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Spatial distribution of earthworms in acid-soil savannas of the eastern plains of Colombia

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Abstract

The horizontal spatial distribution of earthworms in a native savanna and in a 17 year introduced pasture from the area of Carimagua (Colombia) was assessed by means of aggregation indices and geostatistical analysis. Morisita's index (I_d) and Taylor's index (b) were calculated from field data obtained through 1 m² quadrats during a sampling period of 17 months and both semivariograms and contour maps were obtained at three different dates through a rapid sampling of 64, $40 \text{ cm} \times 40 \text{ cm} \times 15 \text{ cm}$ soil blocks, regularly distributed in the nodes of a 8×8 grid (i.e. $70 \text{ m} \times 70 \text{ m}$). Earthworms tended to exhibit contagious distribution in both land use systems and a multivariate relationship appeared between the size of the earthworm and aggregation indices employed. Land use had no significant effect on the spatial distribution of earthworms. Earthworms belonging to different ecological categories and from diverse sizes were spatially distributed in patches of several tens of meters. Semivariograms reflected, occasionally, the size of these patches, although, no spatial structure was perceived at some dates. This might be due to the intersample distance employed (10 m) so that changing the scale of observation could reflect unseen patterns.

In this study our purpose was first to determine the strength of spatial aggregation in some Neotropical earthworm species and then to observe the movement of population patches at the scale of the studied area. The use of these types of spatial analyses in conjunction may be very useful to establish the dynamics of earthworm populations in the field. © 2001 Elsevier Science B.V. All rights reserved.

Keywords: Spatial distribution; Geostatistics; Principal component analysis; Aggregation index; Population ecology; Savannas

1. Introduction

There have been many recent studies on the abundance and spatial distribution of several animal taxa in soils. Geostatistical analysis has been used to assess the spatial distribution of microorganisms and Collembola (Fromm et al., 1993), nematodes (Wallace

and Hawkins, 1994; Robertson and Freckman, 1995; Delaville et al., 1996; Rossi et al., 1996; Ettema et al., 1998) and earthworms (Poier and Richter, 1992; Stein et al., 1992; Rossi et al., 1995, 1997).

Although, earthworm distribution is known to be irregular and aggregated (Guild, 1955; Satchell, 1955; Svendsen, 1957), a first attempt to quantify the spatial pattern on earthworm populations was initiated by Phillipson et al. (1976) and it has been followed by Lavelle (1978, 1983a, 1988), Fragoso (1993), Boag et al. (1994), Rossi (1998), Rossi et al. (1995, 1997),

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Rossi and Lavelle (1998), and Decaëns and Rossi (in press) in more detail. These studies indicated that soil fauna was generally spatially structured at small scales (<100 m) (Robertson, 1994).

Little is known about the factors that control or influence the observed spatial patterns of soil macrofauna. Presumably, abiotic factors are responsible at least partly, for the spatial pattern of these soil macroinvertebrates. However, several studies have shown a lack of relationship between soil abiotic factors and the spatial pattern of various taxa of invertebrates. For example, Rossi et al. (1997) showed that the distribution of the endogeic megascolecid earthworm Polypheretima elongata Perrier was independent of a strong gradient in soil organic matter, their food resource. Ettema et al. (1998), in a study on spatiotemporal dynamics of bacterivorous nematodes, demonstrated the lack of correlation between nematode and soil resource patterns, although, they explained it by the young age of the investigated plots. At a given scale, organisms may appear to be distributed in relation to some abiotic factors (e.g. soil texture and plant-parasitic nematodes (Wallace et al., 1993)) though Rossi and Quénéhervé (1998) showed that these kind of correlations might be spurious in some cases. Generally, species distributional patterns are likely to be controlled by many factors acting at different scales both in time and space.

