
Introduction

Data: set of n attribute measurements $\{z(\mathbf{s}_i), i = 1, \dots, n\}$, available at n sample locations $\{\mathbf{s}_i, i = 1, \dots, n\}$

Objectives:

- quantify spatial *auto-correlation*, or attribute dissimilarity typically expressed as: $\frac{1}{2}[z(\mathbf{s}_i) - z(\mathbf{s}_j)]^2$ as a function of separation distance between sample pairs \mathbf{s}_i and \mathbf{s}_j
- introduce the sample semivariogram, its characteristics, and provide some examples
NOTE: *Spatial auto-correlation is a second-order characteristic of spatial variation, and hence the sample semivariogram should be computed from data whose spatial variation is not explained by first-order effects*
- justify the need of going beyond the sample semivariogram to a semivariogram model
- introduce parametric functions of distance that can be used as formal theoretical semivariogram models
- discuss issues of fitting semivariogram models to sample semivariogram values

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Semivariogram Cloud

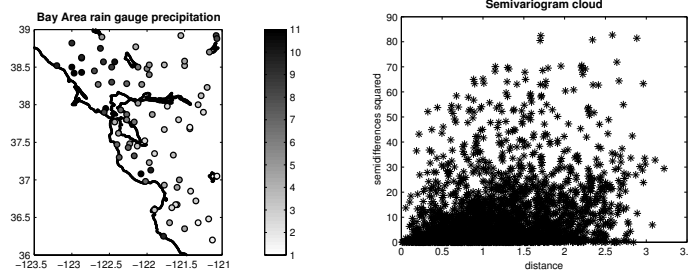
Definition: A scatter-plot of *attribute* squared semidifferences between all possible pairs of samples measured at different locations, versus their separation distance

Computational procedure:

1. construct Euclidean distance matrix $\mathbf{D} = [d_{ij}, i = 1, \dots, n, j = 1, \dots, n]$ between all n^2 pairs of data locations, where d_{ij} is defined as: $d_{ij} = \|\mathbf{h}_{ij}\| = \|\mathbf{s}_i - \mathbf{s}_j\|$
2. construct squared semidifference matrix $\mathbf{E} = [e_{ij}, i = 1, \dots, n, j = 1, \dots, n]$ between all n^2 pairs of attribute values, where e_{ij} is defined as: $e_{ij} = \frac{1}{2}[z(\mathbf{s}_i) - z(\mathbf{s}_j)]^2$
3. plot each distance value d_{ij} against the corresponding squared semidifference e_{ij} ; in other words, plot $\mathbf{e} = \text{vec}(\mathbf{E})$ versus $\mathbf{d} = \text{vec}(\mathbf{D})$. The plot of all pairs $\{d_{ij}, e_{ij}\}$ is termed a semivariogram cloud

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Semivariogram Cloud Example



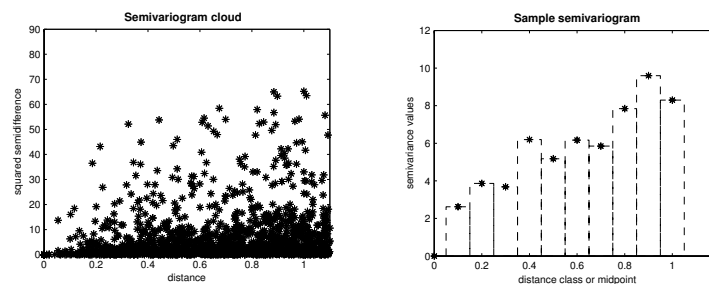
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A measure of dissimilarity between attribute values measured at different locations, i.e., a spatial measure of attribute dissimilarity

Expected graph pattern: As the distance d_{ij} between sample pairs increases, the corresponding squared semidifference e_{ij} should also increase

Difficult to interpret, so we consider groups of sample pairs separated by similar distances i.e., average squared semidifferences within distance classes (x-axis bins in the right graph above)

Semivariogram Cloud Versus Plot



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Going from the first to the second:

- define a set of L distance classes; the l -th class has limits: $(d_l - t_l, d_l + t_l]$, where d_l is the class midpoint and t_l is half the class width (or distance tolerance)
- for a given distance class $(d_l - t_l, d_l + t_l]$, the semivariogram value $\hat{\gamma}(d_l)$ is the average of $n(d_l) \ll n^2$ squared attribute semidifferences computed from sample pairs whose inter-distances d_{ij} satisfy: $d_l - t_l < d_{ij} \leq d_l + t_l$
- in other words, the semivariogram plot can be regarded as a summary of the semivariogram cloud, according to some distance-based grouping of samples

Computing Sample Semivariograms

- compute distance matrix $\mathbf{D} = [d_{ij}, i = 1, \dots, n, j = 1, \dots, n]$ and squared semidifference matrix $\mathbf{E} = [e_{ij}, i = 1, \dots, n, j = 1, \dots, n]$ between n^2 data pairs

$$\mathbf{D} = \begin{bmatrix} 0 & d_{12} & d_{13} & d_{14} & d_{15} \\ d_{12} & 0 & d_{23} & d_{24} & d_{25} \\ d_{13} & d_{23} & 0 & d_{34} & d_{35} \\ d_{14} & d_{24} & d_{34} & 0 & d_{45} \\ d_{15} & d_{25} & d_{35} & d_{45} & 0 \end{bmatrix} \quad \mathbf{E} = \begin{bmatrix} 0 & e_{12} & e_{13} & e_{14} & e_{15} \\ e_{12} & 0 & e_{23} & e_{24} & e_{25} \\ e_{13} & e_{23} & 0 & e_{34} & e_{35} \\ e_{14} & e_{24} & e_{34} & 0 & e_{45} \\ e_{15} & e_{25} & e_{35} & e_{45} & 0 \end{bmatrix}$$

