

# Quebec Pension Plan (QPP) multi-population data analysis

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Fourteenth International Longevity Risk and Capital Market Solutions  
Conference

Amsterdam, Netherlands



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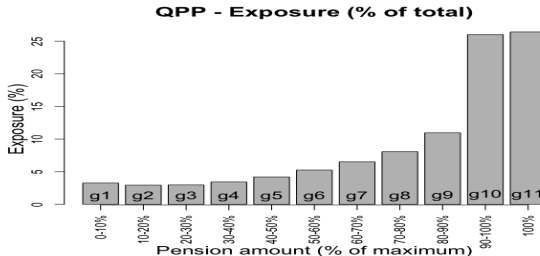
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# QPP data overview

- 11 sub-populations ordered by increasing cohort pension amount in 10% bands.
- Only contains Quebec pensioners.
- Age over 65-89, and year over 1991-2015. ( $11 \times 25 \times 25$ )

# QPP data overview (cont.)

- Males

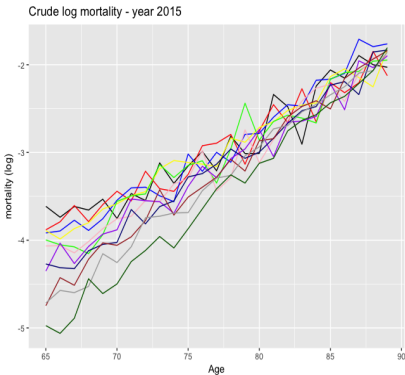
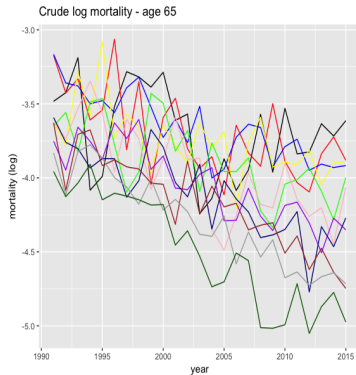


- Females



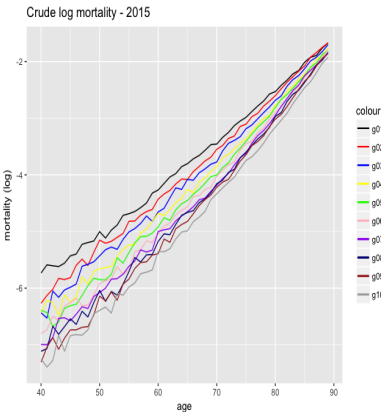
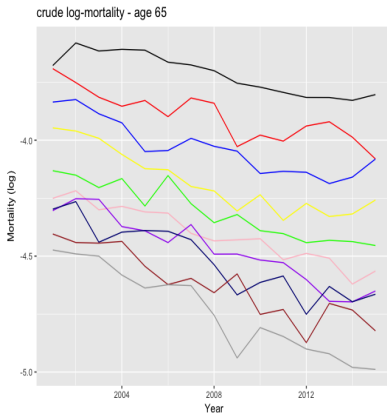
# QPP data overview (cont.)

## - Group-wise crude death rates (log-scale):



# QPP data overview (cont.)

- Comparison with **England IMD** (larger sample size, with groups evenly splited)



## - Age-Standardized Mortality Rate (ASMR)

- ASMR is a weighted average of the crude death rates over a defined age range, for certain specific calendar year  $t$ .
- $E_x^s$  is the 'standard population' at age  $x$  (from European Standard Population, calibrated in 2013).
- $m_{tx}$  is crude death rate.
- 

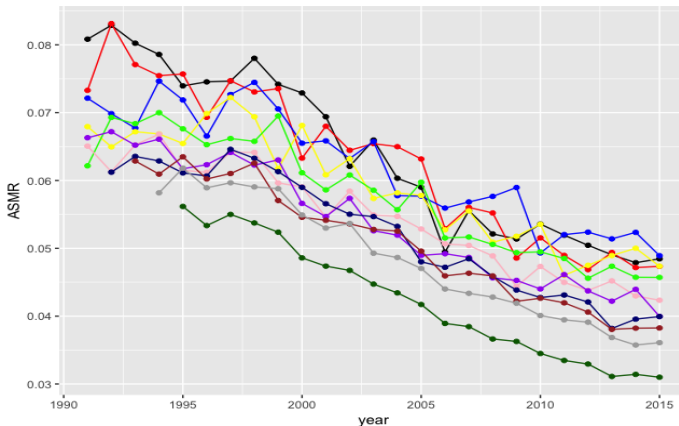
$$ASMR(t) = \frac{\sum_x m_{tx} E_x^s}{\sum_x E_x^s}$$

- Use of ASMR:
  - comparison of mortality over different populations;
  - assessment of mortality term structure;
  - assessment of signal-to-noise ratio.

