

Spatial statistics in public health research: methodological opportunities and computational challenges

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Outline

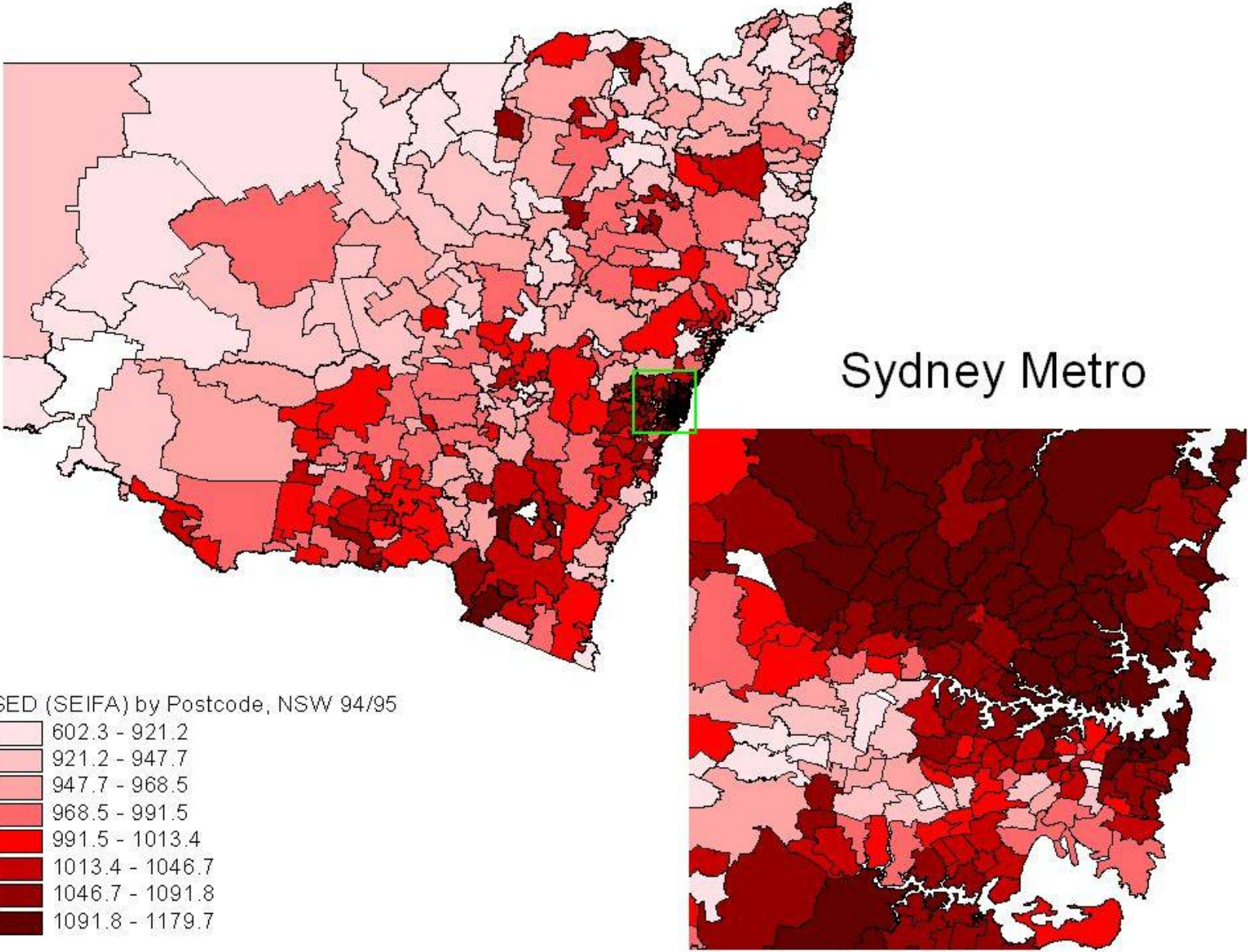
- explosion of spatial data in health research
- examples of spatial health data
- modelling spatial risk in a case-control study
 - focus on computational efficiency
- methodological and research challenges

Increased attention to spatial analysis in public health

- areal data:
 - public databases and geocoding of individuals to areas
 - interest in health disparities and social science questions
 - focus is on covariates, not spatial structure
- point data
 - geocoding and GPS are mainstream
 - * health outcomes can be assigned point locations
 - GIS software
 - * easy data management and manipulation
 - * graphical presentation
 - * spatially-varying covariate generation
 - strong applied interest in kriging and related smoothing methods
 - opportunities for more sophisticated spatio-temporal modelling, particularly Bayesian models

- environmental exposure modelling
 - * spatial smoothing and additive modelling of monitoring data
- mixed point and area data
 - individual locations plus area-level covariates
- multivariate responses
 - multiple pollutants, multiple health endpoints
 - latent variable modelling, causal relationships

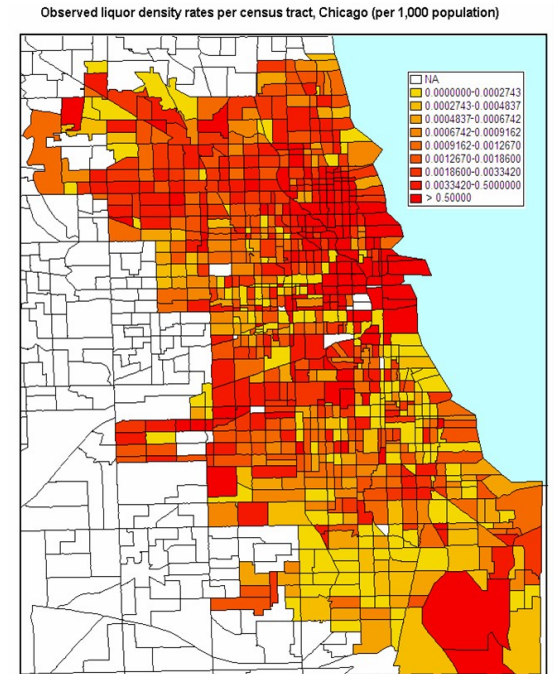
Socioeconomic factors in health outcomes in NSW, Australia



- challenges
 - areal (postcode) units vary drastically in size
 - computational challenge
 - * 650 units, 5 years daily data, 2 sexes, 9 age groups
 - spatial effect and spatially-varying covariates hard to tease apart
 - data misalignment
 - * outcome at postcode, covariate at census analogue
- relate areal data to a latent smooth process (Kelsall & Wakefield, Rathouz)