The spatial distribution of soil organisms is influenced, among other factors, by the plant cover, resulting in a horizontal mosaic of areas subjected to gradients of nutrient availability and microclimatic conditions (Lavelle, 1983b). Such a pattern is probably more pronounced in savanna ecosystems than in any other, because of the strong seasonality affecting both temperature and moisture gradients. On the other hand, land use systems and agricultural practices, have been shown to directly affect soil resource patchiness (Robertson et al., 1993). In turn, a change in soil resource heterogeneity is supposed to have huge potential effect on plant community structure and the distribution of soil living organisms (Tilman, 1988). Plant community composition may be itself determined by the type of agricultural practice and soil nutrient status (Miles, 1985; Wardle and Lavelle, 1997). Furthermore, if plant community influences, for example, termite and earthworm assemblage composition (Lavelle et al., 1997), it should also be affecting their spatial distribution. The impact of these organisms in soil functioning is obviously conditioned by the per capita effect but also by the spatial location of the individuals. This is a relevant feature when the effects of ecosystems engineers (sensu Jones et al., 1994) are assessed.

The aim of this work was to examine the potential effect of changing a natural savanna into improved pasture by planting African grass and tropical herbaceous legume species, combined with grazing, upon spatial distribution of earthworms and their morphological traits. We also tested the relationship between spatial distribution and morphological parameters according to land use, and the suitability of geostatistical techniques for description of earthworm distribution with the intersample distance used.

We estimated different aggregation indices for earthworm species in a natural and a disturbed savanna in order to determine how disturbance could affect spatial distribution of populations. However, aggregation indices do not take into account the actual location of the sampling units and, thus, provide no information on the spatial distribution at scales larger than the sampling unit size (Rossi et al., 1995). We, therefore, intended to fully characterize the earthworm spatial distribution by designing a spatially explicit sampling scheme and processing the data using geostatistics. This method allows assessment of consistency of spatial patterns as well as the scale at which they are expressed (Robertson, 1994).

2. Materials and methods

2.1. Site description

The study was undertaken at the CIAT-CORPOICA Carimagua field station, in the well-drained isohyperthermic savannas of the Eastern Plains of Colombia (4°37′N and 71°19′W) at 175 m altitude. Average yearly rainfall is 2280 mm with a mean temperature of 26°C, and a dry season from December-March. Soils are of two types: low-fertility oxisols in the upland ("altos") and ultisols in the low-lying ("bajos") savannas. The former are characterized by their acidity (pH 4.5, water), a high Al saturation (>90%) and low values of exchangeable Ca, Mg and K.

Two different and contrasting paddocks were investigated: A native savanna without grazing nor burning

where Andropogon bicornis, Gymnopogon sp., Panicum sp., Trachypogon sp. and Imperata sp. are the most abundant plant species, and a 17 years old grazed pasture, in a two ha plot, that combines an exotic African grass, Brachiaria decumbens cv. Basilisk, and a tropical herbaceous legume, Pueraria phaseoloides CIAT 9900 ("Kudzu").

2.2. Earthworm biology background

In the investigated paddocks, the earthworm community is comprised of eight species, all natives and present in the introduced system (Jiménez et al., 1998b). Most species belong to family Glossoscolecidae, the most widespread earthworm family in the neotropical region, Andiodrilus sp. (endogeic mesohumic), Andiorrhinus sp. (endoanecic), Andiorrhinus sp. 2 (endoanecic), Aymara sp. (epigeic), Glossodrilus sp. (endogeic polyhumic) and Martiodrilus carimaguensis (anecic), and two new genera belonging to the families Ocnerodrilidae (endogeic polyhumic) and Acanthodrilidae (epigeic) (Jiménez and Moreno, unpublished data). There are major differences in the structure of the earthworm communities in the savanna and pasture plots. In the savanna endogeic species make up 83.2% of the total biomass, ca. 3.26 g fresh weight per m², Glossodrilus sp. being the most abundant species, whereas in the pasture, several endogeic species represent only 12% of the total earthworm biomass, 57.1 g fresh weight m², and the anecic species accounts for 88% of total biomass (Jiménez et al., 1998b).

2.3. Earthworm sampling

Two sampling strategies were used in this study.