# QPP data overview (cont.)

## - ASMR of QPP males over age 65-89:

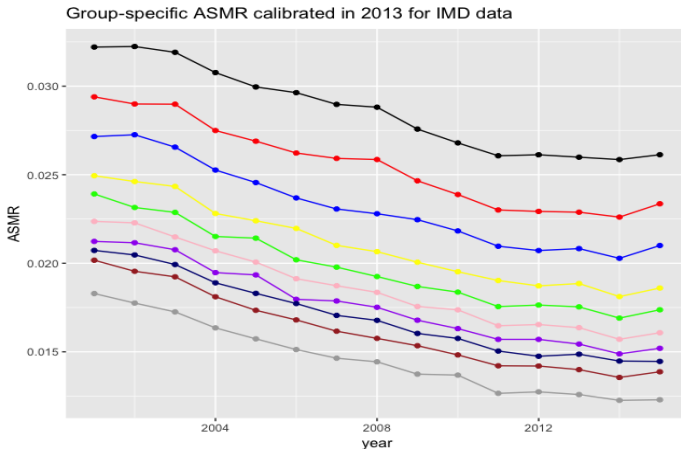
Group-specific ASMR calibrated in 2013 for QPP males data





# QPP data overview (cont.)

- (For comparison) ASMR of England IMD:



# QPP data overview (cont.)

- ASMR is smoother than the crude death rates, but still quite volatile for QPP males.
- Group 10 and 11 (larger size) are smoother than others.
- Groups with higher pension tends to have lower mortality.
- QPP applies different grouping methodology (pension level) from England IMD (deprivation index) - less powerful predictor.

# Model specification

m1  $\log m_{xti} = \alpha_{xi} + \beta_{xi}^1 \kappa_{ti}^1 + \beta_{xi}^2 \kappa_{ti}^2$  (Renshaw and Haberman, 2003)

m2  $\log m_{xti} = \alpha_{xi} + \beta_{xi}^1 \kappa_{ti}^1 + \beta_x^2 \kappa_{ti}^2$

m3  $\log m_{xti} = \alpha_{xi} + \beta_x^1 \kappa_t^1 + \beta_{xi}^2 \kappa_{ti}^2$  (Li and Lee, 2005)

m4  $\log m_{xti} = \alpha_{xi} + \beta_{xi}^1 \kappa_{ti}^1$  (Lee-Carter, 1992)

m5  $\log m_{xti} = \alpha_{xi} + \beta_x^1 \kappa_{ti}^1 + \beta_x^2 \kappa_{ti}^2$  (CAE model by Kleinow, T, 2014)

m6  $\log m_{xti} = \alpha_x + \beta_x^1 \kappa_{ti}^1 + \beta_x^2 \kappa_{ti}^2$  (CAE model with common  $\alpha_x$ )

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- $\alpha$ ,  $\beta$  and  $\kappa$  are stochastic parameters capturing age/period effect.
- $\alpha$  provides a form of base mortality table (while  $\kappa$  is zero).
- $\beta$  determines the relative rates of mortality improvement at different ages.