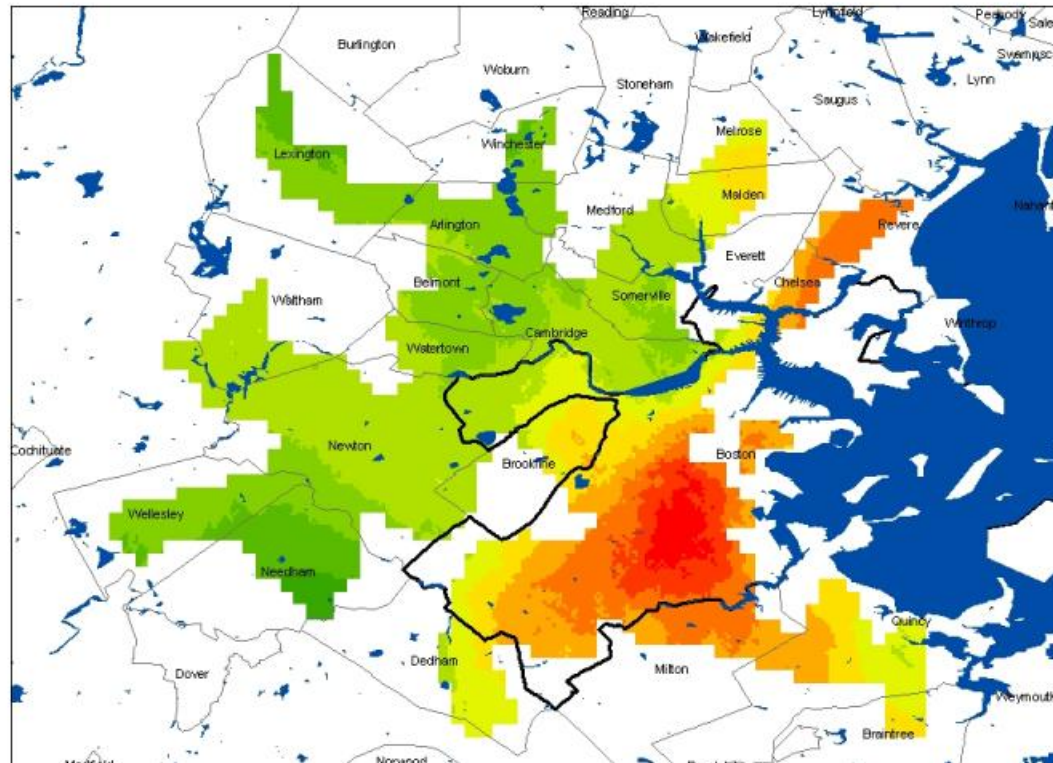
Combining area and individual-level information

- area-level covariates based on point process data
 - access to contraception at health clinics in Malawi
 - accessibility of liquor retail outlets in Chicago
- spatial scale of interest is based on outcome
- consider two-stage Bayesian model so smoothing is informed by the health outcome



Spatial variation in allergenic response

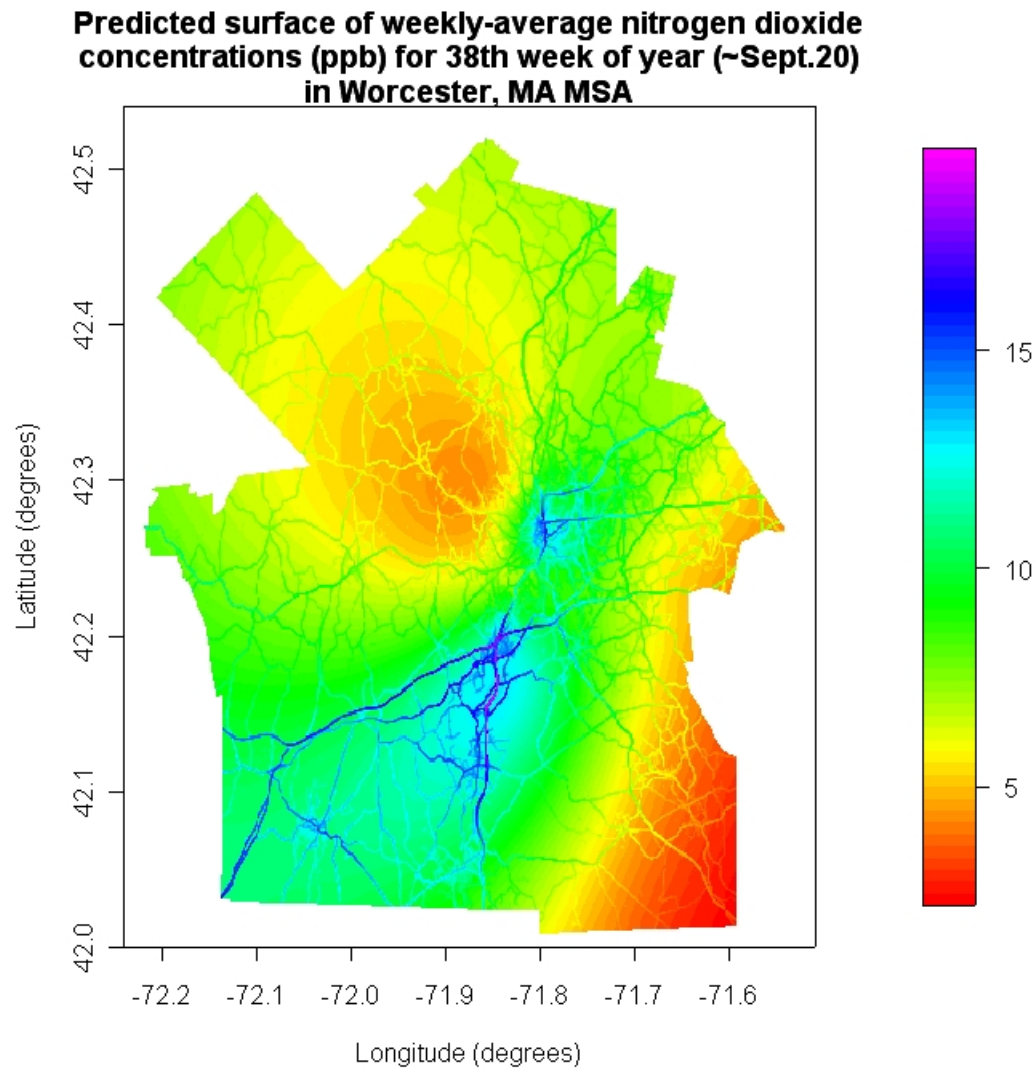
- geocoding of new mothers' residences
- measurement of blood serum IgE immune response
- interest in variance partitioning



