INTRODUCTION

Thraustochytrids are fungoid, Heterokonta protocysts (Cavalier Smith et al. 1994) common in marine and estuarine environments. These globular, saprobic organisms, ranging between 5 and 20 µm in size, undergo asexual reproduction through biflagellate zoospores and are characterized by an ectoplasmic net that penetrates solid particles and absorbs organic matter (Porter 1990, Raghukumar 1996). Thraustochytrids can be isolated from various substrata and appear to be abundant in the sediments of coastal and estuarine environments (Ulken 1981, Raghukumar & Schaumann 1993, Bongiorni 1999). Such widespread occurrence, together with their ability to feed on diverse complex organic substrata, suggests that these organisms play an important ecological role within the microbial loop. Recently, higher biomass and C/N ratios for thraustochytrids than for bacterioplankton were reported by Kimura et al. (1999), who concluded that thraustochytrids could play a more important role in the carbon cycle of marine ecosystems than has been previously recognized. Through degradation and decomposition, thraustochytrids promote the turnover of organic matter and establish trophic and competitive interactions with bacteria, ciliates and other protists. The biomass of thraus-
thraustochytrids in detritus should be high enough to represent a significant food source for detritivores and ciliates (Fenchel 1980, Raghukumar & Balasubramian 1991). Despite their likely ecological importance, these microorganisms are poorly understood because they cannot be preserved and stained directly in sediments, making the collection of quantitative data a trying task. To date, nearly all previous studies on thraustochytrid field distributions have relied upon techniques for culturing live samples mainly through the pine-pollen baiting method (Gaertner 1968). These culturing techniques clearly underestimate the actual numbers, as a direct method based on epifluorescence microscopy has recently yielded values some orders of magnitude higher (Raghukumar & Schaumann 1993, Naganuma et al. 1998, Kimura et al. 1999).

Ciliated protozoa are among the main components making up the micro-communities in temperate, marine, medium- to fine-grained sandy shores (Fenchel 1967, Epstein 1997a, b). Dense ciliate populations, feeding on diatoms and other protists, should control their populations and, feeding on bacteria, promote bacterial mineralization processes in the sediment (Fenchel et al. 1998). Ciliates, in turn, are a significant food source for many suspension feeders (Ribes et al. 1998, 1999). Ciliate density in Mediterranean sandy shores may exceed several hundred cells ml\(^{-1}\) (Santangelo & Lucchesi 1995).

Marine, interstitial ciliates have been the object of several quantitative studies based on direct counts, a partial review of which has been conducted by Finlay & Fenchel (1996). Recently, attempts have been made to describe the community structure and spatial distribution pattern of ciliates in Mediterranean sandy shores (Lucchesi & Santangelo 1995, 1997, Santangelo & Lucchesi 1995). In those studies, ciliates were extracted from the sand following Uhlig’s ice-water filtration method (Uhlig 1964), and live-sample counts were performed just after collection. Under these limiting conditions, a small number of replicated samples were examined. Only in recent years have ciliates been enumerated via fixation and staining by fluorochromes (Alongi 1993, Epstein 1995, 1997a,b, Tao & Taghon 1997). In particular, Epstein (1995) has set forth a new method for enumerating all nano-, micro- and meio-benthos from marine sediments by extraction from glutaraldehyde-seawater washed sand samples and the separation from detritus by centrifugation in a non-linear silica-gel gradient. In order to be counted efficiently, the fluorochrome-stained protists must be separated from sediment and detrital particles, which, due to background fluorescence, would mask them. This method enables extraction of nearly all ciliates from the sediment, and, by preserving them, also allows collection and examination of a higher number of samples, thus providing substantial quantitative data. According to Epstein (1995) this method collects nearly all nano-benthos and should therefore also be effective in isolating thraustochytrids. These fungi-protists were not identified by the protocol described by Epstein (1995) as it lacks a suitable staining procedure. A new method to directly identify thraustochytrids from several marine environments by fixing with formalin and staining them with acrylalvaine was developed by Raghukumar & Schaumann (1993). This fluorescent dye stains thraustochytrid cells, revealing orange-to-red fluorescing multilayered cell walls and yellow-green fluorescing nuclei. It seemed worthwhile to design a protocol allowing for simultaneous fixation, extraction, separation from detritus, and staining of these 2 major components of the interstitial microcommunity.

