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Modelling Pre- and Protohistorical Iconographic Compositions. The R package decorr

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Abstract

By definition, Prehistorical societies are characterised by the absence of a writing system. Writing is one of the most rational semiographical system with a clear distinction between signified and signifier – specially in alphabetic and binary writings – and the development of the signified on a horizontal, vertical or boustrophedon axis. Prehistorical times cover more than 99% of the human living. Even if it is being discussed, first symbolic manifestations start around 200,000 BC (d'Errico and Nowell 2000). The duration from first symbolic expressions to start of writing represents 97% of the human living. In illiterate societies, testimonies of symbolic systems mostly come from iconography (ceramic decorations, rock-art, statuary, etc.) and signs are displayed mostlty a discontinuous figures which can have different relationships one with another. An iconographical composition can be "read" as a spatial distribution of features having intrinsic values possibily having meaningful relationships one with another depending on their pairwise spatial proximities.

To understand meaningful associations of signs, geometric tools, graph analysis and statistical analysis offer great tools to recognize iconographical patterns and to infer collective conventions. We present the **decorr** R package which ground concepts, methods and tools to analyse ancient iconographical systems.

Keywords: Iconography, Prehistory, Graph Theory, Graph Drawing, Spatial Analysis, R.

1. Introduction: Count data regression in R

The introduction is in principle "as usual". However, it should usually embed both the implemented *methods* and the *software* into the respective relevant literature. For the latter both competing and complementary software should be discussed (within the same software environment and beyond), bringing out relative (dis)advantages. All software mentioned should be properly \cite{}d. (See also Appendix B for more details on BibTeX.)

For writing about software JSS requires authors to use the markup \proglang{} (programming languages and large programmable systems), \pkg{} (software packages), \code{} (functions, commands, arguments, etc.). If there is such markup in (sub)section titles (as above), a plain text version has to be provided in the LATEX command as well. Below we also illustrate how abbrevations should be introduced and citation commands can be employed. See the LATEX code for more details.

For decades, study of ancient iconography was linked to history of religion because closely linked to believes, etc. Unlike other aspects of the material culture – a flint blade for cutting, a pottery for containing, a house for living –, the function of an iconographic composition cannot be drawn directly from itself. Since the *New Archaeology* developpement during the 60's (Clarke 2014), symbolic expressions start to be studied with the same formal methods (statistics, seriations, distribution maps, etc.) as any another aspect of social organisation: settlement patterns, tools *chaine opÃlratoire*, susbsitence strategies, etc. (Renfrew and Bahn 1991), (Leroi-Gourhan 1992). Study of ancient iconography had undergone significative improvements at site scale – with GIS, database, paleoclimatic restitutions, etc. – and at the sign scale with the development of archaeological sciences – radiocarbon dating, use-wear analysis, elemental analysis, etc. – but these improvement do not necessarly help to understand the semantic content of the iconography.

Formal methods to study ancient iconography semantcis, has been mostly been grounded (explicitly or not) on the prime principle of Saussurian linguistic: the 'linearity of the signifier' (De Saussure 1989). This principle states that the meaning of a linguistic or writing pattern is linear and directed.

Applying this principle to any other graphical content than a writing, allows to considered as the organisation of graphical as a relationship of figures grouping graphical units, patterns grouping figures, motives grouping patterns, etc., until the entire decorated support is described and can be compared to another decoration (XXX). But during this *decomposition* process, the groups and relationships are often defined empirically, their level of significance are often implicit and the iconographical and spatial proximities between graphical units and categories of graphical units are not quantified. Furthermore, due to the inherent variability of iconography, most of the studies developp proper descriptive vocabularies, singular relationships of categories, idosyncratic methods in a site-dependend or period-dependend scales. This limits drastically the possibility to conduct cross-cultural comparisons and to draw a synthesis of humankind's symbolism at a large scale and over the long-term.