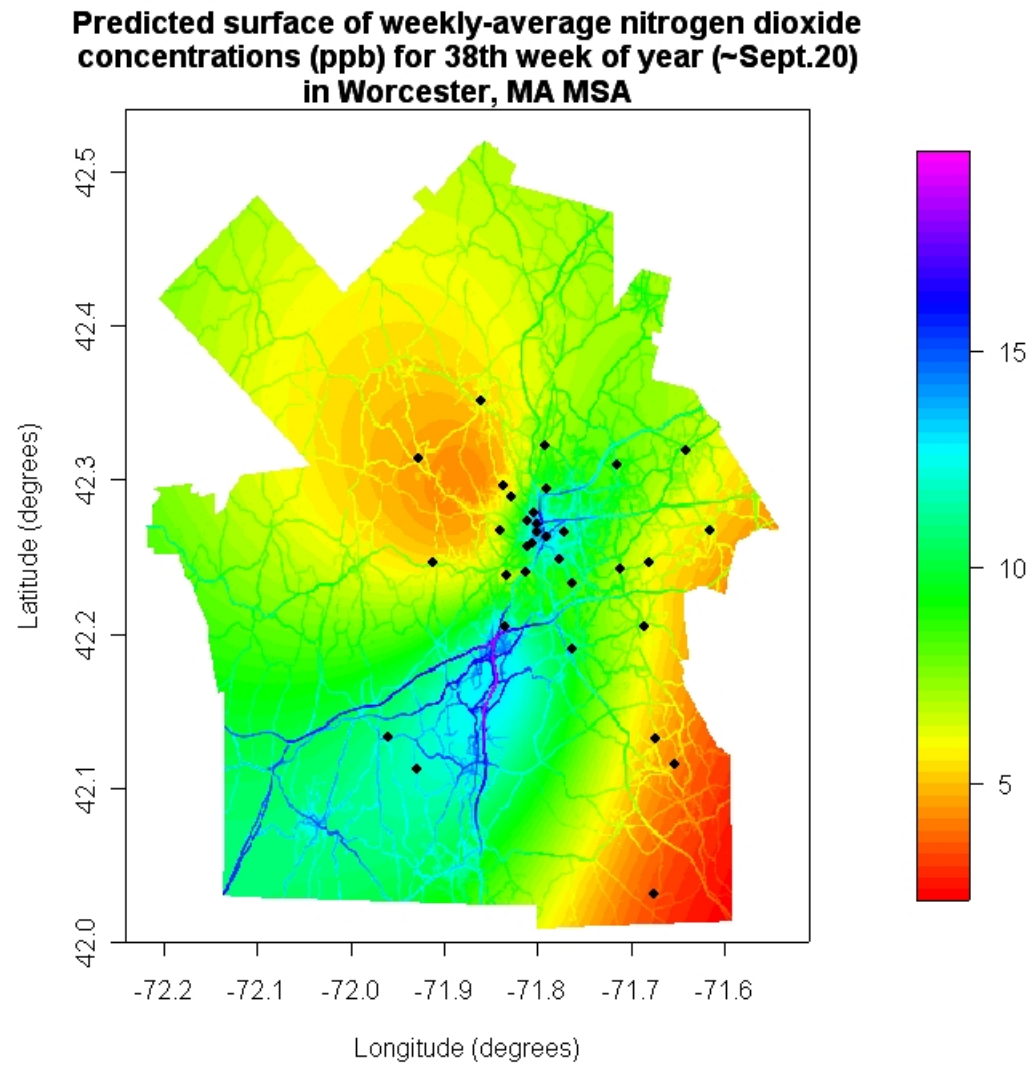
Exposure estimation in the Nurses' Health Study

- spatial estimation of individual environmental exposures
 - often air pollution
- particulate matter (PM) exposure in large cohort of nurses
 - estimate individual exposure, 1985-2003
 - EPA monitoring for large-scale spatio-temporal heterogeneity
 - spatially-varying covariates for local heterogeneity
 - * distance to roads, climate variables, local land use, ...
 - * generated using GIS
 - geocoding of individual residences every two years
 - * relate estimated exposure to health outcomes (chronic heart disease)

- geocoding and GIS make this possible; spatial statistics provides a rigorous framework



- geocoding and GIS make this possible; spatial statistics provides a rigorous framework for estimation

