) + \mu_1 + Z_{1i}(t) && \text{(random walk)} \\ &\quad - \psi \left(\kappa_1^{(i)}(t-1) - \bar{\kappa}_1(t-1) \right) && \text{(gravity between groups)}\end{aligned}$$

$$\begin{aligned}\kappa_2^{(i)}(t) &= \kappa_2^{(i)}(t-1) + \mu_2 + Z_{2i}(t) \\ &\quad - \psi \left(\kappa_2^{(i)}(t-1) - \bar{\kappa}_2(t-1) \right)\end{aligned}$$

where

$$\bar{\kappa}_1(t) = \frac{1}{n} \sum_{i=1}^n \kappa_1^{(i)}(t) \quad \text{and} \quad \bar{\kappa}_2(t) = \frac{1}{n} \sum_{i=1}^n \kappa_2^{(i)}(t)$$

A specific model

$$\kappa_1^{(i)}(t) = \kappa_1^{(i)}(t-1) + \mu_1 + Z_{1i}(t) - \psi \left(\kappa_1^{(i)}(t-1) - \bar{\kappa}_1(t-1) \right)$$

$$\kappa_2^{(i)}(t) = \kappa_2^{(i)}(t-1) + \mu_2 + Z_{2i}(t) - \psi \left(\kappa_2^{(i)}(t-1) - \bar{\kappa}_2(t-1) \right)$$

- $(\bar{\kappa}_1(t), \bar{\kappa}_2(t)) \sim$ bivariate random walk
- Each $\kappa_1^{(i)}(t) - \bar{\kappa}_1(t) \sim AR(1)$ reverting to 0
- Each $\kappa_2^{(i)}(t) - \bar{\kappa}_2(t) \sim AR(1)$ reverting to 0

A specific model

$$\kappa_1^{(i)}(t) = \kappa_1^{(i)}(t-1) + \mu_1 + Z_{1i}(t) - \psi \left(\kappa_1^{(i)}(t-1) - \bar{\kappa}_1(t-1) \right)$$

$$\kappa_2^{(i)}(t) = \kappa_2^{(i)}(t-1) + \mu_2 + Z_{2i}(t) - \psi \left(\kappa_2^{(i)}(t-1) - \bar{\kappa}_2(t-1) \right)$$

The $Z_{i,j}$ are multivariate normal, mean 0 and

$$\text{Cov}(Z_{ki}, Z_{lj}) = \begin{cases} v_{kl} & \text{for } i = j \\ \rho v_{kl} & \text{for } i \neq j \end{cases}$$

ρ = cond. correlation between $\kappa_1^{(i)}(t)$ and $\kappa_1^{(j)}(t)$ etc.

Comments

- Model is very simple
 - One gravity parameter, $0 < \psi < 1$
 - One between-group correlation parameter,
 $0 < \rho < 1$
- Many generalisations are possible
- But more parameters + more complex computing
- This simple model seems to fit quite well.
- Nevertheless \Rightarrow work in progress

