Variogrammes

Ça se discute

A. Boyé

$$\hat{P} = \frac{x_{1} + x_{2}}{n_{1} + n_{1}} \quad \hat{X} = \frac{x_{1} + x_{2} + x_{3} + ... + x_{N}}{\sigma} \quad A = \hat{y} - b \hat{x}$$

$$H_{o} : P = P_{o} \quad S_{\hat{x}} \rightarrow \sqrt{n} \quad N$$

$$SE = \sqrt{\frac{\hat{P}(1 - \hat{p})}{n}} \quad Z = \sqrt{\frac{\hat{P} - P_{o}}{R(1 - P_{o})}} \quad \begin{pmatrix} n \\ k \end{pmatrix} = \frac{n!}{n!(n-k)!}$$

$$SE = \sqrt{\frac{\hat{P}(1 - \hat{p})}{n}} \quad Z = \sqrt{\frac{\hat{P}_{c}(1 - \hat{p}_{o})}{R(1 - \hat{p}_{o})}}$$

$$ME \cdot z^{*} \frac{\sigma}{\sqrt{n}}$$

$$\hat{Y} = A + b \times \mu = \frac{nP}{z} = \frac{x - \mu}{\sigma}$$

Spatial (or temporal) correlation may be described by ¹

Correlograms

A correlogram is a graph in which *spatial correlation* values are plotted, on the ordinate, against *distance classes* among sites on the abscissa.

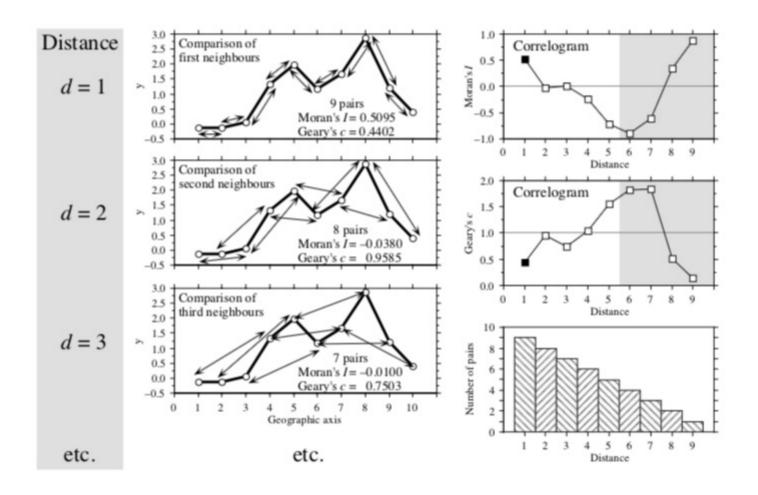
Variograms (semi-variograms)

A variogram is a graph in which *semi-variance* is plotted, on the ordinate, against *distance classes* among sites on the abscissa

Periodograms (structure assumed to consist of a combination of sine waves)

Fit sines and cosines of various periods, one period at a time, and determine the proportion of the series' variance (\mathbb{R}^2) explained by each period. The abscissa is either a period or its inverse, a frequency; the ordinate is the proportion of variance explained.

Principle



Correlograms

Moran's I:
$$I(d) = \frac{\frac{1}{W} \sum_{h=1}^{n} \sum_{i=1}^{n} w_{hi} (y_h - \bar{y}) (y_i - \bar{y})}{\frac{1}{n} \sum_{i=1}^{n} (y_i - \bar{y})^2} \quad \text{for } h \neq i$$
 (13.1)

Geary's c:
$$c(d) = \frac{\frac{1}{2W} \sum_{h=1}^{n} \sum_{i=1}^{n} w_{hi} (y_h - y_i)^2}{\frac{1}{(n-1)} \sum_{i=1}^{n} (y_i - \bar{y})^2} \quad \text{for } h \neq i$$
 (13.2)

Moran's I

Moran's I related to the *Pearson*correlation coefficient. *Usually* takes

values between -1 and +1

Numerator: covariance;

Denominator: maximum-likelihood

estimator of the variance

• Geary's c

Geary's c coefficient is a *distance-type* function : varies from 0 to some unspecified value larger than 1.