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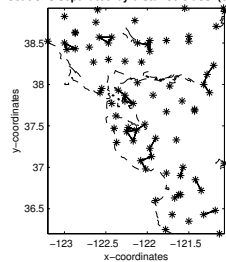
- for a given distance class $(d_l - t_l, d_l + t_l]$, find entries of \mathbf{E} that correspond to entries of \mathbf{D} falling in that distance class, e.g.:

$$\begin{bmatrix} 0 & \boxed{d_{12}} & d_{13} & d_{14} & d_{15} \\ \boxed{d_{12}} & 0 & d_{23} & \boxed{d_{24}} & d_{25} \\ d_{13} & d_{23} & 0 & d_{34} & \boxed{d_{35}} \\ d_{14} & \boxed{d_{24}} & d_{34} & 0 & d_{45} \\ d_{15} & d_{25} & \boxed{d_{35}} & d_{45} & 0 \end{bmatrix} \quad \longrightarrow \quad \begin{bmatrix} 0 & \boxed{e_{12}} & e_{13} & e_{14} & e_{15} \\ \boxed{e_{12}} & 0 & e_{23} & \boxed{e_{24}} & e_{25} \\ e_{13} & e_{23} & 0 & e_{34} & \boxed{e_{35}} \\ e_{14} & \boxed{e_{24}} & e_{34} & 0 & e_{45} \\ e_{15} & e_{25} & \boxed{e_{35}} & e_{45} & 0 \end{bmatrix}$$

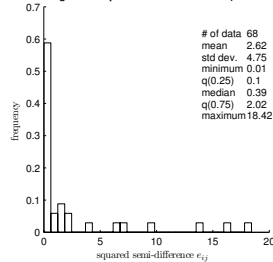
- sample semivariogram $\hat{\gamma}(d_l)$ for that class is the average of the $n(d_l)$ squared semidifferences, e -values, whose corresponding distances, d -values, fall in class $(d_l - t_l, d_l + t_l]$; i.e., the mean of all e -values in boxes in the matrix on the right above

Examples of Semivariogram Computation

Locations separated by distance class (0.05 0.15]

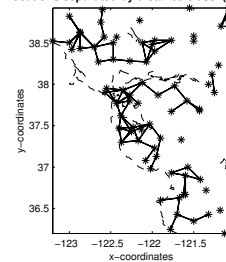


Histogram of squared semi-differences (0.05 0.15]

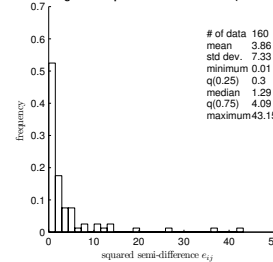


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Locations separated by distance class (0.15 0.25]



Histogram of squared semi-differences (0.15 0.25]



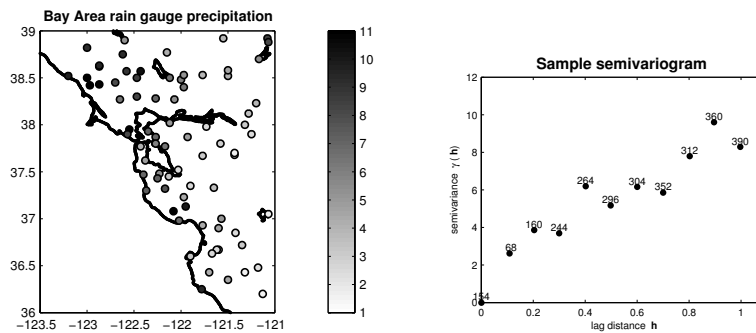
$\hat{\gamma}((0.05 \ 0.15]) = 2.62$, $\hat{\gamma}((0.15 \ 0.25]) = 3.86$ = averages of values displayed in histograms
 Map views linking sample pairs that contribute to such histograms are extremely informative

Sample Semivariogram Plots

Consider a set of L distance classes with midpoints $\{d_l, l = 1, \dots, L\}$ and tolerances $\{t_l, l = 1, \dots, L\}$. The plot of semivariance values $\{\hat{\gamma}(d_l), l = 1, \dots, L\}$ versus the average sample inter-distance for each class is called a sample semivariogram

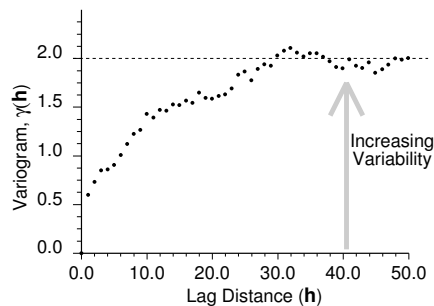
$$\hat{\gamma}(d_l) = \frac{1}{n(d_l)} \sum_{c=1}^{n(d_l)} e_c = \frac{1}{2n(d_l)} \sum_{d_{ij} \in (d_l - t_l, d_l + t_l]}^{n(d_l)} [z(\mathbf{s}_i) - z(\mathbf{s}_j)]^2$$

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numbers above bullets denote # of sample pairs contributing to $\hat{\gamma}(d_l)$ at each lag distance
could also graph variances of e -values within the distance classes; $\hat{\gamma}(0) = 0$, always

