Spatial Models: Point Referenced Data

Point Referenced Models

- Primarily two classes of models
- Bayesian Kriging
 - 2D analog to AR(1)
 - Special case of Gaussian Process model
 - Includes parameter error
- Markov Random Field
 - 2D analog to state space

The basic spatial model

$$Z(s) = \underbrace{\mu(s|\beta) + w(s|\phi) + \epsilon(s)}_{trend} + \underbrace{v(s|\phi) + \epsilon(s)}_{spatial \ error} + \underbrace{\epsilon(s)}_{residual \ error}$$

- $\mu(s)$ = process model (e.g. $X\beta$)
- $\varepsilon \sim N(0,\tau^2) = \text{nugget}$
- W ~ $N(0,C_{\phi})$
- $C_{ij} = Cov[z_i, z_j] = f(|s_i s_j|)$
- Assumes isotropy, second-order stationarity

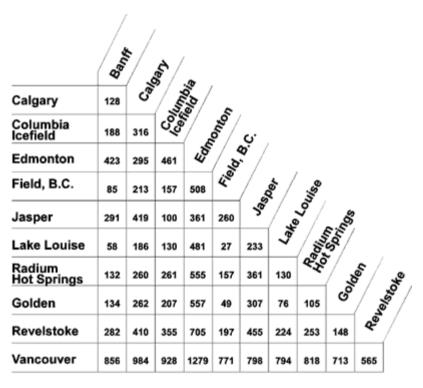
Spatial Covariance Matrix

•
$$C_{ij} = Cov[z_i, z_j] = f(|s_i - s_j|) = \sigma^2 \rho(d_{ij} | \phi)$$

- D = pairwise distance matrix
- Example correlation functions
 - Exponential: $\rho(d|\phi) = \exp(-d_{ij}\phi)$
 - Gaussian: $\rho(d|\phi) = \exp(-d_{ij}^2 \phi)$
 - Matern: $\rho(d|\phi) = \frac{v^{1/2} \phi d_{ij}}{2^{\nu} \Gamma(\nu)} K_{\nu}(2 \phi v^{1/2} d_{ij})$
- Not all functions are valid correlation functions
 - Requirement: Positive Definite covariance matrix

Distance Matrices

- Matrix of all pairwise distances
- "Distance" need not be Euclidean
- Ultimately can generalize TS and Spatial model to any situation where correlation is based on distance
 - Phylogenetic distance
 - Graphs/networks



Distances shown are in Kilometres. To convert to miles multiply by 0.6

Spatial Likelihood

$$Z(s) = \mu(s|\beta) + w(s|\phi) + \epsilon(s)$$

$$\vec{Z} \sim N(\mu, C_{\phi} + \tau^{2} I)$$

```
gp_mle <- function(parm){
  mu <- parm[1]  ## mean
  sigma <- parm[2]  ## spatial variance
  phi <- parm[3]  ## spatial correlation
  tau <- parm[4]  ## nugget variance (optional)
  C <- sigma*exp(-psi*D)  ## exponential corr
  -sum(dmvnorm(Z,rep(mu,n),C + diag(tau,n),log=TRUE))
}</pre>
```

Bayesian Spatial Model

$$\vec{Z} \sim N(\mu, C_{\phi} + \tau^{2} I)$$

$$\mu \sim N(M_{0}, V_{\mu})$$

$$\sigma^{2} \sim IG(s_{1}, s_{2})$$

$$\tau^{2} \sim IG(t_{1}, t_{2})$$

$$\Phi \sim Gamma(f_{1}, f_{2})$$

Bayesian Approach

```
model{
  mu ~ dnorm(0,0.01)
  sigma ~ dgamma(0.01,0.01)
  tau ~ dgamma(0.01,0.01)
  phi ~ dunif(0,100)
  SIGMA <- inverse(1/sigma*exp(-phi*D) + 1/tau)
  Z[] ~ dmnorm(mu,SIGMA)
}</pre>
```

Bayesian Kriging:

Prediction from the Bayesian Spatial Model

- First fit spatial model
- Second compute $D_{pred} = n+m$ pairwise distance matrix of current (n) AND prediction (m) points
- Third, for each stored MCMC iteration

- Calculate
$$\mathbf{C}_{\text{pred}}$$
 Store
$$E[Z_{pred}] = \mathbf{\mu}_{pred} + C_{s',s} C_{s,s}^{-1} (Z - \mathbf{\mu})$$
 Store for CI
$$Var[Z_{pred}] = C_{s',s'} - C_{s',s} C_{s,s}^{-1} C_{s,s'} + \mathbf{\tau}^2 \mathbf{I}$$

