Spatial sampling with SamplingStrata

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Optimization with the spatial method

Let us suppose we want to design a sample survey with k Z target variables, each one of them correlated to one or more of the available Y frame variables.

When frame units are georeferenced or geocoded, the presence of spatial auto-correlation can be investigated. This can be done by executing for instance the Moran test on the target variables: if the null hypothesis is rejected (i.e. the hypothesis of the presence of spatial auto-correlation is accepted) then we should take into account also this variance component.

As indicated by deGruijter et al. (2016) and deGruijter, Wheeler, and Malone (2019), in case Z is the target variable, omitting as negligible the *fpc* factor, the sampling variance of its estimated mean is:

$$V(\hat{ar{Z}}) = \sum_{h=1}^{H} (N_h/N)^2 S_h^2/n_h$$

We can write the variance in each stratum h as:

$$S_h^2 = rac{1}{N_h^2}\sum_{i=1}^{N_{h-1}}\sum_{j=i+1}^{N_h}(z_i-z_j)^2$$

The optimal determination of strata is obtained by minimizing the quantity *O*:

$$O = \sum_{h=1}^{H} rac{1}{N_h^2} \{ \sum_{i=1}^{N_{h-1}} \sum_{j=i+1}^{N_h} (z_i - z_j)^2 \}^{1/2}$$

Obviously, values z are not known, but only their predictions, obtained by means of a regression model. So, in Equation we can substitute $(z_i - z_j)^2$ with

$$D_{ij}^2 = rac{({ ilde z}_i - { ilde z}_j)^2}{R^2} + V(e_i) + V(e_j) - 2Cov(e_i,e_j)$$

where R^2 is the squared correlation coefficient indicating the fitting of the regression model, and $V(e_i)$, $V(e_j)$ are the model variances of the residuals. The spatial auto-correlation component is contained in the term $Cov(e_i, e_j)$.

In particular, the quantity D_{ij} is calculated in this way:

$$D_{ij}^2 = rac{({ ilde z}_i - { ilde z}_j)^2}{R^2} + (s_i^2 + s_j^2) - 2 s_i s_j e^{-k(d_{ij}/range)}$$

where d_{ij} is the Euclidean distance between two units i and j in the frame (calculated using their geographical coordinates, that must be expressed in meters), the s_i and s_j are estimates of the prediction errors in the single points and *range* is the maximum distance below which spatial auto-correlation can be observed among points. The value of *range* can be determined by an analysis of the spatial *variogram*.

To summarize, when frame units can be geo-referenced, the proposed procedure is the following:

- acquire coordinates of the geographic location of the units in the population of interest;
- fit a *kriging* model (or any other spatial model) on data for each Z;
- $\circ\,$ obtain predicted values together with prediction errors for each Z and associate them to each unit in the frame;
- perform the optimization step.

In order to illustrate the "*spatial*" method, we make use of a dataset generally employed as an example of spatially correlated phenomena (in this case, the concentration of four heavy metals in a portion of the river Meuse). This dataset comes with the library "*sp*":

```
library(sp)
# locations (155 observed points)
data("meuse")
# grid of points (3103)
data("meuse.grid")
meuse.grid$id <- c(1:nrow(meuse.grid))
coordinates(meuse)<-c("x","y")
coordinates(meuse.grid)<-c("x","y")
lm_lead <- lm(log(lead) ~ dist,data=meuse)
lm_zinc <- lm(log(zinc) ~ dist,data=meuse)</pre>
```



We analyse the territorial distribution of the *lead* and *zinc* concentration, and model (by using the *universal kriging*) their relations with distance from the river, using the subset of 155 points on which these values have been jointly observed:

```
library(automap)
kriging_lead = autoKrige(log(lead) ~ dist, meuse, meuse.grid)
plot(kriging_lead,sp.layout = NULL, justPosition = TRUE)
```



kriging_zinc = autoKrige(log(zinc) ~ dist, meuse, meuse.grid)
plot(kriging_zinc, sp.layout = list(pts = list("sp.points", meuse)))