Semivariogram Characteristics

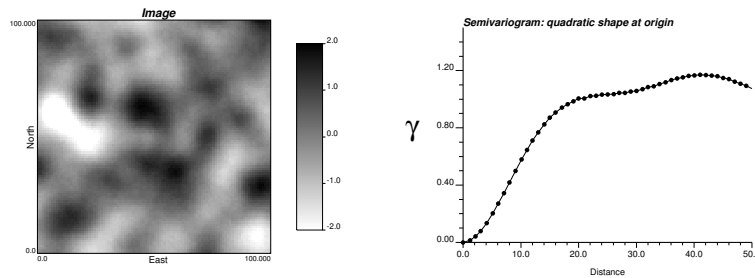


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- **sill**: limit semivariogram value (plateau) is approximately equal to sample variance (for representative sample)
- **range**: distance at which semivariogram reaches (or starts oscillating around) sill = distance of influence of any datum on another
- **nugget effect**: discontinuity at origin ($\hat{\gamma}(\epsilon) > \epsilon$); sum of measurement error and micro-structures (variability at scales smaller than sampling interval)
watch out for sparse data, outliers and positional or attribute errors
- transformation of Euclidean distance into statistical “distance” bearing imprint of specific phenomenon

Sample Semivariogram Shape & Interpretation (1)

Quadratic shape near origin:



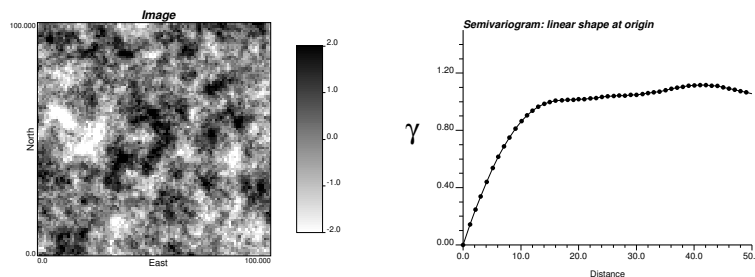
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Interpretation:

- highly continuous (extremely smooth) spatial attribute variability
- spatial attribute is differentiable
- typical variables: elevation, temperature, ...

Sample Semivariogram Shape & Interpretation (2)

Linear shape near origin:



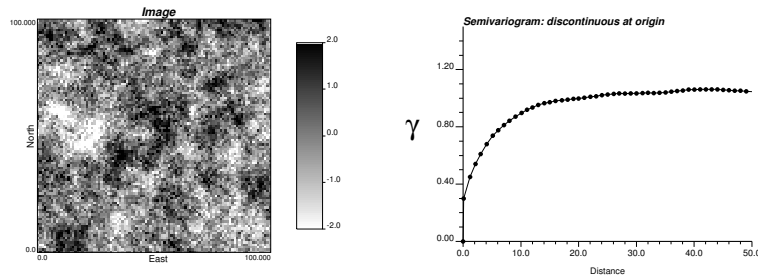
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Interpretation:

- continuous variability (not extremely smooth) of spatial attribute
- attribute is not differentiable
- typical variables: ore grades, ...

Sample Semivariogram Shape & Interpretation (3)

Discontinuous near origin:



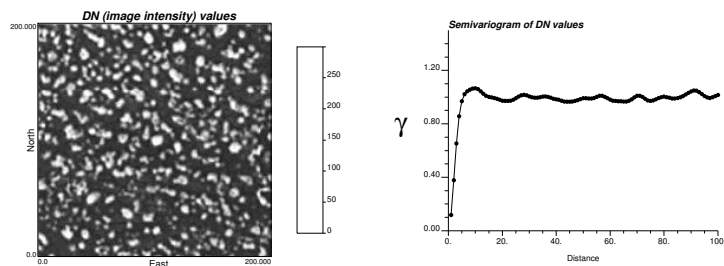
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Interpretation:

- highly irregular (quasi-random) spatial variability at small scales
- typical variables: precipitation, . . .

Sample Semivariogram Shape & Interpretation (4)

Oscillating (around sill):



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Interpretation:

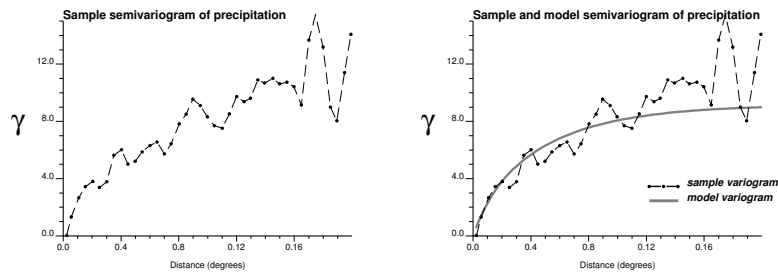
- periodic variability of spatial attribute yields sinusoidal semivariogram
- semivariogram shape possibly due to limited sampling
- need to provide physical evidence for periodicity
- frequently encountered in time series

The Need for Semivariogram Models

Problems: (i) sill, range, and relative nugget, cannot be determined directly from the sample semivariogram plot, (ii) a continuum of semivariogram values $\gamma(d)$ for any distance vector d is required in interpolation, but *sample* semivariogram values $\{\hat{\gamma}(d_l), l = 1, \dots, L\}$ are typically calculated only for few (L) distances $\{d_l, l = 1, \dots, L\}$.

Semivariogram model definition: parametric function $\gamma(d; \theta)$ fitted to sample semivariogram values $\{\hat{\gamma}(d_l), l = 1, \dots, L\}$; θ denotes parameter vector with, e.g., range, and sill (for a given semivariogram function)

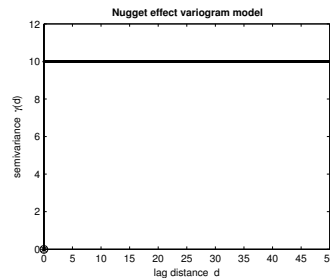
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semivariogram modeling is more than a curve fitting exercise;

Warning: cannot use any curve as semivariogram model !!!

