Block Prediction for a Finite Grid

Jay Ver Hoef

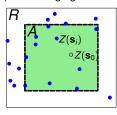
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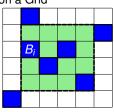
Introduction

Introduction

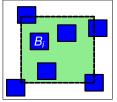
1) Block Kriging



2)Block Prediction for Finite Populations on a Grid



3)Block Prediction for Finite Populations Irregularly Spaced



Ver Hoef, J.M. 2001. Predicting finite populations from spatially correlated data. 2000 Proceedings of the Section on Statistics and the Environment of the American Statistical Association, pgs. 93 - 98.

Ver Hoef, J.M. 2008. Spatial methods for plot-based sampling of wildlife populations. Environmental and Ecological Statistics 15: 3 - 13.



Review of BLUP and Point Prediction

$$\begin{pmatrix} \mathbf{z} \\ Z(\mathbf{s}_0) \end{pmatrix} = \begin{pmatrix} \mathbf{X} \\ \mathbf{x}(\mathbf{s}_0)' \end{pmatrix} \boldsymbol{\beta} + \begin{pmatrix} \boldsymbol{\epsilon} \\ \boldsymbol{\epsilon}(\mathbf{s}_0) \end{pmatrix}$$
$$\operatorname{cov} \begin{pmatrix} \boldsymbol{\epsilon} \\ \boldsymbol{\epsilon}(\mathbf{s}_0) \end{pmatrix} = \begin{pmatrix} \boldsymbol{\Sigma} & \mathbf{c} \\ \mathbf{c}' & \sigma_0^2 \end{pmatrix}$$

Best Linear Unbiased Prediction (BLUP) (or [Universal] Kriging) minimize: $E(\lambda' \mathbf{z} - Z(\mathbf{s}_0))^2$ subject to $E[\lambda' \mathbf{z}] = E[Z(\mathbf{s}_0)] \ \forall \ \boldsymbol{\beta}$.

Unbiasedness $\Rightarrow X'\lambda = x(s_0)$

Minimization yields the solution to:

$$\left(\begin{array}{cc} \Sigma & X \\ X' & 0 \end{array}\right) \left(\begin{array}{c} \lambda \\ m \end{array}\right) = \left(\begin{array}{c} c \\ x(s_0) \end{array}\right)$$



Introduction

Robustness

BLUP for Finite Populations

$$\left(\begin{array}{c} \mathbf{z}_{\scriptscriptstyle S} \\ \mathbf{z}_{\scriptscriptstyle u} \end{array} \right) = \left(\begin{array}{c} \mathbf{X}_{\scriptscriptstyle S} \\ \mathbf{X}_{\scriptscriptstyle u} \end{array} \right) \boldsymbol{\beta} + \left(\begin{array}{c} \boldsymbol{\epsilon}_{\scriptscriptstyle S} \\ \boldsymbol{\epsilon}_{\scriptscriptstyle u} \end{array} \right)$$

$$\operatorname{cov}\left(egin{array}{c} \epsilon_s \ \epsilon_u \end{array}
ight) = \left(egin{array}{cc} \Sigma_{s,s} & \Sigma_{s,u} \ \Sigma_{s,u}' & \Sigma_{s,s} \end{array}
ight)$$

Best Linear Unbiased Prediction (BLUP) (or [Universal] Kriging) minimize: $E(\lambda' \mathbf{z}_s - \mathbf{b}' \mathbf{z})^2$ subject to $E[\lambda' \mathbf{z}_s] = E[\mathbf{b}' \mathbf{z}] \ \forall \ \boldsymbol{\beta}$.

Unbiasedness $\Rightarrow X'_{c}\lambda = X'_{c}b_{s} + X'_{u}b_{u}$

Minimization yields the solution to:

$$\left(\begin{array}{cc} \boldsymbol{\Sigma}_{s,s} & \boldsymbol{X}_s \\ \boldsymbol{X}_s' & \boldsymbol{0} \end{array}\right) \left(\begin{array}{c} \boldsymbol{\lambda} \\ \boldsymbol{m} \end{array}\right) = \left(\begin{array}{cc} \boldsymbol{\Sigma}_{s,s} & \boldsymbol{\Sigma}_{u,u} \\ \boldsymbol{X}_s' & \boldsymbol{X}_u' \end{array}\right) \left(\begin{array}{c} \boldsymbol{b}_s \\ \boldsymbol{b}_u \end{array}\right)$$



