# Bayesian Spatial Analysis

Dr. Jarad Niemi

STAT 615 - Iowa State University

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# Spatial modeling

Three main types of spatial data:

- Point/geo-referenced
- Areal-referenced
- Point process/pattern

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#### Examples

Air quality monitoring

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- Air quality monitoring
- Coastal tide level monitoring

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- Bird point counts

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- Inflation per country

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- Locations of caught Lingcod

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### Assumptions:

Stationarity

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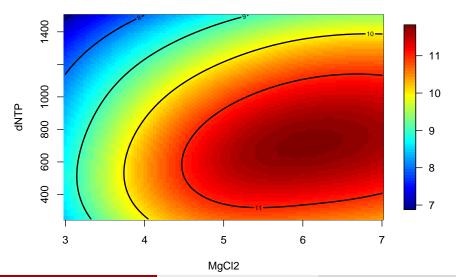
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- Stationarity
  - Intrinsic stationarity
  - Weak stationarity
  - Strong stationarity
- Isotropy
- Gaussian process

### Example spatial process

#### log of DNA amplification rate (KCL=29.77, KPO4=32.13)



# Intrinsic stationarity

#### Definition

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$$E[(Y(s+h) - Y(s))^{2}] = Var[Y(s+h) - Y(s)] = 2\gamma(h)$$

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A process Y(s) is isotropic if the semivariogram function depends only on ||h||, the length of the separation vector. Otherwise the process is anisotropic.

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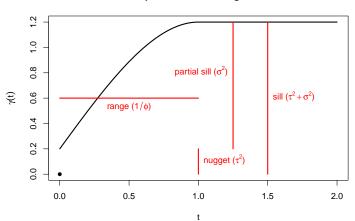
Thus, if the process is ergodic, an intrinsicly stationary process is also weakly stationary.