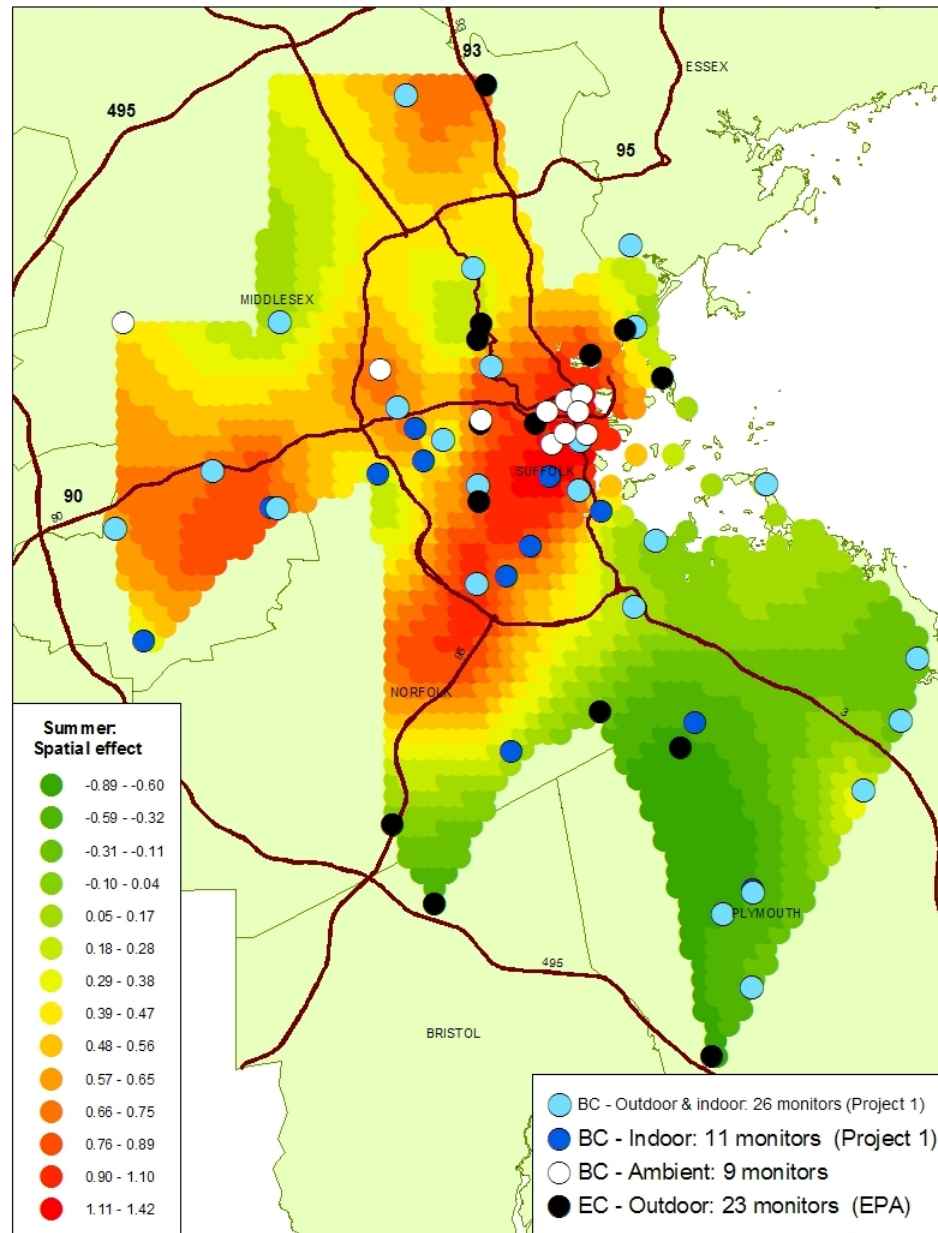


Challenges for spatio-temporal exposure estimation

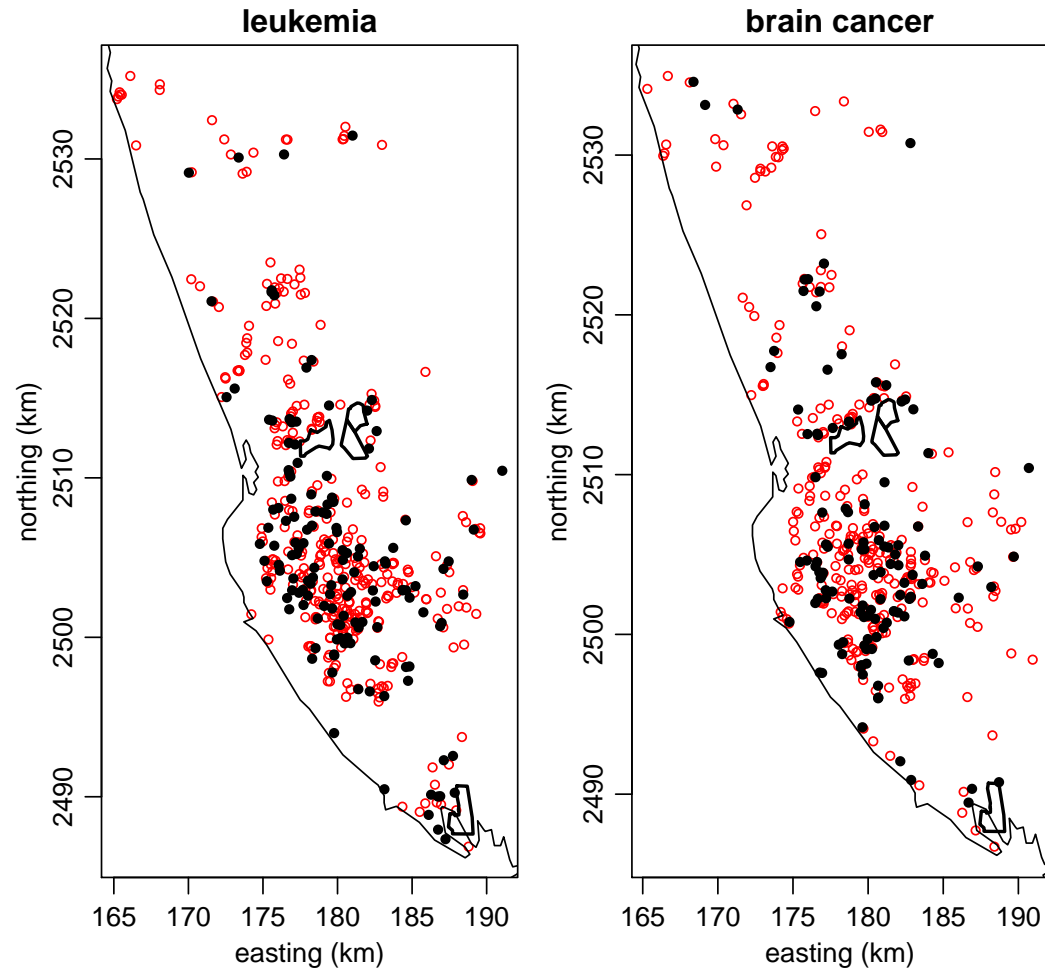
- computations: 50,000 monthly pollution measurements over 20 years at 500 monitoring sites
 - kriging is difficult, particularly Bayesian implementations
 - efficient, user-friendly computation is critical (`gam()` in R)
 - more complicated spatio-temporal structures for better prediction, but ...
 - * Bayesian implementation would require a statistician
 - * more computationally efficient methods needed
- non-standard measurement error results from smoothing
- multivariate, non-Gaussian modelling
 - modelling PM_{2.5} based on PM₁₀ and on airport visibility
 - simple multivariate normality not reasonable

Latent variable modelling

- exposure estimation for PM in the Boston area
- which pollutant sources are responsible for health outcomes?
 - traffic is locally heterogeneous, power plant pollutants (e.g., sulfates) are not
- estimate latent traffic exposure and relate to health outcomes
- two surrogates for traffic, elemental carbon and black carbon
- hierarchical Bayesian model with multiple data sources



Petrochemical exposure in Kaohsiung, Taiwan



$$n = 495$$

$$n_1 = 141$$

$$n = 433$$

$$n_1 = 121$$

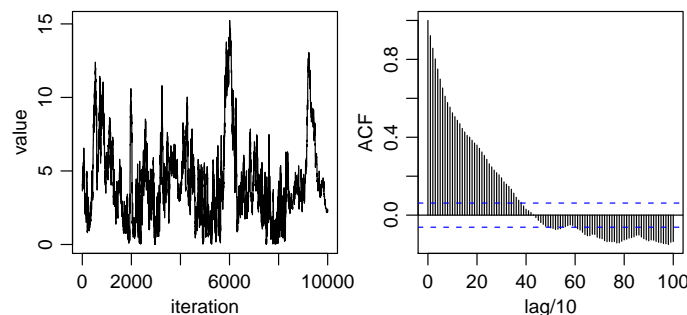
Possible approaches for health analysis

- Explicitly estimate pollutant exposure - difficult retrospectively
- Use distance to exposure source as covariate
- Use a moving window/multiple testing to detect clusters of cases
 - default approach - software available
- **Include space as a covariate to provide a map of risk**

$$Y_i \sim \text{Ber}(p(\mathbf{x}_i, \mathbf{s}_i))$$
$$\text{logit}(p(\mathbf{x}_i, \mathbf{s}_i)) = \mathbf{x}_i^T \boldsymbol{\beta} + g_\theta(\mathbf{s}_i)$$