Valid Semivariogram Models: Pure Nugget Effect



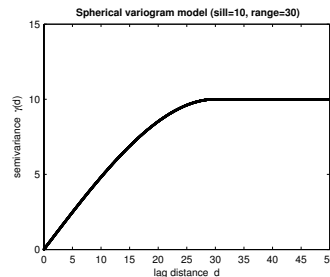
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$$\gamma(d; \theta) = \begin{cases} 0, & \text{if } d = 0 \\ \sigma, & \text{if } d > 0 \end{cases}$$

$\theta = [\sigma]$, where σ denotes attribute variance

- indicates complete absence of spatial correlation
- could occur due to measurement error and microstructure, i.e., features occurring at scales smaller than sampling interval

Valid Semivariogram Models: Spherical



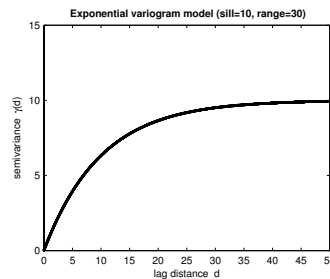
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$$\gamma(d; \theta) = \begin{cases} \sigma \left[\frac{3}{2} \left(\frac{d}{r} \right) - \frac{1}{2} \left(\frac{d}{r} \right)^3 \right], & \text{if } d < r \\ \sigma, & \text{if } d \geq r \end{cases}$$

$\theta = [\sigma \ r]$, where r is the model range

- linear behavior at origin
- clearly defined range parameter r

Valid Semivariogram Models: Exponential



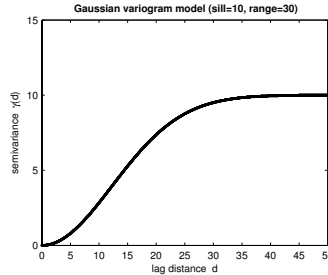
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$$\gamma(d; \theta) = \sigma \left[1 - \exp\left(-\frac{3d}{r}\right) \right]$$

$\theta = [\sigma \ r]$

- linear behavior at origin; rises faster than spherical; reaches sill asymptotically
- effective range parameter r ; distance at which 95% of sill reached

Valid Semivariogram Models: Gaussian



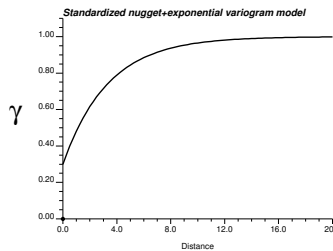
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$$\gamma(d; \theta) = \sigma \left[1 - \exp\left(-\frac{3d^2}{r^2}\right) \right]$$

$$\theta = [\sigma \ r]$$

- quadratic behavior at origin; implies smooth spatial variability of attribute values; reaches sill asymptotically
- effective range parameter r ; distance at which 95% of sill reached

Valid Semivariogram Models: Nugget + Exponential



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$$\gamma(d; \theta) = \begin{cases} 0, & \text{if } d = 0 \\ a + ([\sigma - a][1 - \exp(\frac{3d}{r})]), & \text{if } d \geq \epsilon \end{cases}$$

$$\theta = [\sigma \ a \ r]$$

- discontinuous at origin; reaches sill asymptotically
- practical range parameter r ; distance at which 95% of sill reached
- a/σ = relative nugget contribution = proportion (to total sill) of purely random spatial variability
- more complex models can be built by adding or multiplying valid models

Fitting Semivariogram Models to Sample Data

Or fitting valid semivariogram functions (curves) to sample semivariogram values

Manual fitting:

- select number of semivariograms, their type (functional form), sill, and range
- model behavior at origin (nugget effect, shape of semivariogram at distances smaller than first lag) using prior knowledge about phenomenon

Automatic fitting:

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- least squares fit (ordinary, generalized, weighted): choose semivariogram model parameters (typically iteratively) so as to minimize discrepancy between model and sample semivariogram values over all lags; other methods also available
- treat with caution, especially with sparse data and outliers

Cross-validation:

- given a proposed parameter set, i.e., a semivariogram model, perform cross-validation using geostatistical interpolation, and record resulting error statistics
- repeat with different model parameters, and select as “optimal” model the one whose parameters yield best cross-validation error statistics

Summary

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- Spatial auto-correlation can be quantified by looking at attribute dissimilarity as a function of separation distance
- The semivariogram cloud is “too cloudy” for detecting meaningful patterns
- The semivariogram plot is constructed by averaging squared semidifferences within distance bins to “smooth” out the variability in the semivariogram cloud
NOTE: *Watch out for trends (first-order effects) in the data; a sample semivariogram quantifies second-order effects and might be contaminated by variations due to trends/drifts*
- A quantitative way to encapsulate a sample semivariogram is through a parametric semivariogram model
- Fitting procedures exist for estimating the parameters of semivariogram models, i.e., for fitting model semivariograms to sample semivariograms
- The final semivariogram model can be used for simulation (pattern generation) and geostatistical interpolation

NOTE: *A semivariogram model is a spatial process model, whose parameters are inferred from the sample data through the sample semivariogram*

Introduction

Data: set of n attribute measurements $\{z(\mathbf{s}_i), i = 1, \dots, n\}$, available at n sample locations $\{\mathbf{s}_i, i = 1, \dots, n\}$

Objectives: (i) predict or interpolate unknown attribute value $z(\mathbf{s}_p)$ at location \mathbf{s}_p from the n sample data, and (ii) assess reliability of predicted value

Slide 1 Geostatistical spatial interpolation:

- predicted attribute value = weighted linear combination of sample data values + attribute mean, if known (non-linear methods also exist)
- a semivariogram model is used to determine the weights, which account for:
 - spatial auto-correlation between sample data and unknown value
 - spatial auto-correlation between sample data themselves (data redundancy)
- in addition, and contrary to most interpolation algorithms, geostatistics offers a measure of reliability (prediction error variance) regarding the attribute prediction

Simple Kriging (SK)

SK prediction: $\hat{z}(\mathbf{s}_p) = m + \sum_{i=1}^n w_p(\mathbf{s}_i)[z(\mathbf{s}_i) - m] = \mathbf{w}_p^T \mathbf{r}$

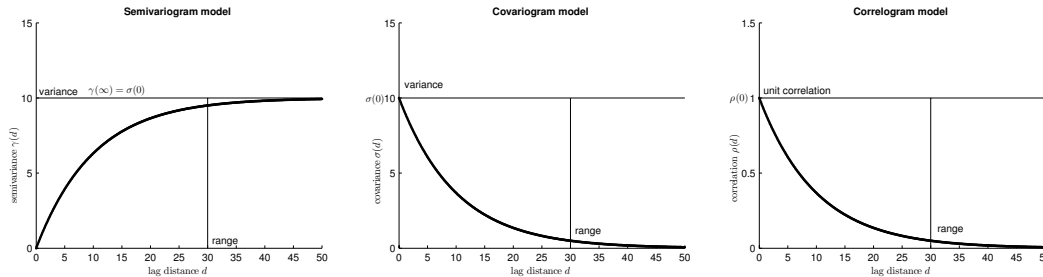
- $\mathbf{w}_p = [w_p(\mathbf{s}_i), i = 1, \dots, n]^T$: $(n \times 1)$ vector of SK-weights assigned to n sample data for prediction at location \mathbf{s}_p ; superscript T denotes transposition
- $\mathbf{r} = [z(\mathbf{s}_i) - m, i = 1, \dots, n]^T$: $(n \times 1)$ vector of residual data from known mean m

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$$\hat{z}(\mathbf{s}_p) = m + \underbrace{\left[w_p(\mathbf{s}_1) \cdots w_p(\mathbf{s}_i) \cdots w_p(\mathbf{s}_n) \right]}_{\mathbf{w}_p^T} \underbrace{\begin{bmatrix} z(\mathbf{s}_1) - m \\ \vdots \\ z(\mathbf{s}_i) - m \\ \vdots \\ z(\mathbf{s}_n) - m \end{bmatrix}}_{\mathbf{r}}$$

use semivariogram model to determine weights at each prediction location; typically, it is the covariogram model that is used due to computational reasons

Semivariogram / Covariogram / Correlogram Model



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Conversion between models, with $\sigma(0) = \gamma(\infty)$ being the sill of the semivariogram model:

- Semivariogram \rightarrow covariogram: $\sigma(d) = \sigma(0) - \gamma(d)$
- Covariogram \rightarrow correlogram: $\rho(d) = \frac{\sigma(d)}{\sigma(0)}$
- Semivariogram \rightarrow correlogram: $\rho(d) = 1 - \frac{\gamma(d)}{\sigma(0)}$
- Covariogram \rightarrow semivariogram: $\gamma(d) = \sigma(0) - \sigma(d)$

Requisites for Geostatistical Interpolation I

Data-to-data and data-to-unknown distances:

$$\mathbf{D} = \begin{bmatrix} 0 & \cdots & d_{1j} & \cdots & d_{1n} \\ \vdots & \ddots & \vdots & \ddots & \vdots \\ d_{i1} & \cdots & 0 & \cdots & d_{in} \\ \vdots & \ddots & \vdots & \ddots & \vdots \\ d_{n1} & \cdots & d_{nj} & \cdots & 0 \end{bmatrix} \quad \text{and} \quad \mathbf{d}_p = \begin{bmatrix} d_{1p} \\ \vdots \\ d_{ip} \\ \vdots \\ d_{np} \end{bmatrix}$$

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Comments:

- as any other interpolation method, one accounts for the proximity of the n sample locations to the prediction location s_p
Note: Vector \mathbf{d}_p changes from one prediction location s_p to another, hence the subscript p
- unlike other interpolation methods, one also accounts for the proximity between sample locations themselves (sample configuration or data layout)
Note: Matrix \mathbf{D} of sample-to-sample distances is the same for all prediction locations

Requisites for Geostatistical Interpolation II

From distance matrices to model covariance matrices: Take any distance value d_{ij} and d_{ip} , i.e., any entry in \mathbf{D} and \mathbf{d}_p , and transform it, via the covariogram model, to a covariance value $\sigma(d_{ij})$ and $\sigma(d_{ip})$

Data-to-data and data-to-unknown model covariances:

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$$\Sigma = \begin{bmatrix} \sigma(0) & \cdots & \sigma(d_{1j}) & \cdots & \sigma(d_{1n}) \\ \vdots & \ddots & \vdots & \ddots & \vdots \\ \sigma(d_{i1}) & \cdots & \sigma(0) & \cdots & \sigma(d_{in}) \\ \vdots & \ddots & \vdots & \ddots & \vdots \\ \sigma(d_{n1}) & \cdots & \sigma(d_{nj}) & \cdots & \sigma(0) \end{bmatrix} \quad \text{and} \quad \sigma_p = \begin{bmatrix} \sigma(d_{1p}) \\ \vdots \\ \sigma(d_{ip}) \\ \vdots \\ \sigma(d_{np}) \end{bmatrix}$$

