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Live benthic foraminiferal faunas from the Bay of Biscay: faunal density, composition, and microhabitats

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Abstract

In the meso-oligotrophic Bay of Biscay, a diminishing downward organic matter flux with depth is accompanied by an important decrease of the live foraminiferal density. Although bottom water oxygenation is not directly influenced by organic matter input, the oxygenation of interstitial waters and the primary redox fronts do change in response to variations of the organic matter flux. The occurrence of deep and intermediate infaunal taxa can be linked to fundamental redox fronts and putative associated bacterial consortia. Our data are in agreement with the TROX-model, which explains the benthic foraminiferal microhabitat as a function of organic flux and benthic ecosystem oxygenation. Both the depth of the principle redox fronts and the microhabitat of deep infaunal species show important increases with depth. At the deepest oligotrophic stations, deep infaunal faunas become relatively poor. Therefore, the exported flux of organic matter appears to be the main parameter controlling the composition and the vertical distribution of benthic foraminiferal faunas below the sediment-water interface. The oxygenation of pore waters plays only a minor role. A species-level adaptation of the TROX-model is presented for the Bay of Biscay. © 2002 Elsevier Science Ltd. All rights reserved.

Keywords: Live benthic foraminifera; Exported organic matter flux; Redox conditions; Microhabitat

1. Introduction

Benthic foraminifera are an important component of the meiofaunal community of deep-sea detritus feeders. In deep-sea environments, they commonly represent more than 50% of the total biomass (Gooday et al., 1992). Thanks to their extraordinary potential of adaptation, benthic

foraminifera are able to survive and proliferate in a wide range of marine environments, including extreme ecosystems, such as oligotrophic abyssal plains (Tietjen, 1971; Coull et al., 1977) or hydrothermal vents (Sen Gupta and Aharon, 1994). Because of their potentially important role in deep ocean environments, they are at present studied intensively for a better understanding of their role in the benthic ecosystem and for a more precise definition of their contribution to the recycling of organic matter at the ocean floor.

There is a general consensus that the faunal composition of heterotrophic benthic foraminiferal faunas is strongly linked to the quantity and

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quality of the organic detritus reaching the ocean floor, and to the oxygenation at the sediment-water interface and of the interstitial waters in the first cm of the sediment (e.g. Van der Zwaan, 1982; Altenbach and Sarnthein, 1989; Loubere et al., 1993; Fariduddin and Loubere, 1997; Jorissen et al., 1998). In this context, this paper describes the variability of live (Rose Bengal stained) benthic foraminiferal assemblages along a transect in the Bay of Biscay comprising five stations from 140 to 2000 m water depth (Table 1, Fig. 1). Since 1997, a number of stations in this depth range have been sampled periodically in order to follow the temporal and spatial succession of the benthic foraminiferal faunas.

The Bay of Biscay is a typical temperate meso-oligotrophic environment with two annual bloom periods. The first one, in late winter or early spring, is associated with a shallowing of the mixed layer and with a strengthening of the rather shallow thermocline. Moreover, internal waves associated with strong spring neap tides cause nutrient injection into the photic zone and subsequent phytoplankton blooms at the shelf break in the northern part of the basin (Pingree et al., 1986). The second main bloom event, less marked than the first one, is commonly recorded in autumn, when the summer thermocline starts to be eroded by a deepening of mixing. The advantage of a study of benthic foraminiferal ecosystems in this mesotrophic-oligotrophic context is the fact that the oxygenation at the sediment-water interface is not seriously influenced by the seasonal variability of the organic matter input. The bottom water oxygen concentrations at the five stations are always relatively high (Table

1), and the seasonal monitoring of the oxygenation indicates that the concentrations do not vary significantly throughout the year (Anschutz et al., 1999; Hyacinthe et al., 2001). Therefore, ecosystem variability is probably caused mainly by changes in the organic flux reaching the ocean bottom. The downward organic flux varies with water depth and in response to the temporal and spatial oscillations of primary production in the surface waters. The five stations discussed in this paper were all sampled in late autumn-early winter (of the years 1997 and 1998; Table 1). The stations were selected in order to better understand the influence of the spatially variable organic matter flux reaching the ocean bottom on benthic foraminiferal faunas in an open slope setting. Our study focuses on the faunal density and composition and on foraminiferal microhabitats. The microhabitat is defined as the vertical distribution of a taxon in the first cms of the sediment, which is controlled by the composite action of all physical, chemical and biological processes (Corliss, 1985). Understanding the foraminiferal microhabitat is important, because it allows the precise trophic and oxic requirements of each species in the total live benthic foraminiferal assemblage to be known.