The protocol, at once an improvement in and a compromise between those expounded by Raghukumar & Schaumann (1993) for thraustochytrids and by Epstein (1995) for ciliates, allows a direct, reliable assessment of the abundance of both these protists. The thraustochytrid distribution in sandy sediments is completely unknown and that of ciliates is still only partly understood (Santangelo & Lucchesi 1995, Jonsson & Johansson 1997, Fenchel & Blackburn 1999). Knowing the scale at which ecological processes operate is highly important to our understanding of such processes. Simple data on abundances of organisms, also new for some protists (as is the case of thraustochytrids), do not however furnish any information about the spatial structure of the populations examined. Such data reveal nothing about distribution gradients and patchiness (Underwood 1997). Thus, to tentatively explore the distribution of both microorganisms, samples were collected following a sampling design for nested ANOVA with 3 successively smaller spatial scales, i.e. sampling units of different size (Fig. 1). Nested sampling designs with different spatial scales should provide reasonably accurate information about the spatial structure and dynamics of sand bottom communities (Morrisey et al. 1992, Underwood 1997). The findings presented here are the first stemming from the direct enumeration and distribution analysis of thraustochytrids in sediment.

**MATERIALS AND METHODS**

Study area and sampling design. Sampling was carried out at Marina di Pisa (43°42’ N, 10°16’ E), just south of the Arno river mouth, Tuscany, Ligurian Sea, Italy (Fig. 1), during the 1998-1999 autumn-winter season in October, November and January. An artificial breakwater with 2 small, north- and south-oriented passes sheltered the sandy shore. The sand was
medium-to-fine grained. Porosity ranged between 19.8 and 21.3%, and total organic matter between 1.17 and 0.72% (w/w %). Temperature and salinity ranged between 18 and 13°C and 34 and 36 psu respectively. Samples were collected during morning low tide in shallow water (0.30 to 0.40 m depth).

A 19.8 m × 6.6 m sandy area (Fig. 1) was divided into 12 3.3 m × 3.3 m squares. Using a randomization program 2 squares were selected at each sampling from within the area. Each square then yielded random two 1 × 1 m plots, and finally two 0.1 × 0.1 m subplots were randomly selected per plot. Four replicate sand samples, 1 ml (±0.04 SE) each, were collected from each subplot by coring the sandy bottom with a glass tube to a depth of 1 cm. Sampling was repeated at 3 different, randomly selected times. The nested factors were therefore: square, plot and subplot (with 2 levels each) with time as a fourth, 3-level random factor, hierarchical to the others.

Each collection yielded 32, 1 ml sand replicated samples (replicates), for an overall number of 96. Replicates were processed as described in the following, and data of both thraustochytrid and ciliate density (cells ml⁻¹) was then examined. The Cochran test was performed to assess variance homogeneity among samples, a basic assumption for ANOVA. As the test yielded significant results for ciliates, the data were log transformed and the test repeated. As the test results on log-transformed data were no longer significant, ANOVA was carried out on the transformed data (Underwood 1997).

**Extraction, separation and staining of protists.** Sand samples were fixed immediately after collection with 5 ml glutaraldehyde-seawater solution (1.5% final concentration) and kept cold. Glutaraldehyde fixation (Epstein 1995) was chosen over formalin fixation (Raghukumar & Schauman 1993) because it is the only fixative compatible with Percoll and works with thraustochytrids as well as ciliates. Glutaraldehyde was diluted in filter-sterilized (0.2 µm pore size membrane) artificial seawater (particle-free seawater = PFSW), kept at the same salinity as that measured in the field. The subsequent phases of the procedure are outlined in Fig. 2. Thraustochytrids and ciliates were extracted from each sample by gently hand washing/shaking them 4 times with 5 ml of 1.5% glutaraldehyde-PFSW solution; each time the 5 ml supernatant was collected yielding a total of 20 ml. A 4 ml subsample of this 20 ml of supernatant was then centrifuged. In this way ⅙ of the original sample volume (0.2 of initial 1 ml) was processed. Epstein (1995) washed sandy samples 5 times with 6 ml of glutaraldehyde-seawater solution, thus collecting 30 ml supernatant of which 2 ml subsamples were centrifuged (Epstein 1995), thereby processing only ⅙ of the original sample volume (0.096 of initial 1.44 ml). According to Epstein (1995), this method enables extraction of over 90% of the nano- and over 95% of the microbenthos from sand samples. We chose to reduce both the number of washings (from 5 to 4) and the volume of the washing solution (from 6 to 5 ml) to reduce the dilution of original samples, achieving an extraction efficiency similar to that obtained by Epstein (1995).