- 1. Monthly sampling: a stratified random sampling procedure was performed during 17 months (from April 1994 to September 1995, except June 1994) in both systems. Earthworms were monthly handsorted from 1 m × 1 m × 0.5 m monoliths (Lavelle, 1978) in each system and carried to the laboratory where length (mm), preclitellar diameter (mm) and weight (g) were measured.
- 2. Spatially explicit sampling: in both plots, 64 points distributed in the nodes of a regular $70 \, \text{m} \times 70 \, \text{m}$ grid were sampled for earthworm analysis. A soil

monolith of $40 \,\mathrm{cm} \times 40 \,\mathrm{cm} \times 15 \,\mathrm{cm}$ was rapidly taken in each point and handsorted in order to count all the earthworm species and their different demographic categories: adults (with clitellum and sexual protuberances), juveniles (subadults, sexual protuberances present but no clitellum and immatures, neither clitellum nor sexual protuberances) and cocoons. The method, following Rossi and Lavelle (1993) was adapted due to the large size of M. carimaguensis. All the earthworms were later replaced into the soil at their original places in the grid. To avoid sampling at the same points in the different dates samples were displaced along a spiral whose origin was represented by the point sampled at the first date. With respect to the size of the paddocks we considered as identical the sampling points from one date to another. The two plots studied were surveyed at three different times: November 1993, November 1994 and May 1995, for the native savanna, and September 1993, October 1994 and June 1995 for the grass-legume pasture.

2.4. Data analysis

2.4.1. Aggregation indices

As a first attempt to determine the spatial pattern of earthworms, two aggregation indices were computed by Morisita (1959) and Taylor (1961) as I_d and b, respectively.

1. Comparing spatial aggregation of populations with very different mean densities requires an index insensitive to mean density. Morisita (1959, 1971) proposed an index of aggregation independent on the sample mean and the total number of individuals in samples $(\sum x)$. However, this index is strongly related to the total number of sampling units. It is defined as

$$I_{d} = \frac{n\left(\sum_{i=1}^{n} (Z_{i}^{2}) - \sum_{i=1}^{n} Z_{i}\right)}{\left(\sum_{i=1}^{n} Z_{i}\right)^{2} - \sum_{i=1}^{n} Z_{i}}$$

where n is the number of sampling units and Z_i the count for the ith sampling unit.

The Morisita's index equals 1 for random distributions, is less than 1 for regular distributions and greater than 1 for clumped distributions. Departure from randomness can be statistically tested

by comparison to a chi-square (χ^2) value (Elliot, 1971).

2. The Taylor's Power Law (Taylor, 1961, 1984; Taylor et al., 1978) is based on the empirical relationship between the mean (m) and the variance s^2 that are related by the exponential function

$$s^2 = am^b$$

The parameters a and b are population parameters. Parameter b is considered to be an intrinsic measure of population aggregation varying continuously from zero for regular distribution ($s^2 = a$ with a < 1) through 1 for random distribution ($s^2 = m$ with a = 1) to ∞ for strongly contagious distribution (Taylor, 1961).

The parameter a is a scaling factor depending on the size of the sample unit (Taylor, 1961; Elliot, 1971) and the exponent b is used as an aggregation index. Parameter b is assumed to be independent of the mean so that it is a value measuring aggregation without being confused by changes in population mean density (Taylor, 1961, 1984; Taylor et al., 1988).

The parameters a and b are determined by linear regression of sample means and variances after the mean and variance estimates are log-transformed.

$$\log s^2 = \log a + b \log m$$

The Taylor's Power Law aggregation index is assumed to be species-specific (Taylor, 1961, 1988) and has been shown to be closely related to life history traits of various African earthworm species (Rossi and Lavelle, 1998).

2.4.2. Principal component analysis (PCA)

A PCA (Webster and Oliver, 1990) was performed to examine the relationships between several biological traits, i.e. adult weight (AW), adult length (AL), immature weight (IW), immature length (IL), length/width ratio (L/W), weight/width ratio (W/W) and mean annual vertical distribution (VD) together with Morisita's and Taylor's Power Law indices. To determine if land use influences these variables, such as aggregation indices and biological traits, that were recorded for earthworm populations from the pasture and the natural savanna plots, they were analysed together.

2.4.3. Geostatistics

Geostatistics or regionalized variable theory (Matheron, 1965, 1971) is a useful tool to describe any spatially structured variable in the soil. Central to geostatistics is the semi-variogram, the function which describes the evolution of the semi-variance with the inter-sample distance (Webster and McBratney, 1989). It is the single most important tool in geostatistical applications to soil (McBratney and Webster, 1986). In the regionalized variable theory the quantity $\gamma(h)$ is known as the semi-variance and it is estimated using the algorithm.

$$\gamma(h) = \frac{1}{2}M(h)\sum_{i=1}^{M(h)} [Z(x_i) - Z(x_i + h)]^2$$

Where M(h) is the number of sample pairs at each distance interval h ("lag") and $Z(x_i)$ and $Z(x_i + h)$ the values of the variable at any two places separated by a distance h.