Modelling challenges from a Bayesian perspective

- thousands of case-control observations - difficult for Bayesian kriging
- non-Gaussian spatial models particularly difficult
 - spatial process cannot be analytically integrated out of the likelihood/posterior
 - MCMC mixing is very slow because of high-level structure
 - * correlation amongst process values and between process values and process hyperparameters



Modelling Framework

$$Y_i \sim \text{Ber}(p(\mathbf{x}_i, \mathbf{s}_i))$$
$$\text{logit}(p(\mathbf{x}_i, \mathbf{s}_i)) = \mathbf{x}_i^T \boldsymbol{\beta} + g_\theta(\mathbf{s}_i)$$

- basic spatial model for $\mathbf{g}_\theta^s = (g_\theta(\mathbf{s}_1), \dots, g_\theta(\mathbf{s}_n))$
 - GAM: $g_\theta(\cdot)$ is a two-dimensional smooth term
 - * basis representation

$$\mathbf{g}_\theta^s = Z\mathbf{u}$$

- * Gaussian process representation:

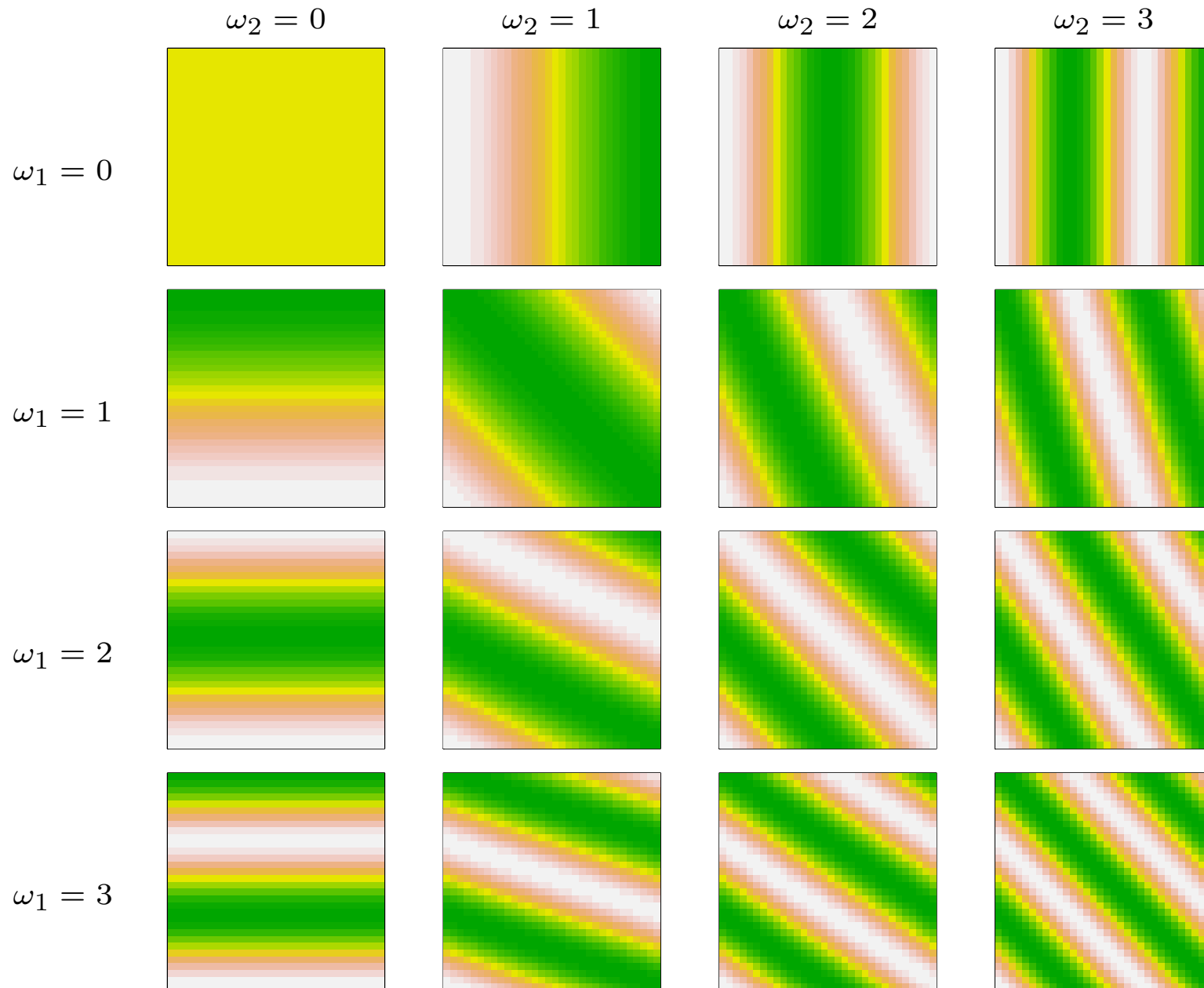
$$g(\cdot) \sim \text{GP}(\mu(\cdot), C_\theta(\cdot, \cdot)) \Rightarrow \mathbf{g}_\theta^s \sim N(\boldsymbol{\mu}, C_\theta)$$

- GLMM: $\mathbf{g}_\theta^s = Z\mathbf{u}$
 - * correlated random effects, $\mathbf{u} \sim N(0, \Sigma)$

Bayesian spectral basis function model

- computationally efficient basis function construction (Wikle 2002)
- $\mathbf{g}^\# = Z\mathbf{u}$ and $\mathbf{g}^s = \sigma P\mathbf{g}^\#$
 - piecewise constant gridded surface on k by k grid
 - P maps observation locations to nearest grid point
- Z is the Fourier (spectral) basis and $Z\mathbf{u}$ is the inverse FFT
- $Z\mathbf{u}$ is approximately a Gaussian process (GP) when...
 - $\mathbf{u} \sim N(0, \text{diag}(\pi_\theta(\boldsymbol{\omega})))$ for Fourier frequencies, $\boldsymbol{\omega}$
 - spectral density, $\pi_\theta(\cdot)$, of GP covariance function defines $V(\mathbf{u})$