The main objectives of this paper are (1) to present and discuss variation of the foraminiferal density with increasing water depth and thus with calculated diminishing organic fluxes, (2) to better explain the compositional changes observed along the five stations bathymetrical/trophic transect, and (3) to explain the microhabitat changes in response to the trophic conditions at the sediment-water interface.

Table 1

Water depth, sampling date, geographical position, bottom water oxygen and depth in the sediment of the zero oxygen level for the five stations used in this paper

Station	Sample	Depth (m)	Date	Latitude	Longitude	Oxygenation (ml/l)	Zero O ₂ boundary (mm)
D	OB1D	140	October 1997	43°41'93N	1°34'10W	4.90	8
B	OB1B	553	October 1997	43°49'98N	2°23'04W	4.80	17
A	OB1A	1012	October 1997	44°09'08N	2°20'27W	4.36	20
F	OB2F	1264	Februar 1998	44°17'10N	2°44'95W	4.70	64
H	OB5H	1993	October 1998	44°32'00N	2°37'00W	5.85	63

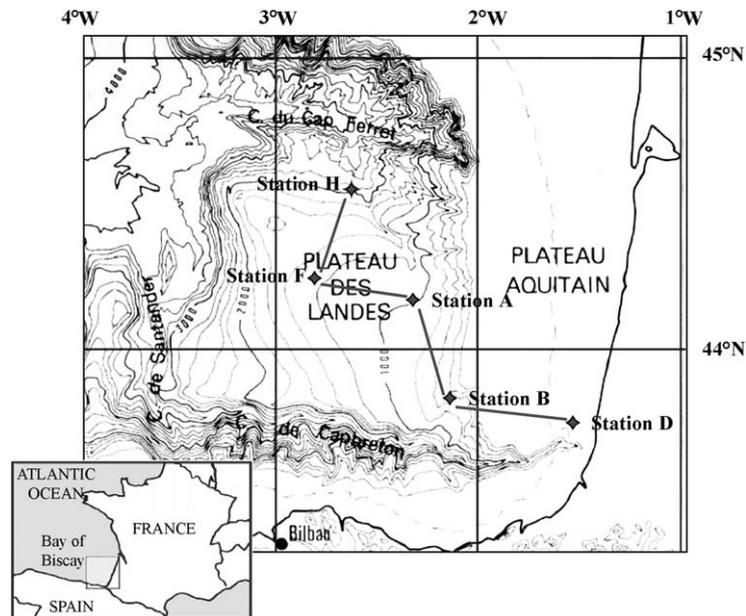


Fig. 1. Study area, bathymetry and geographical position of the five stations.

2. Study area, material and methods

The Bay of Biscay is a semi-enclosed basin at the eastern side of the North Atlantic Ocean, bathed by rather homogeneous oceanic waters belonging to the North Atlantic thermohaline and geostrophic circulation and more precisely, to the North Atlantic Drift. The Bay of Biscay is bordered by the Irish shelf in the north, the Armorican and Aquitaine shelves in the east, and by the Iberian shelf in the south. The hydrographical structure is relatively well known (Ogawa and Tuzin, 1973). The patterns of the surface waters are strongly constrained by seasonal variations of the thermocline and the mixed layer. Below the surface waters (≤ 150 m depth), the Northern Atlantic Central Waters (NACW) are present down to 800 m depth. Between 800 and 1200 m, a branch of Mediterranean outflow waters is present. In comparison with the surrounding water masses, these Mediterranean Waters (MW) are characterised by a high salinity (35.80–35.85, Le Floch, 1968). They have the lowest oxygen values for the Bay of Biscay (3.8 ml/l, Le Floch, 1968; 4.36 ml/l, this paper). Below the MW,

Intermediate Atlantic Water and Polar Atlantic Water, both belonging to Northern Atlantic Deep Waters, fill the deepest parts of the basin. Within the Bay of Biscay, exchanges between the deeper parts and the coastal waters are rather insignificant, especially in spring and summer when a dome of cold and dense water on the continental shelves limits advection of sub-surface water (Ogawa and Tuzin, 1973).

The continental slope bordering the French shelf deepens gradually, and is interrupted by two large canyons (Cap Ferret Canyon, Cap Breton Canyon, Fig. 1). Vertical fluxes represent the main sedimentary component in open slope environments, whereas lateral advection may dominate sedimentary processes in the canyons (Heussner et al., 1999). The present paper concentrates on the open-slope environments, where the impact of laterally advected particles is supposed to be minimal, and where the linkage between surface water primary production, the vertical particle flux, and benthic life should be rather straightforward.