# Model specification

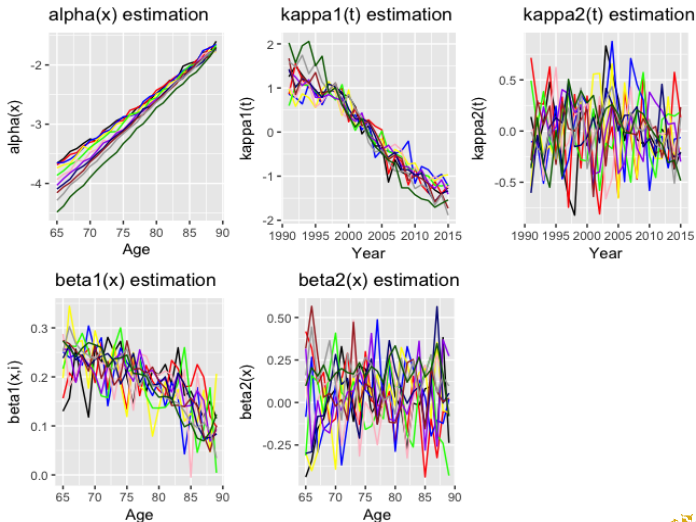
- m7  $\log m_{xti} = \alpha_{xi} + \kappa_{ti}^1 + (x - \bar{x})\kappa_{ti}^2$  (Plat, 2009)
- m8  $\log m_{xti} = \alpha_x + \kappa_{ti}^1 + (x - \bar{x})\kappa_{ti}^2$  (Plat model with common  $\alpha_x$ )
- m9  $\log m_{xti} = \alpha_{xi} + \kappa_t^1 + (x - \bar{x})\kappa_{ti}^2$  (Plat model with common  $\kappa_t^1$ )
- m10  $\log m_{xti} = \alpha_{xi} + \kappa_{ti}^1 + (x - \bar{x})\kappa_t^2$  (Plat model with common  $\kappa_t^2$ )
- m11  $\log m_{xti} = \alpha_{xi} + \kappa_t^1 + (x - \bar{x})\kappa_t^2$  (Plat model with common  $\kappa_t^1$  and  $\kappa_t^2$ )

# Model specification (cont.)

- m1 is the 'basis' with most specified structure among all others.
- All other models are simplifications of m1.
- Parameters are estimated by Poisson assumption on number of deaths with Maximum Log-likelihood Estimation (MLE).

# Parameter estimation and Model selection (cont.)

## - Model m1 - estimated parameters (males)



# Parameter estimation and Model selection (cont.)

- Model m1:

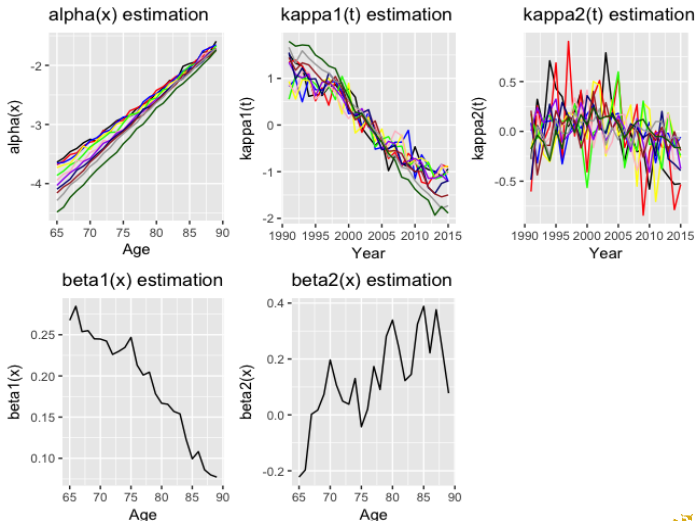
$$\log m_{xti} = \alpha_{xi} + \beta_{xi}^1 \kappa_{ti}^1 + \beta_{xi}^2 \kappa_{ti}^2$$

- Group specific  $\alpha_{xi}$  gives observable group rankings.
- $\kappa_{ti}^1$  and  $\beta_{xi}^1$  have decreasing pattern for all groups.
- $\kappa_{ti}^2$  and  $\beta_{xi}^2$  are quite volatile.



# Parameter estimation and Model selection (cont.)

- Model m5 - estimated parameters (males)



# Parameter estimation and Model selection (cont.)

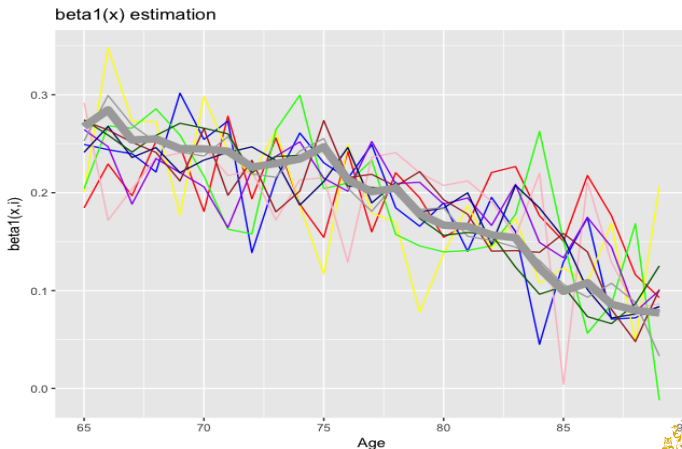
- Model m5:

$$\log m_{xti} = \alpha_{xi} + \beta_x^1 \kappa_{ti}^1 + \beta_x^2 \kappa_{ti}^2$$

- Group specific  $\alpha_{xi}$  gives observable group rankings.
- $\kappa_{ti}^1$  has similar decreasing pattern for all groups.
- $\kappa_{ti}^2$  is quite volatile.
- $\beta_x^1$  decreases over age and is less volatile than  $\beta_x^2$ .