BLUP for Finite Populations

$$\boldsymbol{\lambda}'\mathbf{z} = \mathbf{b}_{\scriptscriptstyle S}'\mathbf{z}_{\scriptscriptstyle S} + \mathbf{b}_{\scriptscriptstyle U}\hat{\mathbf{z}}_{\scriptscriptstyle U}$$

where

Introduction

$$\hat{\mathbf{z}}_{u} = \mathbf{\Sigma}_{u,s} \mathbf{\Sigma}_{s,s}^{-1} (\mathbf{z}_{s} - \hat{\boldsymbol{\mu}}_{s}) + \hat{\boldsymbol{\mu}}_{u},$$

$$\hat{\boldsymbol{\mu}}_{u} = \mathbf{X}_{u} \hat{\boldsymbol{\beta}}_{GLS}, \quad \hat{\boldsymbol{\mu}}_{s} = \mathbf{X}_{s} \hat{\boldsymbol{\beta}}_{GLS}, \quad \hat{\boldsymbol{\beta}}_{GLS} = (\mathbf{X}_{s}' \boldsymbol{\Sigma}_{s,s}^{-1} \mathbf{X}_{s})^{-1} \mathbf{X}_{s}' \boldsymbol{\Sigma}_{s,s}^{-1} \mathbf{z}_{s}$$

and

$$E(\lambda'\mathbf{z}_s - \mathbf{b}\mathbf{z})^2 = \mathbf{b}'\Sigma\mathbf{b} - \mathbf{c}_{\mathbf{b}}'\Sigma_{s,s}^{-1}\mathbf{c}_{\mathbf{b}} + \mathbf{d}_{\mathbf{b}}'(X_s'\Sigma_{s,s}^{-1}X_s)^{-1}\mathbf{d}_{\mathbf{b}}$$

where

$$\mathbf{c}_{\mathbf{b}} = \mathbf{\Sigma}_{s,s} \mathbf{b}_s + \mathbf{\Sigma}_{s,u} \mathbf{b}_u, \quad \mathbf{d}_{\mathbf{b}} = \mathbf{X}' \mathbf{b} - \mathbf{X}'_s \mathbf{\Sigma}_{s,s}^{-1} \mathbf{c}_{\mathbf{b}}$$



Connections to Sampling Theory

Let
$$\mathbf{X} = \left(\begin{array}{c} \mathbf{1}_n \\ \mathbf{1}_{N-n} \end{array} \right) \quad \mathbf{\Sigma} = \left(\begin{array}{cc} \sigma^2 \mathbf{I}_n & \mathbf{0} \\ \mathbf{0}' & \sigma^2 \mathbf{I}_{N-n} \end{array} \right) \quad \mathbf{b} = \left(\begin{array}{c} \mathbf{1}_n(1/N) \\ \mathbf{1}_{N-n}(1/N) \end{array} \right)$$
 Then

$$\lambda' \mathbf{z}_s = \bar{z}$$

and

$$E(\lambda'\mathbf{z}_s - \mathbf{b}\mathbf{z})^2 = \frac{\sigma^2}{n} \left(1 - \frac{n}{N}\right)$$



Connections to Sampling Theory

Let

$$\mathbf{X} = \begin{pmatrix} \mathbf{1}_{n_1} & \mathbf{0} \\ \mathbf{0} & \mathbf{1}_{n_2} \\ \mathbf{1}_{N_1 - n_1} & \mathbf{0} \\ \mathbf{0} & \mathbf{1}_{N_2 - n_2} \end{pmatrix} \quad \mathbf{\Sigma}_{s,s} = \begin{pmatrix} \sigma_1^2 \mathbf{I}_{n_1} & \mathbf{0} \\ \mathbf{0}' & \sigma_2^2 \mathbf{I}_{n_2} \end{pmatrix} \quad \mathbf{\Sigma}_{u,u} = \begin{pmatrix} \sigma_1^2 \mathbf{I}_{N_1 - n_1} & \mathbf{0} \\ \mathbf{0}' & \sigma_2^2 \mathbf{I}_{N_2 - n_2} \end{pmatrix}$$

and

$$\Sigma_{s,u} = \mathbf{0}, \quad \mathbf{z} = (\mathbf{z}_{s,1}', \mathbf{z}_{s,2}', \mathbf{z}_{u,1}', \mathbf{z}_{u,2})', \quad \mathbf{b} = \mathbf{1}_{N_1 + N_2}$$