Few data are available on primary production in general or on the occurrences of bloom events in

the Bay of Biscay. Tréguer et al. (1979) estimate a production between 0.4 and 1.9 g C/m²/day for the spring bloom of 1973. Measurements of primary production in the central part of the Bay of Biscay during the autumn bloom of 1972 yielded values between 0.3 and 0.4 g C/m²/day (Le Corre and Treguer, 1976). These values agree with recent data obtained in the Cap Ferret region during five ECOFER cruises: from 0.7 to 1.2 g C/m²/day in spring (May 1990 and 1991) and 0.3 g C/m²/day in October 1990 (Laborde et al., 1999). Laborde et al. (1999) estimate total annual primary production between 145 and 170 g C/m²/yr. Generally, phytoplankton blooms in the Bay of Biscay are composed of diatoms (mainly *Chaetoceros* spp. and *Nitzschia* spp.) and coccolithophorids (*Gephyrocapsa oceanica*, *Emiliania huxleyi*; Tréguer et al., 1979; Fernandez et al., 1995). Fernandez (1990), however, shows that in the Central Cantabrian Sea (southern part of the Bay of Biscay) high chlorophyll-*a* and primary production rates observed in March consist predominantly of microflagellates. Moreover, in the Celtic Sea, coccolithophore blooms (*Emiliania huxleyi*) may also occur during summer (Fernandez et al., 1993). Finally, Pingree et al. (1986) demonstrate a chlorophyll increase over the shelf break of the northern French shelf, due to internal and tidal waves; these rather exceptional and short bloom events occur during neap tides in spring.

In cases where the particle flux is mainly vertical, and where we have a quantitative estimate of the surface water primary production, we can use the formula proposed by Berger and Wefer (1990), and improved by Herguera (1992), to estimate the annual organic matter flux J_z that reaches the ocean bottom at different water depths:

$$J_z = (2\sqrt{PP(PP/z)}) + ((5/\sqrt{PP})(PP/\sqrt{z})),$$

where PP is the primary production, in g C/m²/yr, z the water depth (in m), J_z the organic flux at a water depth of z meters, in g C/m²/yr.

This formula differentiates between a labile component (first term), which is supposed to represent easily metabolisable organic detritus, and a more refractory component (second term), which decreases much more slowly with depth. We

assume that benthic life will be affected mainly by the first term. When we use a mean primary production value of 150 g C/m²/yr, for all stations, irrespective of water depth and distance from shore, and when we consider only the vertical transport of particles, the total exported flux at the shallowest station (D, 140 m) can be estimated at 31.4 g C/m²/yr, with a labile fraction of 26.6 g C/m²/yr. At the deepest station (H, 1993 m) the estimated total exported flux is 3.2 g C/m²/yr, with a labile component of 1.8 mg C/m²/yr. Estimated total flux values for the intermediate stations B (553 m), A (1012 m) and F (1264 m) are 9.2, 5.6 and 4.6 g C/m²/yr, respectively; whereas the estimates for the labile component are 6.6, 3.6 and 2.9 g C/m²/yr, respectively. These values suggest that the labile organic carbon flux, which should represent a readily available food source for the benthic fauna, is about 15 times higher in the shallowest station D than in the deepest station H.

The five stations D, B, A, F, and H have been selected to compose a rather ideal SE-NW transect, ranging from the outer shelf to bathyal open slope environments (Fig. 1; Table 1). Station D is situated at 140 m depth, at the diffusive boundary between the surface waters and the NACW. According to Ogawa and Tauzin (1973), the NACW and the basal part of surface waters can be characterised by a seasonally constant temperature of 11.9°C and an invariant salinity of 35.60. The temperature and salinity of more superficial surface waters (0–100 m depth) are strongly influenced by the seasonality of mixing, thermocline depth, and continental runoff. The sandy mud deposits found at station D are typical of the sedimentary conditions found in outer shelf environments.

Station B is positioned at 553 m depth. This site represents a mid-slope environment, isolated from possible lateral input of sediment (turbidites and slumps), which could occur in this depth range in areas closer to the two main canyon systems. Station B is positioned within the NACW, which has a salinity of 35.60 and a temperature of about 11°C (Ogawa and Tauzin, 1973). The sediment consists predominantly of fine-grained silty mud.

Station A (1012 m depth) was selected for its strategic position inside the branch of Mediterranean outflow waters (MW). This water mass has a temperature of about 10.5°C and a salinity of 35.80 (Le Floch, 1968; Ogawa and Tauzin, 1973). Sediments at this depth are predominantly silty muds.

