

A General Procedure for Building Mortality Models

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Motivation

- Recently, there has been a proliferation of new mortality models
- Some of these models are “black-box algorithms” such as PCA
 - Involving terms that lack “demographic significance”
- Others have functional terms added without justification and require a priori assumptions about what the model should look like
- We introduce a general procedure which provides structure to the model building process
- This requires an explicit “toolkit” of functions

Age/Period/Cohort Modelling Framework

$$\eta_{x,t} = \alpha_x + \sum_{j=1}^n \beta_x^j \kappa_t^j + \beta_x^0 \gamma_{t-x}$$

Link function between model and data; typically

$$\eta_{x,t} = \ln(\hat{\mu}_{x,t}) = \ln\left(\frac{D_{x,t}}{E_{x,t}^c}\right)$$

$$\eta_{x,t} = \text{logit}(\hat{q}_{x,t}) = \ln\left(\frac{D_{x,t}}{E_{x,t}^0}\right) - \ln\left(1 - \frac{D_{x,t}}{E_{x,t}^0}\right)$$

Static life table – baseline mortality rates at each age

Age functions – which ages mortality is changing at – in non-parametric form.

Smooth age functions $f^j(x; \theta)$ reduce number of parameters to be estimated

Cohort parameters – “lifelong” factors depending upon year of birth

Often set to 1 for simplicity and robustness

Period factors governing trends in time at same age range

Examples of Models in this Form

- Lee and Carter (1992) $\ln(\mu_{x,t}) = \alpha_x + \beta_x \kappa_t$
- Renshaw and Haberman (2006) $\ln(\mu_{x,t}) = \alpha_x + \beta_x^1 \kappa_t + \beta_x^0 \gamma_{t-x}$
 - (modified version of this model without age function on the γ 's found to be more robust)
- Cairns, Blake and Dowd (2006) $\text{logit}(q_{x,t}) = \kappa_t^1 + (x - \bar{x}) \kappa_t^2$
- Plat (2009) $\ln(\mu_{x,t}) = \alpha_x + \kappa_t^1 + (x - \bar{x}) \kappa_t^2 + (x - \bar{x})^+ \kappa_t^3 + \gamma_{t-x}$
- O'Hare and Li (2012) $\ln(\mu_{x,t}) = \alpha_x + \kappa_t^1 + (x - \bar{x}) \kappa_t^2 + \left((x - \bar{x})^+ + \left((x - \bar{x})^+ \right)^2 \right) \kappa_t^3 + \gamma_{t-x}$

Model Selection Criteria

- Adequacy –
 - There should be a sufficient number of terms to capture all significant structure in, and provide a good fit to the data
- Parsimony –
 - Have the smallest number of terms and free parameters necessary (trade off with the adequacy of the model)
- Demographic Significance –
 - Models should be biologically reasonable
 - Terms allow identification with underlying biological and socio-economic processes occurring in the population
- Completeness –
 - Models should span entire age range and not be limited to a subset of ages by construction
 - Models should include allowance for cohort effects and be able to separate these from age/period terms

General Model Building Procedure

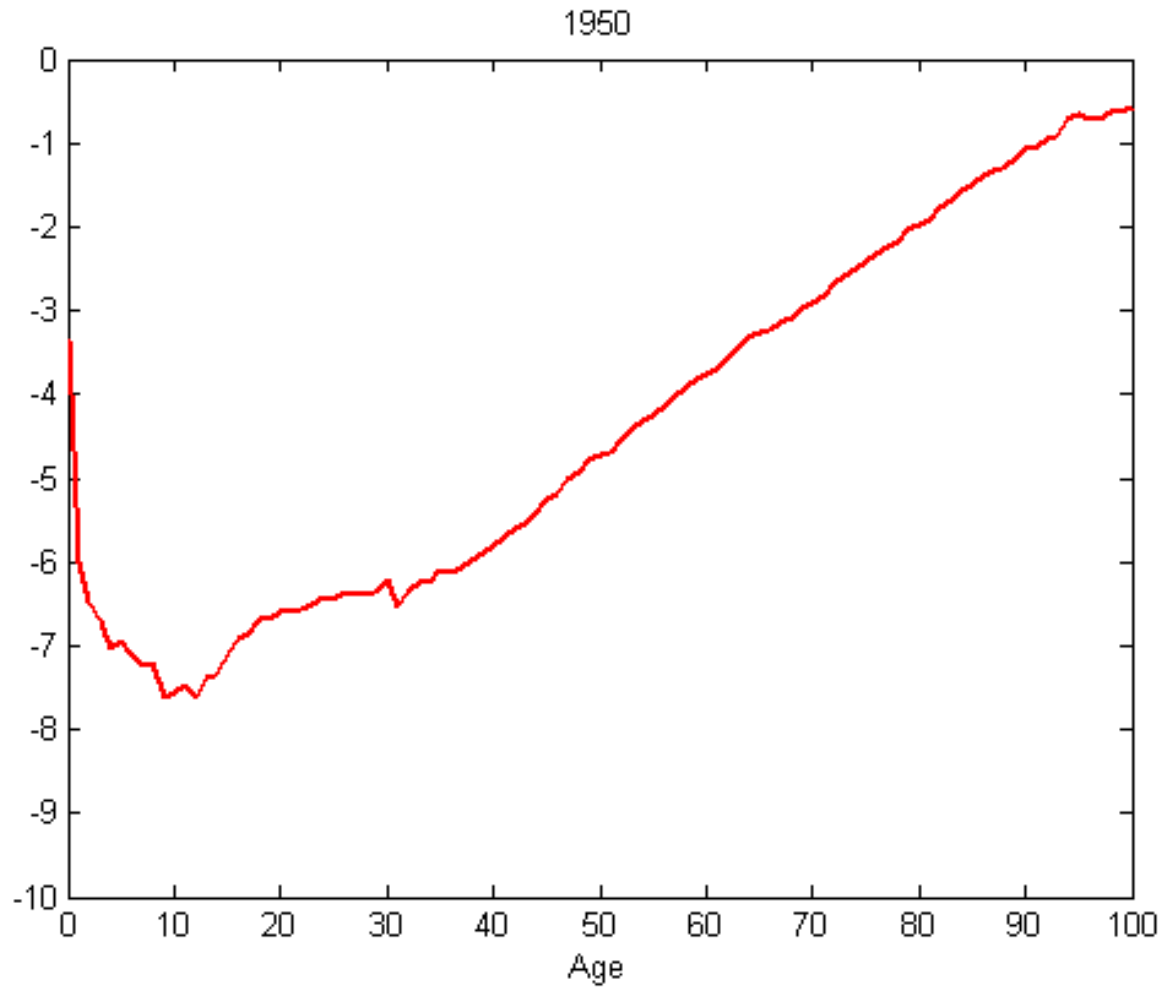
Important to Avoid Over-Parameterisation

- Major risk of procedure is overfitting the model, i.e. adding too many age/period terms.
- We try to avoid this by –
 - Measuring goodness of fit using a metric that penalises the number of parameters used
 - e.g. the Bayes Information Criterion
 - Applying subjective as well as statistical tests on whether new terms are needed – they must be demographically significant as well as statistically significant
 - Requiring age functions to be smooth over a range of adjacent ages to avoid trying to fit statistical noise

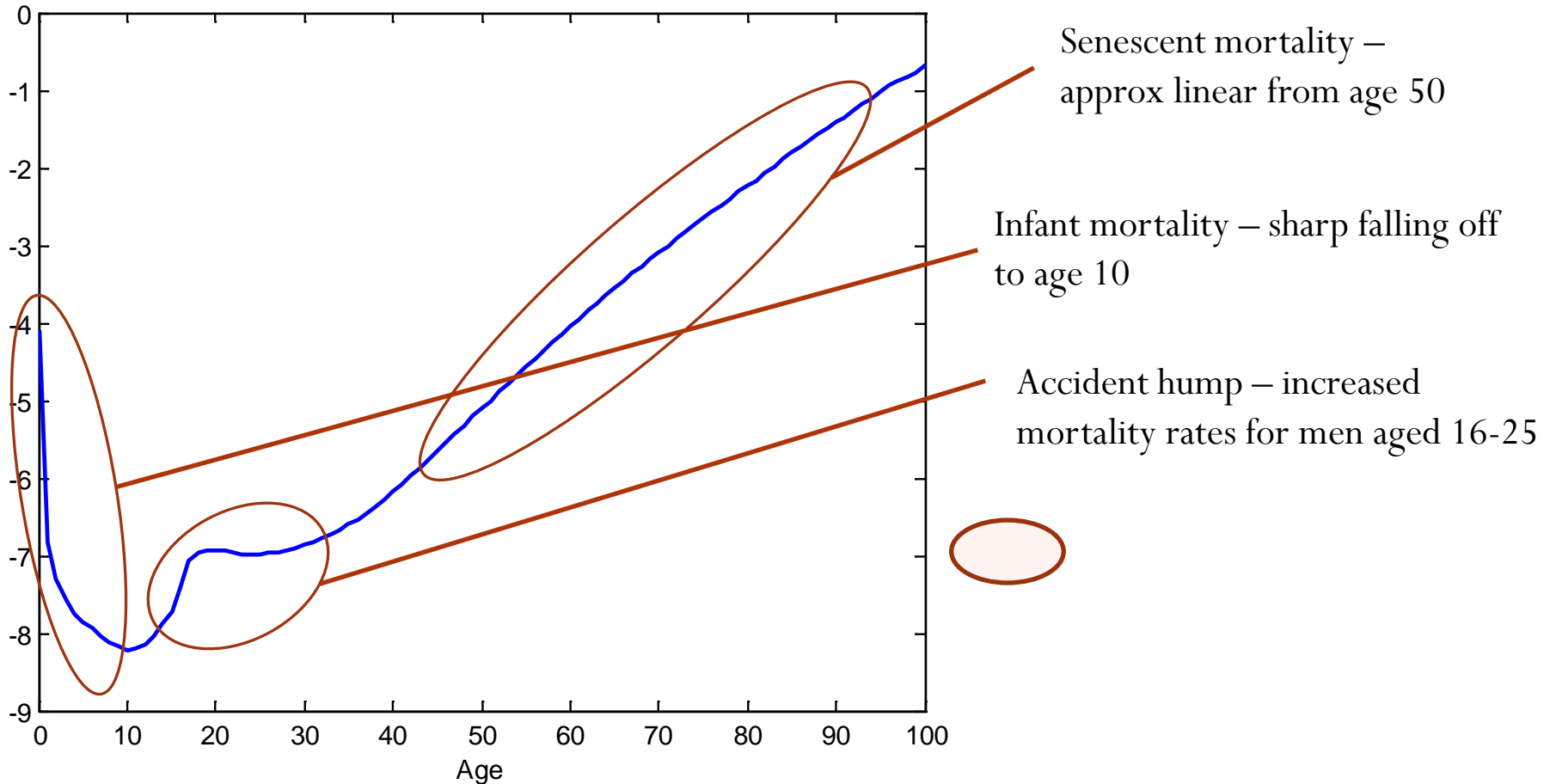
Application

- We apply the general procedure to data for men in the UK from ages 0-100 and years 1950-2009
- Death counts and central exposures to risk from the Human Mortality Database
- We test the final model and its residuals and compare it with existing model fitting procedures
- Need access to a suitable “toolkit” of age functions which we can use to build an appropriate model

