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Sampling strategies for biotope definition: minimal sampling area for selected groups of macroinvertebrates in the rocky subtidal of São Miguel, Azores

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Abstract Azorean rocky shores are mainly characterized by patchy algae-based communities with variable associated macrofauna. Characterization studies should therefore include quantitative information for both algae and macroinvertebrates. Unlike for the algae, minimal sampling areas are undefined for macroinvertebrates in the Azores. The present study defines the minimal area to be used for the assessment of the abundance of conspicuous benthic macroinvertebrate abundance. This study proposes methodologies to be used for a selected group of invertebrates when simultaneously undertaking quantifications of macroalgae.

Keywords Sampling areas · Macroinvertebrates · Rocky subtidal · Biotope definition · Azores

Introduction

Many recent studies have focused their attention on characterising shallow-water marine benthic communities (e.g. Southward and Orton 1954; Russell 1991; Zacharias et al. 1999; Foster et al. 2003). Recently interest has focused on habitat and community characterisation for marine conservation purposes (Mumby and Harborne 1999; Zacharias and Roff 2000). Although coastal ecosystem classifications for management purposes have been developed in the EU and USA, there are only two publications (Tittley et al. 1998; Tittley and

Neto 2000) that outline a provisional benthic biotope classification for Azorean rocky shores following the lines of Hiscock (1995) and Connor et al. (1997). They focused only on stable rocky substrata (not those on mobile stones and cobbles) and were based on descriptive information. Recent studies which are being continued (Macedo 2002) use a quantitative approach to: (a) review the biotope classification of Tittley and Neto (2000); (b) characterise the communities associated with all rocky substrata; (c) define algal-based communities with associated benthic macrofauna; and (d) create a more complete biotope classification of the intertidal zone of São Miguel. These will also set the guidelines for methodologies to be used in further biotope surveys in the Azorean archipelago. The Azorean archipelago (37°40' N and 20°31' W) is distributed unevenly along the Mid Atlantic Ridge. Its nine islands are volcanic in origin and the shores present a convoluted morphology, where high and steep cliffs alternate with rocky beaches of irregular rock sizes (Borges 2004; Morton et al. 1998). Most of the coast of São Miguel Island is subject to medium and high levels of wave action, with low levels restricted to harbours (Macedo 2002). These conditions create different habitat conditions for a wide variety of fauna and flora. Azorean rocky shores are mainly characterised by patchy algal communities with an associated macrofauna of various species. Biotope characterisation studies should therefore encompass quantification of the more conspicuous algae and macroinvertebrates simultaneously. Unlike for algae (Neto 1997), minimal sampling areas have not been calculated for macroinvertebrates in the Azores. These communities are too large to be studied as a whole hence the need to representative samples. Larger samples confer a higher degree of confidence in being a good representative of its population of origin while smaller samples keep sampling effort to a minimum (Weinberg 1978). A compromise is necessary to satisfy both criteria and minimal sampling areas resolve the problem. They are large enough to give accurate qualitative and quantitative information about the composi-

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tion of the community, and small enough to keep sampling effort within reasonable limits (Weinberg 1978).

There are two main methods for the assessment of minimal areas: (1) traditional species-area curves; and (2) similarity/diversity indices-area plots. These methods account for the determination of a minimal area for the whole community, but if the objective is to identify the minimal sampling area for determining abundances of a specific organism or a restricted group of organisms they are not appropriate.

Naturally occurring species assemblages are variable by nature, and spatial variability can occur at different scales ranging from biogeographic to local (Underwood and Chapman 1996, 1998). This applies to species assemblages as well as to single populations. As a consequence, minimal sampling area may vary with population parameters and with geographical location. Therefore it should be determined for each organism/group of organisms and for each place or at least for each region.

As part of developing a methodology for biotope characterization on Azorean rocky shores, the present study defines the minimal area to be used for abundance assessment of conspicuous benthic macroinvertebrates. In areas where communities are mainly characterised by algae for biotope characterization, this study sets the quantification methodologies to be used for a selected group of invertebrates simultaneously with algal quantification.

Methods

The study was carried out on the sublittoral of São Miguel Island, Azores from February to March 2004 on 4 sampling sites (Fig. 1). The study location was selected from 15 possible sampling sites around the island using a table of random values.

The more conspicuous invertebrates were selected using empirical criteria such as representativeness in the habitats studied and low mobility, these were: sea stars [*Ophidiaster ophidianus* (Lamarck 1816), *Marthasterias glacialis* (Linnaeus 1758)]; sea urchins [*Sphaerechinus granularis* (Lamarck 1816), *Paracentrotus lividus*

(Lamarck 1816), *Arbacia lixula* (Linnaeus 1758)]; holothurians (*Holothuria* spp.); fire-worms [*Hermodice carunculata* (Pallas 1766)]; and tube-worms [*Spirographis spallanzani* (Viviani 1805)].