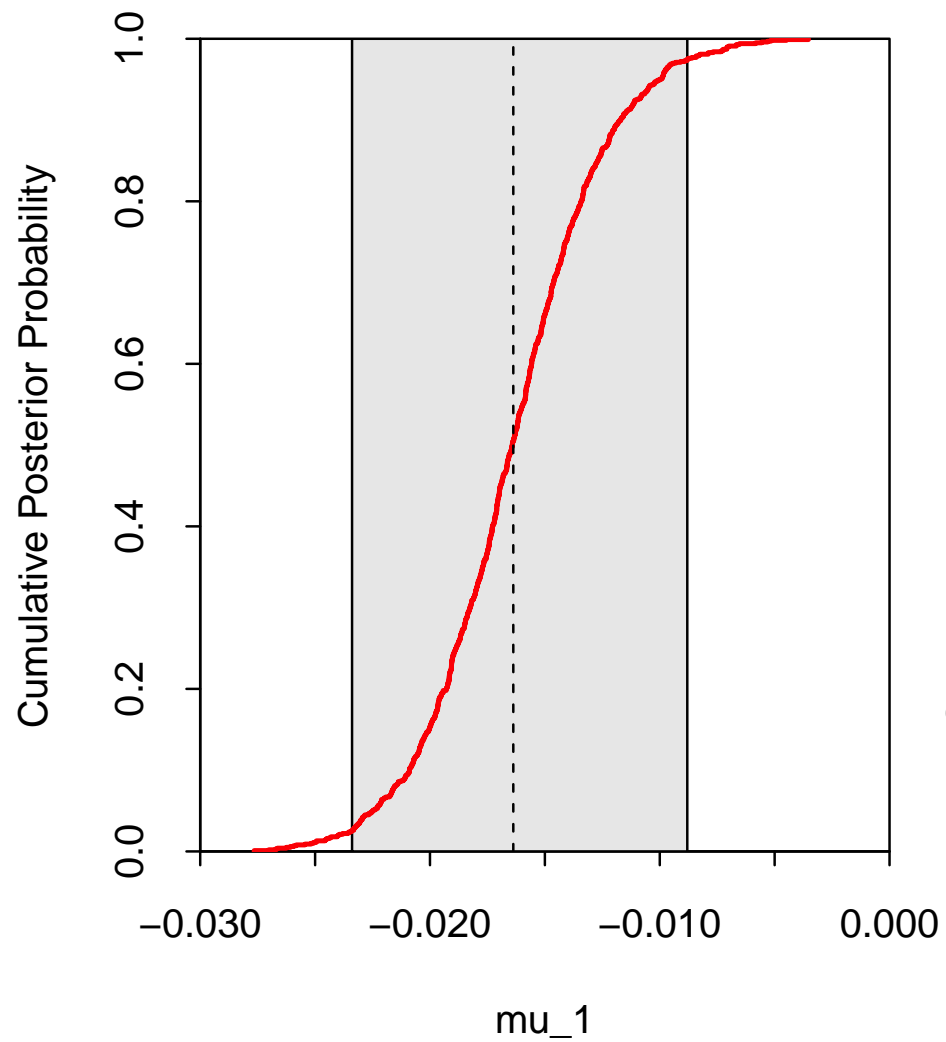
Prior distributions

- As uninformative as possible
- $\mu_1, \mu_2 \sim$ improper uniform prior
- $\{v_{ij}\} \sim$ Inverse Wishart
- $\rho \sim \text{Beta}(2, 2)$
- $\psi \sim \text{Beta}(2, 2)$