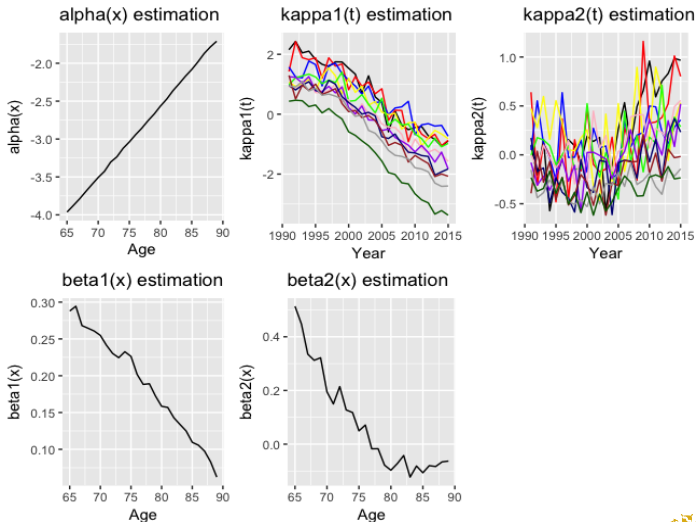
# Parameter estimation and Model selection (cont.)

- Pattern of  $\beta^1$  - model m5 (common - the grey fat solid line) and m1 (group-specific)



# Parameter estimation and Model selection (cont.)

## - Model m6 - estimated parameters (males)



# Parameter estimation and Model selection (cont.)

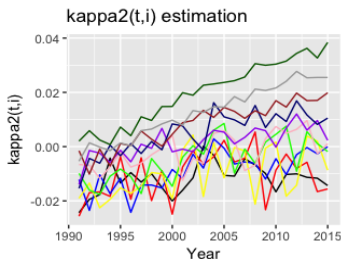
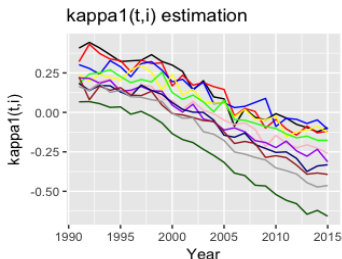
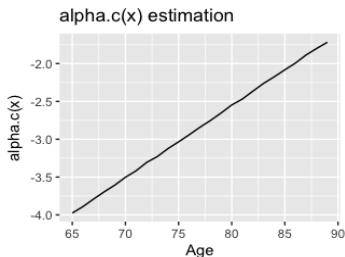
- Model m6:

$$\log m_{x_{ti}} = \alpha_x + \beta_x^1 \kappa_{ti}^1 + \beta_x^2 \kappa_{ti}^2$$

- As  $\alpha_x$  is common, variations between subgroups are captured by  $\kappa_{ti}^1$  and  $\kappa_{ti}^2$ .
- Group 11 stands clear of others in terms of  $\kappa_{ti}^1$ .
- $\beta_x^1$  and  $\beta_x^2$  decreases over age,  $\beta_x^2$  is smoother than under m5.

# Parameter estimation and Model selection (cont.)

## - Model m8 - estimated parameters (males)



# Parameter estimation and Model selection (cont.)

- Model m8:

$$\log m_{x_{ti}} = \alpha_x + \kappa_{ti}^1 + \kappa_{ti}^2(x - \bar{x})$$

- As  $\alpha_x$  is common, variations between subgroups are captured by  $\kappa_{ti}^1$  and  $\kappa_{ti}^2$ .
- Group 11 stands well below and above others for  $\kappa_{ti}^1$  and  $\kappa_{ti}^2$  respectively.

# Parameter estimation and Model selection (cont.)

## - Model selection criteria: log-likelihood and BIC: males

Bayes Information Criterion (BIC) is a statistic based on log-likelihood that penalises over-parameterized models and is used as a purely numerical criterion for selecting out the best model (m8).