Bayesian spectral basis functions



Comparison with usual GP specification

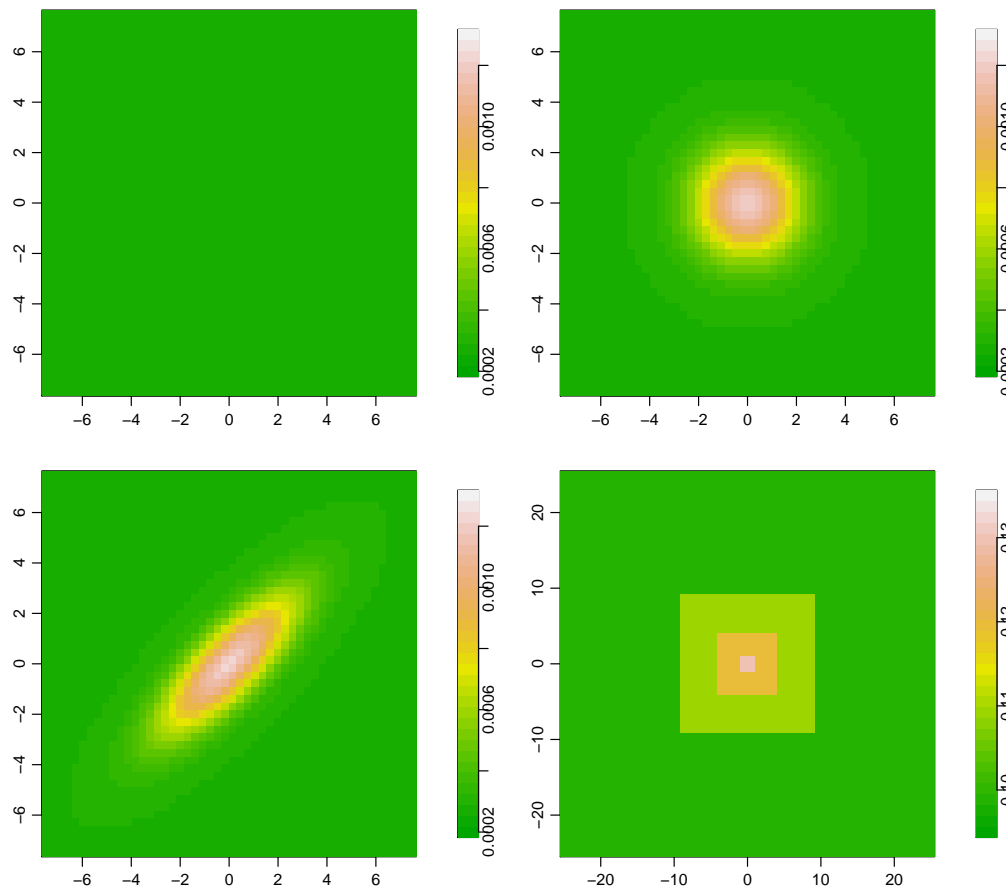
- usual GP model: $\mathbf{g}^s \sim N(\boldsymbol{\mu}, C_\theta)$
 - $O(n^3)$ fitting: $|C_\theta|$ and $C_\theta^{-1}\mathbf{g}$
- spectral basis uses FFT
 - $O((k^2) \log(k^2))$
 - additional observations are essentially free for fixed grid
 - fast computation and prediction of surface given coefficients
 - a priori independent coefficients give fast computation of prior and help with mixing

Other approaches

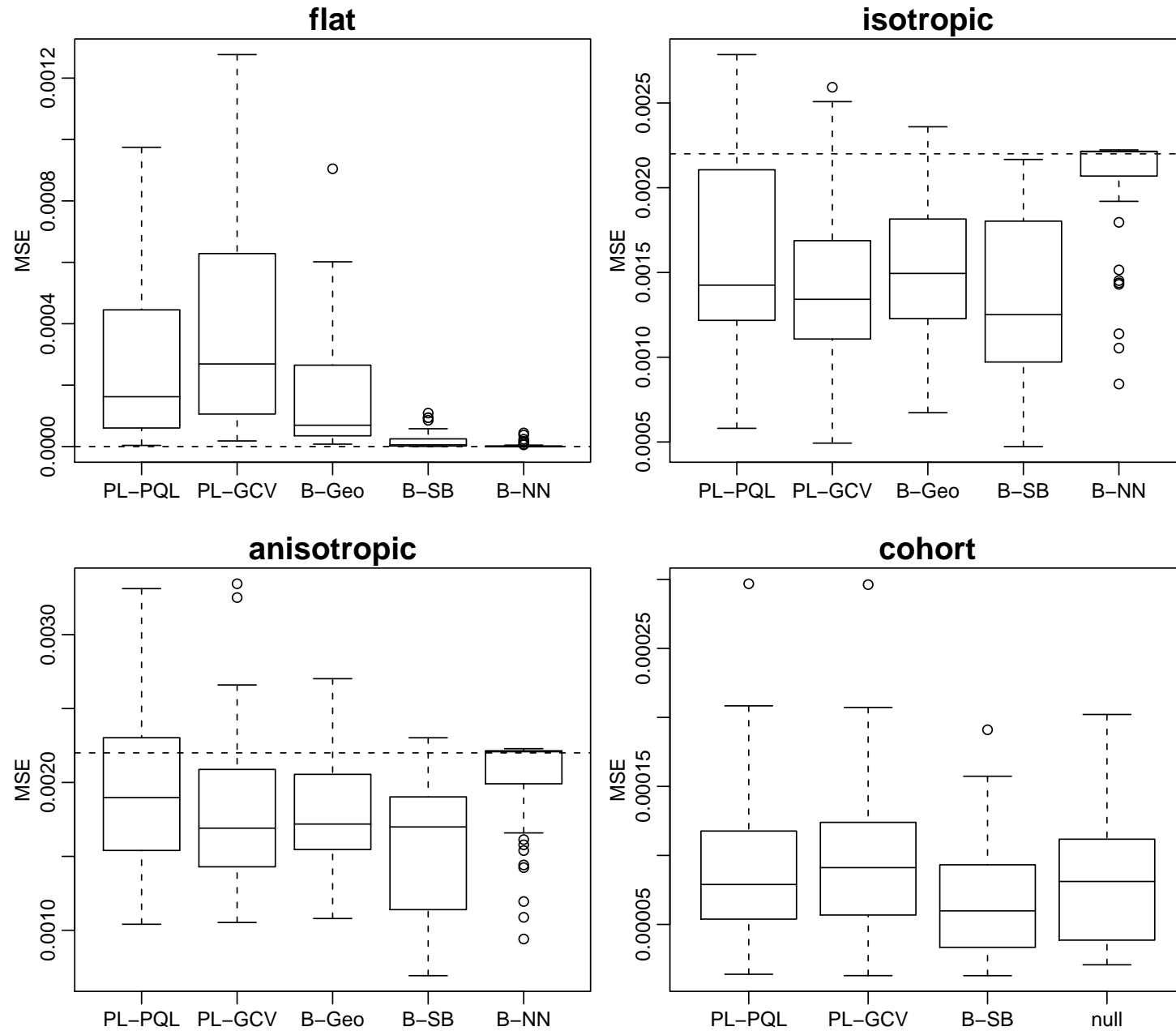
- penalized likelihood based on mixed model (radial basis functions) with REML smoothing
(Kammann and Wand, 2003; Ngo and Wand, 2004) [PL-PQL]
- penalized likelihood with GCV smoothing
(Wood, 2001, 2003, 2004) [PL-GCV]
- Bayesian mixed model/radial basis functions fit by MCMC
(Zhao and Wand 2004) [B-Geo]
- Bayesian neural network model fit by MCMC
(R. Neal) [B-NN]