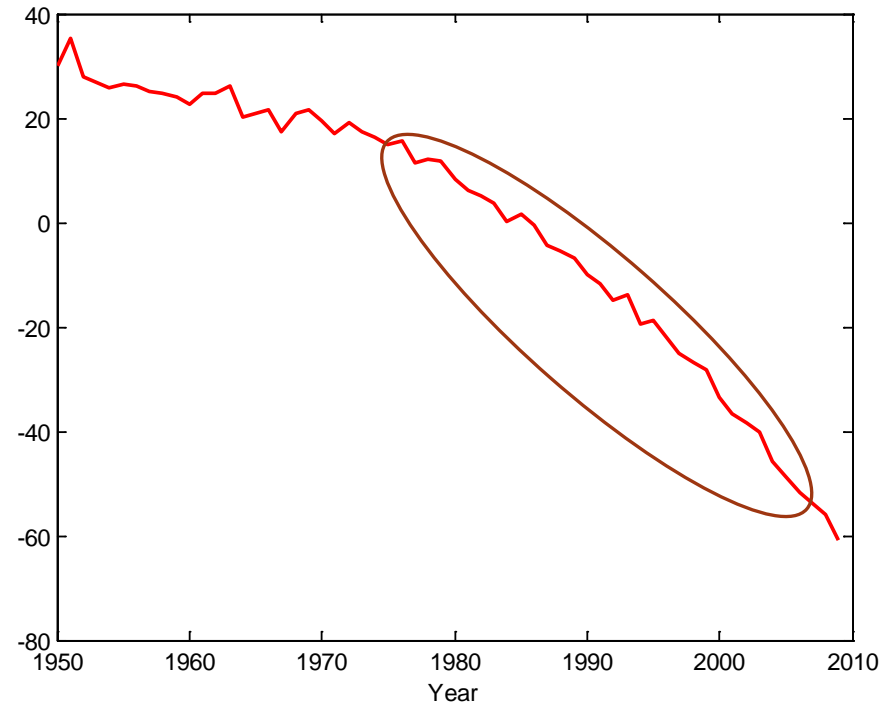
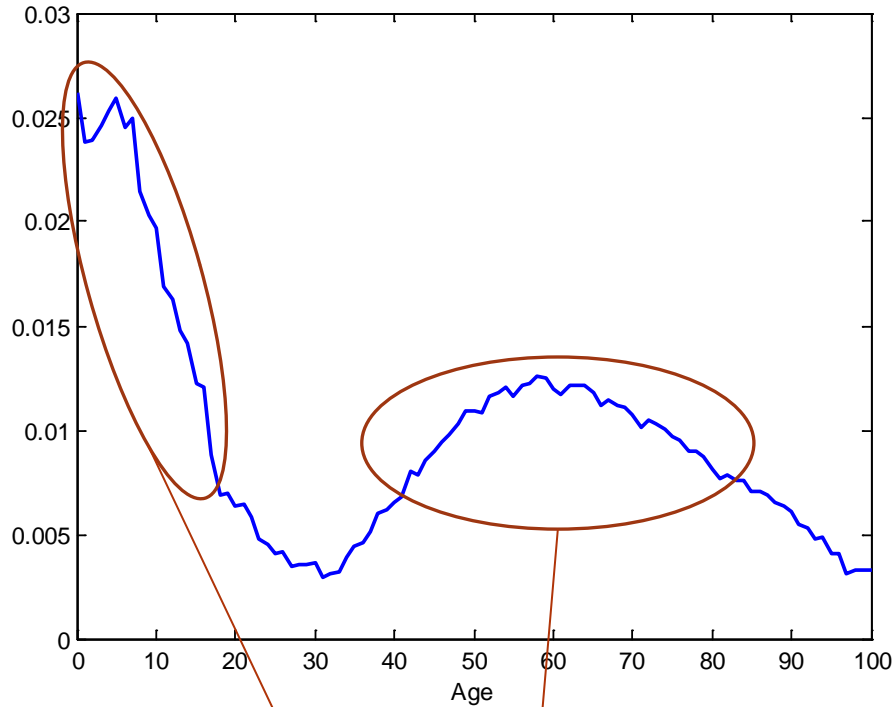
UK Male Data



Stage 0 - Static Life Table

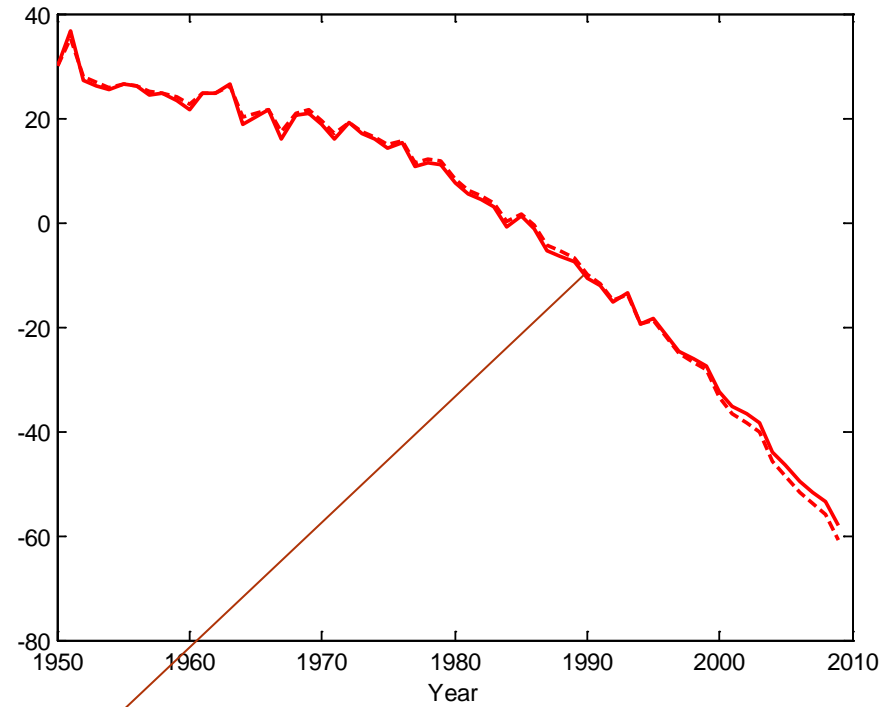
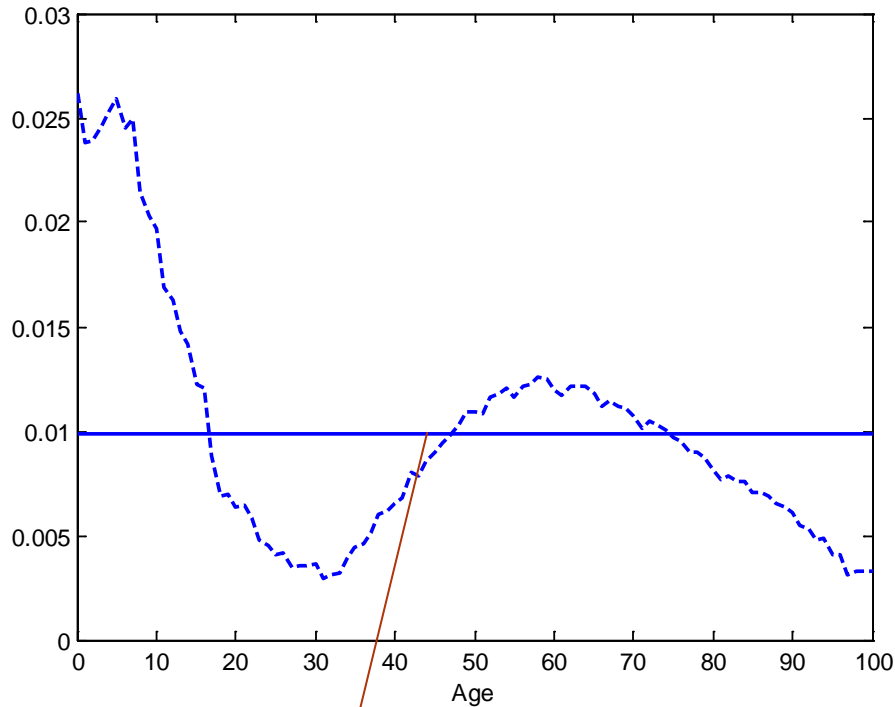


Stage 1 - Add Non-Parametric Age/Period Complementary Pair to Find 1st Dominant Trend



Combines features from across the age range with different underlying causes but which are correlated