# Covariance functions for isotropic models

Model	Covariance function, $C(t)$	Semivariogram, $\gamma(t)$
Linear	C(t) does not exist	$\gamma(t) = \begin{cases} \tau^2 + \sigma^2 t & \text{if } t > 0\\ 0 & \text{otherwise} \end{cases}$
Spherical	$C(t) = \begin{cases} 0 \\ \sigma^{2} \left[ 1 - \frac{3}{2}\phi t + \frac{1}{2}(\phi t)^{3} \right] \\ \tau^{2} + \sigma^{2} \end{cases}$	Semivariogram, $\gamma(t)$ $\gamma(t) = \begin{cases} \tau^2 + \sigma^2 t & \text{if } t > 0 \\ 0 & \text{otherwise} \end{cases}$ $\gamma(t) = \begin{cases} \tau^2 + \sigma^2 & \text{if } t > 0 \\ \tau^2 + \sigma^2 & \frac{1}{2}\phi t - \frac{1}{2}(\phi t)^3 \end{bmatrix} & 0$ $\gamma(t) = \begin{cases} \tau^2 + \sigma^2 \left[1 - \exp(-\phi t)\right] & t > 0 \\ 0 & \text{otherwise} \end{cases}$ $\gamma(t) = \begin{cases} \tau^2 + \sigma^2 \left[1 - \exp(- \phi t ^p)\right] \\ 0 & \text{otherwise} \end{cases}$ $\gamma(t) = \begin{cases} \tau^2 + \sigma^2 \left[1 - \exp(- \phi t ^p)\right] \\ 0 & \text{otherwise} \end{cases}$
Exponential	$C(t) = \begin{cases} \sigma^2 \exp(-\phi t) \\ \tau^2 + \sigma^2 \end{cases}$	$\gamma(t) = \begin{cases} \tau^2 + \sigma^2 \left[1 - \exp(-\phi t)\right] & t > 0 \end{cases}$ oth
Powered exponential	$C(t) = \begin{cases} \sigma^2 \exp(- \phi t ^p) \\ \tau^2 + \sigma^2 \end{cases}$	$\gamma(t) = \begin{cases} \tau^2 + \sigma^2 \left[1 - \exp(- \phi t ^p)\right] \\ 0 \end{cases}$
Gaussian	$C(t) = \begin{cases} \sigma^2 \exp(-\phi^2 t^2) \\ \tau^2 + \sigma^2 \end{cases}$	$\gamma(t) = \begin{cases} \tau^2 + \sigma^2 \left[ 1 - \exp(-\phi^2 t^2) \right] \\ 0 \end{cases}$
Rational quadratic	$C(t) = \begin{cases} \sigma^2 \left( 1 - \frac{t^2}{(1+\phi^2)} \right) \\ \tau^2 + \sigma^2 \end{cases}$	$\gamma(t) = \begin{cases} \tau^2 + \sigma^2 \frac{t^2}{(1+\phi^2)} & t > 0\\ 0 & \text{otherwise} \end{cases}$
Wave	$C(t) = \begin{cases} \sigma^2 \frac{\sin(\phi t)}{\phi t} \\ \tau^2 + \sigma^2 \end{cases}$	$\gamma(t) = \begin{cases} \tau^2 + \sigma^2 \left[ 1 - \frac{\sin(\phi t)}{\phi t} \right] & t > 0 \end{cases}$ other
Power law	C(t) does not exist	$\gamma(t) = \begin{cases} \tau^2 + \sigma^2 t^{\lambda} & t > 0\\ 0 & \text{otherwise} \end{cases}$
Matérn	$C(t) = \begin{cases} \frac{\sigma^2}{2^{\nu-1}\Gamma(\nu)} (2\sqrt{\nu}t\phi)^{\nu} K_{\nu}(2\sqrt{\nu}t\phi) \\ \tau^2 + \sigma^2 \end{cases}$ $C(t) = \begin{cases} \sigma^2(1 + \phi t) \exp(-\phi t) \\ \sigma^2 + \sigma^2 \end{cases}$	$\gamma(t) = \begin{cases} \tau^2 + \sigma^2 \left[ 1 - \frac{\sin(\phi t)}{\phi t} \right] & t > 0 \\ 0 & \text{othe} \end{cases}$ $\gamma(t) = \begin{cases} \tau^2 + \sigma^2 t^{\lambda} & t > 0 \\ 0 & \text{otherwise} \end{cases}$ $\gamma(t) = \begin{cases} \tau^2 + \sigma^2 \left[ 1 - \frac{(2\sqrt{\nu}t\phi)^{\nu}}{2^{\nu - 1}\Gamma(\nu)} (2\sqrt{\nu}t\phi)^{\nu} \right] \\ 0 & \text{otherwise} \end{cases}$ $\gamma(t) = \begin{cases} \tau^2 + \sigma^2 \left[ 1 - (1 + \phi t) \exp(-\phi t) \right] \\ 0 & \text{otherwise} \end{cases}$
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## Spherical semivariogram





## Matérn

Perhaps the most important isotropic process is the Matérn process with covariance

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- ullet  $\phi$  is a spatial scale parameter.

Special cases are the exponential  $(\nu=1/2)$  and Gaussian  $(\nu\to\infty)$ .

## Strong stationarity

#### Definition

A process Y(s) is strongly (or strictly) stationary if, for any set of  $n \ge 1$  sites  $\{s_1, \ldots, s_n\}$  and any  $h \in \mathbb{R}^d$ ,

$$(Y(s_1),\ldots,Y(s_n))^{\top} \stackrel{d}{=} (Y(s_1+h),\ldots,Y(s_n+h))^{\top}$$

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The reverse is not necessarily true.

## Gaussian process

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Y(s) is a Gaussian process if, for any  $n \geq 1$  and any set of sites  $\{s_1, \ldots, s_n\}$ ,  $Y = (Y(s_1), \ldots, Y(s_n))^{\top}$  has a multivariate normal distribution.

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For a Gaussian process, weak stationarity and strong stationarity are equivalent.

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where  $\Sigma$  is constructed from the parameters  $\nu$ ,  $\phi$ ,  $\tau^2$ , and  $\sigma^2$  and the distances between locations, e.g.  $||s_1 - s_2||$ .