- data-to-data covariance matrix Σ : ($n \times n$) matrix with *model* covariance values $\sigma(d_{ij})$ between any two sample locations separated by distance d_{ij}
- data-to-unknown covariance vector σ_p : ($n \times 1$) vector with *model* covariance values $\sigma(d_{ip})$ between the n sample locations and the prediction location \mathbf{s}_p

Note: Vector σ_p changes from one prediction location \mathbf{s}_p to another, hence the subscript p

Requisites for Geostatistical Interpolation III

Data-to-data and data-to-unknown model covariances:

$$\Sigma = \begin{bmatrix} \sigma(0) & \cdots & \sigma(d_{1j}) & \cdots & \sigma(d_{1n}) \\ \vdots & \ddots & \vdots & \ddots & \vdots \\ \sigma(d_{i1}) & \cdots & \sigma(0) & \cdots & \sigma(d_{in}) \\ \vdots & \ddots & \vdots & \ddots & \vdots \\ \sigma(d_{n1}) & \cdots & \sigma(d_{nj}) & \cdots & \sigma(0) \end{bmatrix} \quad \text{and} \quad \sigma_p = \begin{bmatrix} \sigma(d_{1p}) \\ \vdots \\ \sigma(d_{ip}) \\ \vdots \\ \sigma(d_{np}) \end{bmatrix}$$

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Comments:

- data-to-data covariance matrix Σ : encapsulates the redundancy between the sample data; for positive spatial auto-correlation, the more clustered is the sample layout, the more redundant are the sample data (less information content); a clustered sample layout typically translates into larger entries in Σ
- data-to-unknown covariance vector σ_p : encapsulates the statistical proximity (correlation) between the sample data and the unknown attribute value $z(\mathbf{s}_p)$ at the prediction location \mathbf{s}_p ; that correlation is a function of distance between sample and prediction locations, not of the actual (unknown) value $z(\mathbf{s}_p)$; The larger the entries of vector σ_p , the stronger the predictive power of sample data

Simple Kriging (SK) System & Weights

$$\begin{bmatrix} \sigma(0) & \cdots & \sigma(d_{1n}) \\ \vdots & \ddots & \vdots \\ \sigma(d_{n1}) & \cdots & \sigma(0) \end{bmatrix} \begin{bmatrix} w_p(\mathbf{s}_1) \\ \vdots \\ w_p(\mathbf{s}_n) \end{bmatrix} = \begin{bmatrix} \sigma(d_{1p}) \\ \vdots \\ \sigma(d_{np}) \end{bmatrix}$$

$$\Sigma \mathbf{w}_p = \boldsymbol{\sigma}_p$$

Comments:

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- the SK system is a (disguised) version of the normal equations for the case of regression with no intercept term: $\mathbf{X}^T \mathbf{X} \mathbf{b} = \mathbf{X}^T \mathbf{y}$, where \mathbf{X} is the design matrix and \mathbf{y} is the vector of data on the dependent variable; in regression, the data-to-data covariance is estimated as $\mathbf{X}^T \mathbf{X}/n$, and the data-to-unknown covariance as $\mathbf{X}^T \mathbf{y}/n$
- the weights vector \mathbf{w}_p is obtained by solving the SK system, as $\mathbf{w}_p = \Sigma^{-1} \boldsymbol{\sigma}_p$, anew at each prediction location \mathbf{s}_p since the entries of $\boldsymbol{\sigma}_p$ change
- entries of \mathbf{w}_p do not depend on data values or on sill, $\sigma(0)$, of covariogram model:

$$\sigma(0) \begin{bmatrix} \rho(0) & \cdots & \rho(d_{1n}) \\ \vdots & \ddots & \vdots \\ \rho(d_{n1}) & \cdots & 1 \end{bmatrix} \begin{bmatrix} w_p(\mathbf{s}_1) \\ \vdots \\ w_p(\mathbf{s}_n) \end{bmatrix} = \sigma(0) \begin{bmatrix} \rho(d_{1p}) \\ \vdots \\ \rho(d_{np}) \end{bmatrix}$$

Interpreting the Simple Kriging Weights

$$\begin{bmatrix} w_p(\mathbf{s}_1) \\ \vdots \\ w_p(\mathbf{s}_n) \end{bmatrix} = \frac{1}{\sigma(0)} \begin{bmatrix} 1 & \cdots & \rho(d_{1n}) \\ \vdots & \ddots & \vdots \\ \rho(d_{n1}) & \cdots & 1 \end{bmatrix}^{-1} \sigma(0) \begin{bmatrix} \rho(d_{1p}) \\ \vdots \\ \rho(d_{np}) \end{bmatrix} \Rightarrow \mathbf{w}_p = \Sigma^{-1} \boldsymbol{\sigma}_p$$

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- if sample interdistances d_{ij} are larger than correlogram range, then $\rho(d_{ij}) = 0$, and $\Sigma = \sigma(0)\mathbf{I}$, the $(n \times n)$ identity matrix; this entails that $w_p(\mathbf{s}_i) = \rho(d_{ip})$, i.e., weights are equal to correlogram values
- but in general, $\Sigma \neq \sigma(0)\mathbf{I}$, i.e., sample interdistances are within correlation range, in which case Σ^{-1} modulates $\boldsymbol{\sigma}_p$: influence of samples in clusters is downplayed
- the closer the sample data to the prediction location, and the more spread out the data over the study region, the better the SK prediction is expected to be
- for sample data far away (beyond correlation range) from the prediction location \mathbf{s}_p , $\rho(d_{ip}) = 0$ and $w_p(\mathbf{s}_i) = 0$: all weights are equal to 0
- for prediction at a sample location $\mathbf{s}_p \equiv \mathbf{s}_i$, data-to-unknown covariance vector $\boldsymbol{\sigma}_p = \boldsymbol{\sigma}_i$ is same as i -th column of Σ ; this yields $w_p(\mathbf{s}_i) = 1$ if $\mathbf{s}_i = \mathbf{s}_p$, 0 otherwise: only sample co-located with prediction location receives non-zero (= 1) weight

