# Fitting spatial + space-time models

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# What we've learned so far

- Time series can be useful for identifying structure, improving precision, and accuracy of forecasts
- Modeling multivariate time series
  - e.g. MARSS() function, with each observed time series mapped to a single discrete state
- Using DFA
  - Structure determined by factor loadings

### Response generally the same variable



Making inference about population as a whole involves modeling pink as well as blue time series

# Multivariate approach with MARSS

- Time series modeled as independent (allowed to have different states/trends)
- But we estimate the correlation with **Q** matrix
- Q = "unconstrained"

$$Q = \begin{bmatrix} q_{1,1} & q_{1,2} & q_{1,3} & q_{1,4} \\ q_{2,1} & q_{2,2} & q_{2,3} & q_{2,4} \\ q_{3,1} & q_{3,2} & q_{3,3} & q_{3,4} \\ q_{4,1} & q_{4,2} & q_{4,3} & q_{4,4} \end{bmatrix}$$

Obvious problem: (m \* (m+1))/2 parameters!

# Constraints on constraints

- We'd like to be able to reduce # parameters in Q matrix, and can specify alternate structures, but we have to follow rules
- This can cause problems in MARSS

$$Q = \begin{bmatrix} q_{1,1} & q_{1,2} & 0 & 0 \\ q_{2,1} & q_{2,2} & q_{2,3} & 0 \\ 0 & q_{3,2} & q_{3,3} & q_{3,4} \\ 0 & 0 & q_{4,3} & q_{4,4} \end{bmatrix}$$

If (1,2) are correlated and (2,3) correlated,
 (1,3) have to be correlated

# Multivariate approach with DFA

 Estimate how individual observed time series load onto 'trends'





# Potential problems with both the MARSS and DFA approach

 Sites separated by large distances may be grouped together

 Sites close to one another may be found to have very different dynamics



# Are there biological mechanisms that may explain this?

- Puget Sound Chinook salmon
  - 21 populations generally cluster into 2-3 groups based on genetics
  - Historically large hatchery programs





# Motivation of explicitly including spatial structure

• Adjacent sites can be allowed

to covary

• Estimated parameters greatly reduced to 2-5



# Types of spatial data

- Point referenced data
  - Aka geostatistical data
  - Typically 2-D, but could be 1-D or 3-D (depth, altitude)
  - May be fixed station or random (e.g. trawl surveys)
- Point pattern data
  - Spatially referenced based on outcomes
    - Presence of a species
  - Inference focused on describing clustering (or not)
- Areal data
  - Locations occur in blocks
  - counties, management zones, etc.

# Computationally convenient approaches

• CAR (conditionally autoregressive models)

Better suited for Bayesian methods

- Goal of both is to write the distribution of a single prediction analytically in terms of the joint (y1, y2)
- SAR (simultaneous autregressive models)
  - Better suited for ML methods
  - Simultaneously model distribution of predicted values
- 'Autoregressive' in the sense of spatial dependency / correlation between locations

# CAR models (Besag 1991)

- $Y_i = BX_i + \phi_i i + \varepsilon_i$
- BX Regression coefficients

 $\phi_i i$  Spatial component (aka Markov Random Field)

- $\boldsymbol{\mathcal{E}}_i$  Residual error term
- Create spatial adjacency matrix W, based on neighbors, e.g.
- W(i,j) = 1 if neighbors, 0 otherwise
- W often row-normalized (rows sum to 1)



|   | 1 | 2 | 3 | 4 | 5 |
|---|---|---|---|---|---|
| 1 | 0 | 1 | 0 | 0 | 0 |
| 2 | 1 | 0 | 1 | 0 | 0 |
| 3 | 0 | 1 | 0 | 1 | 0 |
| 4 | 0 | 0 | 1 | 0 | 1 |
| 5 | 0 | 0 | 0 | 1 | 0 |

• Besag et al. 1991

$$\phi_k \mid \phi_{-k} \sim N\left(\frac{\sum_{i=1}^n w_{ki}\phi_i}{\sum_{i=1}^n w_{ki}}, \frac{\sigma^2}{\sum_{i=1}^n w_{ki}}\right)$$

• Leroux et al. 1999

$$\phi_{k} \mid \phi_{-k} \sim N \left( \frac{\rho \sum_{i=1}^{n} w_{ki} \phi_{i}}{\rho \sum_{i=1}^{n} w_{ki} + 1 - \rho}, \frac{\sigma^{2}}{\rho \sum_{i=1}^{n} w_{ki} + 1 - \rho} \right)$$

• In matrix form *y* 

$$y \sim N\left(0, \left(I - \rho W\right)^{-1} \tilde{D}\right)$$
$$\tilde{D}_{ii} = \sigma_i^2$$

