Modeling age-space mortality dynamics in small areas

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SCOR chair on mortality research meeting, Paris

8 November 2024





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8 November 2024 1 / 24

Motivation



Figure: Observed mortality. Females, 2015. Left: London Metropolitan Area. Center: single London borough. Right: single Lower layer Super Output Area. *Note:* vertical bars at the bottom of the graphs indicate zero observed deaths.

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Motivation



Figure: Life expectancy at birth by Lower layer Single Output Area calculated from observed mortality. Females, 2015.

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- Model mortality in a very large number of very small areas.
- For example, 4835 sub-municipal small areas (Lower layer Super Output Areas (LS0As)) in a metropolitan area (London) and 91 age categories → 439.985 observations.

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Understanding fine-grain inequalities in mortality

- Small populations lead to substantial fluctuations in observed death counts, creating very noisy mortality signal.
- At the same time, underlying mortality risk can differ substantially in space.
- Difficult to distinguish between real differences and random variation in risk of death.

Existing approaches

- Bayesian "**principal components**" or "**SVD** (Singular Value Decomposition)" model (Alexander, Zagheni, and Barbieri 2017):
 - Model area-specific mortality schedule as a linear combination of several principal components of a matrix of standard mortality schedules (e.g. mortality rates over age and time for the whole country).
 - Pool information with hierarchical structure (counties within a state in US) or across space (autoregressive prior).

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- **TOPALS** model (Gonzaga and Schmertmann 2016; Schmertmann and Gonzaga 2018):
 - Model area-specific mortality schedule as piecewise-linear deviations from a standard schedule.
 - Difference penalty on coefficients of deviations to ensure stability of estimates.

Existing approaches

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 - Model area-specific mortality schedule as a linear combination of several principal components of a matrix of standard mortality schedules (e.g. mortality rates over age and time for the whole country).
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- **D-Splines** (Schmertmann 2021):
 - Model area-specific mortality schedule as a linear combination of cubic splines, penalized with a custom-made penalty derived from a standard schedule of mortality.

- Presence of a standard schedule that must be estimated from real data, but uncertainty surrounding the estimation and incorporation of the standard is not taken into account (Alexander, Zagheni, and Barbieri 2017; Schmertmann and Gonzaga 2018; Gonzaga and Schmertmann 2016).
- Spatial structure of data is not exploited (Schmertmann and Gonzaga 2018; Gonzaga and Schmertmann 2016; Schmertmann 2021).

Data requirements

- Deaths and exposures for a single year for multiple spatial units.
- Enough deaths in the sum of spatial units to distinguish a mortality ٠ signal.

Model assumptions

- Mortality in a single area:
 - resembles the mortality schedule of the total of all area being studied,
 - deviates from the standard schedule smoothly in age and space,
 - can show breaks from the overall smooth pattern in space.
- Standard schedule and deviations from standard are estimated simultaneously \rightarrow uncertainty in standard is accounted for.
- Spatial structure exploited to borrow strength across areas.

Figure: Data inputs



 Data: two m × n matrices, m ages, n spatial units: deaths Y, exposures E. Centroids of territorial units serve as spatial information.

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Methodology

• We model deaths in a Poisson setting:

$$\ln E[\mathbf{y}] = \ln(\mathbf{e}) + \ln(\mu) = \ln(\mathbf{e}) + \eta.$$
(1)

Goal: model

$$\eta = \begin{bmatrix} \eta^{0} \\ \eta_{1} \\ \eta_{2} \\ \vdots \\ \eta_{j} \\ \vdots \\ \eta_{n} \end{bmatrix} = \mathbf{X}\theta \text{ where } \begin{cases} \eta^{0} \text{ a common age schedule,} \\ \eta_{j} = \eta^{0} + \delta_{j} + \gamma_{j} \\ = \text{area } j \text{ schedule} \end{cases}$$
(2)

- δ_j deviations from standard that vary smoothly in age and space
- γ_j area-specific intercepts that allow for unsmooth variation.

