Application of Spatially-Varying Coefficient Models

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Spatially-Varying Coefficient Models

- Extension of spatial regression approaches that allow regression coefficients to vary across space, and not just the intercept
- SVC models are random slopes models, with spatial structure given to the random slopes

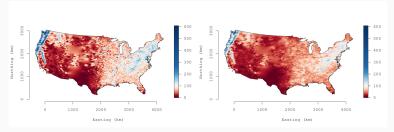
SVC GLMMs

- We can model $\mathbf{w}(\mathbf{s}_i)$ using a GP, predictive process, or NNGP
- We can envision modeling $\mathbf{w}(\mathbf{s}_i)$ in two ways:
 - 1. Multivariate NNGP (see previous forest biomass example)
 - 2. Independent NNGPs
- Here we focus on the latter
- Pros and cons to both approaches, similar to correlations between random slopes and intercepts in mixed models

Potential benefits of SVC models

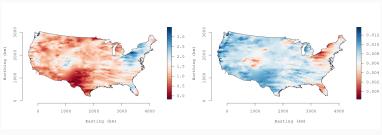
- Improved predictive performance
 - Tremendous flexibility to accommodate spatial variability in effects
- Hypothesis testing and generation
- Accommodate highly non-linear relationships
- Model spatial variability in trends over time

Improved predictive performance





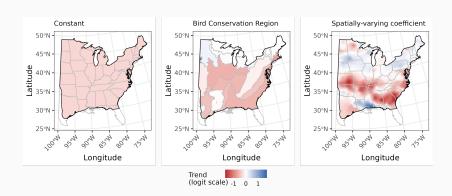
Fitted biomass



 $\beta_0(s)$ $\beta_{NDVI}(s)$

More flexibility to accommodate spatial variability in effects

Gray Catbird occurrence trend across the eastern US from 2000-2019



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 - Further, negative covariances can arise, which may drive σ^2 to be very close to zero.
- Recommendation following Gelfand et al. (2003) JASA: only use positive covariate values. Can lead to identifiability problems.
- For modeling spatially-varying trends, standardiziation is recommended.

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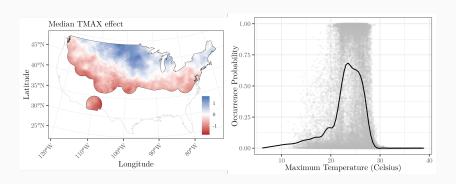
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- Recommend plotting spatial maps of SVC effects together with a plot of the covariate vs. the predicted mean for all observed locations
- Interpretation can be difficult when both the effect and the covariate vary over space. More straightforward with something like a trend.

Example: Effect of max temperature on Bobolink occurrence



Software

- spBayes: univariate Gaussian SVC with full GPs
- sp0ccupancy: univariate Binomial SVC with NNGPs (multivariate on its way)
- varycoef: maximum likelihood Gaussian SVCs (Dambon et al. 2021 Spatial Stats.)
- sdmTMB: penalized likelihood and Bayesian SVC GLMMs (Anderson et al. 2022 bioRxiv)

Exercise: 10-year occurrence trend of Wood Thrush

- Data come from USGS North American Breeding Bird Survey
- We desire to account for observational biases in detection of the birds (i.e., false negatives).
- Add on an additional observational layer to our hierarchical model



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- We model $z_t(\mathbf{s}_i)$ just as before with a Bernoulli GLM, with a SVC for trend

$$z_t(\mathbf{s}_i) \sim \mathsf{Bernoulli}(\psi_t(\mathbf{s}_i))$$

 $\mathsf{logit}(\psi_t(\mathbf{s}_i)) = \tilde{\beta}_0(\mathbf{s}_i) + \tilde{\beta}_1(\mathbf{s}_i) \cdot \mathsf{YEAR}_t$

• $\tilde{\beta}_0(\mathbf{s}_i)$ and $\tilde{\beta}_1(\mathbf{s}_i)$ are modeled as independent SVCs with NNGPs

Exercise: Observation model

- Let $y_{t,k}(\mathbf{s}_i)$ denote the observed detection (1) or nondetection (0) of the bird at site i during year t and survey $k = 1, \ldots, 5$.
- We model $y_{t,k}(\mathbf{s}_i)$ conditional on the true presence/absence of the species $z_t(\mathbf{s}_i)$

$$y_{t,k}(\mathbf{s}_i) \mid z_t(\mathbf{s}_i) \sim \mathsf{Bernoulli}(p_{i,t,k} \cdot z_t(\mathbf{s}_i))$$

 $\mathsf{logit}(p_{i,t,k}) = \alpha_{0,t} + \alpha_1 \cdot \mathsf{DAY}_{i,t,k} + \alpha_2 \cdot \mathsf{DAY}_{i,t,k}^2$

• A key assumption for identifiability is that $z_t(\mathbf{s}_i)$ does not change across the 5 replicate surveys at site i during year t.