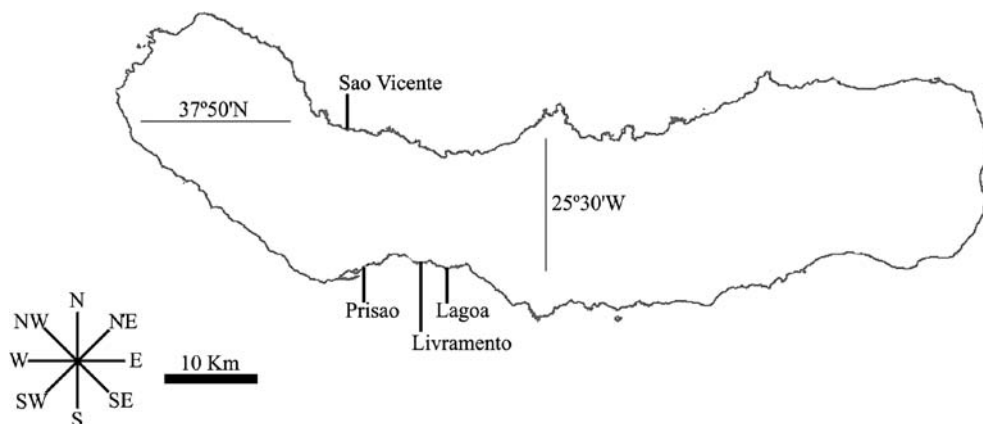
Three different depth levels were considered according to Neto (2001): (1) shallow water—5 m; (2) intermediate transition zone—15 m; and (3) deep water—25 m. At each depth level the total number of individuals was counted in four different areas (7.5, 15, 22.5, and 30 m²)—1.5 m wide transects associated to four length classes: 5, 10, 15 and 20 m. Transect width was chosen based on preliminary tests counting organisms along 10 m long transects of varying width (1, 1.5, and 2 m wide) with the help of PVC bars; counting easiness was the sole factor considered, and 1.5 m width chosen. Transect length categories were chosen to represent four levels of sampling effort, and 20 m considered the maximum effort possible. This decision was based on the time spent in counting organisms along three replicate transects (20 m long and 1.5 m wide) at the maximum depth of 25 m. Three replicates was the number chosen in the experimental design for invertebrate quantification procedures in the biotope characterization study. Invertebrate and algae quantification methodologies were planned to be executable together during only one dive and thus methodologies kept to a minimum of time consumption.

Transect location and geographic orientation were randomly chosen using a table of random values, and two replicate transects of each length were sampled.

Sampling strategy followed a fully orthogonal experimental design with depth (3 levels) and area (4 levels) as fixed factors.

Data were subject to analysis of variance (2-factor ANOVA), homogeneity of variances assessed using Cochran's *C* test and data transformation applied when necessary (Underwood 1997), using the statistical software package GMav 5.0 (University of Sydney). Non metric multidimensional scaling was applied to data for trend identification in samples, SIMPER analysis used to identify organism contributions for sample grouping trends, and ANOSIM to test for differences between samples using the software package PRIMER (Clarke and Warwick 2001).

Fig. 1 Sampling sites on São Miguel Island



Results

Univariate analysis

Analysis of variance showed a significant variability associated with depth regarding sea stars, tubeworms and fire-worms (Table 1). The interaction between depth and area is significant in the case of sea urchins. No significant differences were identified for holothurians.

Figure 2a and b show that both sea stars and tubeworms present higher density at the shallower depth-D1—contrary to what was observed for fire-worms (Fig. 2c) that showed higher density at higher depths-D3. Figure 2d shows great variability of sea urchin density: (i) within depth classes—with less variability in the intermediate depth level D2; and (ii) between areas—with less variability in the longer transects of 15 m and 20 m. Highest and lowest density values for sea urchins were found in the lowest and highest depths sampled (D1 and D3) and for the transect lengths of 5 m and 10 m, respectively, tending to stabilize for the transect lengths of 15 m and 20 m. A relatively constant density of sea

urchins is observed at the intermediate depth (D2) independent of transect length. Density of sea urchins was constant with respect to depth for transect lengths