The subsamples (4 ml) were centrifuged in a non-linear silica gel gradient (Percoll) at 5000 × g for 15 min (the latter was prepared by centrifuging 12 ml of a 50% PFSW-Percoll solution at 30000 × g in a 18.5 ml tube for 15 min). As the uppermost fraction was never found to contain any ciliates or thraustochytrids, and the dense, lowermost layer held only detritus (undesired fluorescence source), these 2 layers were discarded (0.5 cm thick each, for an overall volume of 1.54 ml). The remaining Percoll gradient was then vacuum filtered (35 ml Hg) on a black polycarbonate
membrane (25 mm diameter, 1 µm pore size, Poretics). The membranes with filtered material were then rinsed with PFSW, flooded with fluorochromes and vacuum drained again. Ciliates and thraustochytrids were stained consecutively with 3 fluorochromes: acriflavine hydrochloride (Sigma A-8251), DAPI (4',6'-diamino-2-phenylindole hydrochloride; Sigma D-9542) and FITC (fluorescein isothiocyanate; Sigma F-1522). The staining solutions were prepared as described in Raghukumar & Schaumann (1993) for acriflavine and in Sherr et al. (1993) for DAPI and FITC. None of the fluorochromes interferes with another or has a jelling effect on Percoll.

The stains were vacuum drained after 4, 12 and 10 min, respectively, rinsed each time with PFSW and post-stained with 0.025% of the optical brightener Calcofluor White (Sigma F6259) for 1 min. Each filter was placed on a microscope slide between 2 drops of immersion oil (Cargille type A) and examined under an epifluorescence microscope (Jena-Zeiss) equipped with a 50 W mercury lamp. A filter for DAPI (excitation filter 360 nm, barrier filter 420 nm) and one for acriflavine/FITC (excitation filter 450 to 490 nm, barrier filter 520 nm) were used.

According to Raghukumar & Schaumann (1993), acriflavine simultaneously stains thraustochytrid sulphated polysaccharide cell walls red and their nuclei yellow-green, thus distinguishing these organisms from other similar-sized protists which fluoresce greenish only and lack red fluorescing cell walls. Finally, to distinguish thraustochytrids from other similar-sized photosynthetic protists, natural chlorophyll fluorescence was checked under UV excitation light: the photosynthetic protists were easily distinguished because of their peculiar red fluorescing cell content (Raghukumar & Schaumann 1993).

Independent enumeration for ciliates and thraustochytrids was carried out on randomly selected half-fractions of the polycarbonate membrane. Membranes were scanned at 100 200 and 400 x. The method was tested for efficiency on sterilized sand (collected in the sampling area) onto which known densities of pine-pollen cultured thraustochytrids (Thraustochytrium motivum, Schizochytrium octosporeum) isolated in the same area (Bongiorni 1999) and lab cultured ciliates (Euplotes crassus, Euplotes minuta) had been dispersed. It is worth mentioning that the Euplotes species may be more resistant to this procedure than other ciliates (therefore yielding higher extraction efficiency) and that some ciliates (e.g. long, contractile Karyorelicida) are not well fixed with glutaraldehyde-seawater solution. Moreover, many ciliates cannot be identified at high taxonomic level by this method.

RESULTS AND DISCUSSION

Extraction efficiency was checked by examining sterilized sand samples onto which known densities of cultured thraustochytrids and ciliates had been dispersed. A nearly complete recovery of both protists was achieved.