Stage 1 – Find Simplest Parametric Term That Does the Same Job

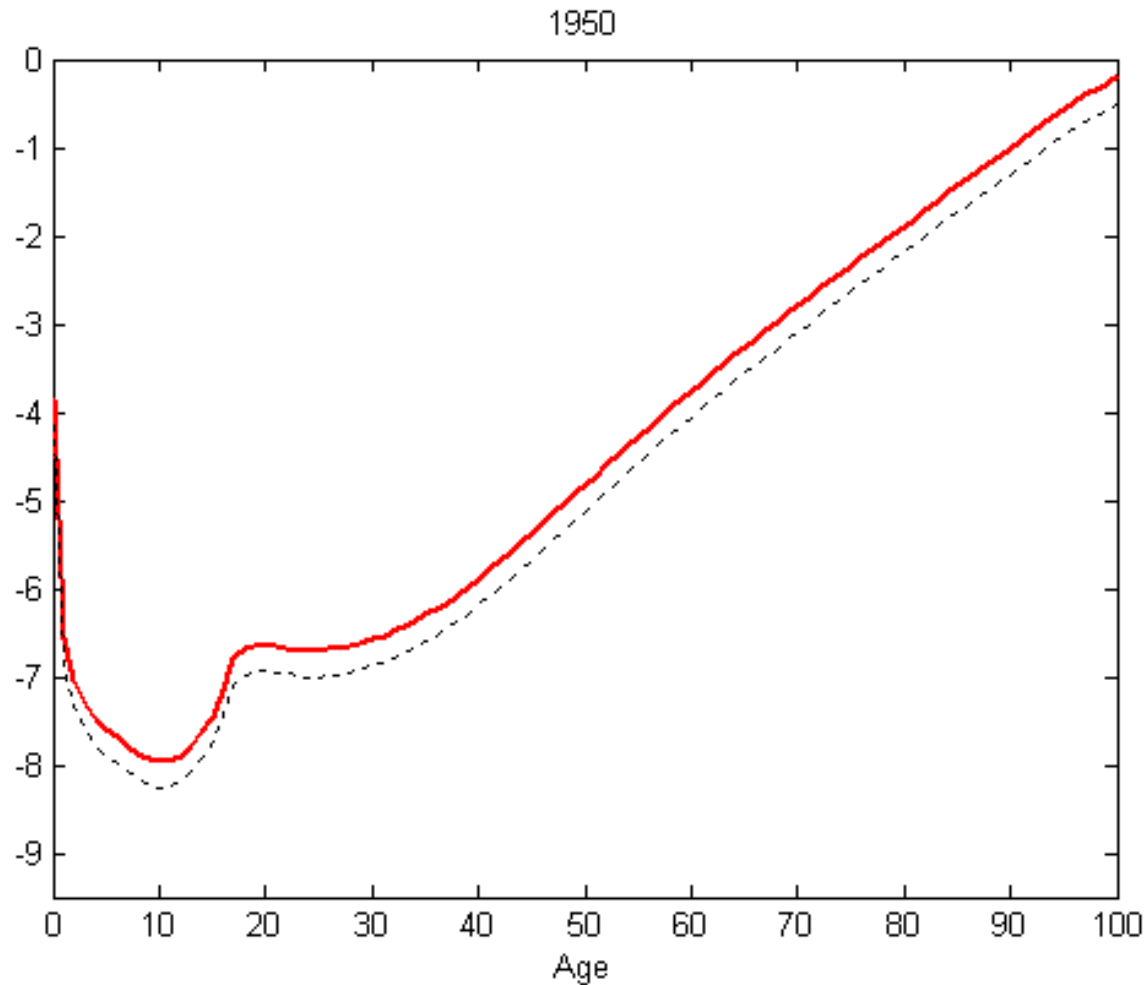


Constant age function – changes in overall level of mortality due to accidents, hygiene, etc

Finds same trend in the data – general fall in mortality rate with time



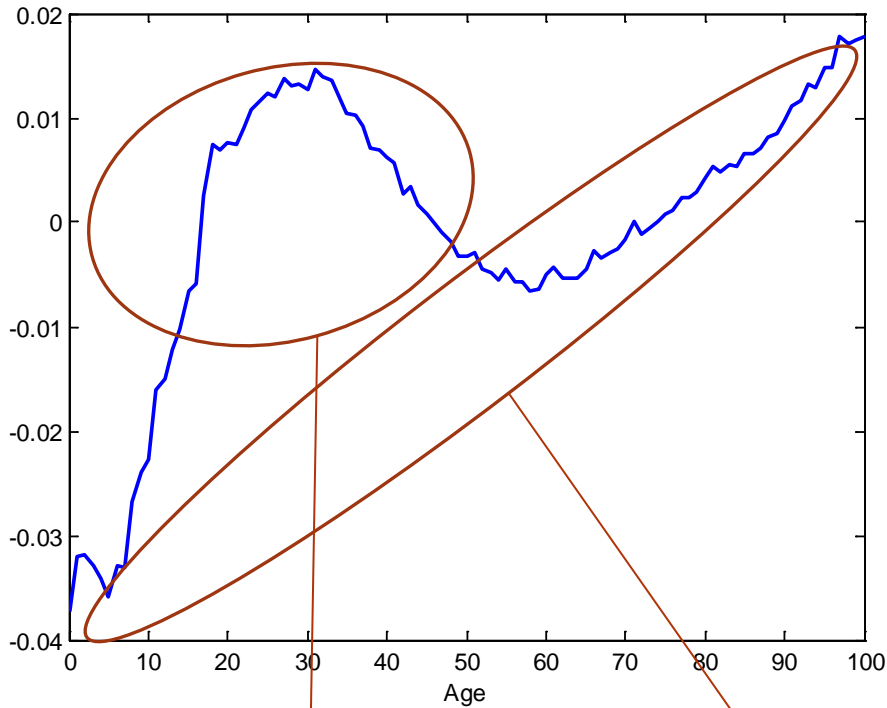
Stage 1 – Effect of 1st Term



Stage 1 - Test Goodness of Fit

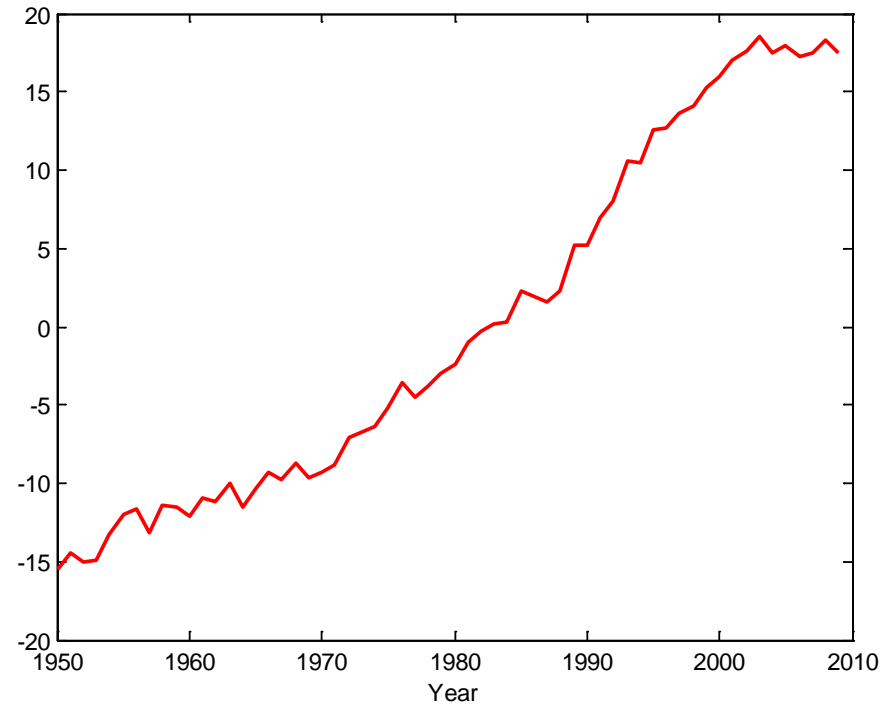


Stage 2 - Add Non-Parametric Age/Period Complementary Pair to Find 2nd Dominant Trend

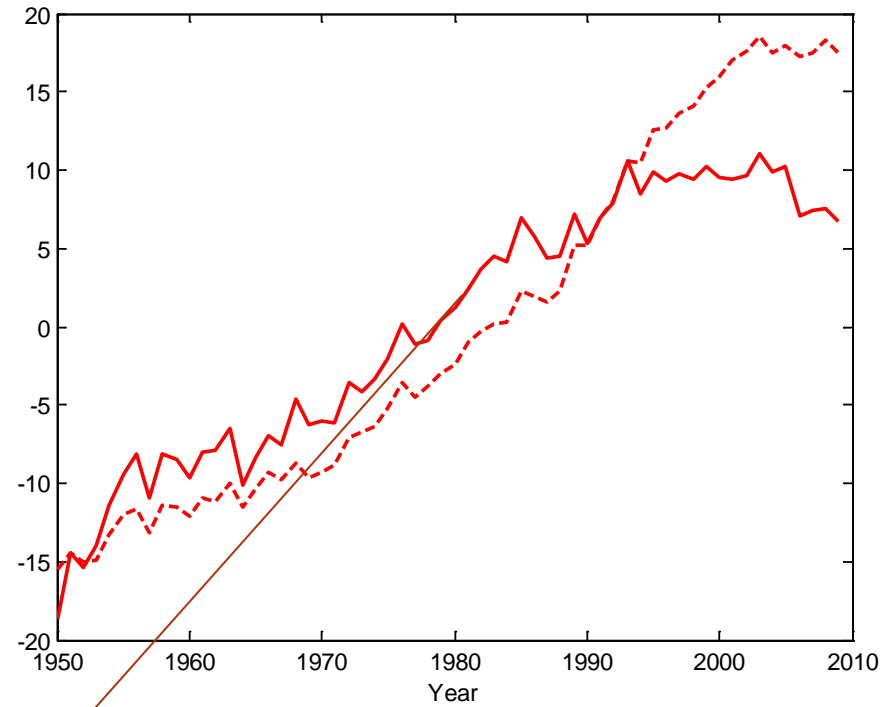
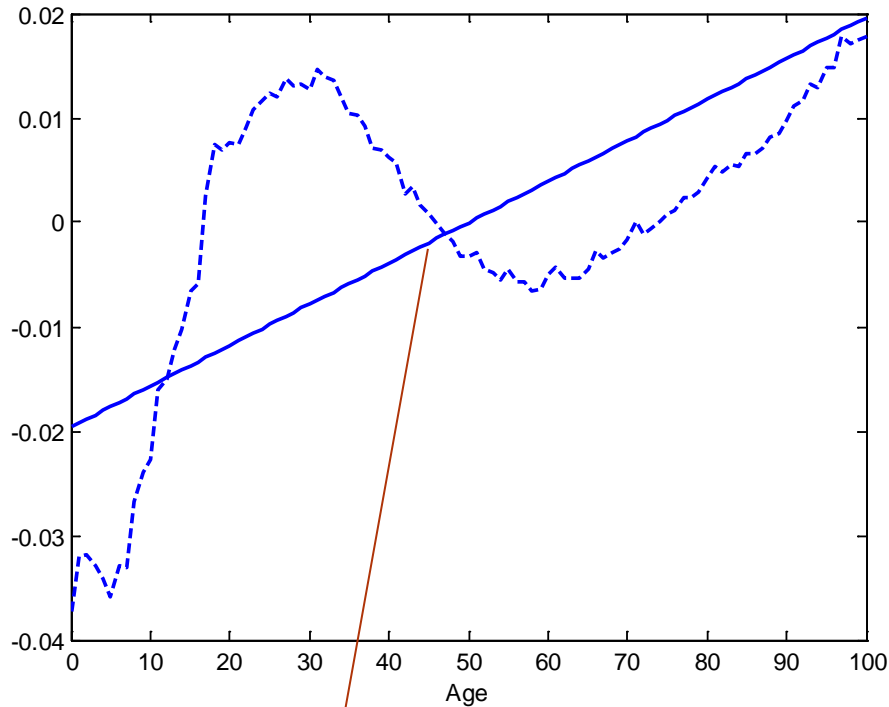


“Hump” for low ages?

Linear trend across entire age range?