The semi-variogram is the plot of the semi-variance against the distance (lag). Its shape indicates whether the variable is spatially dependent or not, e.g. there is spatial autocorrelation. The more alike the values separated by a given distance, the lower the semi-variance. Thus, generally the semi-variance increases with increasing distance as similarity between points decreases. If the variable is not spatially dependent, the semi-variance fluctuates around the sample variance and the semi-variogram is "flat".

Estimated values of the semivariance are adjusted to a theoretical model (McBratney and Webster, 1986; Webster and Oliver, 1990; Rossi et al., 1995), which is fitted to the semi-variogram calculated from sample values. Generally, only several authorized functions are used in semi-variogram modeling (see Webster, 1985 and Rossi et al., 1995 for a review of the models normally used). In the case of spatial autocorrelation, when the lag distance is increased the semi-variance gets to a maximum value (the "sill" variance) for a given distance interval (the range). In the semivariogram the range marks the limit of the spatial dependence or autocorrelation in the variable concerned. For distances larger than the range the semi-variance remains fixed leading to a plateau in the semi-variogram. The intercept of the semi-variogram is generally a non-zero value called the nugget variance (C_0) that expresses the variability due to unseen patterns

(occurring at scales shorter than minimum inter-sample distance). The sill variance minus the nugget variance is the spatial variance (C). This term accounts for the part of the total variance that can be modeled by the spatial structure (McBratney and Webster, 1986).

Semivariograms were estimated using the software VAR 5 (Yost et al., 1989), which allows semivariogram estimation and model fitting by a modified least square differences procedure (Cressie, 1985).

2.5. Contour maps

Classical contour maps were plotted using the command "contour plot" of the software Sigma plot 4.0.

3. Results

Density of earthworms varied both among species and dates in relation to geostatistical sampling. In both systems the highest abundance was obtained for *Glossodrilus* sp. In the savanna numbers ranged from 18.9 (November 1994) to 46.3 individuals per

m² (November 1993), and from 36.9 (1994) to 102.5 individuals per m² (1995) in the pasture. The epigeic species Aymara sp. was only obtained in 1995 sampling where the density of this species was 7.81 and 13.5 individuals per m² in the savanna and pasture, respectively. The density obtained for M. carimaguensis and Andiorrhinus sp. was less than 1 individual per m², except for the former in 1995 in the pasture plot (14.5 individuals per m²). Values of density for Ocnerodrilidae n. gen n. sp. were also very low; in the savanna the abundance of this species varied from 2 individuals per m² in 1993 to 7 individuals per m² in 1995 and in the pasture from 6.6 to 18.8 individuals per m² in 1994 and 1995, respectively. Finally, the numbers of Andiodrilus sp. varied from 1.97 individuals per m² in 1993 to 2.7 individuals per m^2 in 1995 in the savanna and from 3.5 (1993) to 6.2 (1995) in the pasture. Tables 1 and 2 summarize the data obtained for the different demographic stages of all species in the savanna and pasture, respectively.

Earthworm species tended to exhibit clumped spatial distributions. Indices I_d and b showed a strong aggregation for all species in both systems (Table 3).

Table 1 Summary statistics of abundance of earthworms (N m⁻²) of the 64 points regular grid during three dates in the savanna^a