Model	log-likelihood	# parameters	df	BIC
m1	-22,252.44	1375	1331	56,265.12
m5	-22,628.04	875	851	52,775.22
m6	-22,771.52	625	621	51,029.98
m8	-22,867.36	575	573	50,797.55



# Parameter estimation and Model selection (cont.)

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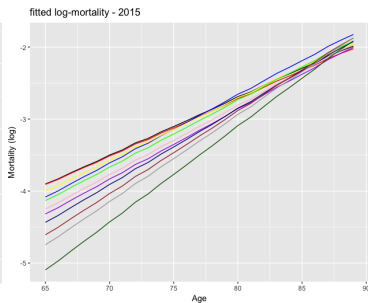
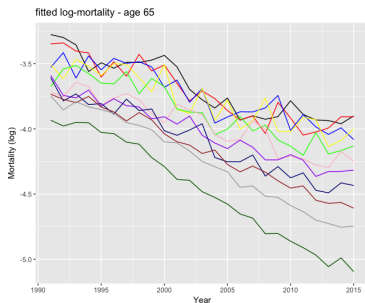
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# Diagnostic on fitting results

- m8 has fewest parameters and better BIC than other three.
- More parameters improves log-likelihood but is also penalized for over-parameterization.
- Greater complexity does not necessarily improve fitting significantly.
- Additional diagnostic is also required for selection.

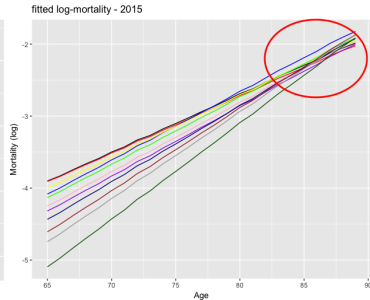
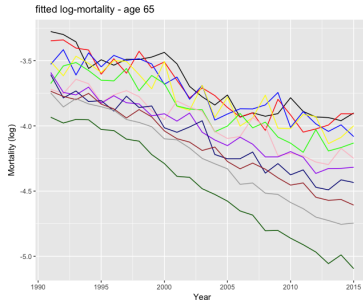
# Diagnostic on fitting results

- Fitted mortalities (log-scale) from model m8 (males)



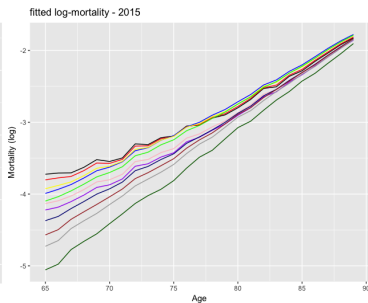
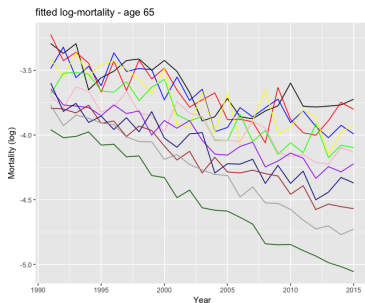
# Diagnostic on fitting results

- Fitted mortalities (log-scale) from model m8 (males)



# Diagnostic on fitting results

- Fitted mortalities (log-scale) from model m6 (males)



# Diagnostic on fitting results

## - Standardized Residuals

$$Z_{txi} = \frac{D_{txi} - E_{txi} \hat{m}_{txi}}{\sqrt{E_{txi} \hat{m}_{txi}}}$$

- Measures standardized difference between crude and estimated figures.
- Not affected by absolute scale of observations.
- Well-fitted model is expected to have random standardized residuals.

# Diagnostic on fitting results

- Standardized residuals from m6: QPP males



# Diagnostic on fitting results

- Standardized residuals from m8: QPP males





# Diagnostic on fitting results

- Both m6 and m8 have quite random standardized residuals.
- There is no significant non-random cluster along x-axis (year), y-axis (age) or diagonal (cohort).
- m6 doesn't have significant crossover in fitted mortality curves. m8 has crossovers at high ages.
- m6 is selected as the most suitable model for QPP males.

# Cluster analysis on QPP data

- QPP has relatively small population size.
- Subpopulations are not evenly grouped.
- Crude mortalities are quite volatile.
- Some adjacent groups typically have quite similar levels of mortality.
- We consider to re-cluster the QPP dataset.