Simulated datasets

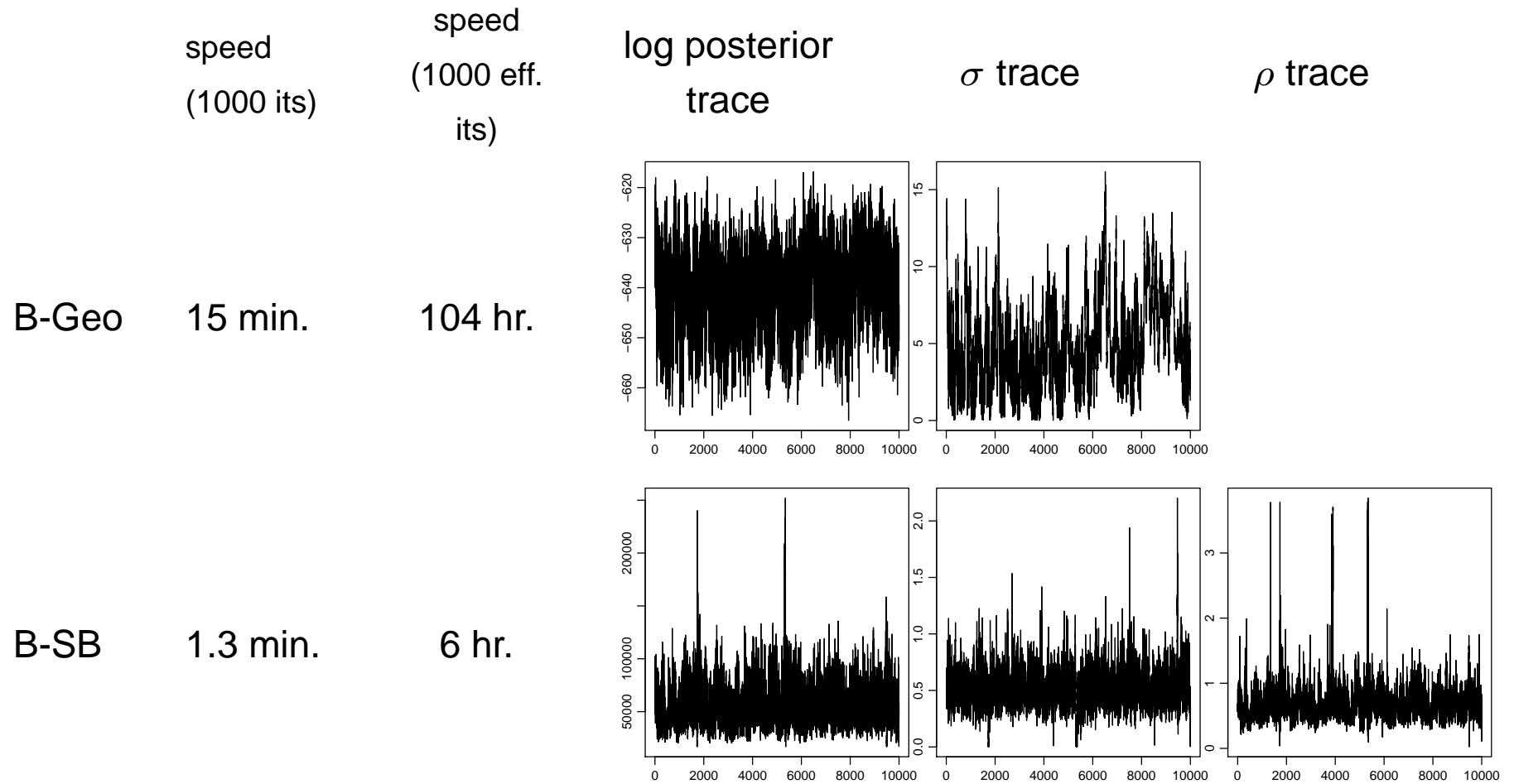
- 3 case-control scenarios: $n_0 = 1,000$; $n_1 = 200$; $n_{\text{test}} = 2500$ on 50 by 50 grid
- 1 cohort scenario: $n = 10,000$; $n_{\text{test}} = 2500$ on 50 by 50 grid