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$$\rho(u;\nu,\phi) = \frac{(u/\phi)^{\nu} K_{\nu}(u/\phi)}{2^{\nu-1} \Gamma(\nu)}$$

as defined in geoR:matern.

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as defined in geoR:matern. The overall mean is modeled separately and uses covariates x(s) via

$$\mu(s) = x(s)^{\top} \beta.$$

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where

$$Y|\theta \sim N(X\beta, \sigma^2 H(\phi) + \tau^2 I).$$

Let  $\theta=(\beta,\sigma^2,\tau^2,\phi)$ , then parameter estimates may be obtained from the posterior distribution:

$$p(\theta|y) \propto p(y|\theta)p(\theta)$$

where

$$Y|\theta \sim N(X\beta, \sigma^2 H(\phi) + \tau^2 I).$$

Typically, independent priors are chosen so that

$$p(\theta) = p(\beta)p(\sigma^2)p(\tau^2)p(\phi).$$

As a general rule, non-informative priors can be chosen for  $\beta$ , e.g.  $p(\beta) \propto 1$ .

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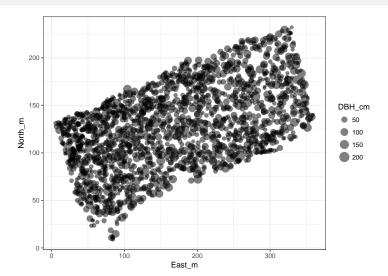
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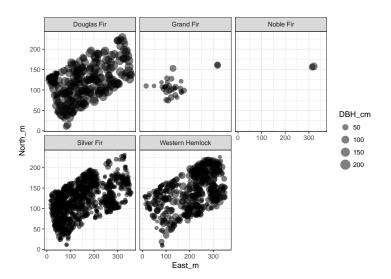
Typically, independent priors are chosen so that

$$p(\theta) = p(\beta)p(\sigma^2)p(\tau^2)p(\phi).$$

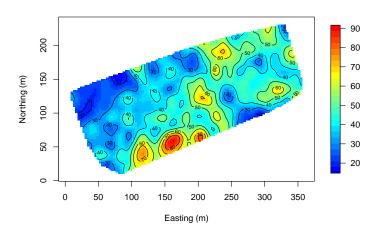
As a general rule, non-informative priors can be chosen for  $\beta$ , e.g.  $p(\beta) \propto 1$ . However, improper (or vague proper) priors for the variance parameters can lead to improper (or computationally improper) posteriors.

# Diameter at breast height (DBH) for an experimental forest





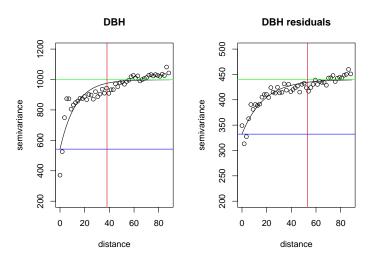
# Interpolation of mean DBH (ignoring species)



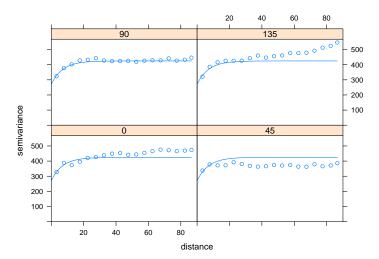
## Regression

```
## variog: computing omnidirectional variogram
## variofit: covariance model used is exponential
## variofit: weights used: equal
## variofit: minimisation function used: nls
##
## Call:
## lm(formula = DBH_cm ~ Species, data = d)
## Residuals:
          10 Median
      Min
                              30
                                    Max
## -78.423 -9.969 -3.561 10.924 118.277
##
## Coefficients:
##
                        Estimate Std. Error t value Pr(>|t|)
## (Intercept)
                          89.423
                                    1.303 68.629 <2e-16 ***
## SpeciesGrand Fir
                       -51.598
                                    4.133 -12.483 <2e-16 ***
                     -5.873 15.744 -0.373 0.709
## SpeciesNoble Fir
## SpeciesSilver Fir
                    -68.347
                                 1.461 -46.784 <2e-16 ***
## SpeciesWestern Hemlock -48.062
                                     1.636 -29.377 <2e-16 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 22.19 on 1950 degrees of freedom
## Multiple R-squared: 0.5332, Adjusted R-squared: 0.5323
## F-statistic: 556.9 on 4 and 1950 DF, p-value: < 2.2e-16
```