Stage 2 – Find Simplest Parametric Term That Does the Same Job

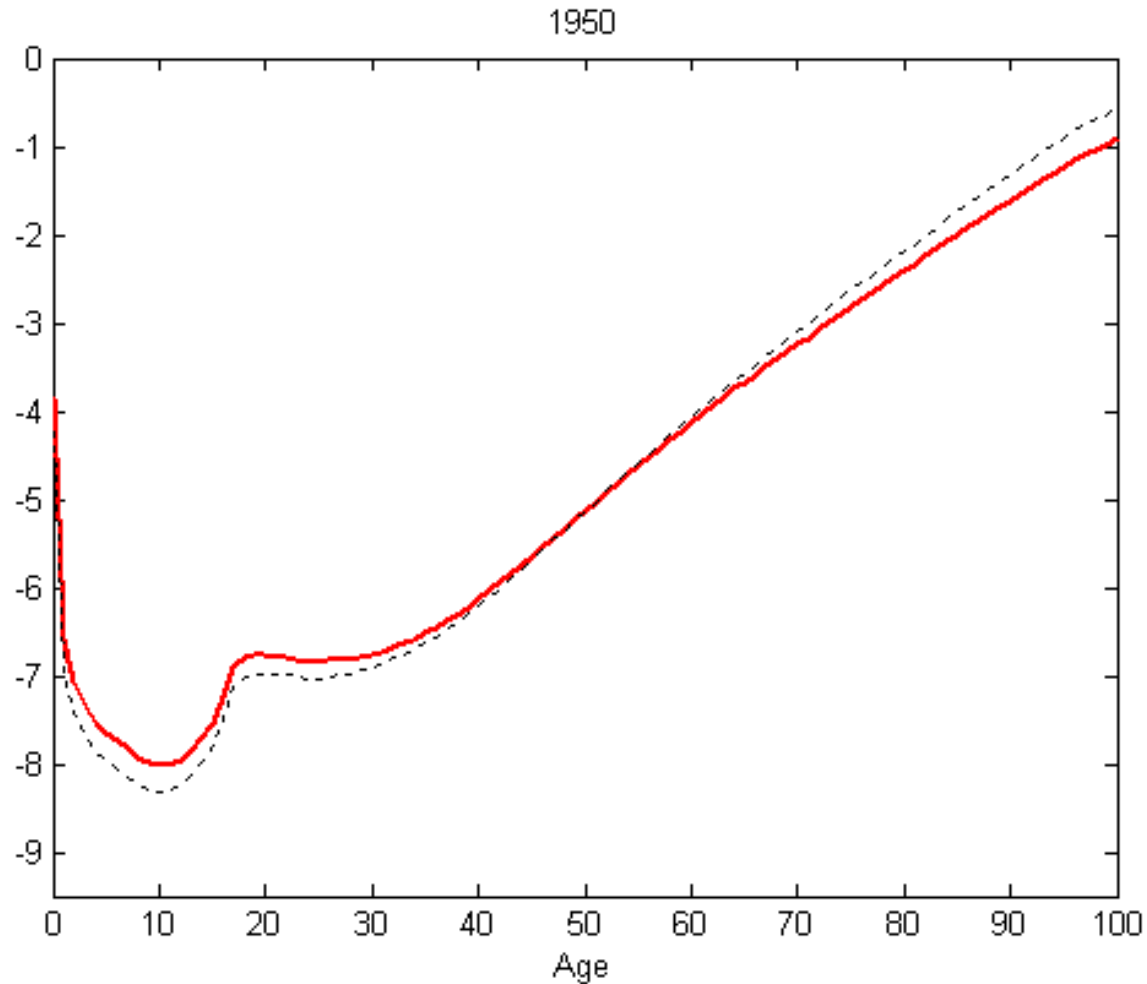


Linear age function – governs changes in the Gompertz slope for given year

“Rectangularisation” of mortality curve with time



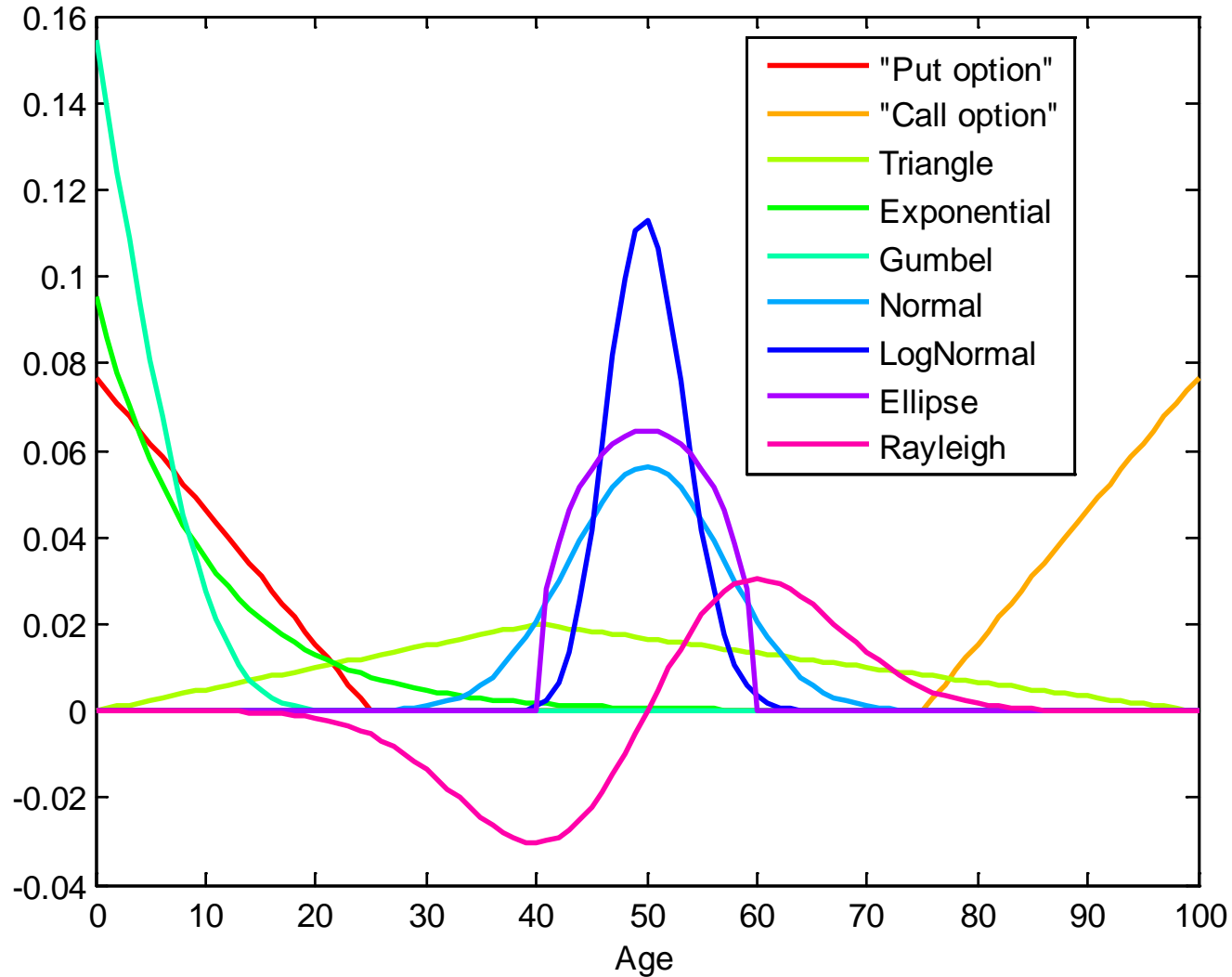
Stage 2 – Effect of 2nd Term



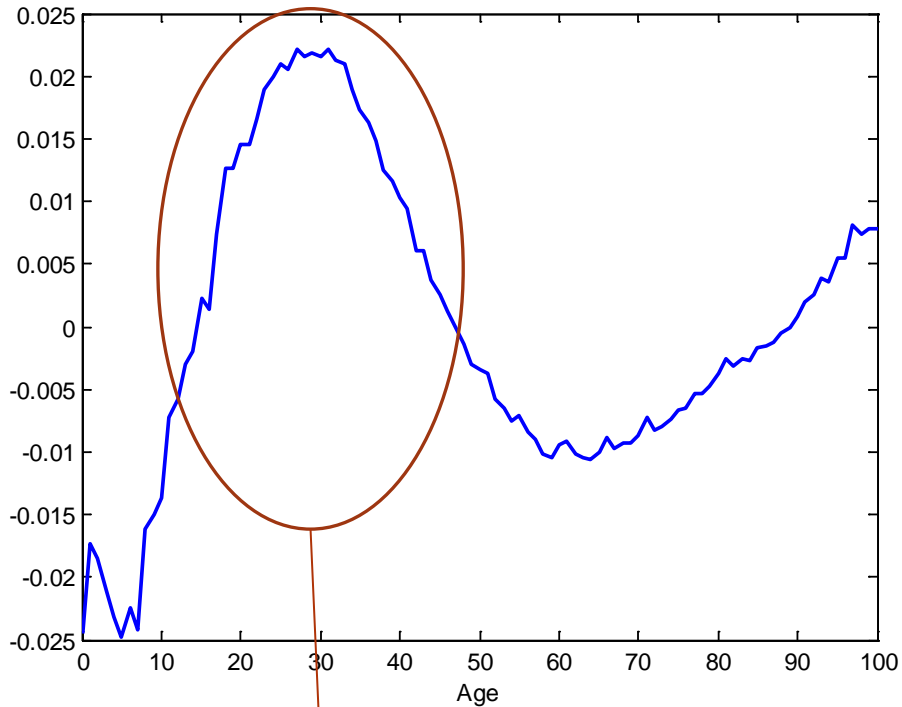