• Implemented in R: spdep, CARBayes

# SAR models

• Simultaneous autoregressive model

$$y \sim N \left( 0, \left( I - \rho W \right)^{-1} \tilde{D} \left( I - \rho W \right)^{-1} \right)$$
$$\tilde{D}_{ii} = \sigma_i^2$$

Remember that the CAR was

$$y \sim N\left(0, \left(I - \rho W\right)^{-1} \tilde{D}\right)$$
$$\tilde{D}_{ii} = \sigma_i^2$$

# Commonalities of both approaches

- Adjacency matrix W can also instead be modified to include distance
- Models spatial dependency as a function of single parameter  $\rho$

- Models don't include time dimension in spatial field
  - One field estimated for all time steps

# Problems with these approaches

• Wall (2004) "A close look at the spatial structure implied by the CAR and SAR models"



Fig. 5. Lines in both plots represent the implied model correlations between first order neighbors in the US lattice based on the SAR model (left) and CAR model (right) as functions of the respective parameters  $\rho_s$  and  $\rho_c$ .

Alternative to CAR & SAR: model elements of Q as functions

• Create matrix D, as pairwise distances

This can be 1-D, or any dimension – We'll use Euclidian distances for 2-D

$$D = \begin{bmatrix} 0 & |x_1 - x_2| & |x_1 - x_3| & |x_1 - x_4| \\ |x_1 - x_2| & 0 & |x_2 - x_3| & |x_2 - x_4| \\ |x_1 - x_3| & |x_1 - x_3| & 0 & |x_3 - x_4| \\ |x_1 - x_4| & |x_1 - x_4| & |x_3 - x_4| & 0 \end{bmatrix}$$

# Next specify spatial covariance function (aka "kernel")

• Exponential

 $Cov(x_1, x_2) = \sigma^2 \exp(-d_{1,2}/\theta)$   $Cov(x_1, x_2) = \sigma^2 \exp(-\theta \cdot d_{1,2})$ 

Squared exponential\*

 $Cov(x_1, x_2) = \sigma^2 \exp(-d_{1,2}^2 / \theta)$ 

• Matern family

$$Cov(x_1, x_2) = \sigma^2 \frac{1}{\Gamma(\nu) 2^{\nu-1}} \left(\sqrt{2\nu} \frac{d_{1,2}}{\rho}\right) \mathbf{K}_{\nu} \left(\sqrt{2\nu} \frac{d_{1,2}}{\rho}\right)$$

\*Also referred to as radial basis function, or Gaussian

#### Squared exponential



Distance



Theta = 1





Response



Theta = 1



Response

### Considerations for time series models

• Should spatial dependency be included

- How to model it
  - Constant
  - Time varying
    - Autoregressive
    - Random walk
    - Independent variation?

# Model-based geostatistical approaches

• 1. Generalized least squares

• 2. Bayesian methods in spBayes

• 3. INLA models

• 4. Spatial GAMs

# Method 1: using gls()

- Generalized least squares function
   Similar syntax to lm, glm, etc.
- Flexible correlation structures
  - corExp()
  - corGaus()
  - corLin()
  - corSpher()
- Allows irregularly spaced data / NAs

– unlike Arima(), auto.arima(), etc.



the first reading of the day (typically 00:00).

Portland, Oregon http://www.wcc.nrcs.usda.gov

- We'll use Snow Water Equivalent (SWE) data in Washington state
- 70 SNOTEL sites
   We'll focus only on Cascades
- 1981-2013
- Initially start using just the February SWE data
- 1518 data points (only 29 missing values!)



# We can use AIC to evaluate different correlation structures

mod.exp = gls(Feb ~ elev, correlation =
corExp(form=~lat+lon,nugget=T), data = y[which(is.na(y
\$Feb)==F & y\$Water.Year==2013),])
AIC(mod.exp) = 431.097

mod.gaus = gls(Feb ~ elev, correlation =
corGaus(form=~lat+lon,nugget=T), data =
y[which(is.na(y\$Feb)==F & y\$Water.Year==2013),])
AIC(mod.gaus) = 433.485

# Diagnostics: fitting variograms

var.exp <- Variogram(mod.exp, form =~ lat+lon)
plot(var.exp,main="Exponential",ylim=c(0,1))</pre>

var.gaus <- Variogram(mod.gaus, form =~ lat+lon)
plot(var.gaus,main="Gaussian",ylim=c(0,1))</pre>

# Exponential variogram

Exponential



### Gaussian variogram

Gaussian



# Extensions of these models

- corExp and corGaus spatial structure useful for wide variety of models / R packages
- Linear/non-linear mixed effect models
- Ime() / nlme() in nlme package Generalized linear mixed models
- glmmPQL() in MASS package
- Generalized additive mixed models
- gamm() in mgcv package

# Method 2: spBayes

- Slightly more complicated syntax:
- Specify:
- a. Priors on parameters
- b. Tuning parameters for Metropolis sampling (jumping variance)
- c. Starting / initial values
- d. Covariance structure ("exponential", "gaussian", "matern", etc)
- e. Number of MCMC iterations / burn-in, etc.