# Variogram (exponential model)



# Isotropy?



## spBayes

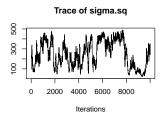
```
p = nlevels(d$Species)
r = spLM(DBH_cm ~ Species,
        data = d,
        coords = as.matrix(d[c('East_m','North_m')]),
        knots = c(6,6,.1), # for spatial prediction
        cov.model = 'exponential'.
        starting = list(tau.sq = fit.DBH.resid$nugget,
                        sigma.sq = fit.DBH.resid$cov.pars[1],
                                 = fit.DBH.resid$cov.pars[2]),
                        phi
        tuning = list(tau.sq = 0.015,
                      sigma.sq = 0.015,
                              = 0.015),
                      phi
        priors = list(beta.Norm = list(rep(0,p), diag(1000,p)),
                      phi.Unif = c(3/1,3/0.1),
                      sigma.sq.IG = c(2,200),
                      tau.sq.IG = c(3,300).
        n.samples = 10000.
        n.report = 200,
        verbose=TRUE)
```

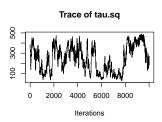
Example

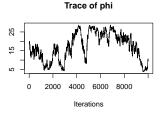
```
General model description
## Model fit with 1955 observations.
## Number of covariates 5 (including intercept if specified).
##
## Using the exponential spatial correlation model.
##
## Using modified predictive process with 36 knots.
##
## Number of MCMC samples 10000.
##
## Priors and hyperpriors:
   beta normal:
   mu: 0.000 0.000 0.000 0.000 0.000
   cov:
   1000.000 0.000 0.000 0.000 0.000
   0.000 1000.000 0.000 0.000 0.000
   0.000 0.000 1000.000 0.000 0.000
   0.000 0.000 0.000 1000.000 0.000
   0.000 0.000 0.000 0.000 1000.000
##
   sigma.sq IG hyperpriors shape=2.00000 and scale=200.00000
   tau.sq IG hyperpriors shape=3.00000 and scale=300.00000
   phi Unif hyperpriors a=3.00000 and b=30.00000
##
   Sampling
## Sampled: 200 of 10000, 2.00%
## Report interval Metrop. Acceptance rate: 36.50%
## Overall Metrop. Acceptance rate: 36.50%
## Sampled: 400 of 10000, 4.00%
## Report interval Metrop. Acceptance rate: 36.50%
```

## Traceplots

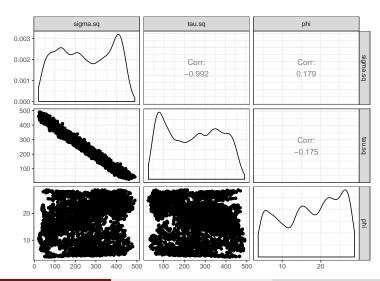
plot(r\$p.theta.samples, density=FALSE)







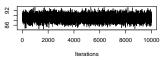
```
r$p.theta.samples[burnin:nreps,] %>%
as.data.frame %>%
GGally::ggpairs() +
theme_bw()
```



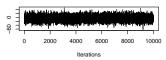
# Traceplot 2s

plot(r\$p.beta.samples, density=FALSE)

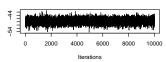
#### Trace of (Intercept)