Stage 2 - Test Goodness of Fit



Toolkit of Age Functions



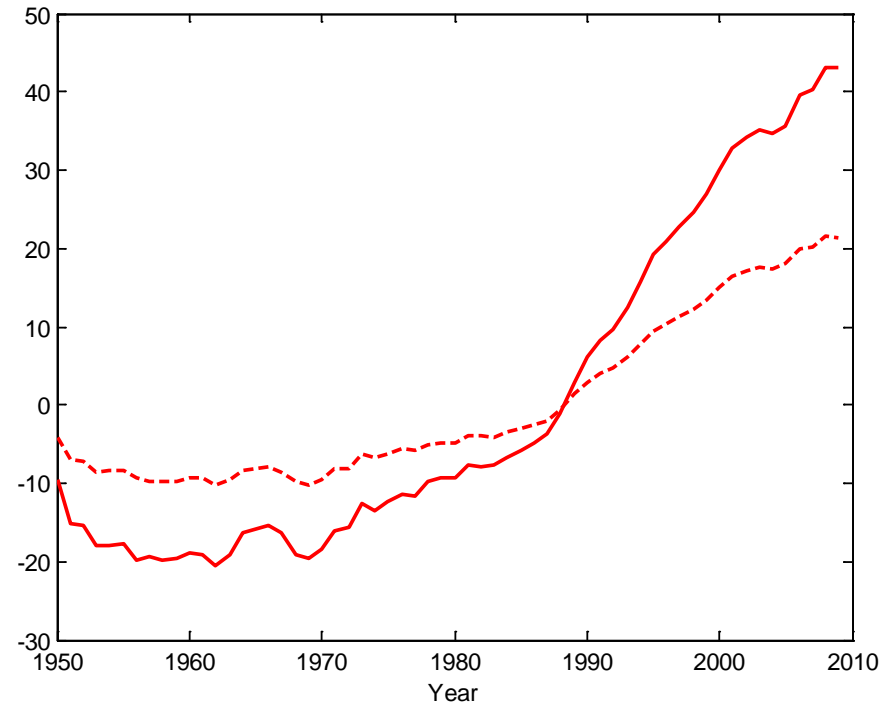
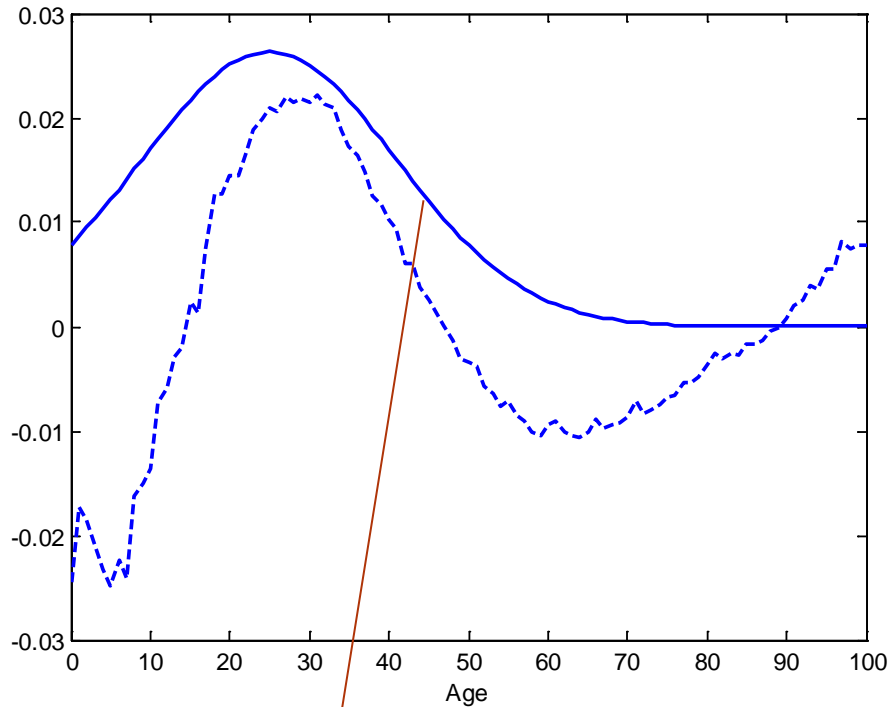
Stage 3 - Add Non-Parametric Age/Period Complementary Pair to Find 3rd Dominant Trend



“Hump” for low
ages



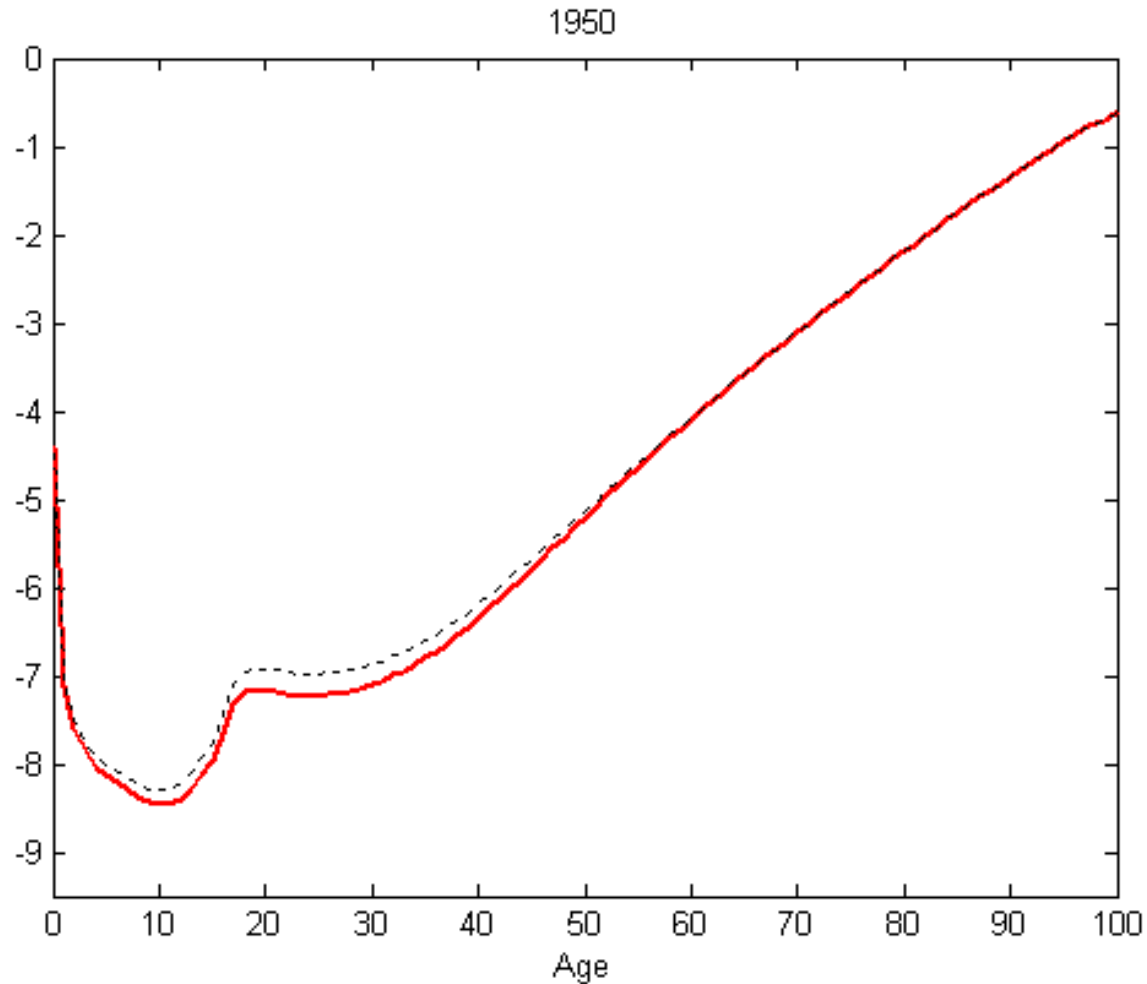
Stage 3 – Find Simple Parametric Term That Does the Same Job