Numerator: sums the squared

differences

Varies as the reverse of a Moran's I correlogram

Correlograms

Moran's I:
$$I(d) = \frac{\frac{1}{W} \sum_{h=1}^{n} \sum_{i=1}^{n} w_{hi} (y_h - \bar{y}) (y_i - \bar{y})}{\frac{1}{n} \sum_{i=1}^{n} (y_i - \bar{y})^2} \quad \text{for } h \neq i$$
 (13.1)

Geary's c:
$$c(d) = \frac{\frac{1}{2W} \sum_{h=1}^{n} \sum_{i=1}^{n} w_{hi} (y_h - y_i)^2}{\frac{1}{(n-1)} \sum_{i=1}^{n} (y_i - \bar{y})^2} \quad \text{for } h \neq i$$
 (13.2)

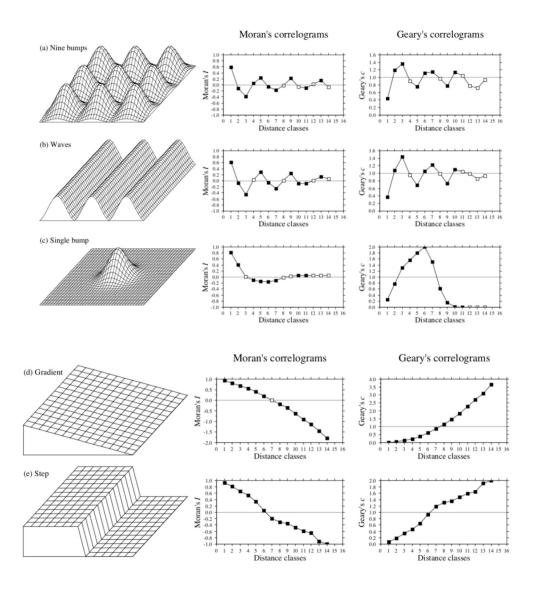
Positive spatial correlation : positive values of I.

Negative correlation : negative values.

 \Rightarrow Strong spatial correlation produces high values of I and low values of c

Sensitive to extreme values and asymmetry in distribution: normalize data

Absence of auto-correlation : Moran's I = 0 & Geary's c = 1



Correlograms

Moran's I:
$$I(d) = \frac{\frac{1}{W} \sum_{h=1}^{n} \sum_{i=1}^{n} w_{hi} (y_h - \bar{y}) (y_i - \bar{y})}{\frac{1}{n} \sum_{i=1}^{n} (y_i - \bar{y})^2} \quad \text{for } h \neq i$$
 (13.1)

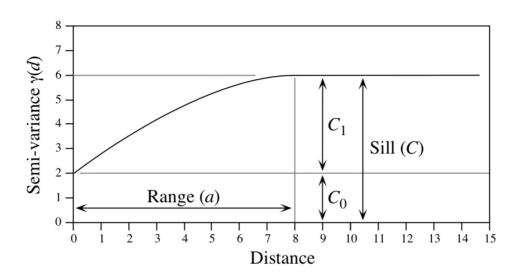
Geary's c:
$$c(d) = \frac{\frac{1}{2W} \sum_{h=1}^{n} \sum_{i=1}^{n} w_{hi} (y_h - y_i)^2}{\frac{1}{(n-1)} \sum_{i=1}^{n} (y_i - \bar{y})^2} \quad \text{for } h \neq i$$
 (13.2)

Semi-variograms

$$\gamma(d) = \frac{1}{2W(d)} \sum_{h=1}^{n} \sum_{i=1}^{n} w_{hi} (y_h - y_i)^2 \quad \text{for } h \neq i$$
 (13.9)