Assessment on 50 simulated datasets



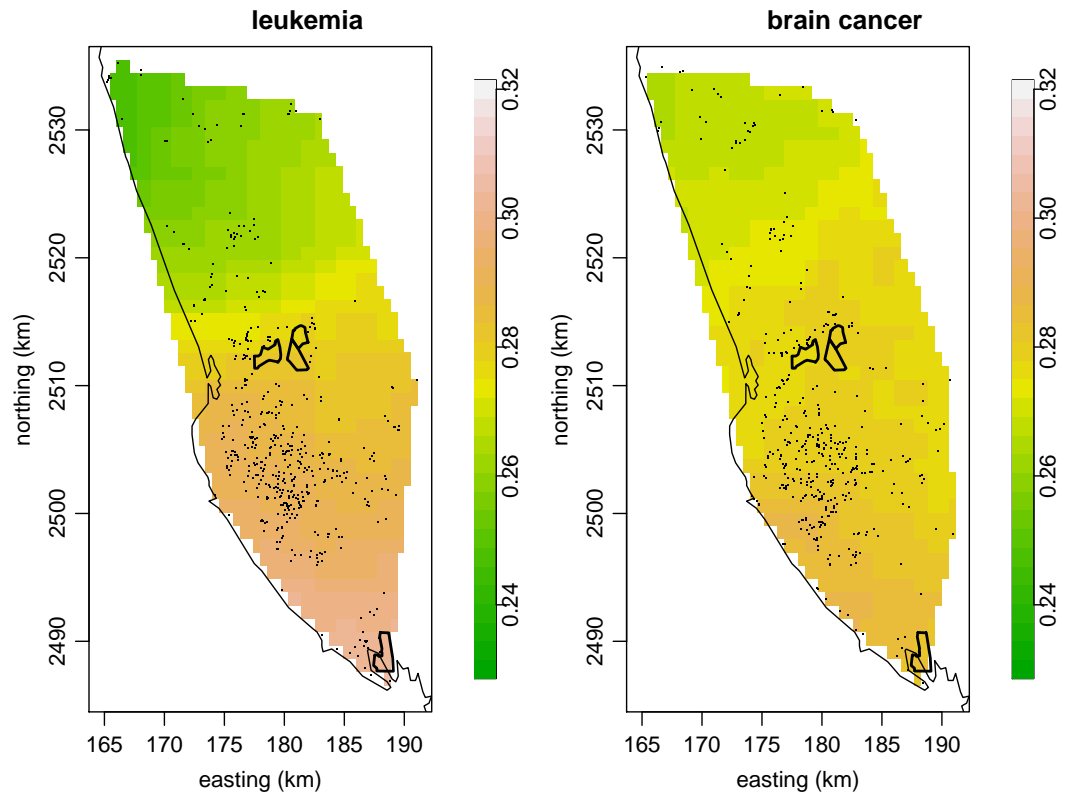
Mixing and speed of Bayesian methods



Taiwan revisited - assessment

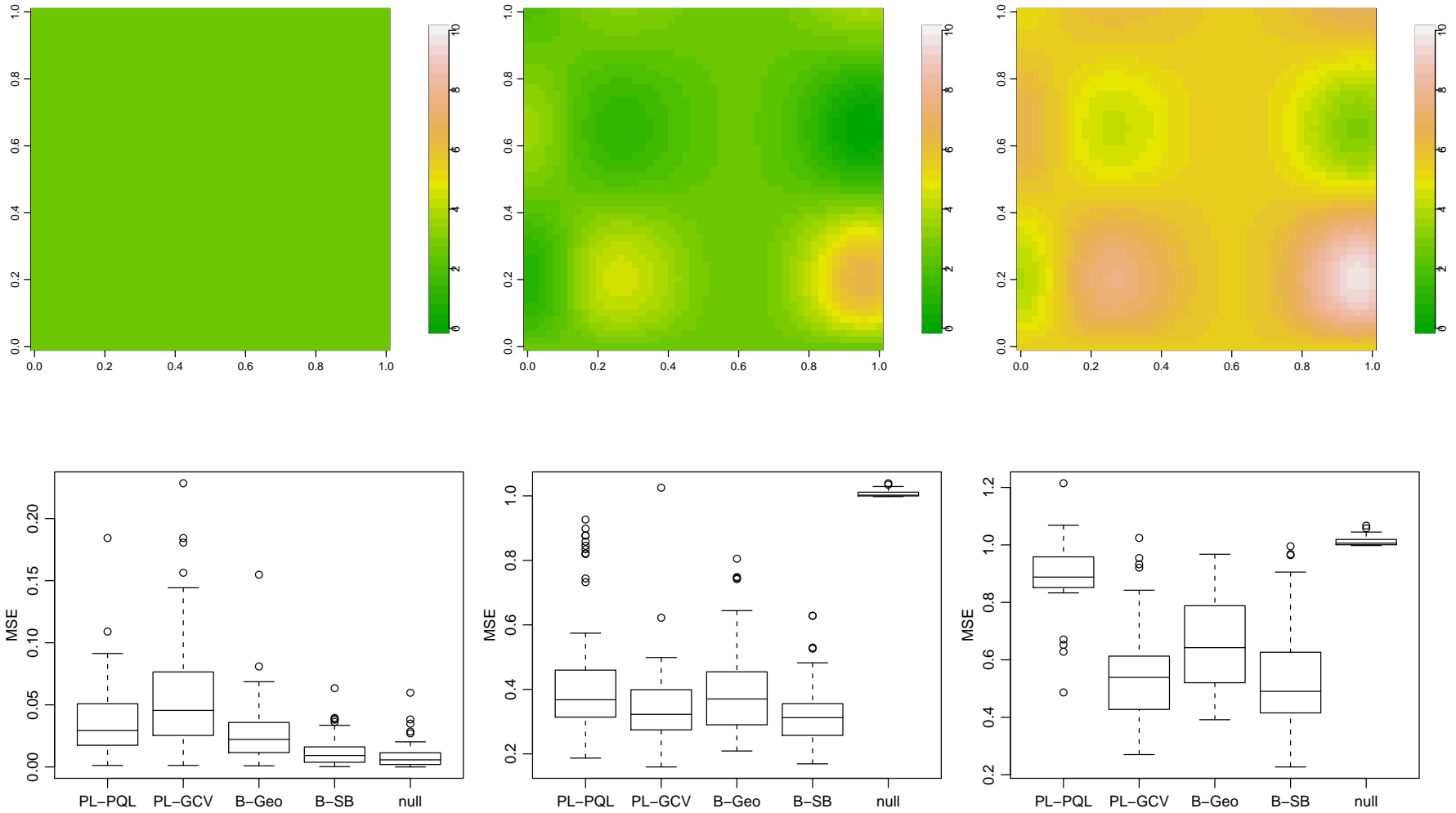
Summed test deviance
over 10-fold C-V sets

	leukemia	brain cancer
PL-GCV	590.1	529.8
PL-PQL	585.6	529.5
B-Geo	583.3	525.7
B-SB	582.1	525.1
null	581.6	525.5



Assessment on count simulations

$n = 225$, $n_{\text{test}} = 2500$ on 50 by 50 grid



Evaluation of methods

- Effective process parameterization = effective Bayesian estimation
 - feasible for spatial models with thousands of observations
- Natural Bayesian complexity penalty works well
 - GP representation zeroes out high-frequency coefficients as appropriate
- Implementation requires MCMC, not very accessible to practitioners
- Power is a real issue with spatial data in general, but particularly with binary observations
- Focused cluster-hunting or distance-based assessment of health risk may provide more power, but without full spatial assessment

Methodology challenges in spatial statistics related to public health

- design and power
 - how do we choose monitoring sites?
 - when we have enough power to estimate spatial features?
 - how do we model spatial processes when monitoring data is at lower resolution than the true surface?
- surveillance and hotspot detection
 - do Bayesian methods have a place in biosurveillance and cluster detection?
 - * current applied work focuses on testing not modelling
 - surveillance likely to benefit from a decision theoretic approach that carefully considers both false positives and false negatives
- assigning one location to an individual is problematic
- variance partitioning between spatial terms and spatially-varying covariates
- confidentiality restrictions with respect to point locations and individual privacy

General challenges for spatial statistics in public health research

- computational: big datasets and fitting of complicated models
- collaborative: developing expertise among applied researchers
- leadership
 - statisticians should be at the forefront of analyzing geographically-indexed health data
 - we shouldn't leave this area to GIS analysts/geographers
 - necessity of providing and publicizing software for rigorous statistical methods
 - * e.g., success of mixed model software – PROC MIXED, lme()
 - * evidence of mgcv: public health researchers will learn R if useful model-building tools exist

- reproducibility: difficult to replicate analyses with complicated models, particularly MCMC implementations
 - posting code and releasing software with papers
 - standardized MCMC in R
 - * many models, particularly new methods, can't be implemented in BUGS
 - e.g., complicated spatio-temporal models
 - * library of MCMC sampling functions with random variable classes
 - Jouni Kerman (Columbia) has an initial implementation for Gibbs and Metropolis sampling (umacs)
 - contributed sampling functions (e.g., slice sampling, Langevin sampling) would make this very powerful
 - * reduce bugs, increase portability and reproducibility, optimize mixing