Then

$$\boldsymbol{\lambda}'\mathbf{z}_s = N_1\bar{z}_{s,1} + N_2\bar{z}_{s,2}$$

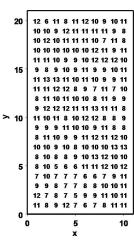
and

$$E(\lambda'\mathbf{z}_s - \mathbf{b}\mathbf{z})^2 = N_1^2 \frac{\sigma_1^2}{n_1} \left(1 - \frac{n_1}{N_1}\right) + N_2^2 \frac{\sigma_2^2}{n_2} \left(1 - \frac{n_2}{N_2}\right)$$

Ecoscience paper 2002

Simulations

Number of plant species in 70 cm × 70 cm plot



1000 samplings

N = 200n = 100



Fixed Pattern, Random Samples

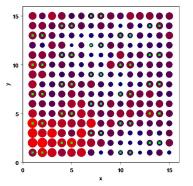
1000 random samples of size 100 Ver Hoef, J.M. 2002. Sampling and geostatistics for spatial data. *Ecoscience* 9: 152 - 161.

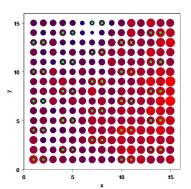
Validation Statistics	Simple Random Sample	Finite Block Pred
Bias	-0.002	-0.001
RMSPE	0.121	0.106
RAEV	0.122	0.105
80%CI	0.802	0.806



Random Pattern, Fixed Samples

Ver Hoef, J.M. 2002. Sampling and geostatistics for spatial data. Ecoscience 9: 152 - 161.





Ecoscience paper 2002

Random Pattern, Fixed Samples

Validation Statistics	SRS_{random}	$FPBK_{random}$	FPBK _{fixed}
Bias	0.522	-0.181	0.127
RMSPE	28.0	20.7	17.3
RAEV	28.0	20.3	17.5
80%CI	0.801	0.791	0.796



Simulation With Covariates

Ver Hoef, J.M. and Temesgen, H. 2013. A comparsion of the spatial linear model to nearest neighbor (k-NN) methods for forestry applications. *PloS* ONE 8(3): e59129. doi:10.1371/journal.pone.0059129.

$$\mathbf{w}_1 = \mathbf{z}_1 + \boldsymbol{\epsilon}_1$$

$$\mathbf{w}_{\eta} = \phi_{\eta - 1} \mathbf{w}_{\eta - 1} + \mathbf{z}_{\eta} + \boldsymbol{\epsilon}_{\eta}$$

$$\mathbf{x}_{\eta} = \boldsymbol{\mu}_{\eta} + \mathbf{w}_{\eta}$$

$$\mathbf{y} = \mathbf{X}\boldsymbol{\beta} + \mathbf{z}_{y} + \boldsymbol{\epsilon}_{y}$$

$$\mathbf{V}[j,j';\sigma^2,\rho] = \sigma^2 \left(1 - \frac{3}{2} \frac{d_{j,j'}}{\rho} + \frac{1}{2} \frac{d_{j,j'}}{\rho^3} \right) I\left(\frac{d_{j,j'}}{\rho} \le 1 \right)$$

8 spatially-patterned and cross-correlated covariates 2 excluded, 2 with $\beta_i = 0$ included



Prediction Methods

- mah1 k-NN that uses Mahalanobis distance with k=1.
- mah5: k-NN that uses Mahalanobis distance with k=5.
- msn1: k-NN that uses most significant neighbor (MSN) with k = 1.
- msn5: k-NN that uses MSN with k = 1.
- best: k-NN that uses both Mahalanobis distance and MSN, and tries $k = 1, 2, \dots, 30$, and then chooses the distance matrix and k with the smallest cross-validation RMSPF from the observed data.
- slm: a spatial linear model using the same covariates as all k-NN methods as main effects only, with an exponential autocovariance model estimated by REML, and using FPBK prediction and variance equations.
- Im: multiple regression like slmMain but assuming all random errors are independent.



Performance Measures

Root-mean-squared-prediction error (RMSPE):

RMSPE =
$$\sqrt{\frac{1}{m} \sum_{j=1}^{m} (\hat{\theta}_j - \theta_j)^2},$$

SRB: signed relative bias.

$$SRB = sign(\tau) \sqrt{\frac{\tau^2}{MSPE - \tau^2}},$$

where

$$\tau = \frac{1}{m} \sum_{j=1}^{m} (\hat{\theta}_j - \theta_j),$$

and sign(τ) is the sign (positive or negative) of τ .