Species	Category	Date 1			Date 2			Date 3		
		Mean	S.D.	S.E.	Mean	S.D.	S.E.	Mean	S.D.	S.E.
Andiodrilus sp.	Adults	0.94	2.9	0.36	0.7	2.2	0.3	2.1	4.2	0.53
	Juveniles	1.03	2.9	0.37	2.1	3.7	0.5	0.4	1.5	0.18
•	Cocoons	0	0	0	0.4	1.8	0.2	0.5	1.6	0.20
Andiorrhinus sp.	Adults	0	0	0	1.6	12.5	1.6	0.09	0.75	0.09
_	Juveniles	0	0	0	0	0	0	0	0	0
	Cocoons	0	0	0	0	0	0 ,	0	. 0	0
Aymara sp.	Adults	0	0	0	0	0	0	4.5	6.4	0.80
	Juveniles	0	0	0	1.36	3.6	0.4	3.3	5.3	0.67
	Cocoons	0	0	0	0	0	0	0	0	0
Glossodrilus sp.	Adults	19.3	19.6	2.5	14.3	17.2	2.2	0	0	0
	Juveniles	27.0	29.8	3.7	4.7	6.1	0.7	30.45	25.9	3.2
	Cocoons	3.9	7.2	0.9	0.3	1.8	0.2	0	0	0
M. carimaguensis	Adults	0	0	0	0.09	0.75	0.09	0.3	1.78	0.22
	Juveniles	0	0	0	0	0	0	0.39	1.92	0.24
	Cocoons	0	0	0	0	0	0	0	0	0
Ocnerodrilidae sp.	Adults	1.9	6.0	0.75	3.13	6.23	0.78	4.8	1.63	0.2
	Juveniles	0.09	0.75	0.09	0	0	0	2.1	8.3	1.03
	Cocoons	0	0	0	0	0	0	0	6.1	0.76

^a Data from the spatially explicit sampling.

Table 2
Summary statistics of abundance of earthworms (N m⁻²) of the 64 points regular grid during three dates in the pasture^a

Species	Category	Date 1		Date 2			Date 3			
	· · · · · · · · · · · · · · · · · · ·	Mean	S.D.	S.E.	Mean	S.D.	S.E.	Mean	S.D.	S.E.
Andiodrilus sp.	Adults	0.6	2.1	0.26	1.5	3.9	0.48	2.4	5.4	0.67
	Juveniles	2.9	5.6	0.70	2.9	6.5	0.82	3.8	9.6	1.2
	Cocoons	0	0	0	0.9	3.0	0.37	1.9	4.2	0.53
Andiorrhinus sp.	Adults	0.56	1.76	0.22	` 0	0	0	0.4	1.5	0.18
	Juveniles	0.84	2.36	0.29	0.2	1.0	0.13	0	0,	0
	Cocoons	0	0	0	0	0	Ó	0	0	0
Aymara sp.	Adults	0 -	0	0	0.2	1.05	0.13	10.5	12.7	1.6
1	Juveniles	Ò	0	0	0.3	1.28	0.16	3.0	5.8	0.7
	Cocoons	0	0	0	0.7	3.34	0.42	0.8	6.2	0.8
Glossodrilus sp.	Adults	6.9	9.4	1.18	34.3	22.0	2.7	0	0	0
	Juveniles	59.7	54.2	6.78	2.5	6.4	0.8	102.5	68.4	8.5
	Cocoons	0.2	1.0	0.13	0	0 .	0	0	0	0
M. carimaguensis	Adults	0.94	2.20	0.27	1.8	4.6	0.57	6.0	7.1	0.89
	Juveniles	0.09	0.75	0.09	0.1	0.7	0.09	8.5	10.4	1.30
	Cocoons	0	0	0 -	0	0	0	0 , 1,	. ,0	0
Ocnerodrilidae sp.	Adults	0 -	0 ;	0	6.44	10.6	1.32	14.6	17.7	2.22
	Juveniles	0	.0	0	0.2	1.62	0.20	4.2	8.2	1.02
	Cocoons	. 0	0	. 0	0	0	0	0	0	0

^a Data from the spatially explicit sampling.

All regression coefficients were significant at 1% in both systems except for the anecic M. carimaguensis (P < 0.05). Owing to $I_{\rm d}$ index Andiodrilus sp., M. carimaguensis and Aymara sp. were the most aggregated species in the savanna, whereas in the pasture the highest aggregation appeared for Andiodrilus sp., Andiorrhinus sp. and Aymara sp. All the values were statistically different from 1, as given by the χ^2 -test.