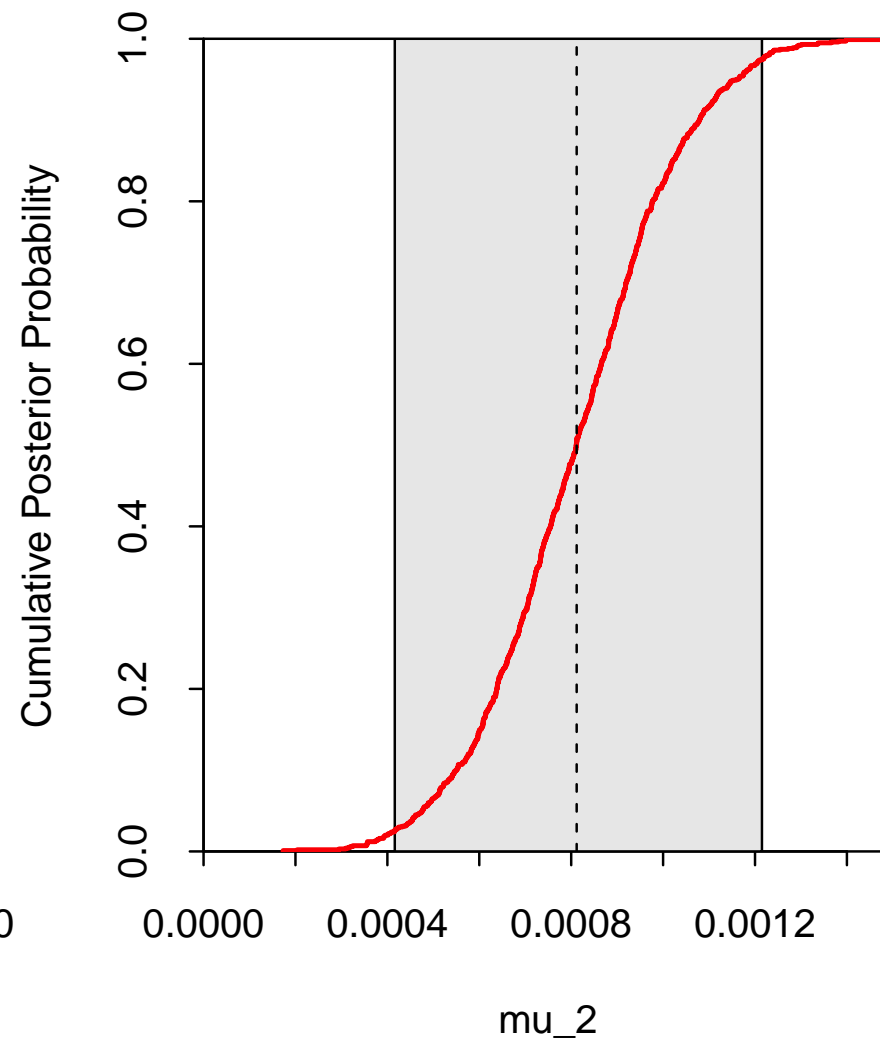
State variables and process parameters estimated using
MCMC (Gibbs + Metropolis-Hastings)