PIC90: 90% prediction interval coverage

$$\text{PIC90} = \frac{1}{m} \sum_{j=1}^{m} I\left(\left(\hat{\theta}_{j} - 1.645 \hat{\text{se}}(\hat{\theta}_{j})\right) < \theta_{j} \& \theta_{j} < \left(\hat{\theta}_{j} + 1.645 \hat{\text{se}}(\hat{\theta}_{j})\right)\right),$$

where $\hat{se}(\hat{\theta}_i)$ is the estimated standard error of $\hat{\theta}_i$.



Table: Performance summaries from 2000 simulated spatial Gaussian data sets with 100 samples and 300 predictions each. Prediction methods form the columns and are described in Section 3.4. Performance measures form the rows and are described in Section 3.3.

	mah1	mah5	msn1	msn5	best	lm	slm	
Point								
RMSPE ^a	9.329	7.451	5.379	4.423	4.456	3.892	2.443	
SRB^b	-0.006	-0.009	0	-0.004	-0.004	-0.002	0.001	
PIC90 ^c	0.897	0.9	0.887	0.889	0.88	0.896	0.892	
	Total							
RMSPE	262.6	289.8	174.3	153.3	154.5	139.3	87.8	
SRB	-0.058	-0.067	-0.003	-0.034	-0.034	-0.02	0.009	
PIC90	0.952	0.87	0.914	0.886	0.874	0.88	0.887	

^a Root-mean-squared-prediction error



^b Signed relative bias

^c 90% prediction interval coverage

Poisson Simulations

Table: Performance summaries from 2000 simulated Poisson data sets with 100 samples and 300 predictions each. Prediction methods form the columns and are described in Section 3.4. Performance measures form the rows and are described in Section 3.3.

	mah1	mah5	msn1	msn5	best	lm	slm
			Tot	al			
RMSPE	320	295.9	296.3	262.3	272.2	283.1	226.1
SRB	-0.137	-0.188	-0.047	-0.135	-0.182	-0.033	-0.005
PIC90	0.912	0.842	0.9	0.867	0.83	0.86	0.858



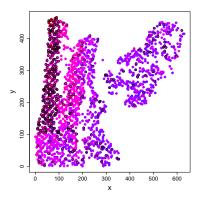
Bernoulli Simulations

Table: Performance summaries from 2000 simulated binary data sets with 100 samples and 300 predictions each. Prediction methods form the columns and are described in Section 3.4. Performance measures form the rows and are described in Section 3.3.

	mah1	mah5	msn1	msn5	best	lm	slm
			Propo	rtion			
RMSPE	0.0395	0.0394	0.0387	0.0334	0.0343	0.0329	0.0298
SRB	0.072	0.09	0.003	0.019	0.079	0.018	0.014
PIC90	0.919	0.841	0.913	0.882	0.84	0.886	0.884



Forestry Data Resampling PMAI



- response: maximum potential mean annual increment (PMAI)
- covariates: 1) temperature, 2) precipitation, 3)Climate Moisture Index, 4) an indicator variable for shade tolerance based on Western Hemlock trees, and 5) elevation



PMAI Resampling

Table: Performance summaries for 500 resamplings of PMAI forest data with 386 samples and 1500 predictions each. Prediction methods form the columns and are described in Section 3.4. Performance measures form the rows and are described in Section 3.3.

	mah1	mah5	msn1	msn5	best	lm	slm
			Tot	al			
RMSPE	219.1	230.7	243.3	200.9	223.2	197	180.4
SRB	0.437	0.712	0.064	-0.019	0.446	0.082	0.058
PIC90	0.944	0.838	0.948	0.922	0.834	0.904	0.904



Robustness

PMAI Resampling with Unbalanced Sampling

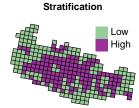
Table: Performance summaries for 500 resamplings of PMAI forest data with 386 spatially unbalanced samples and 1500 predictions each.

	mah1	mah5	msn1	msn5	best	lm	slm
			Tota	al			
RMSPE	637.9	853.6	457.1	442.2	576.1	608.1	269
SRB	2.635	4.055	1.418	1.77	1.651	2.86	0.369
PIC90	0.248	0.010	0.626	0.438	0.308	0.128	0.92