The PCA showed general multivariate relationships between the size of earthworm and aggregation (Fig. 1). The first two axes accounted for 57.6 and 22.1% of the total inertia, respectively. Axis 1 was defined by the yearly average vertical distribution and biometric variables, except L/W ratio. Axis 2, on the other hand, separates the former ratio and both aggregation indices. Axis 1 segregated larger species living at great depth (M. carimaguensis) from those smaller living near the soil surface, whereas axis 2 separated those species with higher L/W ratio (Glossodrilus sp.) from those species with a

Table 3
Aggregation indices obtained for several earthworm species in the savanna and pasture plots

Species	Morisita's index (I _d)		Taylor's inde	x (b)	Average value of b	S.E.
	Savanna	Pasture	Savanna	Pasture		
Andiodrilus sp.	2.903 ^a	3.396 ^a	1.889	1.704	1.797	0.131
Andiorrhinus sp.	ND^{b}	3.759 ^a	ND^b	1.801	1.801	ND^b
Aymara sp.	3.722 ^a	3.505 ^a	1.634	1.385	1.510	0.176
Glossodrilus sp.	1.833 ^a	2.160 ^a	1.476	1.615	1.546	0.098
M. carimaguensis	4.857 ^a	1.137 ^a	1.542	1.405	1.474	0.097
Ocnerodrilidae sp.	2.604 ^a	2.234 ^a	1.579	1.603	1.591	0.017

^a Values statistically different from randomness (Chi-square test).

^b ND: not determined.

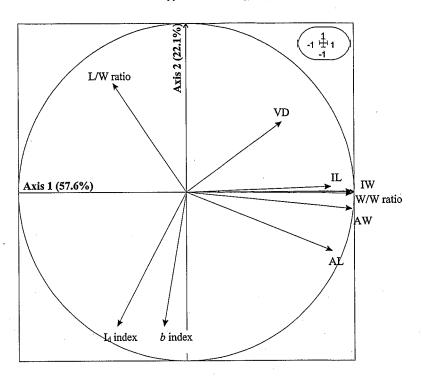


Fig. 1. Factorial map of the principal component analysis (PCA) performed with some biometric variables and both aggregation indices employed.

more contagious distribution and lesser L/W ratio (Andiodrilus sp.). The effect of land use was not significant except for the variable annual vertical depth (t-test, P=0.0123). The pattern of species upon

factorial axes 1 and 2 (Fig. 2) revealed that land use was of no consistent effect so that the distribution of these species is to be explained by biological features.

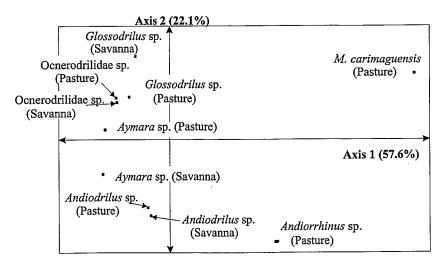


Fig. 2. Projection of the objects (species in plots) onto the first two axes of the PCA.

Table 4
Models of the semivariograms and range (m) obtained for each species during the whole study period^a

Species name	Demographic stage	Native savanna				Introduced pasture		
		1993	1994	1995	1993	1994	1995	
Andiodrilus sp.	Adults		· _	Spherical 44.7	_	Spherical 28.7	Spherical 54.5	
	Juveniles	_	4 , • • • • •			Spherical 27.1	Spherical 30.3	
	Cocoons	-	- . **				Spherical 32.1	
Aymara sp.	Adults		_	Spherical 41.2	<u>.</u>		_	
	Juveniles	_	· _	Linear	_ `.	-		
	Cocoons	_		_	-	-:	· - (,	
Glossodrilus sp.	Adults	Spherical 37.2	_		Linear	-	<u></u>	
	Juveniles	Spherical 29.8	Spherical 36.2			Spherical 57.4	Spherical 41.8	
	Cocoons	Linear		_	_	<u>-</u>	_	
Ocnerodrilidae sp.	Adults		Spherical 30.9		_	- Spring	Spherical 30.8	
	Juveniles			_				
* • •	Cocoons	ND ^{b'}	ND^b	NDb	ND^b	ND_p	ND^b	

^a Symbol (-): nugget variogram.