Gaussian “hump” governs changes in mortality rates for young adults



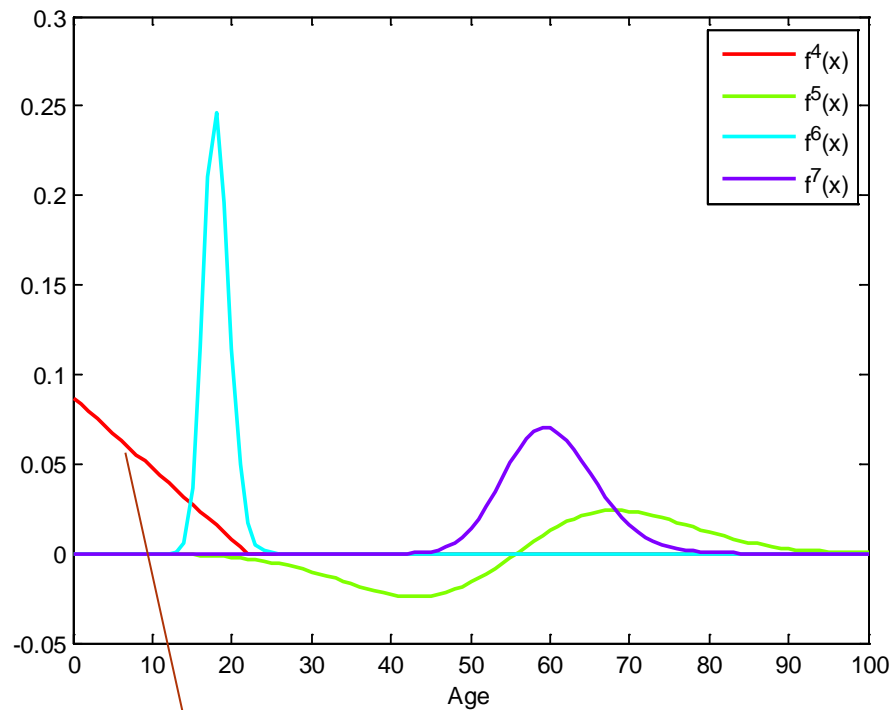
Stage 3 – Effect of 3rd Term



Stage 3 - Test Goodness of Fit



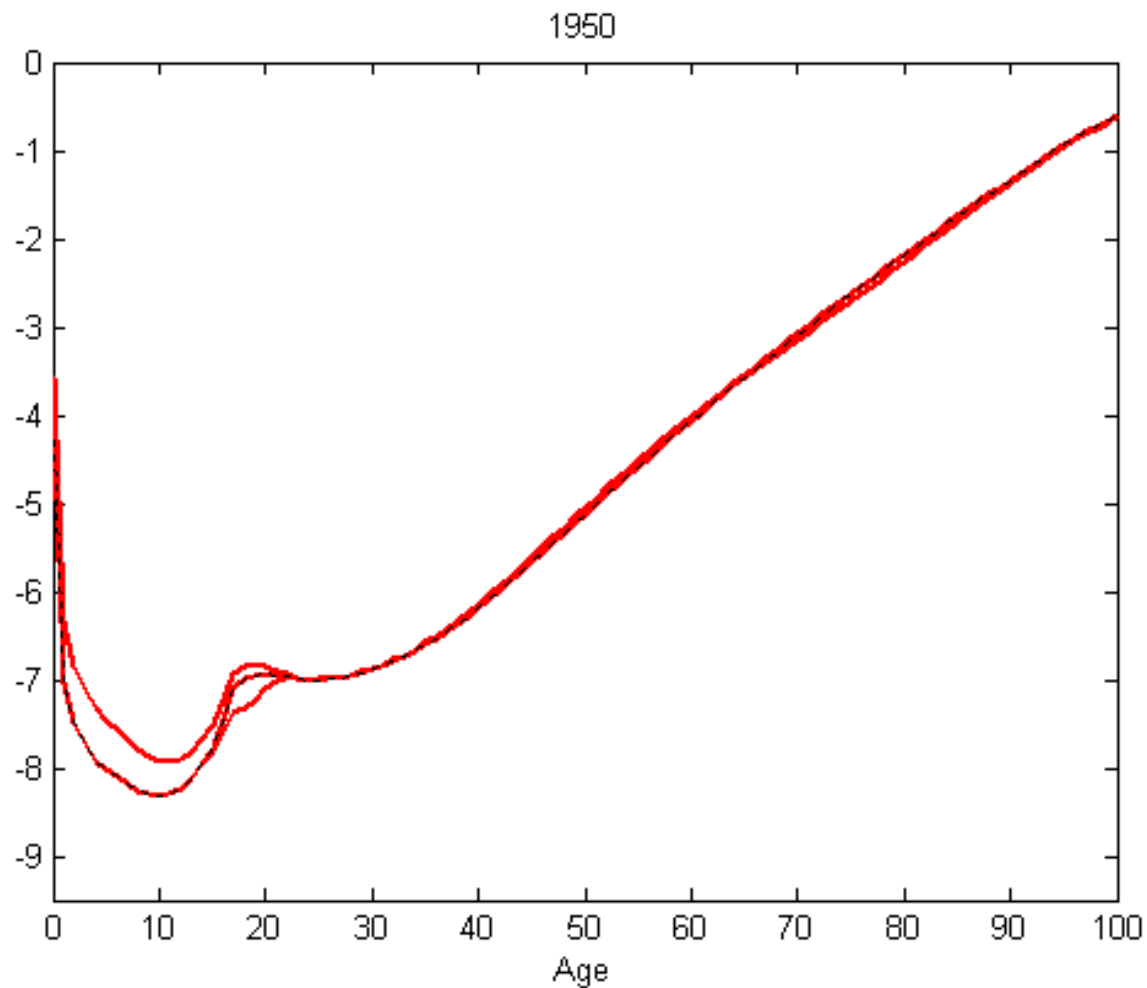
Stage 4 Onwards



“Call option”
function –
childhood
mortality



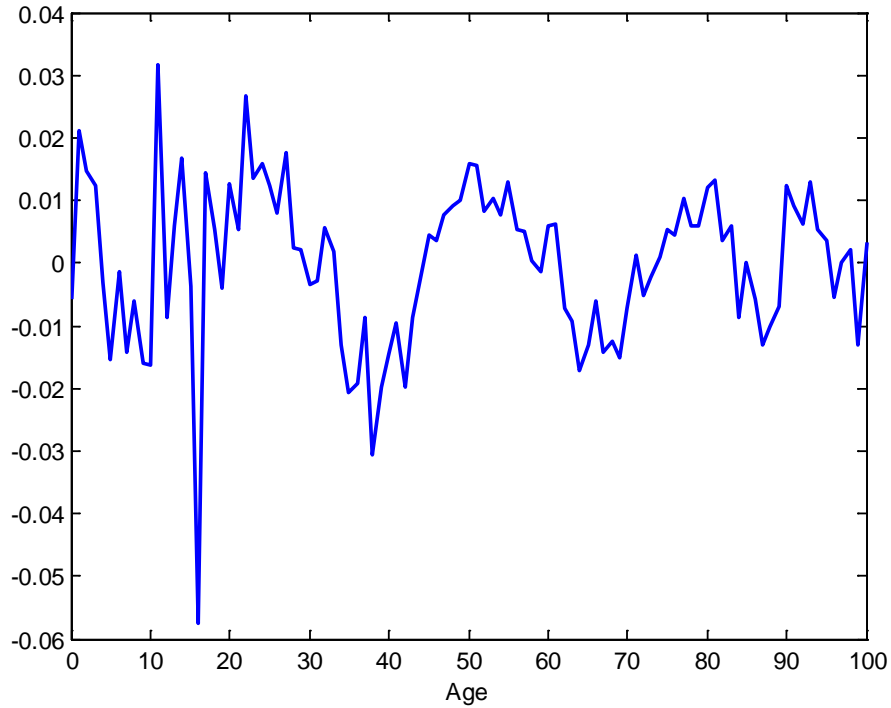
Stage 4 Onwards – Effect of Terms



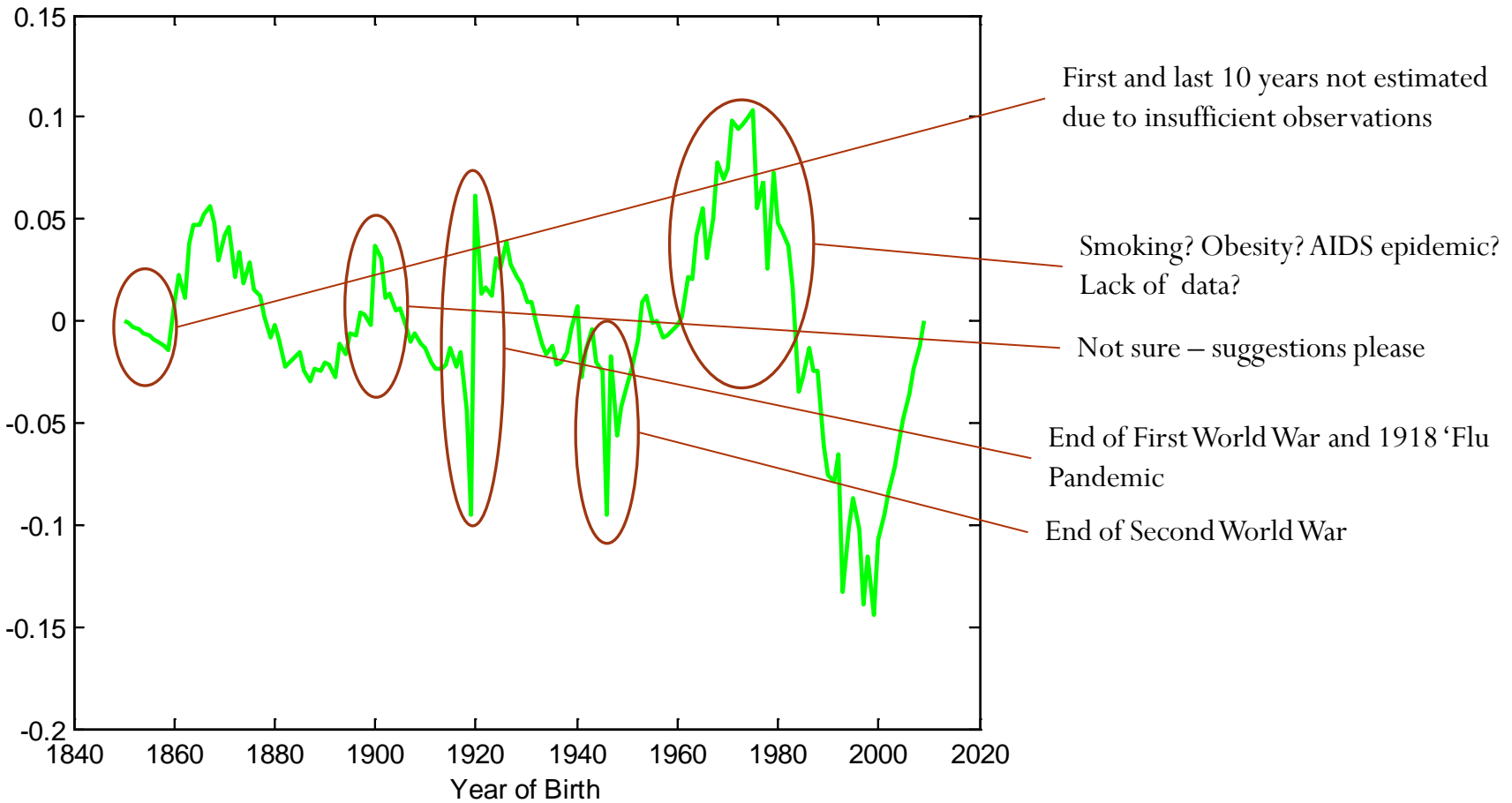
Stage 7 - Test Goodness of Fit



Stopping the Loop

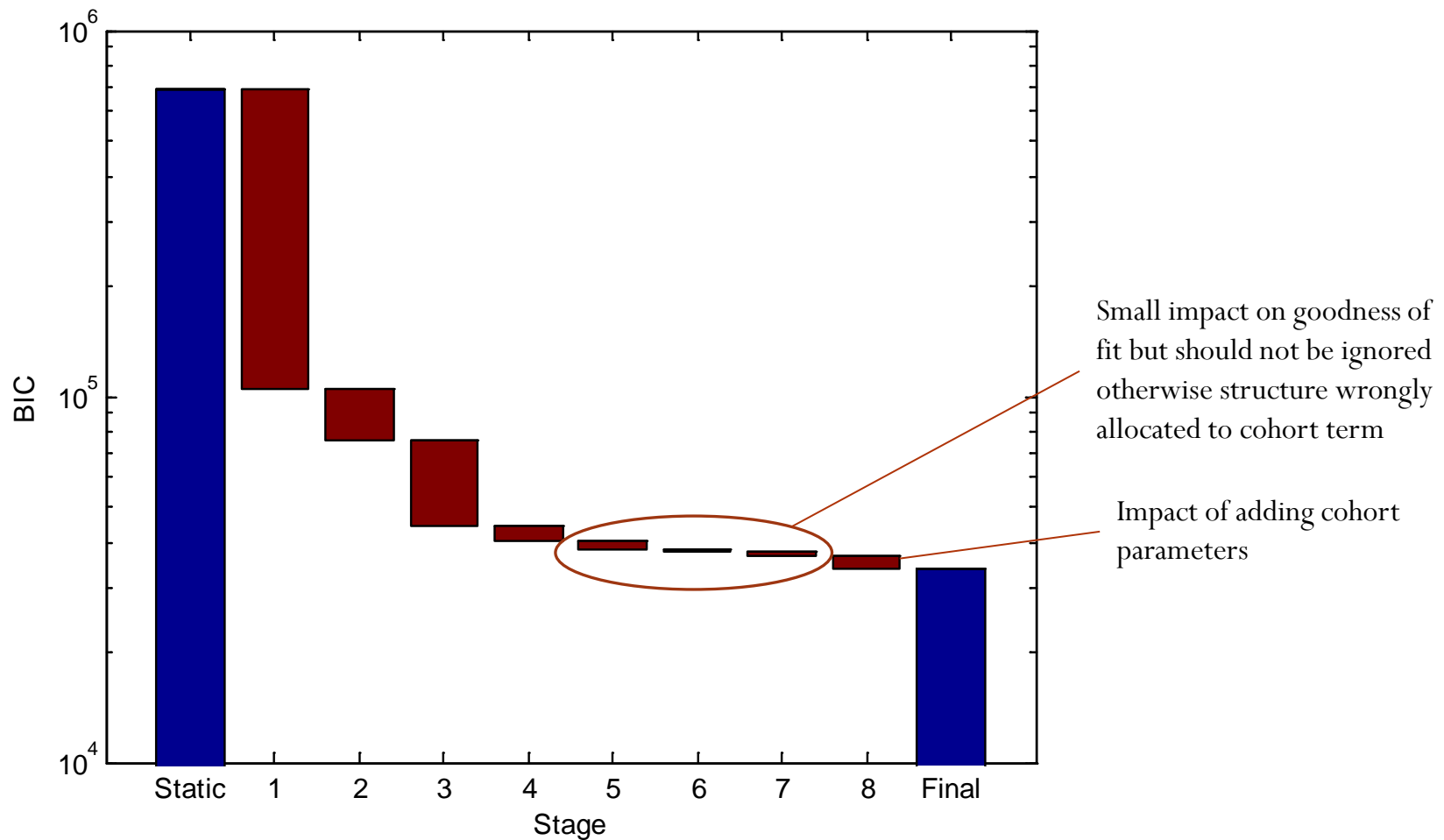


Stage 8 – Add Cohort Parameters

