# Validation of (digital) soil maps

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## Map validation

- No map is perfect. All maps, including soil maps, are representations of reality that are often based on an underlying model.
- This means that there will always be a **deviation** between the phenomenon depicted on the map and the phenomenon observed in the real world, i.e. each map will contain **errors**.
- Validation is defined an activity in which the soil map predictions are compared with observed values.
- From this comparison, the map quality can be quantified and summarized using map quality measures.

## Why validate?

- Validation is an important step in the soil mapping workflow.
- Why do we want to validate: soil maps are not perfect!
  - One should want to check the quality of ones work before this is made public
  - Compare the performance of methods
  - End users must know the quality of maps to judge their usability for specific purposes

### Validation data

- Internal versus external accuracy.
- Validation should be done with independent data, i.e. data not used for the production of the soil map.
- Validation provides summary global measures of accuracy: how accurate the map is on average for the mapping area (i.e. what is the expected error at a randomly selected location in the mapping area)
- Uncertainty assessment provides local measures of accuracy (i.e. pixel specific)

Prediction error:

$$e(\mathbf{s}) = \hat{z}(\mathbf{s}) - z(\mathbf{s})$$

Mean error (bias: systematic over- or underestimation):

$$ME = \bar{e} = \frac{1}{N} \sum_{i=1}^{N} e(\mathbf{s}_i)$$

Mean absolute error and (root) mean squared error:

$$MAE = |\bar{e}| = \frac{1}{N} \sum_{i=1}^{N} |e|(\mathbf{s}_i)$$

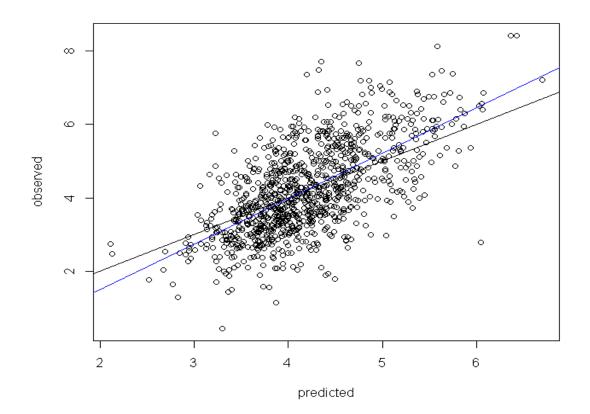
$$MSE = \overline{e^2} = \frac{1}{N} \sum_{i=1}^{N} e^2(\mathbf{s_i})$$

Amount of variance explained:

$$AVE = 1 - \frac{\sum_{i=1}^{N} (\hat{z}(\mathbf{s}_i) - z(\mathbf{s}_i))^2}{\sum_{i=1}^{N} (z(\mathbf{s}_i) - \bar{z})^2}$$

• It is not good practice in validation to do a regression between the observed and predicted value and use the  $R^2$  as measure for the amount of variance explained.

- Black: 1:1 line; Blue: regression line
- AVE = 0.40;  $R^2$  = 0.42



 Mean square deviation ratio: measures how well the prediction model estimates the prediction uncertainty:

$$MSDR = \frac{1}{N} \sum_{i=1}^{N} \frac{\left(\hat{z}(\mathbf{s}_i) - z(\mathbf{s}_i)\right)^2}{\sigma^2(\mathbf{s}_i)}$$

- MSDR should be 1; median SDR 0.455
- MSDR < 1: model overestimates the uncertainty</li>
- MSDR > 1: model underestimates the uncertainty

## Estimating map quality measures

- We estimate the population means of the map quality measures from a sample taken from a limited number of locations in the mapping area.
- Because we estimate, we are uncertain about the estimations.
- When the validation data are collected properly with a probability sampling design - we can quantify this uncertainty.
- We use the same equations as before but now N is replaced by n:

$$ME = \hat{e} = \frac{1}{n} \sum_{i=1}^{n} e(\mathbf{s}_i)$$

## Sampling types

- Purposive sampling: locations are selected purposively (e.g. representative, good spatial coverage)
- Haphazard sampling: locations are selected arbitrarily, 'more-or-less random'. Be ware: this is **NOT** a random sample.
- Probability sampling:
  - locations are selected randomly (based on a probability mechanism).
     Inclusions probabilities are known.
  - Design-based estimation of the accuracy statistics.
  - Sampling data are independent: spatial correlation does not have to be accounted for.