The spatial structure of the demographical stages of the different earthworm species was only assessed in some semivariograms. This means that, in most cases, the intersample distance employed, i.e. 10 m, was too large to allow a proper assessment of the spatial structure for this specific earthworm community. The aggregated patterns showed by aggregation indices are likely to be expressed beyond the scale of 10 m in most of the cases we studied. When significant, most of the variograms were of spherical type (Fig. 3) and

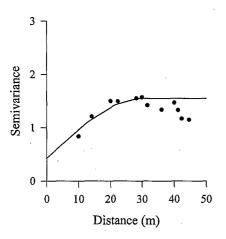


Fig. 3. Spherical semivariogram obtained for Andiodrilus sp. in the pasture plot in June 1995.

indicated that spatial variability was expressed at scales ranging from the 27.1 to 57.4 m. (Table 4).

In both systems earthworm population was spatially distributed in patches of ranging sizes from date to date. Species were aggregated in several patches of an average size of 30–40 m, sometimes greater. As an example, in September 1993 Glossodrilus sp. was mostly composed of juveniles in the pasture (Fig. 4). Two patches could be observed, one of them large placed on the left side of the plot and the other one located at the right corner somewhat smaller than the former. By June 1995 the whole population was made up of juveniles, since adults begin to appear at the end of the rainy season (Jiménez et al., 1998b). It was as if the large patch would have moved to the left and in the right corner two new patches were distinguished.

4. Discussion

Earthworms are organisms with a highly contagious spatial distribution (Rossi and Lavelle, 1998). Our results agree with Piearce (1982) who found that, with independence of the aggregation index employed, earthworms were highly aggregated in soils of the north western coasts of England. Values for the Taylor b index were very similar to those obtained by Taylor et al. (1978) who used data from Gerard (1960). In

^b ND: not determined.

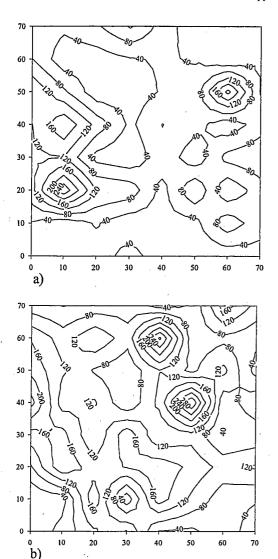


Fig. 4. Contour maps of abundance $(N \, m^{-2})$ of juveniles of Glossodrilus sp. at two different dates in the pasture plot.

our study, the large value of b index for Andiorrhinus sp. could be explained by means of its abundance in the systems studied. Hairston (1959, quoted in Lloyd, 1967) concludes that aggregation indices are higher for rare species and Andiorrhinus sp. is a rare species in Carimagua (Jiménez et al., 1998b).

The evaluation of aggregation indices depends on the measurement scale and applies to all methods that quantify the spatial variance of the distribution of any organism (Horne and Schneider, 1995). Curtis and Mcintosh (1950) and later Pielou (1969) and Greig-Smith (1983) demonstrated the dependence of several indices on the measurement scale. Hurlbert (1990) concluded, on the contrary, that it is an aggregation measure without interpretation. So, comparison of aggregation indices among populations are only valid when evaluated at the same spatial scale.

M. carimaguensis is active and in the same burrow during at least the first week after cast deposition began (Jiménez et al., 1998a; Decaëns, 2000). This species inhabits semi-permanent U-shaped vertical burrows (Jiménez and Decaëns, unpublished data) and is strongly influenced by seasonal patterns in climate. The burrow is normally abandoned after rain saturation that results in a depletion of O_2 in the soil (Jiménez et al., 1998a). Maybe this kind of behavior leads to the low values of I_d index obtained in the pasture, and assessment of the distribution pattern of the different demographic stages, i.e. adults versus immatures, would reveal unseen patterns.

Both aggregation indices employed in this study provided useful tools to assess spatial pattern and its correlation with biological features, i.e. earthworm size. This type of relationship was also established in the case of earthworm community from an African savanna (Rossi and Lavelle, 1998). These authors found that smaller species were strongly aggregated. At Carimagua species of medium to large size were more aggregated, so conclusions must be drawn carefully and other earthworm communities should be included in this type of analysis in order to establish the patterns that define the aggregation of species. A plausible explanation may be the absence of a true anecic species, like M. carimaguensis, in Lamto. We also agree with these authors that in order to define more precisely the ecological categories of earthworms this kind of analysis should be used, especially if b index is species-specific as Taylor et al. (1988) stated.

Several factors may be responsible for the unsuccessful performing of geostatistical analysis.