As the method allows for sample preservation, we were able to examine a high number of replicated samples and follow an ANOVA sampling design in which different spatial scales (sampling units of different size) were included. The abundance values for thraustochytrids are reported in Fig. 3A. Abundance in field samples ranged between 10 and 170 cells ml⁻¹ and the average was 42.4 ± 35.2 cells ml⁻¹, which according to previous estimations (Naganuma et al.
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1998, Kimura et al. 1999) corresponds to the biovolume of 22,200 µm³ ml⁻¹ and an average biomass of 7000 pg C ml⁻¹. Average October, November and January abundances were, respectively, 59.3 ± 41.9, 45.6 ± 26.1 and 21.9 ± 25.2 cells ml⁻¹. On the basis of nested ANOVA, no significant difference in thraustochytrid abundance exists between either the different spatial scales or time of sampling (Table 1). The main variance components (hereafter expressed as % of the overall variance) were those of residual (56.87%) and time (24.67%). The variance components of all 3 spatial factors, mainly those of plot and subplot, were extremely low (Table 1).

The abundances of ciliates in the different samples are reported in Fig. 3B. Their average abundance was 44 ± 26 cells ml⁻¹, ranging between 10 and 130 cells ml⁻¹. Such abundances are comparable with those previously assessed in Mediterranean sandy shores of the same geographic area during autumn-winter, which is characterized by a remarkably lower ciliate density than spring-summer and by a random distribution of these protists (Lucchesi & Santangelo 1995). The average estimated biovolume of ciliates was 9 × 10⁵ µm³ ml⁻¹, corresponding to a biomass of 13 × 10⁴ pg C ml⁻¹ (Putt & Stoeckter 1989). On the basis of multifactorial ANOVA (Table 2), time was the only significant factor (p = 0.014), due to a remarkable decrease in abundance from autumn to winter (October 64.2 ± 30.3, November 45.9 ± 15.6 and January 21 ± 11.1 cells

Table 1. Multifactorial ANOVA calculated for thraustochytrid densities. Nested factors: Square (S), Plot (P), Subplot; orthogonal, random factor: Time (T) with 3 levels. No factor was significant. The main variance components (%) are those of residual and time. Cochran test: not significant (no data transformation)

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<tr>
<th>Source of variation</th>
<th>df</th>
<th>SS</th>
<th>F</th>
<th>p</th>
<th>F vs</th>
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<tr>
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<td>Total</td>
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Table 2. Multifactorial ANOVA calculated for ciliate densities. Nested factors: Square (S), Plot (P), Subplot; orthogonal and random factor: Time (T) with 3 levels. The only significant factor was Time. The main variance components (%) are those of residual and time. Cochran test: significant. Data transformation: logE (x + 1); C = 0.139, p > 0.05

<table>
<thead>
<tr>
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<th>F</th>
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found to contain high levels of polyunsaturated fatty acids (PUFAs) (Kimura et al. 1999). Thraustochytrids should provide a further source of essential fatty acids, which are not synthesized by bacteria nor by animals, to their consumers, such as amoebae, ciliates, filter feeders, detritivores, and larvae of invertebrates and vertebrates (Ulken 1981, Raghukumar 1986, Raghukumar & Balasubramian 1991, Zhukova & Kharlamenko 1999). This additional PUFAs source may be extremely important in the microbial loop.

No variability in abundance was found at any of the different spatial scales explored for either protist, and the main variance component was residual. In other words, at that time and at the spatial scales examined, no spatial structure was found in the sampled area and the major differences in density were those between 1 ml replicated sand samples, regardless of the distances between which they were collected. Such a finding could be due to the time at which samples were collected: during autumn and winter, hydrodynamic disturbance may inhibit patchiness of protists (Santangelo & Lucchesi 1995). Clearly, protists could show some patchiness at a spatial scale smaller than those examined.

Some differences in the trend over time of ciliate and thraustochytrid abundance were found. In contrast with thraustochytrids, ciliates showed a significant decrease from autumn to winter, varying with temperature. Ciliates are thought to be predators of thraustochytrids (Fenchel 1980, Raghukumar 1996, Raghukumar & Balasubramian 1991, Bongiorni 1999). As some degree of correlation between the densities of the 2 protists was found in the field, this hypothesis is strengthened.

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