## Spatial Leave One Out

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The purpose of the code is to evaluate the effectiveness of the spatial leave one out method for determining the predictive power of each model given. In order to acomplish this, the model given will be trained with data that has been reduced by **one sample**. **The sample that** is left out will then be used as a new data point to be predicted by the given model. Let's first start by generating some spatial autocorrelated data.

```
# Import Statement(s)
library(RandomFields)
# Create Grid
n = 100
# Set distance between points (Must be derived from data)
spat_range = 5
# Create a model with spatial dependence
mod_spat_dep = RMexp(var=1, scale=spat_range)
# Create spatially autocorralated predictor variables
x1 = RFsimulate(mod_spat_dep, x=1:n, y=1:n, grid=T)
x2 = RFsimulate(mod_spat_dep, x=1:n, y=1:n, grid=T)
x3 = RFsimulate(mod_spat_dep, x=1:n, y=1:n, grid=T)
# Create spatail error term
spat_err = RFsimulate(mod_spat_dep, x=1:n, y=1:n, grid=T)
# Convert objects to vectors
spat_err = as.vector(spat_err)
x1 = as.vector(x1)
x2 = as.vector(x2)
x3 = as.vector(x3)
y = 2*x1 + x2 + 3*x3 + spat_err
# Create coordinates
coords = cbind(rep(x=1:n, times=n), rep(x=1:n,each=n))
# Organize coordinates, predictors, and responce in data frame
dataset = data.frame(y, x1, x2, x3, coords)
head(dataset)
```

```
##
                     x1
                                 x2
                                            x3 X1 X2
            У
## 1 10.464906 0.6481868
                         1.46388265
                                    2.2667846
## 2 11.183313 1.2908259 1.43218933
                                    1.7588525
## 3 5.838396 0.6052640 1.00718886
                                    0.5743281
## 4 5.460486 1.0873799 0.14793471
                                    0.5645594
     3.845845 0.5288988 0.20043316
                                    0.5279037
## 6 2.348367 1.1348261 -0.04346441 -0.2177105 6 1
```

Now that we have our spatially autocorrelated data, lets take a sample of our data and verify that the results generated by the model are reasonable estimates of the true model's coefficients (Remebering that they should be close to the exspected values of: x1=2, x2=1, and x3=3).

```
# Create sample population
n = 2000
samp = sample(nrow(dataset), size=n)
# Create model(s)
mods = list()
mods$x1234_mod = glm(y \sim x1 + x2 + x3, data = dataset[samp,])
model = glm(y \sim x1 + x2 + x3, data = dataset[samp,])
summary(mods$x1234 mod)
##
## Call:
## glm(formula = y \sim x1 + x2 + x3, data = dataset[samp,])
## Deviance Residuals:
##
       Min
                 10
                      Median
                                   30
                                           Max
## -3.3753 -0.6498 -0.0104
                                         3.1780
                               0.6643
##
## Coefficients:
##
               Estimate Std. Error t value Pr(>|t|)
                           0.02249
                                     4.625 3.99e-06 ***
## (Intercept) 0.10402
                1.94477
                           0.02389
                                    81.409 < 2e-16 ***
## x2
                1.01779
                           0.02253
                                    45.178 < 2e-16 ***
## x3
                2.93773
                           0.02273 129.267 < 2e-16 ***
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for gaussian family taken to be 0.9974151)
##
##
       Null deviance: 29479.2 on 1999
                                        degrees of freedom
## Residual deviance: 1990.8 on 1996 degrees of freedom
## AIC: 5676.6
## Number of Fisher Scoring iterations: 2
```

Looking at the model summary, it appears that calculated coefficients are fairly close to actual coefficients. Now that we feel cofident about the sample taken, lets see how good the models generated will be at predicting new data. In order to do this, lets apply the spatial leave one out method to the data.

```
# Function(s)
get_training_rows = function(coords, dist_thres, longlat=FALSE) {
    ## Computing the distance matrix from the data:
    if (longlat) {
        require(sp)
        dist_matrix = spDists(coords, longlat=TRUE)
    }
    else
```

```
dist_matrix = as.matrix(dist(coords))
    ## Initializing the row indices of the dataset to be used in training
    training_rows = list()
    ## Creating the sets of training indices
    for (i in 1:nrow(dist_matrix)) {
        # Keeping only the observations far enough of the i-st observation by
        # using the threshold distance
        num_cell = which(dist_matrix[i, ] > dist_thres)
        training_rows[[i]] = num_cell
    return(training_rows)
}
sloo_simple = function(model,training){
  intercept = NULL
 x1 = NULL
  x2 = NULL
  x3 = NULL
  predicted = NULL
  observed = NULL
  logLik = NULL
  # Creating the response variable name
  y <- as.character(x=formula(x=model)[2])</pre>
  # Initializing the 'logLik' object
  # logLik <- vector(mode="numeric", length=nrow(x=model$data))</pre>
    # Calculating the SLOO logLikelihoods for each observation i
    for(i in 1:nrow(x=model$data)){
        # Extracting the i-st training set:
       training_data <- model$data[training[[i]],]</pre>
        # Calculating the model parameters from the i-st training set:
        m <- glm(formula=formula(model),data=training_data,family= model$family)</pre>
    intercept[i] = coef(m)[1]
    x1[i] = coef(m)[2]
    x2[i] = coef(m)[3]
    x3[i] = coef(m)[4]
        # Predicting the i-st observation from the i-st training set:
        predicted[i] <- predict(object=m,newdata=model$data[i,],type="response")</pre>
    observed[i] <- residuals(m) + predict(m)</pre>
        # Calculating the probability of the i-st observed value according
    # to the predicted one by the i-st training set:
    logLik[i] <- dnorm(x=model$data[i,y],mean=predicted[i],</pre>
                       sd=sqrt(sum(residuals(m)^2)/nrow(training_data)),
                      log=T)
  finalMatrix = data.frame(intercept, x1, x2, x3, predicted, observed, logLik)
    # Calculating the overall SLOO logLikelihood:
    # Sum.logLik <- sum(logLik)</pre>
    return(finalMatrix)
}
training = get_training_rows(coords[samp, ], dist_thres = spat_range)
```

```
# evalutate the four models using log likelihood
# with the SLOO and a non-spatial approach
sloo_ll = sloo_simple(model, training)
#ll = sapply(mods, logLik)
head(sloo_ll)
```

```
## intercept x1 x2 x3 predicted observed logLik

## 1 0.11528982 1.953732 1.023969 2.928731 0.3366165 1.975940 -2.2423216

## 2 0.09513063 1.954122 1.007003 2.938309 -0.2376246 -1.282299 -3.3876483

## 3 0.10395412 1.944242 1.017989 2.934739 4.2300400 -1.282299 -0.9864654

## 4 0.10130567 1.943705 1.018483 2.940407 3.8182245 -1.282299 -0.9708725

## 5 0.10829345 1.945668 1.016436 2.937313 -0.1077664 -1.282299 -1.1648178

## 6 0.10249711 1.945144 1.013993 2.936559 -1.2009444 -1.282299 -1.0184229
```