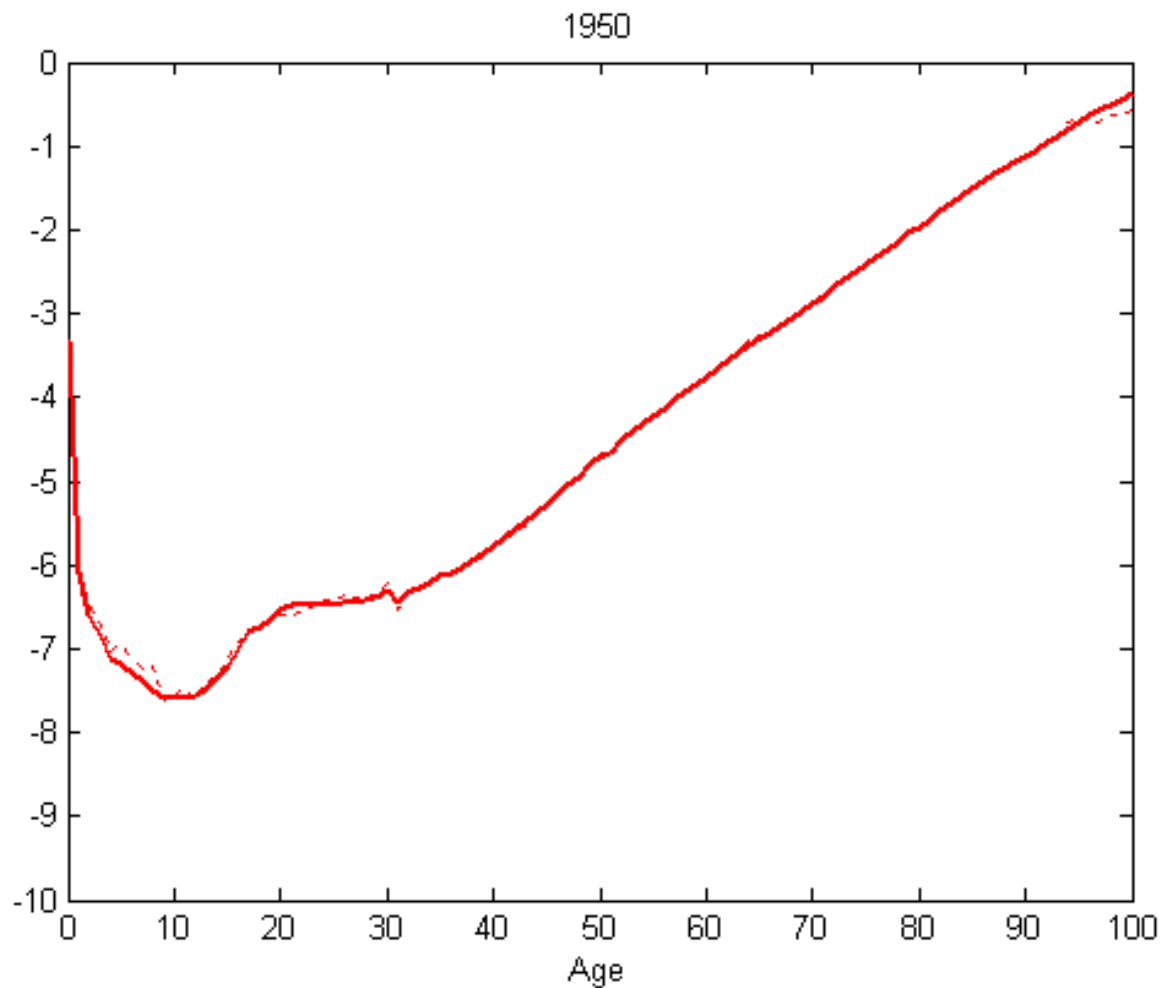




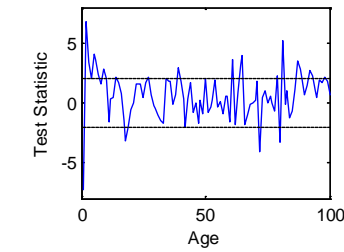
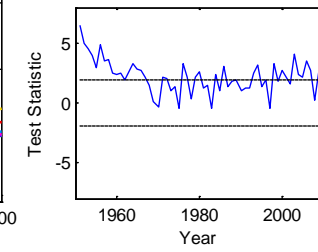
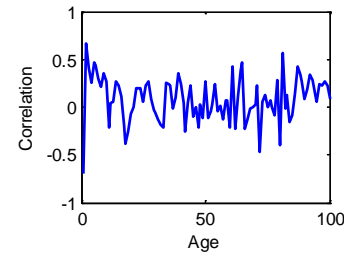
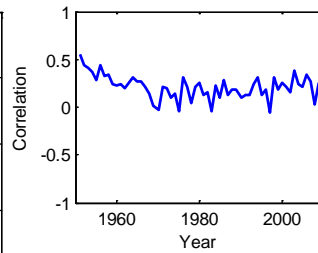
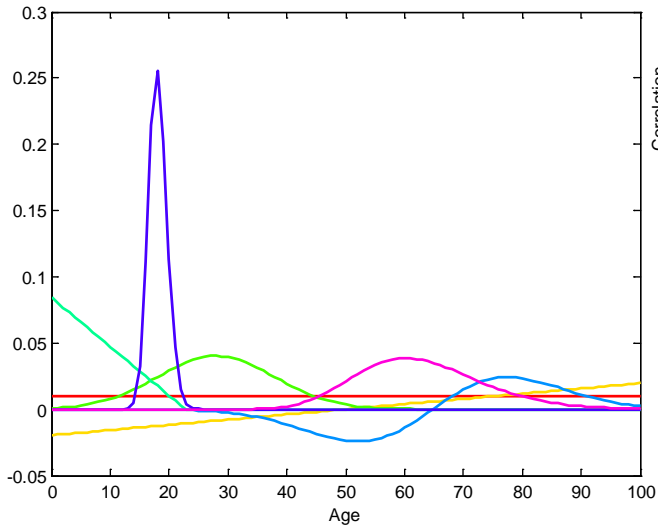
Improvements in Goodness of Fit



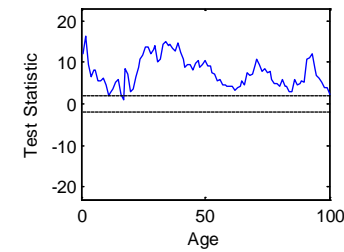
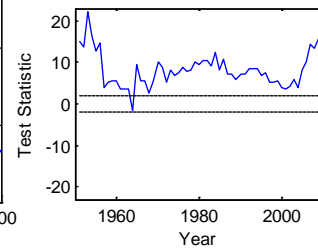
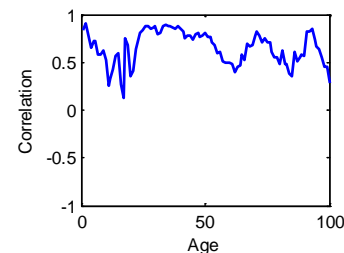
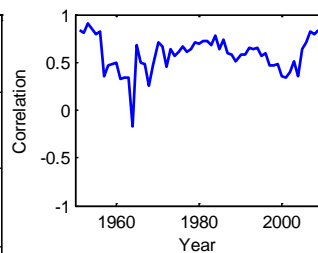
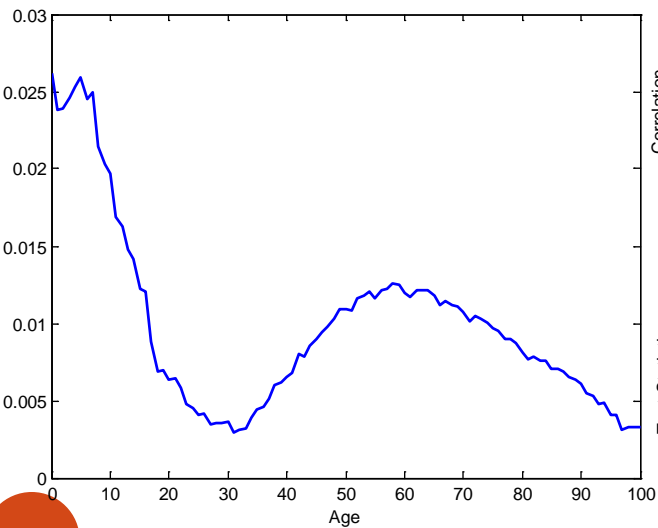
Fitted vs. Observed Mortality