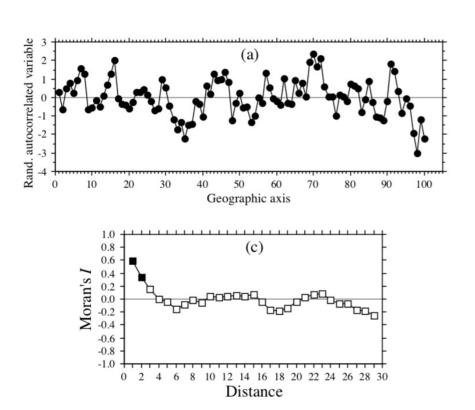
⇒ Numerator of Geary's c

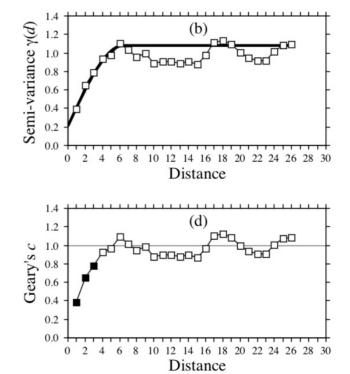
Semi-variograms



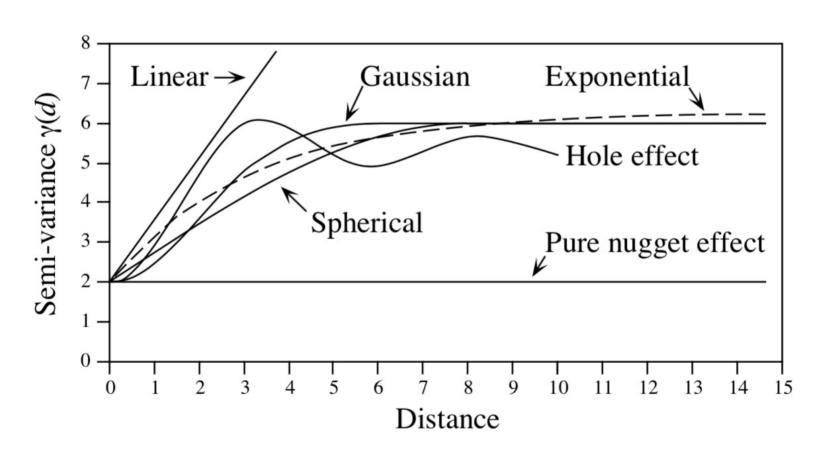
- *sill*: variance of the variable
- range: beyond that distance, the sampling units are not spatially correlated
- C_0 (nugget effect): local variation occurring at scales finer than the sampling interval
 - ex: sampling error, fine-scale spatial variability, and measurement error
- C_1 : spatially structured component

Example





Fit models to the sample variogram (kriging)



Computation in R

Autocorrelation

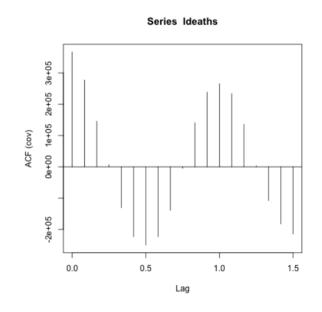
Cross-correlation/cross-variogram

```
ccf(x, y, lag.max = NULL, type = c("correlation", "covariance"),
  plot = TRUE, na.action = na.fail, ...)
```

Monthly Deaths from Lung Diseases in the UK, 1974–1979

acf(ldeaths,
type = "correlation")

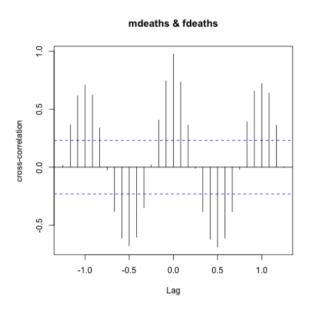
acf(ldeaths,
type = "covariance")