Station F, at 1264 m deep, is positioned in the upper layers of Northern Atlantic Deep Waters (Intermediate Atlantic Waters), commonly characterising the bottom waters of the Celtic Sea (8°C, 35.65, Ogawa and Tauzin, 1973). The sediments are silty muds.

Station H, at 1993 m depth, is typical for benthic environments found in the Northern Atlantic Deep Waters. Although this station is geographically rather close to the Cap Ferret Canyon, it is still an open slope environment with muddy sediments. The water temperature is about 4°C and the salinity is about 35.00 (Ogawa and Tauzin, 1973).

All cores were sampled with a classic Barnett multi-tube corer (Barnett et al., 1984). Each tube has a surface area of about 72 cm², and accordingly, all our faunal density data are standardised for this surface area. The multi-corer allows sampling of the first dm of the sediment, the overlying bottom waters, and an undisturbed sediment-water interface. Free waters were collected immediately after core recovery for dissolved O₂ measurements by the Winkler titration method (Strickland and Parsons, 1972). Profiles of pore water O₂ were obtained on board with a cathode-type mini-electrode (Revsbech and Jørgensen, 1986; Helder and Bakker, 1985; Revsbech, 1983). The temperature was maintained by using an insulating device. This operation was completed in duplicate within 30 min after core recovery. Subsequently, the core used for O₂ profiling was sliced in thin horizontal sections (every 0.5 cm for the top 2 cm, 1 or 2 cm below) within 1½ h. For every level a sub-sample was immediately sealed in a pre-weighed vial and frozen under inert atmosphere (N₂) for further analyses of porosity and chemistry of the solid fraction. Another sub-sample was centrifuged under N₂ at 5000 rpm for 20 min for collection pore waters. Two aliquots of water were filtered

(0.2 µm) and frozen at –25°C for nutrient analyses. In the laboratory, porosity was determined by comparison of the weights of wet and freeze-dried sediment. Interstitial water compounds were analysed by techniques adapted for small volumes (Anschutz et al., 1999; Hyacinthe et al., 2001). Nitrate and nitrite were measured by flow injection analysis (FIA) according to Anderson (1979). Sulphate was measured by a nephelometric method (Stookey, 1970).

For faunal analysis, one entire core was sliced horizontally for each station; every 0.25 cm for the first cm of sediment, every half cm between 1 and 4 cm depth, and every cm between 4 and 10 cm. Exceptionally, for station A, the core was cut into 0.5 cm slices between 1 and 4.5 cm, and, for station D, the first cm was cut into 3 slices (0–0.35, 0.35–0.75 and 0.75–1.0 cm). Sediments were stored in 500 cm³ bottles, which were filled with 95% ethanol containing 1 g/l Rose Bengal stain. Rose Bengal staining is commonly used to identify live foraminifera (Walton, 1952). All samples were gently shaken for several minutes in order to get a homogeneous mixture. In the laboratory, they were sieved through 63 and 150 µm mesh screens, and the sieve residues were stored in 95% ethanol. Stained foraminifera belonging to the 150 µm fraction were sorted from wet samples, and stored in Chapman slides. The 63–150 µm fraction was preserved for future studies. The interpretation of staining of benthic foraminifera is rather subjective. One problem of this technique is the fact that Rose Bengal may stain the protoplasm of dead foraminifera, which may be relatively well preserved for a considerable period of time under the anoxic conditions that generally prevail deep in the sediment (Bernhard, 1988; Corliss and Emerson, 1990). As a consequence, a strict application of the staining criteria is most times easy in superficial samples, but may become more critical in the deeper levels. In all cases, we applied our staining criteria (all chambers except the last one stained brightly pink) very strictly, and compared doubtful individuals with perfectly stained individuals of the same species found in superficial sediment layers. Non-transparent agglutinated and miliolid taxa were broken on many occasions for inspection of the interior of the test. We tried to

identify most of live foraminifera at specific level. We used taxonomic references which are presented in Appendix A. Fragments of branch-like agglutinating foraminifera (such as *Hyperamina*) and *Glomospira charoides* and *Glomospira gordialis* (which, because of the orange–reddish colour of their test are very difficult to confirm as living), were not included in the quantitative analyses.

In order to describe the vertical distribution of the total faunas or individual taxa, we use the average living depth (ALD, Jorissen et al., 1995), which allows a rapid description of the microhabitat patterns. The ALD is calculated with the following formula:

$$ALD_x = \sum_{i=0,x} (n_i D_i) / N,$$

where x is the lower boundary of deepest sample, n_i the number of individuals in interval i , D_i the midpoint of sample interval i , N the total number of individuals for all levels.