Posterior Distributions

Kappa_1 Drift, μ_1

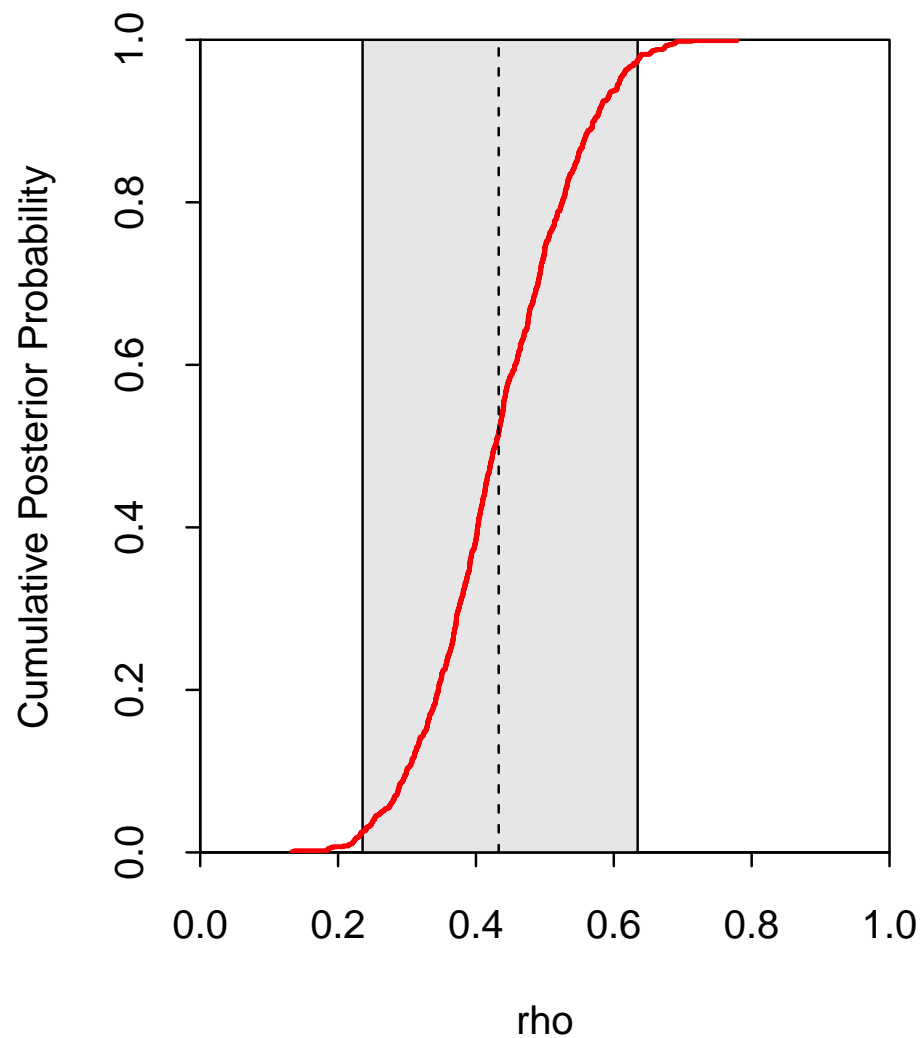