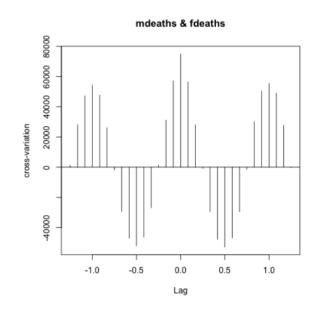


Covariation male/female

```
ccf(mdeaths, fdeaths,
  type = "correlation",
ylab="cross-correlation")
```

```
ccf(mdeaths, fdeaths,
    type = "covariance",
    ylab="cross-variation")
```





Another example from marine systems ²

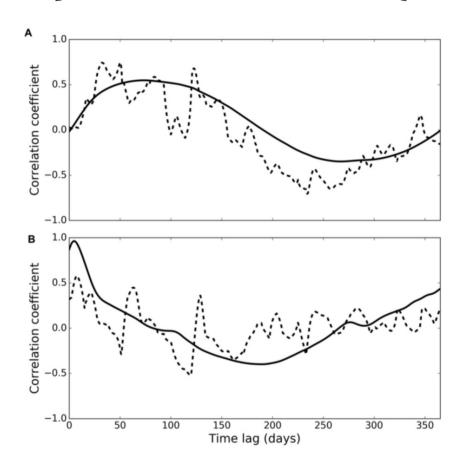


FIGURE 5 | Cross-correlation of phytoplankton with observed (dashed line) and modelled (solid line) deposit feeders (A) and suspension feeders (B).

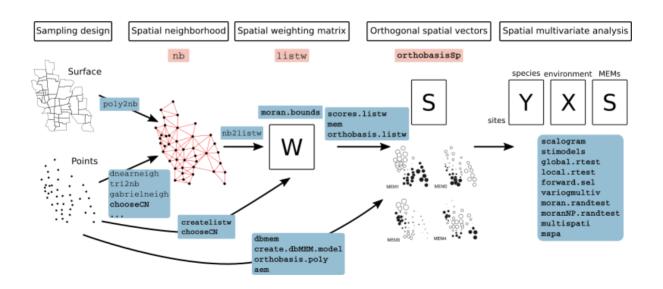
[2] Lessin, G., Bruggeman, J., McNeill, C. L., and Widdicombe, S. (2019). Time scales of benthic macrofaunal response to pelagic production differ between major feeding groups. *Frontiers in Marine Science*, 6:15.

Other approaches (MEM)

Principle

MEM³ (Moran's eigenvector maps): spatial variables representing structures of all relevent scales

Geographic distances among sites or spatial weighting matrix ⇒ spatial explanatory variables



[3] Dray, S., Legendre, P., and Peres-Neto, P. R. (2006). Spatial modelling: a comprehensive framework for principal coordinate analysis of neighbour matrices (PCNM). Ecological Modelling, 196(3):483 – 493.

Examples ⁴

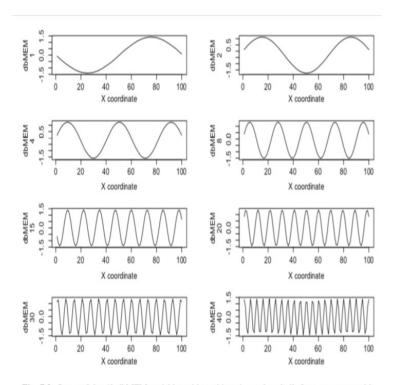


Fig. 7.3 Some of the 49 dbMEM variables with positive eigenvalues built from a transect with 100 equispaced points