- Density: population density fluctuates seasonally due to environmental conditions. The number of earthworms at the end of the rainy season (1993 and 1994) is somewhat low for some species as they have already entered the inactivity period.
- 2. Size of species: sampling was carried out manually and not all the individuals are caught when

hand-sorting method is applied. For several species, *Aymara* sp., *Glossodrilus* sp. and Ocnerodrilidae n. gen. n. sp., almost 50% of the whole population is not collected when employing this method (Jiménez, unpublished data).

- 3. Size of sampling units: the size employed may not be so optimum as to get large anecic species as *M. carimaguensis* which rapidly retreats deeper into its vertical burrow.
- 4. Intersampling distance: since populations display a clear aggregated spatial structure, at least at the scale of the sampling units, it is likely that the minimum inter-sample distance is too large to allow a clear assessment of the spatial distribution at scales larger than this value. Our results suggest that some earthworm species are aggregated in local spots less than 10 m diameter.
- 5. Mosaic structure: Martin (1991) hypothesised about the presence of subpopulations at different demographic phases for the species *Millsonia anomala* Omodeo. Maybe, the observed patterns in the earthworm community of both systems studied correspond to a spatiotemporal mosaic of population patches. But this hypothesis needs to be further tested.

The horizontal distribution patterns of earthworms are difficult to assess since they have been described on various scales, from local (Phillipson et al., 1976; Lavelle, 1978) to regional (Evans and Guild, 1948; Németh, 1981) and geographical scale (Bouché, 1972; Fragoso, 1993). The spatial variability in the horizontal distribution of earthworms ranges from 1 (Rossi, 1998) to 50 m (Poier and Richter, 1992). The tropical endogeic P. elongata Perrier is distributed in clumps of 20-30 m diameter, with a marked opposite pattern amongst adults and juveniles (Rossi et al., 1997). None of the demographic stages of this species were spatially autocorrelated with some soil variables, e.g. soil organic matter and clay gradients. In the study previously mentioned the authors explain the observed patterns on the basis of demographic processes, such as intraspecific competition that leads to decreasing fecundity in areas of high adult density. The size of patches coincides with those reported by Rossi et al. (1997) and Rossi and Lavelle (1998) for some savanna species from Lamto (Ivory Coast). Besides, the limits of the spatial distribution obtained in this study are

within those described by Robertson (1994), below 100 m.

Our sampling strategy employed was not very useful to assess adequately using geostatistics the spatial pattern of earthworm distribution, even though when species have been shown to be spatially aggregated (aggregation index values different from random). Aggregation is a characteristic of earthworm populations that was easy to demonstrate with both I_d and b indices, but a finer resolution must be carried out to show actual significant spatial patterning with geostatistics.

From our results we cannot conclude if the observed distribution is due to inner or external driving factors within the population. Rossi et al. (1997) assessed the spatial distribution of *P. elongata* in market-flower gardens of Martinique Island by means of inner processes within the population. In other study Rossi and Quénéhervé (1998) found no significant relationship between both plant-feeding and free-living nematodes and several soil parameters in vertisols from the same site, except for the density of the plant-feeding nematode *Helicotylenchus retusus* Siddiqui and Brown which was negatively correlated with clay content.

Disturbance plays a major role in structuring ecological systems as they create a spatiotemporal mosaic of patches. Moloney and Levin (1996) indicated that the spatial pattern of disturbance has in general been ignored, with attention only restricted to understanding the effects of both rate and intensity of disturbance. Studies on disturbance hardly have considered the spatial structure of the disturbance regime. In addition to this, the spatiotemporal dynamics of soil variables in disturbed systems, i.e. density of earthworms, have not been deeply studied despite of Rossi (1998), Decaëns and Rossi (in press) and Jiménez (1999) who provide information on this subject.

In this paper we have showed the usefulness of the different statistical tools employed in the spatial analysis of earthworm populations. On the other hand, and interestingly enough, land-use practices in the disturbed ecosystem had no clear impacts on the horizontal spatial distribution of earthworms. This suppports the hypothesis that spatial distribution is determined by factors acting within the population itself, but it needs further research. In a next paper (Jiménez et al., unpublished) a brand new method to detect common spatial patterns across several

sampling dates will show the stability and opposite distributions among different earthworm species.

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