# Example with WA SWE data

# This syntax is dependent on model parameters. See vignette
priors <- list("beta.Norm"=list(rep(0,p), diag(1000,p)), "phi.Unif"=c(3/1,
3/0.1), "sigma.sq.IG"=c(2, 2), "tau.sq.IG"=c(2, 0.1))</pre>

# Phi is spatial scale parameter, sigma.sq is spatial variance, tau.sq = residual starting <- list("phi"=3/0.5, "sigma.sq"=50, "tau.sq"=1)</pre>

# variance of normal proposals for Metropolis algorithm tuning <- list("phi"=0.1, "sigma.sq"=0.1, "tau.sq"=0.1)</pre>

m.1 <- spLM(y~X-1, coords=cords,n.samples=10000, cov.model =
"exponential", priors=priors, tuning=tuning, starting=starting)</pre>

# Coefficients need to be extracted

##recover beta and spatial random effects burn.in <- 5000

m.1 <- spRecover(m.1, start=burn.in, verbose=FALSE)</pre>



# Standard MCMC diagnostics

Output is of class 'mcmc'



# Other spatial models in spBayes

Non-Gaussian errors spGLM

<u>Multivariate</u> spMvLM() spMvGLM()

Dynamic linear model (DLM) spDynLM

# Method 3: models in INLA

• INLA = Integrated Nested Laplace Approximation

• R-INLA software = interface between R and the GMRFLib C library, etc.

• Get it here:

http://www.r-inla.org/

# Motivation of INLA

- Some problems contain simple spatial structure
  - e.g. the harbor seal data in MARSS() with only 5 7 time series
- Others are much more complex
  - WA SWE data
  - Fisheries survey data (1000s of points)
- Including time-varying spatial fields becomes very computationally difficult
- Doing all of the above in a Bayesian setting can be prohibitive, but we can use Laplace approximation

### Varying spatial fields example: eulachon-shrimp bycatch



### INLA's approximation: SNOTEL data

**Constrained refined Delaunay triangulation** 



**Constrained refined Delaunay triangulation** 



**Constrained refined Delaunay triangulation** 



**Constrained refined Delaunay triangulation** 



How many points fall on vertices? Is the boundary area large enough? Choosing this must be done very carefully!

### Estimation done via maximum likelihood

- Estimates seems similar to those from gls() and spBayes()
- Year included as numeric here (not significant)

#### > res2pos\$summary.fixed 0.025quant 0.5quant 0.975quant k1d mode sd mean Intercept -14.576549539 2.788559e+01 -69.319288831 -14.579598755 40.13298938 -14.583354523 2.924022e-16 0.010432282 6.838504e-13 elev 0.010464086 8.931255e-04 0.008740241 0.010453876 0.01224473 -0.004192103 5.540561e-16 -0.004199109 1.409759e-02 -0.031884876 -0.004197175 0.02345057 year >

- Alternatively, we can include year in spatial field
- Year can also be included as factor

# Projecting INLA estimates to surface



### Fixed effect (factor) estimates



# Spatial SWE fields by year

-10 -20 -30 -40 -50

-10 -20 -30 -40













-10 -20 -30 -40 -50















# Additional output / diagnostics

• Plots of observed v fitted values

• Prediction to new locations

 Samples from posterior distribution via Laplace approximation

# Approach 4: Spatial GAMs

 Previous approaches modeled errors as correlated

 Spatial GAMs generally model mean as spatially correlated

• Somewhat analogous to AR vs MA models

# Examples of 2D GAMs

- g = gam(y ~ s(lat, lon)) # basis spline
- g = gam(y ~ te(lat, lon)) # tensor product spline

g = gam(y ~ te(lat, lon, by = as.factor(year)) # smooths by year

# Examples of recent NOAA work

Ward, E.J., G.R. Pess, K. Anlauf-Dunn, and C.E. Jordan. 2012. Applying time series models with spatial correlation to identify the scale of variation in habitat metrics related to threatened coho salmon (Oncorhynchus kisutch) in the Pacific Northwest. Canadian Journal of Fisheries and Aquatic Sciences, 69(11): 1773-1782

Shelton, A.O., J.T. Thorson, E.J. Ward, and B.E. Feist. 2014. Spatial, semi-parametric models improve estimates of species abundance and distribution. Canadian Journal of Fisheries and Aquatic Sciences, 71: 1655-1666

Thorson, J.T., H. Skaug, K. Kristensen, A.O. Shelton, E.J. Ward, J. Harms, and J. Benante. 2015. The importance of spatial models for estimating the type and strength of density dependence. Ecology, In press

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Ward, E.J., J.H. Anderson, T.J. Beechie, G.R. Pess, and M.J. Ford. 2015. Increasing hydrologic variability threatens depleted anadromous fish populations. *In press, Global Change Biology*