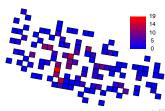
Robustness

Nome Moose Survey



Elevation 638 479 319 160 0

Counts



```
library (spPlotSampCourse)
library (maptools)
path <- system.file("rawdata/moose", package = "spPlotSampCourse")</pre>
samplesFile <- paste(path, "/", "Samples", sep = "")
samples <- readShapePoly(samplesFile)
samples@data[, "x"] <- LLtoUTM(samples@data[, "CENTRLAT"], samples@data[, "CENTRLON"])[,</pre>
samples@data[, "v"] <- LLtoUTM(samples@data[, "CENTRLAT"], samples@data[, "CENTRLON"])[,</pre>
samples@data[, "TOTAL"] <- as.numeric(as.character(samples@data[, "TOTAL"]))</pre>
samples@data[, "b"] <- rep(1, times = length(samples@data[, 1]))</pre>
sdata <- samples@data
coordinates(sdata) <- c("x", "v")</pre>
```

Nome Moose Survey

```
moFit <- splmm(TOTAL ~ ELEVMEAN + STRAT, spdata = sdata, varComps = "exponential")
# summary(moFit)
summary (moFit) $coefficients
##
               Estimate Std. Error t value Pr(>|t|)
## (Intercept) 2.592622
                           0.711248 3.6452 0.0004013
                0.001019 0.003291 0.3097 0.7573331
## ELEVMEAN
## STRATL
               -2.232373
                           0.611141 -3.6528 0.0003908
summary (moFit) $covparms
     Variance Component Parameter Type Estimate
## 1
                                nugget 1.928e-05
                 nugget
## 2
            exponential
                                parsil 1.006e+01
## 3
            exponential
                                range 1.043e+01
```



Nome Moose Survey

```
moFit <- splmm(TOTAL ~ STRAT - 1, spdata = sdata, varComps = "exponential")
summary (moFit) $coefficients
## Estimate Std. Error t value Pr(>|t|)
## STRATH 2.7442 0.5429 5.0547 1.609e-06
## STRATL 0.5289 0.5725 0.9239 3.574e-01
summary (moFit) $covparms
    Variance Component Parameter Type Estimate
## 1
               nuaaet
                      nugget 1.287e-04
         exponential parsil 9.925e+00
## 2
         exponential
## 3
                          range 1.023e+01
FPBK <- predictBlockFinPop(moFit, "b")</pre>
FPBK
## Resp Wts Pred Pred SE
## 1 TOTAL b 554.6 60.62
```

```
samples.H <- samples[samples@data[, "STRAT"] == "H", ]</pre>
samples.L <- samples[samples@data[, "STRAT"] == "L", ]</pre>
sdataH <- samples.H@data
coordinates(sdataH) <- c("x", "v")</pre>
sdataL <- samples.L@data
coordinates(sdataL) <- c("x", "v")</pre>
moFitH <- splmm(TOTAL ~ 1, spdata = sdataH, varComps = "exponential")
summary (moFitH) $coefficients
##
     Estimate Std. Error t value Pr(>|t|)
## (Intercept) 2.78 0.7452 3.731 0.0003782
moFitL <- splmm(TOTAL ~ 1, spdata = sdataL, varComps = "exponential")</pre>
summary (moFitL) $coefficients
       Estimate Std. Error t value Pr(>|t|)
## (Intercept) 0.3082 45.94 0.006709 0.9947
```

```
FPBK.H <- predictBlockFinPop (moFitH, "b")
FPBK.L <- predictBlockFinPop(moFitL, "b")</pre>
rbind (FPBK.H, FPBK.L)
## Resp Wts Pred Pred SE
## 1 TOTAL b 501.26 38.40
## 2 TOTAL b 51.38 14.08
FPBK.strat <- data.frame(Pred = FPBK.H[, "Pred"] + FPBK.L[, "Pred"], PredSE = sqrt(FPBK.H[,
    "Pred SE"]^2 + FPBK.L[, "Pred SE"]^2))
FPBK.strat
## Pred PredSE
## 1 552 6 40 9
FPBK
## Resp Wts Pred Pred SE
## 1 TOTAL b 554.6 60.62
```

Robustness of Block Prediction Methods

- BLUP is non-parametric
- ▶ REML are estimating equations, so covariance estimation is non-parametric
- Predictions are robust to mis-specification of covariance model
- Both predictor and predictand are linear combinations, so we can appeal to a correlated version of central limit theorem and use normal-distribution for probability statements (e.g., prediction intervals).