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In this article we present the R package **decorr**. Its purpose is to formalise a method based on geometric graphs to analyse any graphical content

magick for image manipulation (Ooms 2018) igraph (Csardi and Nepusz 2006) for graph and network analysis grDevices for colors and font plotting (R Core Team 2019) graphics for graphics (R Core Team 2019)

2. Concepts

As any formal system, iconography can be seen as spatial features related one with the other depending on rules of proximities. Just like in R, the features a, r and t concatenated in this order mean "art", and not "rat", in illetrate societies, spatial relations between graphical features are not necessarly linear and directed but multi-directional and undirected: the direction of the interactions of pairwise graphical units can be in any order (Fig. 1)

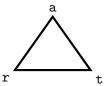


Figure 1: Potential spatial relations between a, r and t graphical units. Edges of the triangle represent the proximity links between the a, r and t graphical features

To resolve this difficulty, we consider the decoration as the tessellation of the support. The undecorated parts of the support, or background, is considerated homogeneous. Exist a link between a couple of graphical units when theses graphical units share a border. A planar graph is constructed from grahical units (nodes) and their proximity links (edges). This model is a Voronoi diagram of the support where the Voronoi seeds are the graphical units. Its geographical equivalent is a Thiessen polygon. This model has a minimal of a priori definitions. Those only concern the graphical units (shape, color, orientation, size, etc.). Nodes and edges are created on a GIS interface where the decoration figure is the basemap and gives the total extent (xmin, xmax,) of the analysis. This decoation figure is open in a GIS without any projection where the decoration. Since, les x and y coordinates of the nodes are relative to the decoration and measured in pixels.

Between two graphical units the links are represented with a plain line. But sometimes a graphical unit can be divided into a main unit (eg, a man) and attribute units (eg, a helmet, a sword). So, the relations between the main unit and the attribute units are directed and displayed with a dashed line.

The spatial levels of the graphical units can be retrieve by a planar graph (Graph Theory) and a spatial (GIS) analysis.

2.1. Graph Theory

Graph theory offers a conceptual framework and indices (global at the entire graph scale, local at the vertex scale) to deal with notions of networks, relationships and neighbourhoods. A system can be represented by vertices connected (or not) to each other with edges. Let us take the example of the word "art" contains three vertices (a, r, t) and two edges (one between a and r, the other between r and t).

The r graphical unit receives two edges, so its centrality degree is 2, r is also central in the graph so its betweeness degree is 1.

2.2. GIS interface

3. The R package decorr

The **decorr** can be downloaded on GitHub.

3.1. Function index

list_nds_compar and list_eds_compar allow to compare respectively the common nodes and the common edges between two graphs

3.2. Function listdec()

This function store graphs in a list.

3.3. Function xxx

xxx

The basic Poisson regression model for count data is a special case of the GLM framework? It describes the dependence of a count response variable y_i (i = 1, ..., n) by assuming a Poisson distribution $y_i \sim \text{Pois}(\mu_i)$. The dependence of the conditional mean $\mathsf{E}[y_i \mid x_i] = \mu_i$ on the regressors x_i is then specified via a log link and a linear predictor

$$\log(\mu_i) = x_i^{\top} \beta, \tag{1}$$

where the regression coefficients β are estimated by maximum likelihood (ML) using the iterative weighted least squares (IWLS) algorithm.

Note that around the {equation} above there should be no spaces (avoided in the LATEX code by % lines) so that "normal" spacing is used and not a new paragraph started.

R provides a very flexible implementation of the general GLM framework in the function glm() (?) in the stats package. Its most important arguments are

```
glm(formula, data, subset, na.action, weights, offset,
  family = gaussian, start = NULL, control = glm.control(...),
  model = TRUE, y = TRUE, x = FALSE, ...)
```

where formula plus data is the now standard way of specifying regression relationships in R/S introduced in ?. The remaining arguments in the first line (subset, na.action, weights, and offset) are also standard for setting up formula-based regression models in R/S. The arguments in the second line control aspects specific to GLMs while the arguments in the last line specify which components are returned in the fitted model object (of class 'glm' which inherits from 'lm'). For further arguments to glm() (including alternative specifications of starting values) see ?glm. For estimating a Poisson model family = poisson has to be specified.