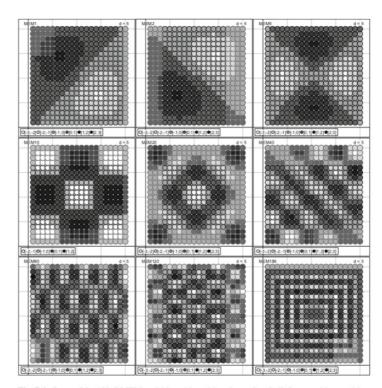


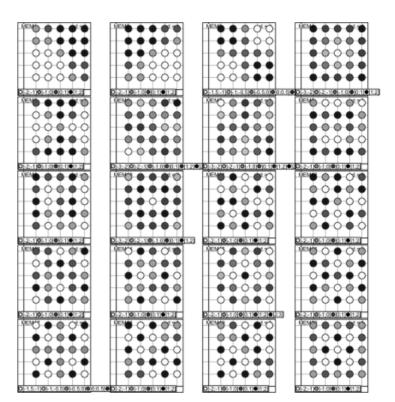
Fig. 7.4 Some of the 189 dbMEM variables with positive eigenvalues built from a grid comprising 20 by 20 equispaced points

[4] Borcard, D., Gillet, F., & Legendre, P. (2018). Numerical ecology with R. 2nd edition. Springer.

MEM and dbMEM can be computed using adespatial

```
library(adespatial)
library(adegraphics)
# Generate grid point coordinates
xvqrid2 <- expand.grid(1:5, 1:5)</pre>
# Creation of the dbMEM eigenfunctions v
xvqrid2.dbmem.tmp <-</pre>
  dbmem(xvgrid2,MEM.autocor="non-null")
xygrid2.dbmem <-
  as.data.frame(xygrid2.dbmem.tmp)
# Plot some dbMEM variables using s.vali
somedbmem2 <- c(1:20)
s.value(xvgrid2.
        xvgrid2.dbmem[.somedbmem2].
  method = "color",
  symbol = "circle".
  ppoints.cex = 0.5
```

Information: Square regular grid; multi



Analyse de codépendance

Ecology, 91(10), 2010, pp. 2952–2964 © 2010 by the Ecological Society of America

Multiscale codependence analysis: an integrated approach to analyze relationships across scales

Guillaume Guénard, 1,3 Pierre Legendre, 1 Daniel Boisclair, 1 and Martin Bilodeau 2

¹Département de Sciences Biologiques, Université de Montréal, C.P. 6128, Succursale Centre-ville, Montréal, Québec H3C3J7 Canada ²Département de Mathématiques et de Statistique, Université de Montréal, C.P. 6128, Succursale Centre-ville, Montréal, Québec H3C3J7 Canada

Analyse de codépendance

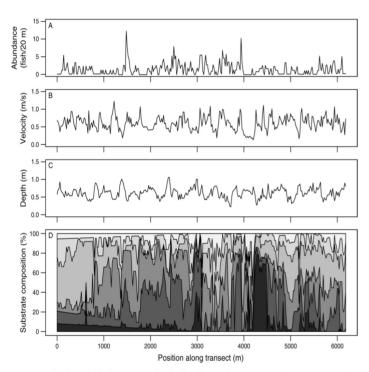


Fig. 4. (A) Abundance of Atlantic salmon parr after the effect of among-day temperature differences and linear trend were removed by regression. Also shown are (B) flow velocity, (C) channel depth, and (D) substrate composition after removal of linear trends. The six levels of shading represent the contributions of the six grain size classes of the river bed, from fine substrate (light) to metric boulder (dark).

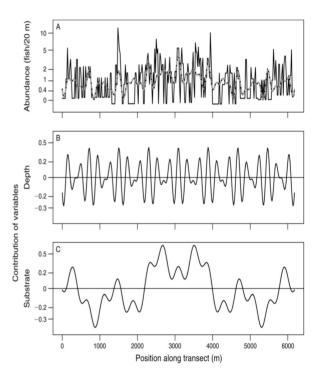


Fig. 5. (A) Observed (solid) and fitted (dotted) parr abundances, on a $\log_c(y+1)$ scale, along the river transect; (B) the fraction of parr abundance explained by channel depth; and (C) substrate composition obtained from Eq. 9. The model explains 18.9% of the $\log(x+1)$ -transformed variation in parr abundance.