### Validation methods

#### Methods:

- Additional probability sampling
- Data-splitting
- Cross-validation (*n*-fold, leave-one-out)
- Data-splitting and cross-validation if there is only one dataset available for calibration and validation.

#### Literature:

 Brus, Kempen, Heuvelink, 2011. Sampling for validation of digital soil maps. European Journal of Soil Science 62, 394-407.

## Additional probability sampling

- For validation it is preferred to use data collected from randomly selected locations, because:
  - no model is needed for estimating map quality estimates. We can apply design-based estimation, meaning that model-free unbiased and valid estimates of the map quality measures can be obtained;
  - discussions on the **validity** of the estimated map quality are avoided;
  - model-free, valid estimates of the variance of the map quality measures can be obtained that allow for hypothesis testing, e.g. for comparison of model performance.

# Additional probability sampling

- All sampling units have probability >0 of being selected, but the probabilities need not be equal
- Inclusion probabilities must be known for all sampling units in the population
- Inclusion probabilities are known by design and are used to estimate the quality measures: design-based estimation
- Various designs: sample, stratified, clustered, two-stage, systematic random sampling

## Data-splitting

- Sample data set is split in two subsets.
- One subset is used to calibrate the prediction model. The other subset is used for validation.
- A frequently used splitting criterion is 70-30, where 70% of the sample data are used for calibration and 30% for validation.
- For sparse data sets, data-splitting can be inefficient since the information in the data set is not fully exploited for both calibration and validation.

## Data-splitting

- A random subsample of a non-probability sample is not a probability sample of the study area -> design-based estimation of quality measures impossible.
- If the validation subset is a non-probability sample of the study area: one must account for possible spatial autocorrelation of prediction (classification) error, i.e. modelbased estimation.
- Often, spatial autocorrelation is not accounted for. Map quality measures cannot be considered unbiased and valid estimates of the population means.
- In this case, the map quality measures are only valid at the validation locations.

### **Cross-validation**

- n-fold cross-validation
- Procedure:
  - Data is split in n subsets of equal size.
  - Model is calibrated using data from n-1 subsets.
  - Model is used to predict at the subset left out.
  - Repeated n times: each time setting aside a different subset.
- Special case: leave-one-out (n=number of samples)
- More efficient than data-splitting
- Problem of spatially correlated errors remains

## Tools in R

• Kriging *n*-fold cross-validation: **krige.cv** function of the **gstat** package.

```
rfk.cv <- krige.cv(formula = resid \sim 1, nfold = 10, locations = d, model = vmf)
```

• *n*-fold cross-validation: **train** function of the **caret** package.

```
## set cross-valiation parameters
cvPar <- trainControl(
  method = "cv",
  number = 10,
  verboseIter = TRUE,
  savePredictions = TRUE
)

## cross-validation with caret package
rf.cv <- train(x = covar, y = tval, method = "rf", tuneGrid = data.frame(mtry = 15), trControl = cvPar)</pre>
```

Subsetting data: createFolds function of the caret package.

```
d$fold <- createFolds(d$ocs1, k=5, FALSE)
```

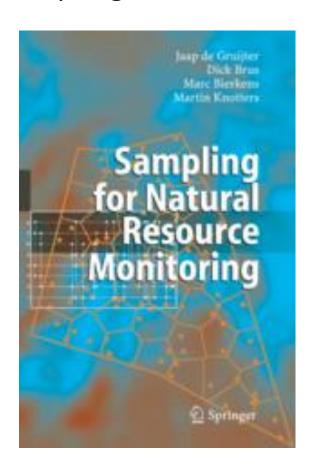
Note: using categorical covariates might give problems for cross-validation

# Sampling for mapping

- Does not have to be a probability (random) sampling (can be sub-optimal)
- Different designs are possible
  - Minimizing prediction variance: spatial coverage sampling
  - Covering the attribute space: Latin hypercube sampling
  - Variogram estimation: spatial coverage + small clusters
- When designing a sampling scheme: start with careful planning of the entire project to avoid mismatch between data acquisition and analysis

### Good resource

• Sampling for Natural Resource Monitoring; De Gruijter, Brus, Bierkens, Knotters; Springer, ISBN 978-3-540-33161-2.



# Thank you for listening