Comparison with Lee-Carter

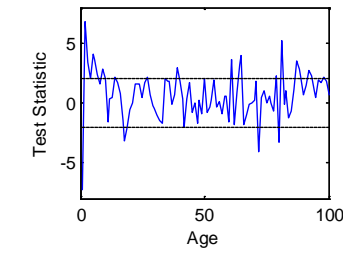
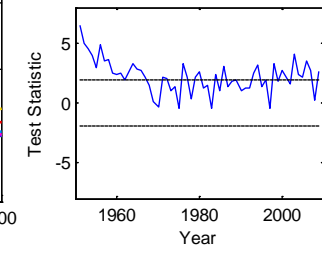
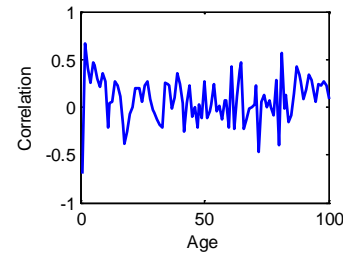
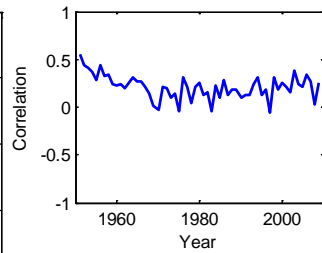
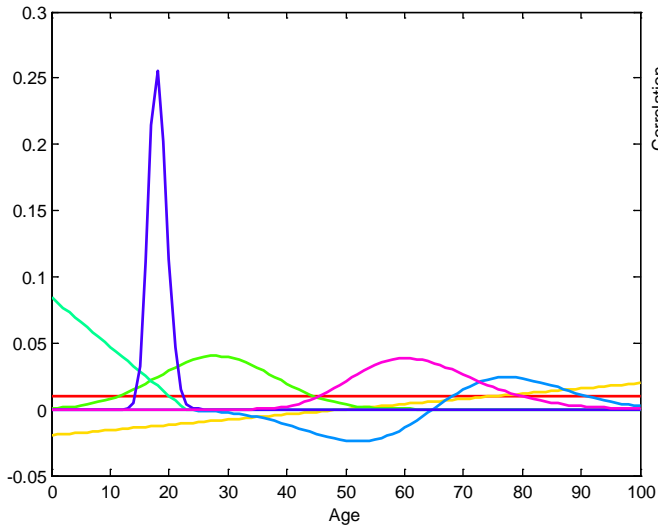


Model BIC	-3.38e+04
Res Mean	-0.01
Res SD	0.94
Res Skew	-0.03
Res Kurt	3.38

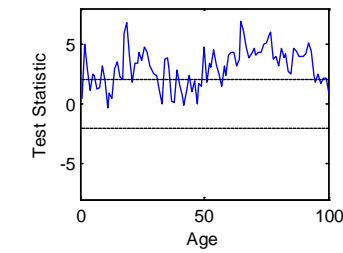
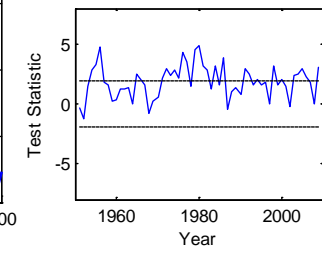
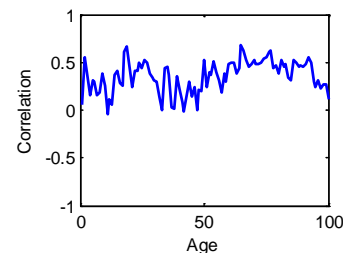
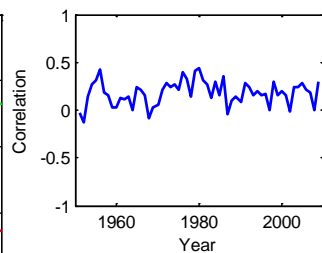
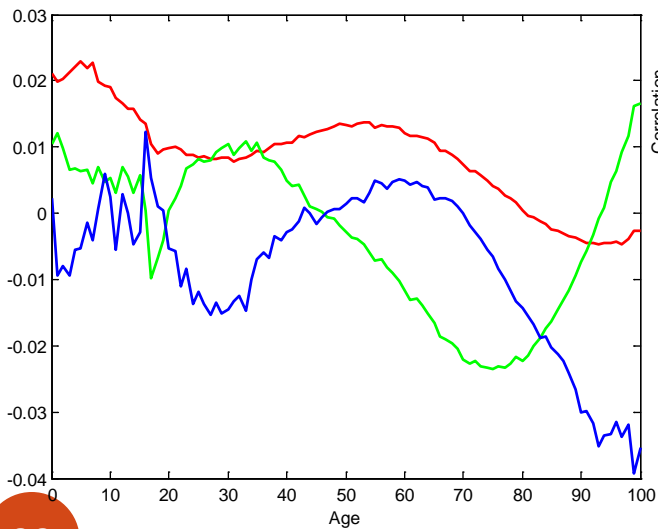


Model BIC	-5.25e+04
Res Mean	-0.02
Res SD	0.98
Res Skew	0.47
Res Kurt	9.75

Comparison with PCA

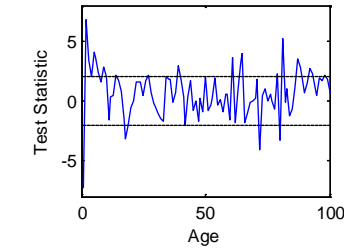
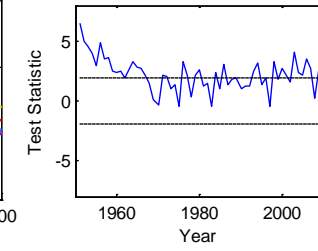
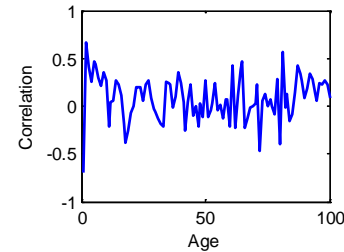
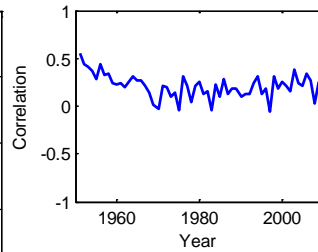
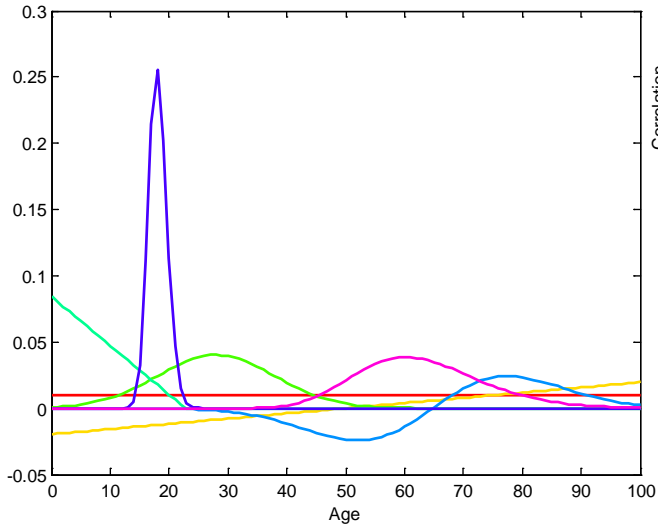


Model BIC	-3.38e+04
Res Mean	-0.01
Res SD	0.94
Res Skew	-0.03
Res Kurt	3.38

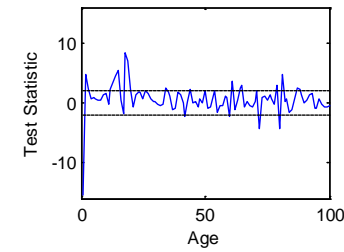
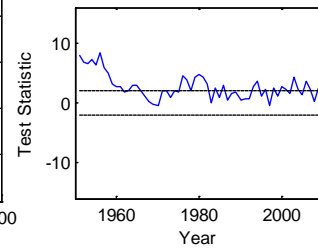
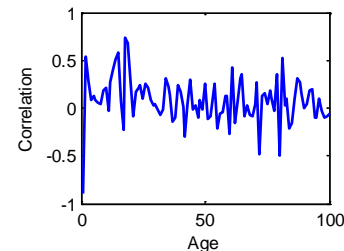
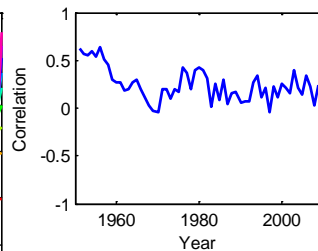
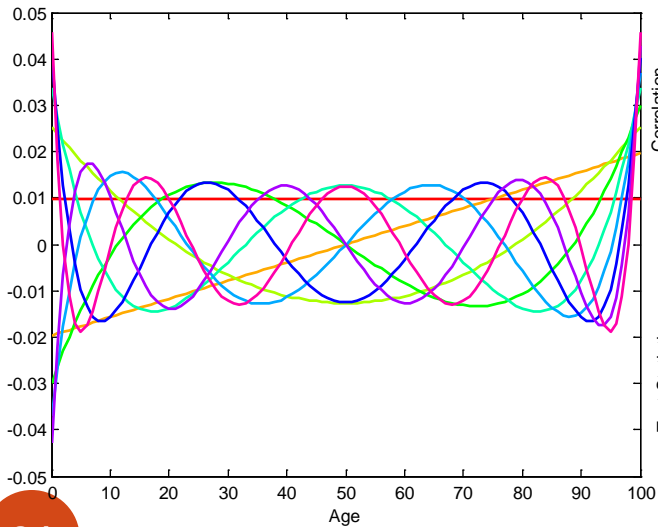


Model BIC	-3.39e+04
Res Mean	0.00
Res SD	0.94
Res Skew	0.02
Res Kurt	3.20

Comparison with Legendre Polynomials



Model BIC	-3.38e+04
Res Mean	-0.01
Res SD	0.94
Res Skew	-0.03
Res Kurt	3.38



Model BIC	-3.44e+04
Res Mean	-0.00
Res SD	0.93
Res Skew	-0.17
Res Kurt	3.69

Assessing the Model Selection Criteria

- Adequacy –
 - Each age/period term has been justified statistically as improving the fit to data
- Parsimony –
 - Fewest terms needed to capture information in the data
 - Each age function requires no more than two free parameters
 - Fewer parameters than alternative PCA and Legendre polynomial procedures
- Demographic Significance –
 - Each term we can demonstrate demographic significance in terms of causes of death at relevant ages
- Completeness –
 - Models spans entire age range
 - Models includes allowance for cohort effects and we can link features of this to history of the population

Other Data Sets and Robustness

- Have followed procedure for a female UK data since 1950
- Also obtain mode with seven age/period functions
 - Shape of age/period functions similar to those for male data
 - Sequence of selected functions different (same processes, different importance for men and women?)
- Female data slightly less easy to apply procedure for as some of the trends of comparable size and highly correlated
- Have also tested for parameter uncertainty by changing range of data, using residual bootstrapping technique (Koissi et al (2006)) and by systematically removing ages/years from dataset
- Final model gives parameter estimates which are robust under all approaches

Next Steps

- Remaining structure in the residuals:
 - Geostatistical techniques to analyse correlation structures - Débon et al. (2010)
- Projections:
 - No reliable way of projecting model with multiple time series consistently
 - Nielsen and Nielsen (2010) show that suitable time series depend upon identification constraints
 - Intuition on cohort parameters implies mean-reverting structure
 - Structural breaks and changes of trend need to be allowed for
 - Modelling two populations (e.g. UK men and women) coherently with linkages between similar age/period terms

Selected References

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Questions?

- Thank you very much for your attention and your feedback