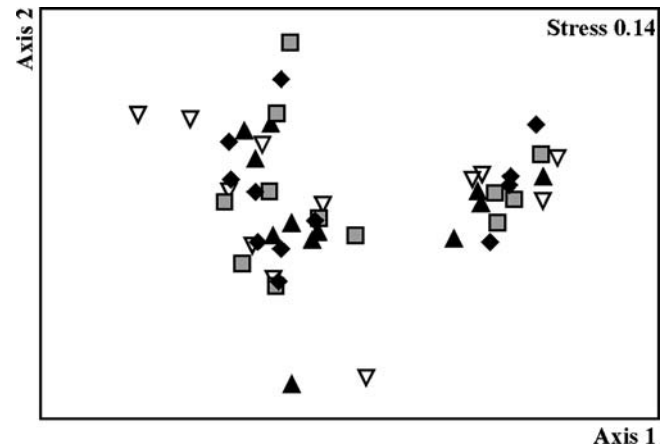


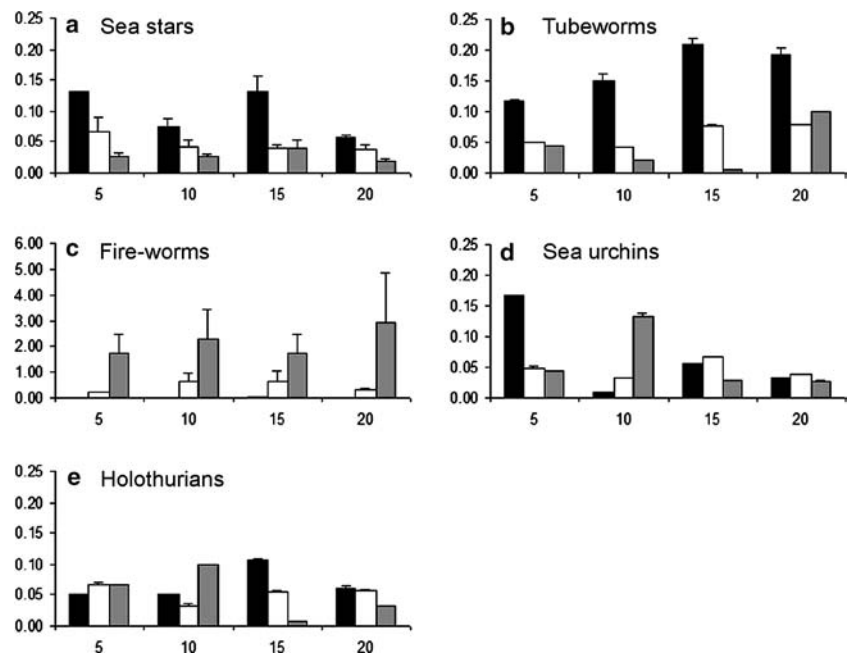
Fig. 3 Non metric Multidimensional scaling (nmMDS) groupings of samples labeled according to the four transect lengths considered for this study (*filled triangle* 5 m, *open triangle* 10 m, *shaded square* 15 m and *filled diamond* 20 m)

Table 1 ANOVA results comparing species abundance of selected macroinvertebrates at three depths and four sampling areas

	Depth (2 Df)		Area (3 Df)		Depth×area (6 Df)		Res (36 Df)
	MS	F	MS	F	MS	F	
Sea stars	0.0209	8.50**	0.0037	1.52	0.0016	0.66	0.0025
Tubeworms	0.0727	5.22*	0.0079	0.56	0.0037	0.27	0.0139
Fire-worms	194.0419	14.51**	1.3502	0.10	1.7758	0.13	13.3767
Sea urchins	22.0045	0.35	105.6977	1.69	157.0370	2.51*	62.5839
Holothurians	0.0010	0.19	0.0003	0.07	0.0049	0.95	0.0052

Bold text = significant P values ($P < 0.05 = *$; $P < 0.01 = **$)

Fig. 2 Mean abundance of macroinvertebrates (+ SE) at three depths (*filled square* 5 m, *open square* 15 m, *shaded square* 25 m), where the x -axis represents transect length categories (m) and the y -axis represents density (nm^{-2})



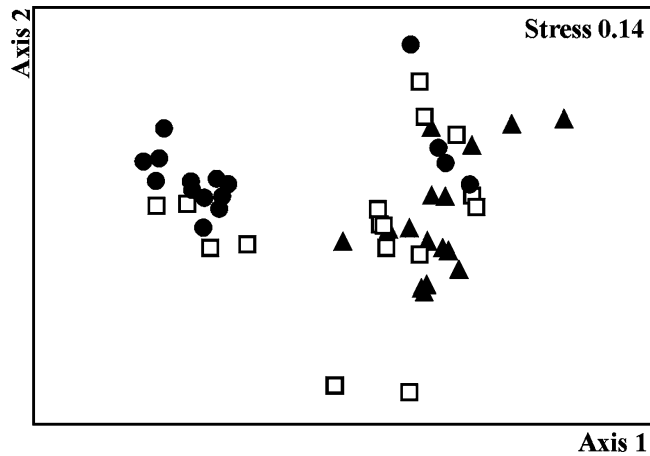


Fig. 4 Non metric Multidimensional scaling (nmMDS) groupings of samples labeled according to the three depth levels considered for this study (filled triangle 5 m, open square 15 m and filled circle 25 m)

of 15 m or more. Holothurian density was relatively constant for both depth and area classes (Fig. 2e).