- For PI directly draw $z_{pred} \sim N(E[z_{pred}], var[z_{pred}])$

C_{pred} submatrices

$$E[Z_{pred}] = \mu_{pred} + C_{s',s} C_{s,s}^{-1} (Z - \mu)$$

$$\mathbf{m}$$

C_{s,s}

m C_{s',s} C_{s',s'}

Words of Warning

Singularities

- There can be no off-diagonal zeros in D
- min(d_{ij}) << max(d_{ij}) can cause numerical singularity

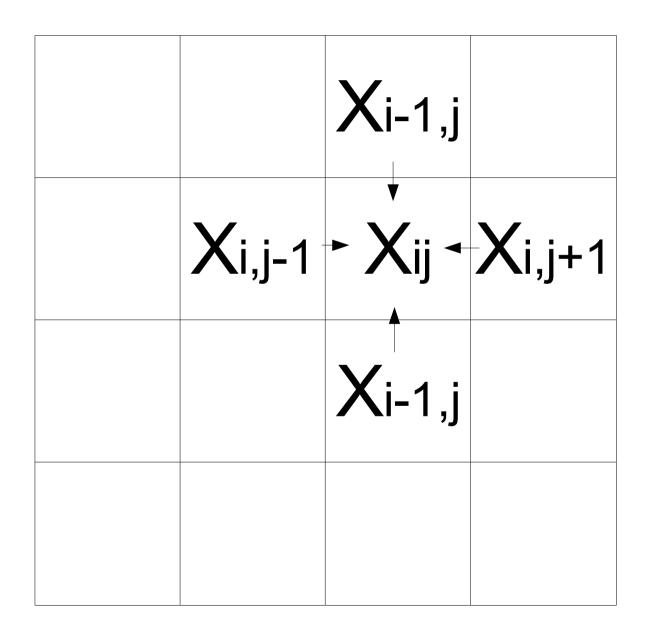
Matrix inversion

- Kriging requires large & computationally expensive matrix multiplications and inversions
- Have to do for each MCMC iteration!

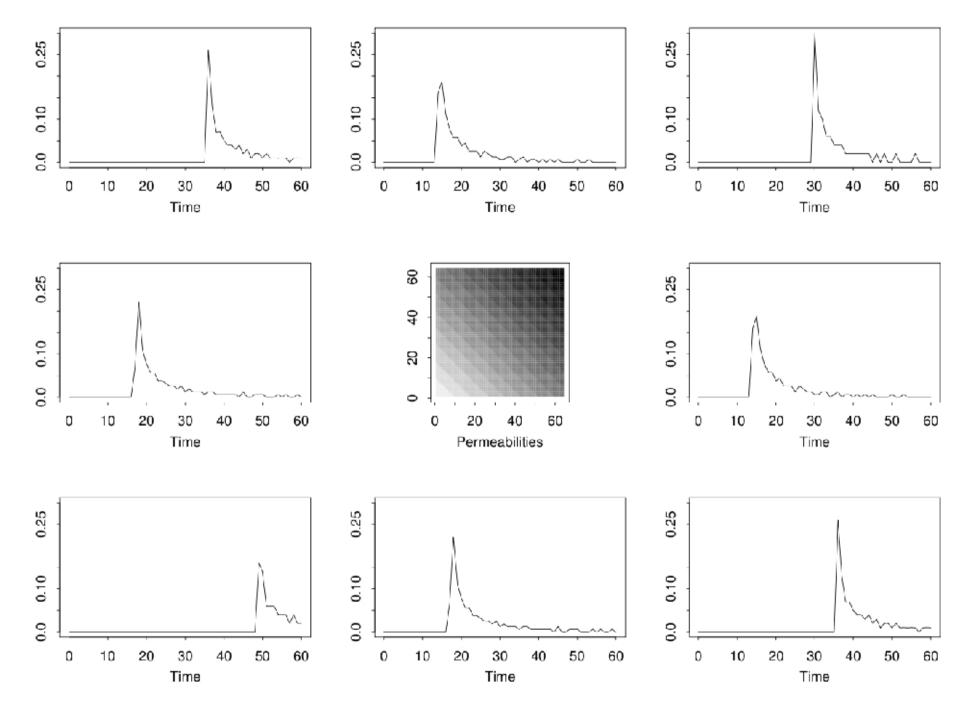
Memory

- Have to store a MAP for every MCMC iteration

Markov Random Fields



- 2D grid of latent variables
 - State-space model
- Process model connects to nearest neighbors
- Data, Y, only observed at a few points



Lee et al. 2000. MARKOV RANDOM FIELD MODELS FOR HIGH-DIMENSIONAL PARAMETERS IN SIMULATIONS OF FLUID FLOW IN POROUS MEDIA