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10 / 24

Building the model matrix

- Smooth standard over age → B_a, m × k_a, a rich cubic B-spline basis over age ([m/5] internal knots) adjusted for infant mortality (Camarda 2019).
- Smooth age-space deviations:
 - **B**_s, $n \times k_{\rm s}$, basis over space:

$$\mathbf{B}_{s} = \mathbf{B}_{\mathrm{lat}} \Box \mathbf{B}_{\mathrm{lon}} = (\mathbf{B}_{\mathrm{lat}} \otimes \mathbf{1}'_{k_{\mathrm{lon}}}) \odot (\mathbf{1}'_{k_{\mathrm{lat}}} \otimes \mathbf{B}_{\mathrm{lon}}), \tag{3}$$

 $\begin{array}{l} \mathbf{B}_{\mathrm{lon}}, \ n \times k_{\mathrm{lon}} \ \text{basis over longitude}, \ \mathbf{B}_{\mathrm{lat}}, \ n \times k_{\mathrm{lat}} \ \text{basis over latitude}. \\ \bullet \ \mathbf{\breve{B}}_{a}, \ m \times \breve{k}_{\mathrm{a}} \ \text{reduced basis over age (} \lfloor m/10 \rfloor \ \text{internal knots)}. \\ \bullet \ \mathbf{B}_{s} \otimes \mathbf{\breve{B}}_{a} \end{array}$

- Unsmooth area specific intercepts $\rightarrow I_n$ identity matrix.
- Model matrix: $\mathbf{1}_n \otimes \mathbf{B}_a : \mathbf{B}_s \otimes \breve{\mathbf{B}}_a : \mathbf{I}_n \otimes \mathbf{1}_m$

Model specification: ensuring identifiablility

- Add sum of areas to ensure convergence \rightarrow n+1 spatial units.
- Model matrix:

$$\mathbf{X} \boldsymbol{\theta} = \left[\begin{array}{c|c} \mathbf{1}_{n+1} \otimes \mathbf{B}_{a} & \mathbf{0}_{m,k_{a}k_{s}+n} \\ \hline \mathbf{B}_{s} \otimes \mathbf{\breve{B}}_{a} & \mathbf{I}_{n} \otimes \mathbf{1}_{m} \end{array} \right] \boldsymbol{\theta} \,. \tag{4}$$

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- Second difference penalty on coefficients for η^0 and δ_j s. Ridge penalty for γ_j .
- Four smoothing parameters $(\lambda_a, \lambda_{lon}, \lambda_{lat}, \check{\lambda}_a)$ and one ridge penalty (κ) .

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Model specification: penalty

- Let D_a , D_{lon} , $D_{lat} \check{D}_a$ be the difference penalties associated with the age standard, longitude, latitude, and the second age basis.
- Penalty matrix

$$\mathbf{P} = \begin{bmatrix} \mathbf{P}_1 & \mathbf{0} & \mathbf{0} \\ \mathbf{0} & \mathbf{P}_2 & \mathbf{0} \\ \mathbf{0} & \mathbf{0} & \mathbf{P}_3 \end{bmatrix},$$

where

$$\mathbf{P}_1 = \lambda_a \mathbf{D}'_a \mathbf{D}_a,$$

$$\begin{split} \mathbf{P}_{3} &= \lambda_{\mathrm{lon}} \mathbf{I}_{k_{\mathrm{lat}}} \otimes \mathbf{D}_{\mathrm{lon}}' \mathbf{D}_{\mathrm{lon}} \otimes \mathbf{I}_{\breve{k}_{\mathrm{a}}} + \\ &\lambda_{\mathrm{lat}} \mathbf{D}_{\mathrm{lat}}' \mathbf{D}_{\mathrm{lat}} \otimes \mathbf{I}_{k_{\mathrm{lon}}} \otimes \mathbf{I}_{\breve{k}_{\mathrm{a}}} + \breve{\lambda}_{a} \mathbf{I}_{k_{\mathrm{lat}}} \otimes \mathbf{I}_{k_{\mathrm{lon}}} \otimes \breve{\mathbf{D}}_{a}' \breve{\mathbf{D}}_{a}, \end{split}$$