As the synopsis above is a code listing that is not meant to be executed, one can use either the dedicated {Code} environment or a simple {verbatim} environment for this. Again, spaces before and after should be avoided.

Type	Distribution	Method	Description
GLM	Poisson	ML	Poisson regression: classical GLM, esti-
			mated by maximum likelihood (ML)
		Quasi	"Quasi-Poisson regression": same mean
			function, estimated by quasi-ML (QML)
			or equivalently generalized estimating equa-
			tions (GEE), inference adjustment via esti-
			mated dispersion parameter
		Adjusted	"Adjusted Poisson regression": same mean
			function, estimated by QML/GEE, inference
			adjustment via sandwich covariances
	NB	ML	NB regression: extended GLM, estimated by
			ML including additional shape parameter
Zero-augmented	Poisson	ML	Zero-inflated Poisson (ZIP), hurdle Poisson
	NB	ML	Zero-inflated NB (ZINB), hurdle NB

Table 1: Overview of various count regression models. The table is usually placed at the top of the page ([t!]), centered (centering), has a caption below the table, column headers and captions are in sentence style, and if possible vertical lines should be avoided.

Finally, there might be a reference to a {table} such as Table 1. Usually, these are placed at the top of the page ([t!]), centered (\centering), with a caption below the table, column headers and captions in sentence style, and if possible avoiding vertical lines.

4. Illustrations

For a simple illustration of basic Poisson and NB count regression the quine data from the MASS package is used. This provides the number of Days that children were absent from school in Australia in a particular year, along with several covariates that can be employed as regressors. The data can be loaded by

R> data("quine", package = "MASS")

and a basic frequency distribution of the response variable is displayed in Figure 2.

For code input and output, the style files provide dedicated environments. Either the "agnostic" {CodeInput} and {CodeOutput} can be used or, equivalently, the environments {Sinput} and {Soutput} as produced by Sweave() or knitr when using the render_sweave() hook. Please make sure that all code is properly spaced, e.g., using y = a + b * x and not y=a+b*x. Moreover, code input should use "the usual" command prompt in the respective software system. For R code, the prompt "R> " should be used with "+ " as the continuation prompt. Generally, comments within the code chunks should be avoided – and made in the regular LATEX text instead. Finally, empty lines before and after code input/output should be avoided (see above).

As a first model for the quine data, we fit the basic Poisson regression model. (Note that JSS prefers when the second line of code is indented by two spaces.)

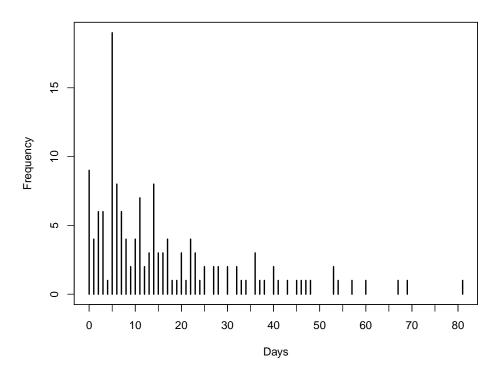


Figure 2: Frequency distribution for number of days absent from school.