Simple Kriging Prediction and Error Variance

Once the SK weights are computed as $\mathbf{w}_p = \Sigma^{-1} \boldsymbol{\sigma}_p$, they are substituted in the following equations to compute the SK prediction $\hat{z}(\mathbf{s}_p)$ and associated error variance $\hat{\sigma}(\mathbf{s}_p)$

SK prediction does not depend on sill $\sigma(0)$ of covariogram model:

$$\hat{z}(\mathbf{s}_p) = m + \mathbf{w}_p^T \mathbf{r} = m + [w_p(\mathbf{s}_1) \cdots w_p(\mathbf{s}_n)] \begin{bmatrix} z(\mathbf{s}_1) - m \\ \vdots \\ z(\mathbf{s}_n) - m \end{bmatrix} = m + \sum_{i=1}^n w_p(\mathbf{s}_i) [z(\mathbf{s}_i) - m]$$

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SK prediction error variance does depend on covariogram model sill $\sigma(0)$:

$$\hat{\sigma}(\mathbf{s}_p) = \sigma(0) - \mathbf{w}_p^T \boldsymbol{\sigma}_p = \sigma(0) - [w_p(\mathbf{s}_1) \cdots w_p(\mathbf{s}_n)] \begin{bmatrix} \sigma(d_{1p}) \\ \vdots \\ \sigma(d_{np}) \end{bmatrix} = \sigma(0) - \sum_{i=1}^n w_p(\mathbf{s}_i) \sigma(d_{ip})$$

which can also be written as: $\hat{\sigma}(\mathbf{s}_p) = \sigma(0) [1 - \sum_{i=1}^n w_p(\mathbf{s}_i) \rho(d_{ip})]$

Interpreting the SK Prediction and Error Variance

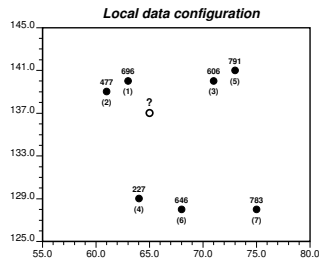
$$\hat{z}(\mathbf{s}_p) = m + \sum_{i=1}^n w_p(\mathbf{s}_i) [z(\mathbf{s}_i) - m] \quad \hat{\sigma}(\mathbf{s}_p) = \sigma(0) - \sum_{i=1}^n w_p(\mathbf{s}_i) \sigma(d_{ip})$$

Comments:

- for sample data far away (beyond correlation range) from the prediction location \mathbf{s}_p , $w_p(\mathbf{s}_i) = 0, \forall i$: *all weighs are equal to 0*. In this case, the SK prediction equals the known mean m and the SK error variance equals the known covariogram sill: $\hat{z}(\mathbf{s}_p) = m$ and $\hat{\sigma}(\mathbf{s}_p) = \sigma(0)$; *away from the sample data, SK yields back the (assumed known) attribute overall mean and variance*
- for prediction at a sample location $\mathbf{s}_p \equiv \mathbf{s}_i$, $w_p(\mathbf{s}_i) = 1$ if $\mathbf{s}_i = \mathbf{s}_p$, 0 otherwise: the SK prediction identifies the known sample datum and the SK error variance is zero: $\hat{z}(\mathbf{s}_i) = z(\mathbf{s}_i)$ and $\hat{\sigma}(\mathbf{s}_i) = 0$; *SK is an exact interpolation algorithm*
- for all other prediction locations, the SK predictions depend on the sample data configuration and their values, while the SK error variances depend only on the sample data configuration; *both SK predictions and error variances depend on the covariogram model $\sigma(d)$ adopted*

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Determining the SK Weights: Step 1



(n x n) matrix of data-to-data inter-distances:

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$$D = \begin{bmatrix} 0.00 & 2.24 & 8.00 & 11.05 & 10.05 & 13.00 & 16.97 \\ 2.24 & 0.00 & 10.05 & 10.44 & 12.17 & 13.04 & 17.80 \\ 8.00 & 10.05 & 0.00 & 13.04 & 2.24 & 12.37 & 12.65 \\ 11.05 & 10.44 & 13.04 & 0.00 & 15.00 & 4.12 & 11.05 \\ 10.05 & 12.17 & 2.24 & 15.00 & 0.00 & 13.93 & 13.15 \\ 13.00 & 13.04 & 12.37 & 4.12 & 13.93 & 0.00 & 7.00 \\ 16.97 & 17.80 & 12.65 & 11.05 & 13.15 & 7.00 & 0.00 \end{bmatrix}$$

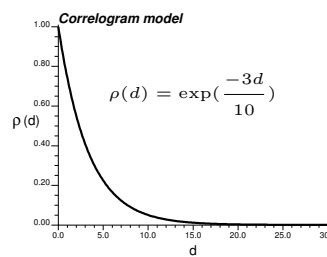
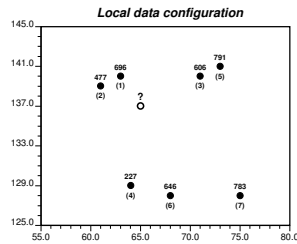
i, j-th element of **D**: $d_{ij} = \|\mathbf{s}_i - \mathbf{s}_j\|$