Kappa_2 Drift, μ_2

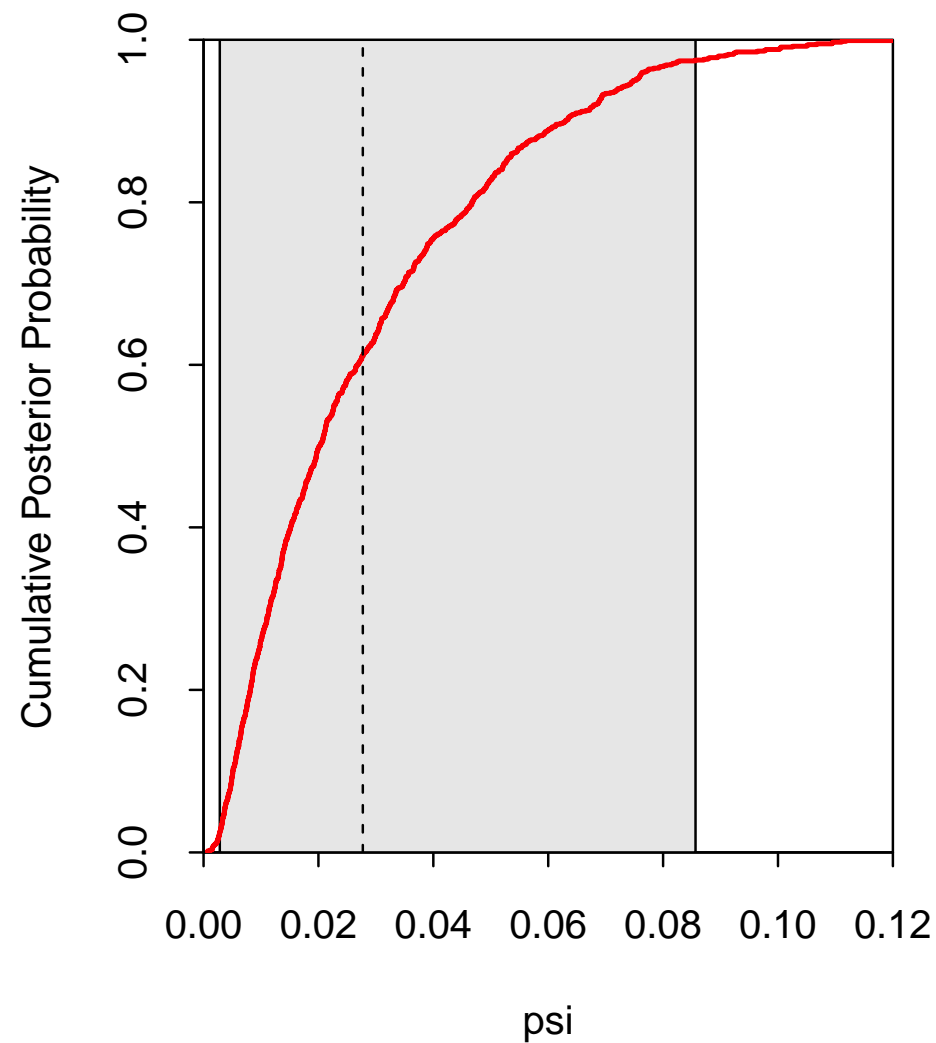


Posterior Distributions