To account for potential overdispersion we also consider a negative binomial GLM.

```
R> library("MASS")
R> m_nbin <- glm.nb(Days ~ (Eth + Sex + Age + Lrn)^2, data = quine)</pre>
```

In a comparison with the BIC the latter model is clearly preferred.

```
df BIC
m_pois 18 2046.851
m_nbin 19 1157.235
```

R> BIC(m_pois, m_nbin)

Hence, the full summary of that model is shown below.

```
R> summary(m_nbin)
```

Call:

```
glm.nb(formula = Days ~ (Eth + Sex + Age + Lrn)^2, data = quine,
  init.theta = 1.60364105, link = log)
```

Deviance Residuals:

Max

3Q

Min

1Q

Median

Theta: 1.604 Std. Err.: 0.214

2 x log-likelihood: -1062.546

```
-3.0857 -0.8306 -0.2620
                           0.4282
                                   2.0898
Coefficients: (1 not defined because of singularities)
           Estimate Std. Error z value Pr(>|z|)
(Intercept) 3.00155
                       0.33709
                                8.904 < 2e-16 ***
EthN
           -0.24591
                       0.39135 -0.628 0.52977
SexM
           -0.77181
                       0.38021
                               -2.030 0.04236 *
AgeF1
           -0.02546
                       0.41615
                               -0.061
                                       0.95121
AgeF2
           -0.54884
                       0.54393
                               -1.009 0.31296
                       0.40558 -0.635 0.52574
AgeF3
           -0.25735
LrnSL
            0.38919
                    0.48421
                               0.804 0.42153
EthN:SexM
                       0.29430
                               1.231 0.21818
            0.36240
EthN:AgeF1 -0.70000
                    0.43646 -1.604 0.10876
                    0.42962 -2.870
                                       0.00411 **
EthN:AgeF2 -1.23283
EthN:AgeF3
            0.04721
                       0.44883
                                0.105
                                       0.91622
EthN:LrnSL
            0.06847
                       0.34040
                               0.201
                                       0.84059
SexM:AgeF1
            0.02257
                       0.47360
                               0.048 0.96198
SexM:AgeF2
                       0.51325
                               3.026 0.00247 **
            1.55330
SexM: AgeF3
            1.25227
                       0.45539
                                2.750 0.00596 **
                    0.40805
SexM:LrnSL
            0.07187
                                0.176 0.86019
AgeF1:LrnSL -0.43101
                       0.47948
                               -0.899 0.36870
AgeF2:LrnSL 0.52074
                       0.48567
                                1.072 0.28363
AgeF3:LrnSL
                 NA
                           NA
                                   NA
                                            NA
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
(Dispersion parameter for Negative Binomial(1.6036) family taken to be 1)
   Null deviance: 235.23 on 145 degrees of freedom
Residual deviance: 167.53 on 128 degrees of freedom
AIC: 1100.5
Number of Fisher Scoring iterations: 1
```

5. Summary and discussion

As usual ...

Computational details

If necessary or useful, information about certain computational details such as version numbers, operating systems, or compilers could be included in an unnumbered section. Also, auxiliary packages (say, for visualizations, maps, tables, ...) that are not cited in the main text can be credited here.

The results in this paper were obtained using R 3.4.1 with the MASS 7.3.47 package. R itself and all packages used are available from the Comprehensive R Archive Network (CRAN) at https://CRAN.R-project.org/.

Acknowledgments

All acknowledgments (note the AE spelling) should be collected in this unnumbered section before the references. It may contain the usual information about funding and feedback from colleagues/reviewers/etc. Furthermore, information such as relative contributions of the authors may be added here (if any).

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- Graphics formatting.
- Naming conventions.
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- Trouble shooting.
- Many other potentially helpful details...

B. Using BibTeX

References need to be provided in a BIBTeX file (.bib). All references should be made with \cite, \citet, \citep, \citealp etc. (and never hard-coded). This commands yield different formats of author-year citations and allow to include additional details (e.g., pages, chapters, ...) in brackets. In case you are not familiar with these commands see the JSS style FAQ for details.

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- JSS-specific markup (\proglang, \pkg, \code) should be used in the references.
- Titles should be in title case.
- Journal titles should not be abbreviated and in title case.
- DOIs should be included where available.
- Software should be properly cited as well. For R packages citation("pkgname") typically provides a good starting point.

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