Multivariate analysis

Non-metric multidimensional scaling of samples (n-MDS) showed no grouping associated to the factor area (Fig. 3). However, when depth is considered, patterns can be identified, namely on the horizontal axis between most of the samples taken at 25 m and 5 m depth, while 15 m depth samples co-occur with both groups (Fig. 4). On the vertical axis no separation can be identified. A higher variability was associated to samples taken at 5 m and 15 m depth.

For the separation of samples according to depth, SIMPER results (Table 2) showed that dissimilarity was lower between 5 m and 15 m (68.94) and between 15 m and 25 m (75.19), than that observed between 5 m and 25 m (84.28). Fire worms were the main cause of the dissimilarity between depth levels, and showed an increasing abundance with depth. Tube worms and sea stars—the second and third contributors for the dissimilarity—showed a decreasing abundance with depth, stabilizing down from the transition zone. Sea urchins and holothurians showed a constant average abundance across all depth levels.

ANOSIM test (Table 3) showed that all depth classes were significantly different from each other, being the most significant difference associated to the depth classes of 5 m and 25 m.

Discussion

Results from univariate and multivariate analysis were consistent: depth was the main factor which contributed for density variation in the selected groups of macroinvertebrates considered by the present study. These showed a similar depth distribution pattern to that reported by Neto (2001) for algae: two distinct commu-

Table 3 ANOSIM analysis results when testing for differences between the three studied depths (5, 15, and 25 m)

Groups	R statistic	Significance level (%)	Actual permutations	Number Observed
5, 15	0.128	0.2	999	1
5, 25	0.578	0.1	999	0
15, 25	0.229	0.6	999	5

Table 2 SIMPER analysis results when testing for differences between macroinvertebrate mean abundances at the three studied depths (5, 15 and 25 m)

Species	Average abundance	Average abundance	Average dissimilarity	% Contribution to dissimilarity	Cumulative % contribution
Average dissimilarity = 68.94					
	Group 5	Group 15			
Fire-worms	0.02	0.45	23.49	34.07	34.07
Tubeworms	0.17	0.06	16.85	24.44	58.51
Seastars	0.10	0.05	10.26	14.89	73.39
Holothurians	0.07	0.05	9.32	13.52	86.91
Sea urchins	0.07	0.05	9.03	13.09	100.00
Average dissimilarity = 84.28					
	Group 5	Group 25			
Fire-worms	0.02	2.18	60.07	71.27	71.27
Tubeworms	0.17	0.04	9.62	11.42	82.69
Seastars	0.10	0.03	5.40	6.40	89.09
Sea urchins	0.07	0.06	4.78	5.67	94.77
Average dissimilarity = 75.19					
	Group 15	Group 25			
Fire-worms	0.45	2.18	58.66	78.02	78.02
Holothurians	0.05	0.05	5.24	6.97	84.99
Sea urchins	0.05	0.06	4.04	5.37	90.36

nities at 5 m and 25 m depth separated by a distinct transition zone at 15 m depth. In the present study multidimensional scaling showed overlap of the samples taken at 15 m with those taken both at 5 m and 25 m. Both ANOSIM R values (Table 3) and SIMPER dissimilarity values (Table 2) are consistent with these observations although the latter are generally high, and similarity between 25 m samples lower than for samples at shallower depths. The areas (transect lengths) used in the present study did not imply any significant variability in the density of the selected groups of macro-invertebrates. This might mean that the transect lengths considered were not sufficiently large to identify the minimum area for a sampling strategy with this kind of organisms. Nevertheless, Table 3 indicates sea urchin average abundance to be stable across depth classes which leads to the conclusion that the main effect of this interaction arises from the area factor (i.e. transect length). From Fig. 2d it was possible to associate such high variability mainly to the transect lengths of 5 m and 10 m. Sea urchin density associated to longer transects (namely 15 m and 20 m) appeared to be quite stable, and these could therefore be considered as the transect lengths to be used in future sampling strategies. In this context, given the diving time constraints and the need to combine sampling strategies for algae and invertebrates simultaneously in the same dive, and to minimize sampling effort, the area chosen for future macroinvertebrate sampling in the Azores was $15 \times 1.5 \text{ m}^2$.

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