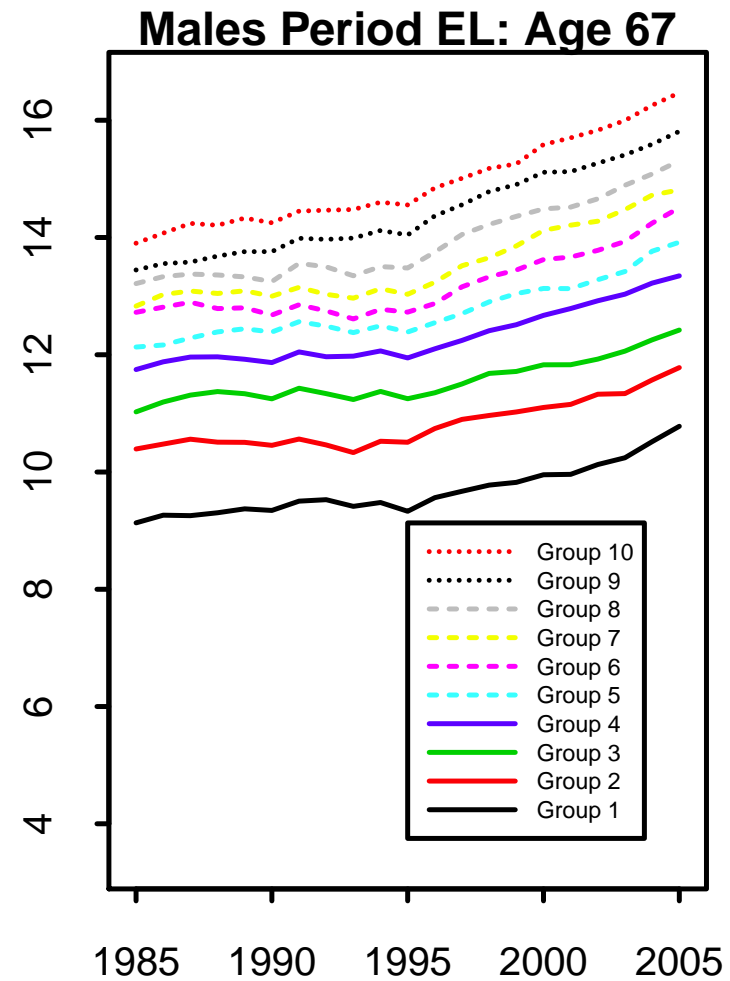
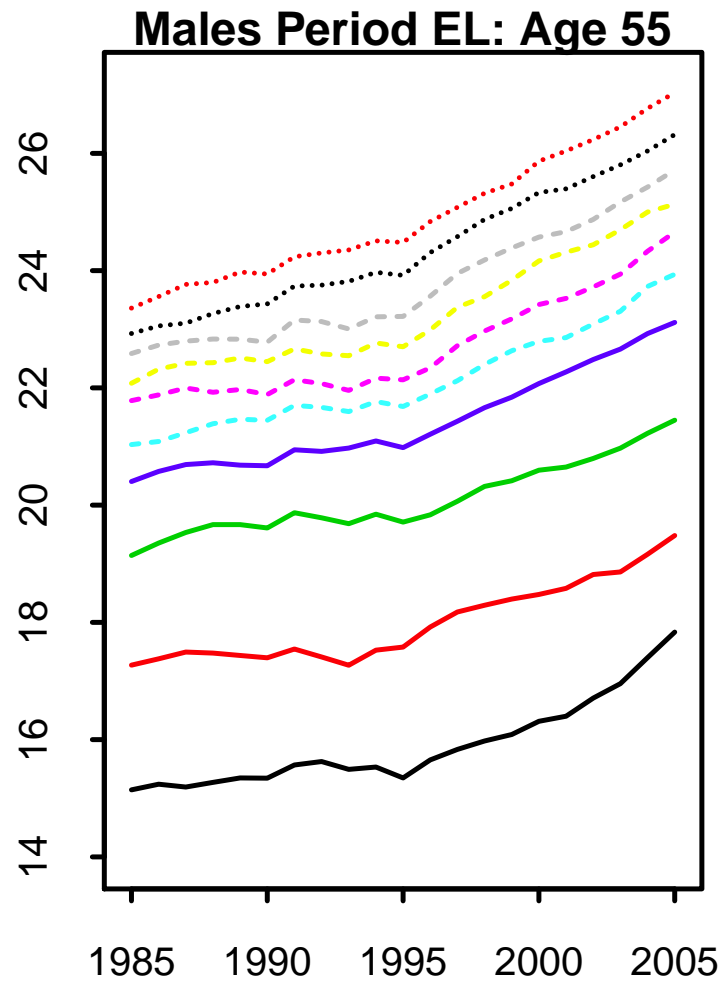
Between Group Correlation, ρ