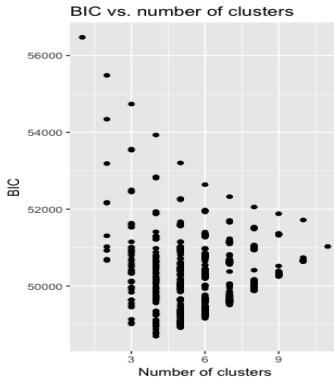
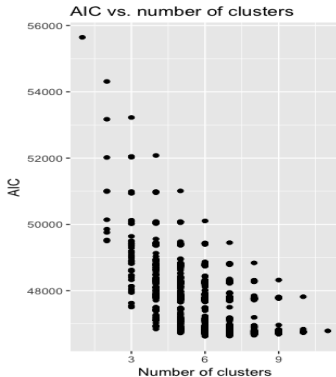
# Cluster analysis on QPP data

## - Algorithm:

- 1 Restructure the data by combining neighbouring groups into clusters. Each cluster could contain  $1, 2, \dots, 11$  groups.
- 2 We obtain new restructured datasets with  $\leq 11$  groups.
- 3 There are 1,024 different combinations in total.  $(\sum_{i=0}^{11-1} C_{11-1}^i)$
- 4 Fit underlying models to each reclustered dataset.

# Cluster analysis on QPP data

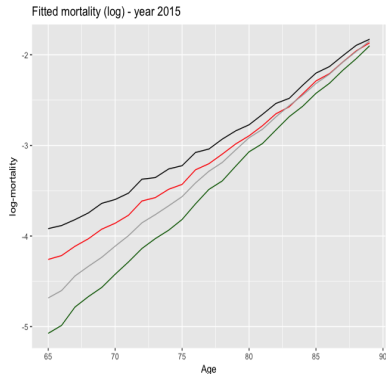
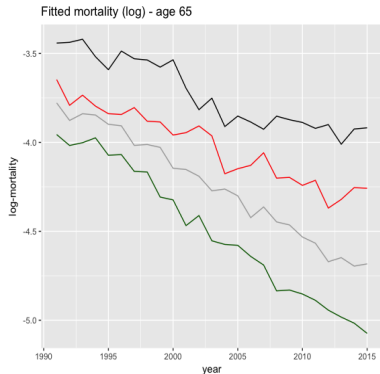
- AIC and BIC for all 1,024 cluster combinations fitted for model m6:



BIC is **48,694.69** under the optimized scenario (used to be 51,029.98).

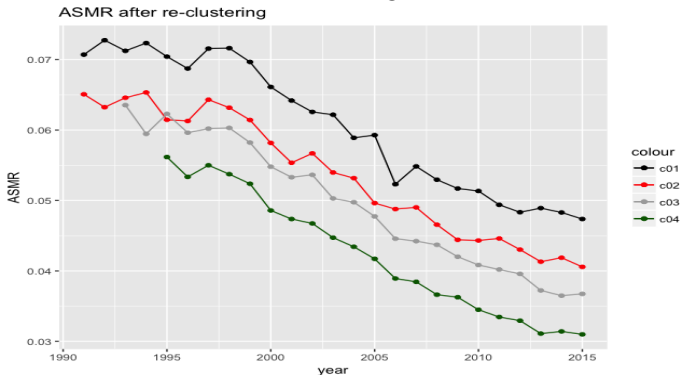
# Cluster analysis on QPP data

- Fitted mortalities (log-scale) from m6 after re-clustered into 4 groups.



# Cluster analysis on QPP data

- ASMR of QPP males after re-clustering:



# Cluster analysis on QPP data

- Conclusion from cluster analysis:
  - All models suggest the same optimal clustering by BIC - with 4 clusters:
    - Cluster 1: group 1-5;
    - Cluster 2: group 6-8;
    - Cluster 3: group 9 and 10;
    - Cluster 4: group 11.
  - Volatilities are reduced significantly.
  - It enables us to see more clearly the different trends of clusters.

# Summary

- For volatile population, models with simpler structure fits better, i.e. model m6 and m8 over m1.
- Besides quantitative criteria, qualitative criteria like graphical diagnostics are the same important.
- Clustering improves fitting quality and signal-to-noise ratio.
- Future researches: Smoothing of modelling results; More detailed cluster analysis; Long-term mortality projection.



# ANY QUESTIONS ?