(n x 1) vector of prediction-to-data-location distances:

$$\mathbf{d}_p = \begin{bmatrix} 3.61 & 4.47 & 6.71 & 8.06 & 8.94 & 9.49 & 13.45 \end{bmatrix}^T$$

i-th element of \mathbf{d}_p : $d_{ip} = \|\mathbf{s}_i - \mathbf{s}_p\|$

Determining the SK Weights: Step 2

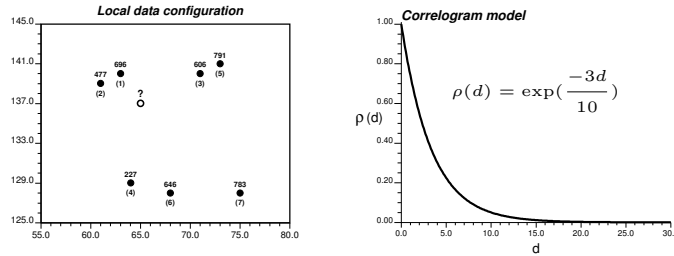


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$$\underbrace{\begin{bmatrix} 3.61 \\ 4.47 \\ 6.71 \\ 8.06 \\ 8.94 \\ 9.49 \\ 13.45 \end{bmatrix}}_{\mathbf{d}_p} \rightarrow 1 \underbrace{\begin{bmatrix} \exp(-3 \times 3.61/10) \\ \exp(-3 \times 4.47/10) \\ \exp(-3 \times 6.71/10) \\ \exp(-3 \times 8.06/10) \\ \exp(-3 \times 8.94/10) \\ \exp(-3 \times 9.49/10) \\ \exp(-3 \times 13.45/10) \end{bmatrix}}_{\sigma_p = \text{sill} \exp(-3\mathbf{d}_p/\text{range})} = \begin{bmatrix} 0.34 \\ 0.26 \\ 0.13 \\ 0.09 \\ 0.07 \\ 0.06 \\ 0.02 \end{bmatrix}$$

These would be the weights if one ignored auto-correlation between sample data

Determining the SK Weights: Step 3



SK system:

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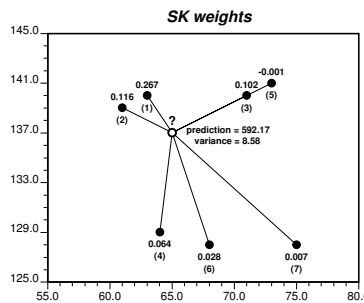
$$\underbrace{\begin{bmatrix} 1.00 & 0.51 & 0.09 & 0.04 & 0.05 & 0.02 & 0.01 \\ 0.51 & 1.00 & 0.05 & 0.04 & 0.03 & 0.02 & 0.00 \\ 0.09 & 0.05 & 1.00 & 0.02 & 0.51 & 0.02 & 0.02 \\ 0.04 & 0.04 & 0.02 & 1.00 & 0.01 & 0.29 & 0.04 \\ 0.05 & 0.03 & 0.51 & 0.01 & 1.00 & 0.02 & 0.02 \\ 0.02 & 0.02 & 0.02 & 0.29 & 0.02 & 1.00 & 0.12 \\ 0.01 & 0.00 & 0.02 & 0.04 & 0.02 & 0.12 & 1.00 \end{bmatrix}}_{\Sigma} \underbrace{\begin{bmatrix} w_p(s_1) \\ w_p(s_2) \\ w_p(s_3) \\ w_p(s_4) \\ w_p(s_5) \\ w_p(s_6) \\ w_p(s_7) \end{bmatrix}}_{\mathbf{w}_p} = \underbrace{\begin{bmatrix} 0.34 \\ 0.26 \\ 0.13 \\ 0.09 \\ 0.07 \\ 0.06 \\ 0.02 \end{bmatrix}}_{\boldsymbol{\sigma}_p}$$

i, j -th element of matrix Σ : $\sigma_{ij} = 1 \times \exp(-3 \times d_{ij}/10)$

Determining the SK Weights: Step 4

$$\underbrace{\begin{bmatrix} w_p(s_1) \\ w_p(s_2) \\ w_p(s_3) \\ w_p(s_4) \\ w_p(s_5) \\ w_p(s_6) \\ w_p(s_7) \end{bmatrix}}_{\mathbf{w}_p} = \underbrace{\begin{bmatrix} 1.36 & -0.69 & -0.09 & -0.02 & 0.00 & 0.00 & -0.01 \\ -0.69 & 1.35 & 0.00 & -0.02 & 0.00 & -0.01 & 0.01 \\ -0.09 & 0.00 & 1.36 & -0.01 & -0.69 & -0.01 & -0.01 \\ -0.02 & -0.02 & -0.01 & 1.09 & 0.00 & -0.32 & -0.01 \\ 0.00 & 0.00 & -0.69 & 0.00 & 1.35 & -0.01 & -0.01 \\ 0.00 & -0.01 & -0.01 & -0.32 & -0.01 & 1.11 & -0.12 \\ -0.01 & 0.01 & -0.01 & -0.01 & -0.01 & -0.12 & 1.02 \end{bmatrix}}_{\Sigma^{-1}} \underbrace{\begin{bmatrix} 0.34 \\ 0.26 \\ 0.13 \\ 0.09 \\ 0.07 \\ 0.06 \\ 0.02 \end{bmatrix}}_{\boldsymbol{\sigma}_p}$$

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original weights vector ($\mathbf{w}_p = \boldsymbol{\sigma}_p$) modified by Σ^{-1} to account for sample redundancy;
 e.g., $w_p(s_1) = 0.27$ instead of $\rho(d_{1p}) = 0.34$