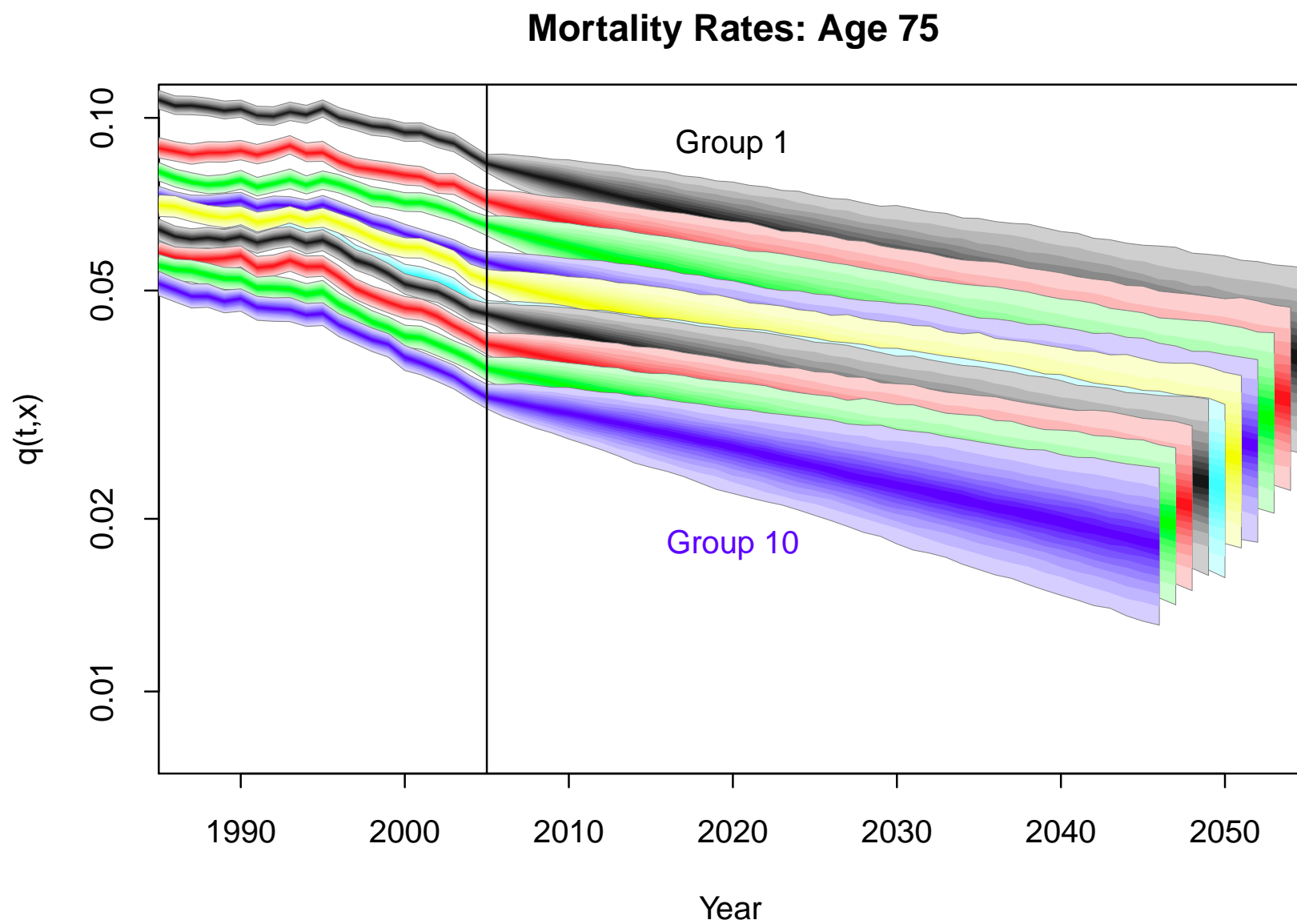
Gravity Parameter, ψ



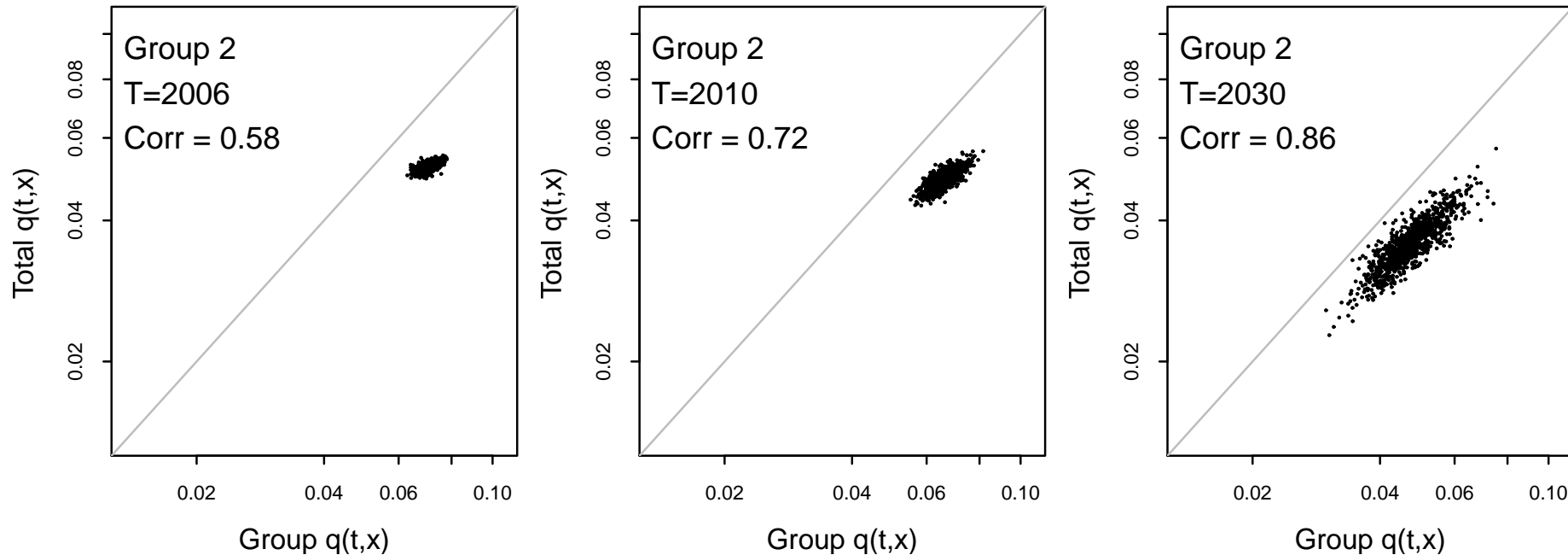
Life Expectancy for Groups 1 to 10



Mortality Fan Charts Including Parameter Uncertainty



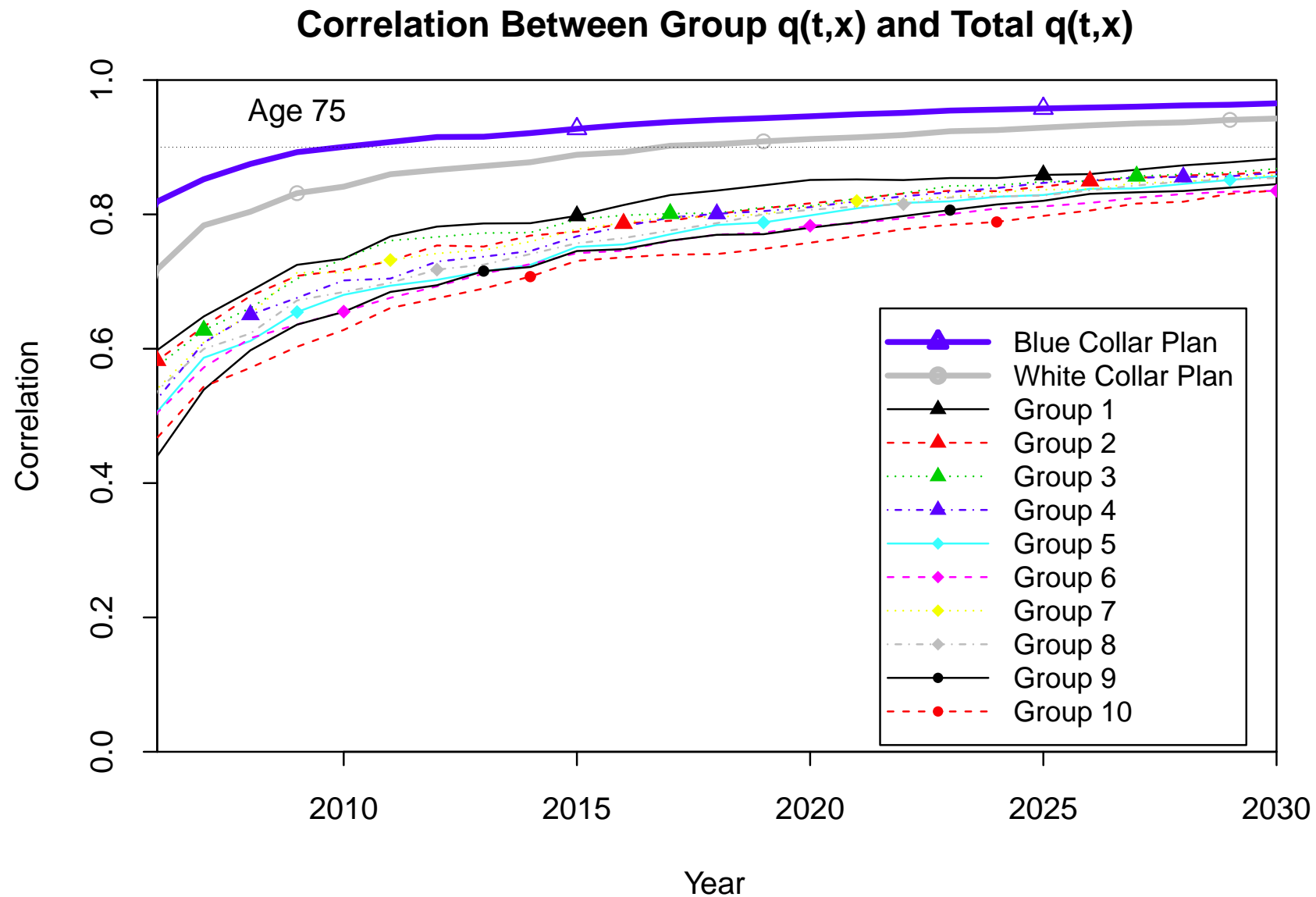
Simulated Group versus Population Mortality



As T increases

- Scatterplots become more dispersed
- Shift down and to the left
- Correlation increases

Forecast Correlations



Conclusions

- Development of a new multi-population dataset for Denmark
strong biologically reasonable group rankings
- Unlike multi-country data
a priori ranking of wealth-related groups
- Proposal for a simple new multi-population model
- Next steps:
 - Females data
 - More general correlation and gravity structures