$$\mathbf{P}_3 = \kappa \mathbf{I}_n.$$

 Model fitting: Iteratively re-Weighted Least Squares (IWLS) using Generalized Linear Array Model (GLAM) (Currie, Durbán, and Eilers 2006) arithmetic:

$$(\mathbf{X}'\tilde{\mathbf{W}}\mathbf{X} + \mathbf{P})\hat{\mathbf{\theta}} = \mathbf{X}'\tilde{\mathbf{W}}\tilde{\mathbf{z}},$$
 (5)

- $\hat{\boldsymbol{ heta}}$ next iteration of coefficients to be estimated
- W diagonal matrix of weights
- z working dependent variable
- Tilde (~) current approximation.

Greater London Authority Data

Data

- *n* = 4835 Lower layer Super Output Areas (LS0As) in Greater London Authority.
- Deaths and mid-year population by m = 91 (0, 1, ..., 90+) ages.
- Total female population between 298 and 4413.
- One year: 2015

Model fitting

- Smoothing parameters equal to 1, ridge penalty equal to 100.
- 7782 coefficients to estimate.
- Effective dimension of 408.

Estimated values of components



Figure: Left: estimated standard schedule and observed mortality of total GLA. Center: estimated values of δ_i . Right: estimated values of γ_i .

Mapping components



Figure: Average over age of δ_j and γ_j .

8 November 2024

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Fitted values



Figure: Fitted log-mortality schedule for four LSOAs. Females, 2015.

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8 November 2024 19 / 24

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Estimating sub-municipal life-expectancy



Figure: Estimated life expectancy at birth. Females, 2015.

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8 November 2024

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Smoothing parameter selection

- Normally we would select the smoothing parameters through a grid search to minimize either Aikaike Information Criterion (AIC) or Bayesian Information Criterion (BIC).
- However, BIC and AIC may be inadequate in the presence of underor over-dispersion.
- We can correct AIC and BIC with ψ^2 :

$$\psi^2 = \frac{\mathsf{Dev}}{\mathsf{df}},$$

where Dev is the deviance of the model and df is the number of observations with nonzero offset minus the effective dimension of the model.

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- Accounting for under- or over-dispersion.
- Allowing the non-smooth effects to vary by age.
- Incorporating time to model multiple years of data.
- Incorporating hierarchical structure in the model.
- Adjusting the distributional assumption to model prevalence of health conditions to produce small-area health expectancy estimates.

Extension: incorporating time

- Now *m* ages, *n* spatial units, *l* years.
- Construct \mathbf{B}_t , $l \times k_a$, basis over time.
- Possible model matrices (omitting zeros and sum for identifiablility):

$$\mathbf{B}_t \otimes \mathbf{1}_n \otimes \mathbf{B}_a : \mathbf{B}_t \otimes \mathbf{B}_s \otimes \breve{\mathbf{B}}_a : \mathbf{I}_I \otimes \mathbf{I}_n \otimes \mathbf{1}_m$$

or

$$\mathbf{B}_t \otimes \mathbf{1}_n \otimes \mathbf{B}_a : \mathbf{B}_t \otimes \mathbf{B}_s \otimes \breve{\mathbf{B}}_a : \mathbf{1}_l \otimes \mathbf{I}_n \otimes \mathbf{1}_m.$$

• Multiple years of data may give enough strength to be able to abandon the standard:

$$\mathbf{B}_t \otimes \mathbf{B}_s \otimes \mathbf{B}_a : \mathbf{I}_I \otimes \mathbf{I}_n \otimes \mathbf{1}_m$$

- Our model borrows strength across age and space in order to produce mortality estimates in data-sparse contexts.
- Our modeling approach can be used to identify spatial patterns in mortality, as well as localized breaks from the general spatial pattern.
- Further work needed for selecting smoothing parameters and modeling multiple years of data.
- Contact: jacob.martin@ined.fr

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