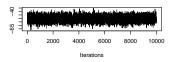
#### Trace of SpeciesNoble Fir



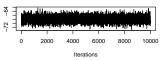
#### Trace of SpeciesWestern Hemlock



#### Trace of SpeciesGrand Fir



#### Trace of SpeciesSilver Fir



# Summary statistics

```
summary(r$p.theta.samples)
##
## Iterations = 1.10000
## Thinning interval = 1
## Number of chains = 1
## Sample size per chain = 10000
##
## 1. Empirical mean and standard deviation for each variable,
     plus standard error of the mean:
##
                       SD Naive SE Time-series SE
##
             Mean
## sigma.sq 246.59 127.324 1.27324
                                          31.224
## tau.sq
           244.94 126.725 1.26725
                                          30.490
## phi
        17.24 7.204 0.07204
                                          2.743
##
## 2. Quantiles for each variable:
##
             2.5%
                     25%
                            50%
                                   75% 97.5%
## sigma.sq 43.200 133.07 238.13 366.61 446.7
## tau.sq 51.208 124.40 252.81 355.24 449.0
## phi 4.593 11.46 17.55 23.58 27.9
```

# Summary statistics 2

```
summary(r$p.beta.samples)
## Tterations = 1:10000
## Thinning interval = 1
## Number of chains = 1
## Sample size per chain = 10000
##
  1. Empirical mean and standard deviation for each variable,
     plus standard error of the mean:
##
                                    SD Naive SE Time-series SE
##
                            Mean
## (Intercept)
                          89.011 1.291 0.01291
                                                       0.01291
## SpeciesGrand Fir
                       -50.361 4.125 0.04125
                                                       0.04125
## SpeciesNoble Fir
                        -4.406 14.034 0.14034
                                                      0.14034
## SpeciesSilver Fir
                    -67.923 1.458 0.01458
                                                      0.01458
## SpeciesWestern Hemlock -47.605 1.631 0.01631
                                                       0.01631
##
## 2. Quantiles for each variable:
##
##
                          2.5%
                                  25%
                                          50%
                                                  75% 97.5%
## (Intercept)
                        86.48 88.13 89.006
                                               89.873 91.53
## SpeciesGrand Fir -58.49 -53.11 -50.306 -47.617 -42.21
## SpeciesNoble Fir -32.44 -13.91 -4.382
                                              5.187 22.83
## SpeciesSilver Fir
                    -70.79 -68.90 -67.917 -66.944 -65.09
## SpeciesWestern Hemlock -50.79 -48.69 -47.615 -46.506 -44.44
```

#### Spatial surface

If interest resides in  $\boldsymbol{w}$ , draws can be obtained using the following relationship

$$p(w|y) = \int p(w|\sigma^2, \phi, y) p(\sigma^2, \phi|y) d\sigma^2 d\phi$$

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which suggests the following strategy:

- 1. Run the MCMC sampler to obtain draws  $(\sigma^2,\phi)^{(g)} \sim p(\sigma^2,\phi|y)$
- 2. After burn-in and for  $g=1,\ldots,G$ , sample  $w^{(g)}\sim p(w|(\sigma^2,\phi)^{(g)},y)$ .

For prediction at points  $s_{01},\ldots,s_{0m}$  and denoting  $Y_0=(Y(s_{01}),\ldots,Y(s_{0m}))^{\top}$  and design matrix  $X_0$  having rows  $x(s_{0j})^{\top}$ ,

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$$p(y_0|y, X, X_0) = \int p(y_0|y, \theta, X_0) p(\theta|y, X) d\theta \approx \frac{1}{G} \sum_{g=1}^{G} p(y_0|y, \theta^{(g)}, X_0).$$

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It is more common to take draws  $y_0^{(g)} \sim p(y_0|y,\theta^{(g)},X_0)$  and estimate the predictive distribution using

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$$\left(\begin{array}{c} y \\ y_0 \end{array}\right) \sim N\left(\left[\begin{array}{c} X\beta \\ X_0\beta \end{array}\right], \left[\begin{array}{cc} \Omega_{11} & \Omega_{12} \\ \Omega_{21} & \Omega_{22} \end{array}\right]\right)$$

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where

$$\Omega_{11} = \sigma^{2} H(\phi) + \tau^{2} \mathbf{I} 
\Omega_{22} = \sigma^{2} + \tau^{2} 
\Omega_{12}^{\top} = \sigma^{2} [\rho(d_{01}; \phi), \dots, \rho(d_{0n}; \phi)]$$

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and  $d_{ij} = ||s_i - s_j||$ .