Using these *kriging* models, we are able to predict the values of lead and zinc concentration on the totality of the 3,103 points in the Meuse territory:

```
df <- NULL
df$id <- meuse.grid$id
df$lead.pred <- kriging_lead$krige_output@data$var1.pred
df$lead.var <- kriging_lead$krige_output@data$var1.var
df$zinc.pred <- kriging_zinc$krige_output@data$var1.pred
df$zinc.var <- kriging_zinc$krige_output@data$var1.var
df$lon <- meuse.grid$x
df$lat <- meuse.grid$y
df$lat <- 1
df <- as.data.frame(df)
head(df)</pre>
```

```
      #
      id Lead.pred Lead.var zinc.pred zinc.var Lon Lat dom1

      #
      1
      5.509360
      0.1954937
      6.736502
      0.2007150
      181180
      333740
      1

      #
      2
      2
      5.546006
      0.1716895
      6.785460
      0.1749260
      181140
      333700
      1

      #
      3
      3
      5.488913
      0.1784052
      6.698883
      0.1826314
      181180
      333700
      1

      #
      4
      4
      5.388320
      0.1855561
      6.558216
      0.1906426
      181220
      333700
      1

      #
      5
      5
      5.584415
      0.1463018
      6.841612
      0.1465346
      181100
      333660
      1

      #
      6
      5.525538
      0.1533757
      6.749216
      0.1549663
      181140
      333660
      1
```

The aim is now to produce the optimal stratification of the 3,103 points under a precision constraint of 1% on the target estimates of the mean *lead* and *zinc* concentrations:

```
library(SamplingStrata)
frame <- buildFrameSpatial(df=df,</pre>
                      id="id",
                      X=c("lead.pred","zinc.pred"),
                      Y=c("lead.pred","zinc.pred"),
                      variance=c("lead.var","zinc.var"),
                      lon="lon",
                      lat="lat",
                      domainvalue = "dom1")
# id
                              Y1
                                       Y2
                                                                         Lat domainvalue
            X1
                     X2
                                               var1
                                                         var2
                                                                  Lon
# 1 1 5.509360 6.736502 5.509360 6.736502 0.1954937 0.2007150 181180 333740
                                                                                        1
# 2 2 5.546006 6.785460 5.546006 6.785460 0.1716895 0.1749260 181140 333700
                                                                                        1
# 3 3 5.488913 6.698883 5.488913 6.698883 0.1784052 0.1826314 181180 333700
                                                                                        1
# 4 4 5.388320 6.558216 5.388320 6.558216 0.1855561 0.1906426 181220 333700
                                                                                        1
# 5 5 5.584415 6.841612 5.584415 6.841612 0.1463018 0.1465346 181100 333660
                                                                                        1
# 6 6 5.525538 6.749216 5.525538 6.749216 0.1533757 0.1549663 181140 333660
                                                                                        1
cv <- as.data.frame(list(DOM=rep("DOM1",1),</pre>
```

```
CV1=rep(0.01,1),

CV2=rep(0.01,1),

domainvalue=c(1:1) ))

CV

# DOM CV1 CV2 domainvalue

# 1 DOM1 0.01 0.01 1
```

To this aim, we carry out the optimization step by indicating the method spatial:

```
set.seed(1234)
solution <- optimStrata (</pre>
 method = "spatial",
 errors=cv,
  framesamp=frame,
 iter = 15,
  pops = 10,
  nStrata = 5,
  fitting = c(summary(lm lead)$r.square,summary(lm zinc)$r.square),
  range = c(kriging_lead$var_model$range[2],kriging_zinc$var_model$range[2]),
  kappa=1)
# *** Domain : 1
                    1
# Number of strata : 3103
# GA Settings
  Population size
#
                        = 10
   Number of Generations = 15
#
```

```
Elitism
#
                       = 2
#
   Mutation Chance
                       = 0.111111111111111
#
#
#
  *** Sample cost: 79.87774
#
  *** Number of strata: 4
#
  *** Sample size : 80
#
   *** Number of strata : 4
#
      #
```



obtaining the following optimized strata:



that can be visualised in this way:

frameres <- SpatialPointsDataFrame(data=framenew, coords=cbind(framenew\$LON,framenew\$LAT))
frameres2 <- SpatialPixelsDataFrame(points=frameres[c("LON","LAT")], data=framenew)
frameres2\$LABEL <- as.factor(frameres2\$LABEL)
spplot(frameres2,c("LABEL"), col.regions=bpy.colors(5))</pre>



We can now proceed with the selection of the sample:

whose units are so distributed in the territory:



References

deGruijter, J. J., A. B. McBratney, B. Minasny, I. Wheeler, B. P. Malone, and U. Stockmann. 2016. "Farm-Scale Soil Carbon Auditing." *Geoderma*, no. 265: 12–130.

deGruijter, J. J., I. Wheeler, and B. P. Malone. 2019. "Using Model Predictions of Soil Carbon in Farm-Scale Auditing - a Sofwtare Tool." *Agricultural Systems*, no. 169(C): 24–30.