Thus  $y_0|y,\theta,X,X_0$  is normal with

$$E[Y(s_0)|y,\theta,X,X_0] = x_0^{\top}\beta + \Omega_{12}^{\top}\Omega_{22}^{-1}(y - X\beta) Var[Y(s_0)|y,\theta,X,X_0] = \sigma^2 + \tau^2 - \Omega_{12}^{\top}\Omega_{22}^{-1}\Omega_{12}$$

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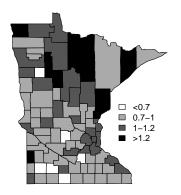
$$Y(s) \sim Po(e^{x(s)^{\top}\beta + w(s)}).$$

For GLMs (other than linear models), w(s) cannot be integrated out and therefore a common MCMC strategy is

- 1. Sample  $\beta | \dots$
- 2. Sample  $w | \dots$
- 3. Sample  $\theta | \dots$  (the spatial parameters [no nugget]).

# Choropleth

#### **MN Lung Cancer SMR**



Let  $Y_i$  represent the SMR for lung cancer in MN county i.

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$$Y_i|y_{-i} \sim N\left(\sum_{j \in n_i} y_j/m_i, \tau^2/m_i\right)$$

#### where

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This defines a Markov Random Field.

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#### Definition

Brook's Lemma states that

$$\frac{p(y_1,\ldots,y_n)}{p(y_1',\ldots,y_n')} = \frac{p(y_1|y_2,\ldots,y_n)}{p(y_1'|y_2,\ldots,y_n)} \cdot \frac{p(y_2|y_1',y_3,\ldots,y_n)}{p(y_2'|y_1',y_3,\ldots,y_n)} \cdot \cdot \cdot \cdot \frac{p(y_n|y_1',\ldots,y_{n-1}')}{p(y_n'|y_1',\ldots,y_{n-1}')}$$

for all  $(y'_1, \ldots, y'_n)$ .

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for all  $(y'_1,\ldots,y'_n)$ .

lf

$$p(y_1',\ldots,y_n') = \int \frac{p(y_1|y_2,\ldots,y_n)}{p(y_1'|y_2,\ldots,y_n)} \cdot \frac{p(y_2|y_1',y_3,\ldots,y_n)}{p(y_2'|y_1',y_3,\ldots,y_n)} \cdots \frac{p(y_n|y_1',\ldots,y_{n-1}')}{p(y_n'|y_1',\ldots,y_{n-1}')} dy_1,\ldots,dy_n < \infty$$

then  $p(y_1, \ldots, y_n)$  is a proper joint distribution.

## Conditionally autoregressive models

More generally, we can consider

$$Y_i|y_{-i} \sim N\left(\sum_{j \neq i} b_{ij}y_j, \tau_i^2\right)$$

Through Brook's Lemma, we have

$$p(y_1, \dots, y_n) \propto \exp\left(-\frac{1}{2}y^{\top}D^{-1}[\mathbf{I} - B]y\right)$$

where

- ullet B has elements  $b_{ij}$
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where

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In order for  $D^{-1}[I-B]$  to be symmetric, we need  $\frac{b_{ij}}{\tau_i^2}=\frac{b_{ji}}{\tau_j^2}$  for all i,j.

#### Proximity matrix

#### Definition

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### Common choices for $w_{ij}$ are

• 1 if i is a neighbor of j and 0 otherwise

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  - neighbors defined by those who share an edge

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  - ullet neighbors defined by those who are within distance  $\delta$

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  - K-nearest neighbors
- "distance"

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This is called the *intrinsically autoregressive* model.

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a prior for  $\rho$  that induces a reasonable amount of spatial association should put most of its mass near 1.

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indicates some issues with this model:

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- $\bullet$   $\rho$  needs to be very close to 1 to obtain a consequential amount of spatial association.

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- $\bullet$  Choose  $\rho=1$  and estimate a mean (remove mean from the fixed effect)
- Let  $\rho \sim Be(18,2)$  (Banerjee pg 164) and estimate it.

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and the conditional distributions are

$$Y_i|y_{-i} \sim N\left(\frac{\rho \sum_{j \neq i} w_{ij} y_j}{rho \sum_{j \neq i} w_{ij} y_j + 1 - \rho}, \frac{\tau^2}{rho \sum_{j \neq i} w_{ij} y_j + 1 - \rho}\right).$$

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This distribution is proper so long as  $0 \le \rho \le 1$ . Lee (2011) argued that this CAR should be preferred for a variety of reasons.

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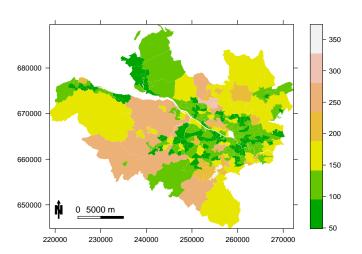
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$$p(\omega_1, \dots, \omega_S) \propto \exp\left(-\frac{1}{2\tau^2}\omega^{\top}[D_w - \rho W]\omega\right)$$



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- $\bullet$   $Y_i$  be the logarithm of the median home price in each Intermediate Geography (IG) to the north of the river Clude in the Greater Glasgow and Clyde health board,
- use explanatory variables
  - crime: crime rate (number of crimes per 10,000 people) in each IG (logged),
  - rooms: median number of rooms in a property in each IG,
  - type: predominant property type in each IG with levels: detached, flat, semi, terrace.
  - sales: percentage of properties that sold in each IG in a year, and
  - driveshop: average time taken to drive to a shopping centre in minutes (logged).

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- $oldsymbol{\omega}_i$  are assumed to come from an intrinsic CAR model with proximity matrix indicating those regions that share a border

```
propertydata.spatial@data$logprice <- log(propertydata.spatial@data$price)</pre>
propertydata.spatial@data$logdriveshop <- log(propertydata.spatial@data$driveshop)
### code chunk number 9: CARBayes.Rnw:495-498
library(splines)
form <- logprice~ns(crime,3)+rooms+sales+factor(type) + logdriveshop</pre>
model <- lm(formula=form, data=propertydata,spatial@data)
### code chunk number 10: CARBayes.Rnw:505-510
library(spdep)
W.nb <- poly2nb(propertydata.spatial, row.names = rownames(propertydata.spatial@data))
W.list <- nb2listw(W.nb, style="B")
resid.model <- residuals(model)
moran.mc(x=resid.model, listw=W.list, nsim=1000)
##
  Monte-Carlo simulation of Moran I
##
## data: resid model
## weights: W.list
## number of simulations + 1: 1001
##
## statistic = 0.2733, observed rank = 1001, p-value = 0.000999
## alternative hypothesis: greater
```

```
Areal-referenced
```

```
##
## Call:
## lm(formula = form, data = propertydata,spatial@data)
##
## Residuals:
       Min
                10 Median
                                  30
                                         Max
## -0.91319 -0.15992 0.00136 0.15647 0.81675
##
## Coefficients:
##
                      Estimate Std. Error t value Pr(>|t|)
                     4.436135
                                0.157971 28.082 < 2e-16 ***
## (Intercept)
## ns(crime, 3)1
                    -0.358967 0.089006 -4.033 7.24e-05 ***
## ns(crime, 3)2
                    -0.617084 0.165152 -3.736 0.000229 ***
                    -0.299454 0.126516 -2.367 0.018670 *
## ns(crime, 3)3
                     ## rooms
                                0.000362 5.619 4.93e-08 ***
## sales
                    0.002034
## factor(type)flat
                  -0.215967 0.066412 -3.252 0.001298 **
## factor(type)semi
                    -0.153610 0.057750 -2.660 0.008301 **
## factor(type)terrace -0.280023   0.072634   -3.855   0.000146 ***
## logdriveshop
                     -0.089084
                                 0.025588 -3.482 0.000585 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 0.2243 on 260 degrees of freedom
## Multiple R-squared: 0.6206, Adjusted R-squared: 0.6075
## F-statistic: 47.26 on 9 and 260 DF, p-value: < 2.2e-16
```

# Bayesian analysis using Leroux CAR