For all stations, ALD_{10} was calculated for the whole fauna as well as for individual taxa, on the basis of the numbers of stained individuals found in the successive sediment slices. Isolated individuals separated from the main population by more than 1 cm of “sterile” sediment (without live individuals of the studied taxa) were not integrated in the calculations of the ALD_{10} . We suppose that such isolated individuals had been transported downward (outside their normal microhabitat) by bioturbation, or correspond to dead organisms that have been counted erroneously. In the data sheets (Tables 2a–e), the latter individuals are indicated in brackets.

After the first classification with four main microhabitats proposed by Corliss and Chen (1988), it was argued that only species living on elevated substrates can be considered as “epifaunal” (Buzas et al., 1993). Therefore, in the soft bottom communities described in this study, we recognise only three different groups: shallow infaunal, intermediate infaunal and deep infaunal species.

3. Results

The total density of the live foraminiferal fauna were determined by integrating the numbers of live individuals picked in all levels from 0 to 10 cm depth. It is expressed as the number of live foraminifera found in and below a 72 cm² surface area (Fig. 2). Concerning vertical profiles, specific foraminiferal densities were normalised for each layer to a 50 cm³ sediment volume; we generally regrouped the four uppermost slices of each core into two 0.5 cm thick samples. The percentages of the various taxa were calculated on the basis of the non normalised densities.

Fig. 2, which presents the foraminiferal densities for the five stations, shows a clear negative correlation between the foraminiferal density and water depth. The shallowest station D, with about 2000 live individuals collected, presents a maximum value, whereas stations H and F, with 179 and 122 individuals, respectively, are much poorer.

At all five stations, perforate foraminifera form the main component of the benthic foraminiferal faunas (75–90%; Fig. 2). At station B, they compose almost 90% of the total fauna. Agglutinated taxa account for 10–20% of the total fauna, whereas miliolids are rare in all stations (between 0.3% and 8.2%). The latter group tends to be richer on the middle and lower slope (between 4.2% and 8.2%) than on the outer shelf and upper slope, where they are almost absent.

Species diversity is highest at station B where about 50 species are found. Station D, with 36 species, is slightly less diverse. Station A contains 41 species; at stations F and H, where agglutinated and porcellaneous taxa represent more than half of all species, 25 and 28 species are found, respectively.

Bottom waters at station D (140 m) have an oxygen concentration of 4.9 ml/l. Within the sediment, there is a rapid decrease, until anoxic conditions are reached at 0.8 cm depth (Fig. 3a). The nitrate + nitrite profile shows a sharp decrease in the first mm of the sediment, suggesting that also the zone of nitrate reduction is extremely close to the sediment–water interface. Sulphate reduction is chemically detected below 5 cm depth, and corresponds to an increase of particulate sulphur

and a decrease of dissolved sulphate. This is the only station where sulphate reduction is significant. Station D, which has the highest foraminiferal density of all five stations (1989 individuals/72 cm²), presents a faunal assemblage that is strongly concentrated in the uppermost cm of the sediment (Fig. 3b). The highest density is recorded in the superficial sample (0.00–0.35 cm), where a value of 1400 live individuals/50 cm³ has been calculated. This value falls abruptly to about 350 individuals/50 cm³ in the 0.75–1.0 cm level, where the oxygen concentration is already close to zero. About 350 live foraminifera/50 cm³ can still be noted in the 1.0–1.5 cm level, in a completely anoxic environment. The ALD₁₀ of the total live fauna is 1.1 cm. The fauna is dominated by *Chilostomella oolina* (30.2%), *Valvulinera bradyana* (17.0%), *Clavulina cylindrica* (15.0%), *Nonion scaphum* (13.0%), *Bolivina subaenariensis* (3.1%), *Hyalinea balthica* (2.7%) and *Bulimina marginata* (2.3%) (Table 2a). *V. bradyana* (ALD₁₀=0.4 cm), *C. cylindrica* (ALD₁₀=0.4 cm), and *C. oolina* (ALD₁₀=1.2 cm) are the dominant taxa in the well-oxygenated first half cm (Fig. 3c). They are accompanied there by less frequent species such as *Bolivina alata*, *B. subaenariensis*, *Uvigerina peregrina* and *Rectuvigerina phlegeri*. *C. oolina* (ALD₁₀=1.2 cm), and *N. scaphum* (ALD₁₀=1.8 cm) dominate the faunas in the anoxic zone from 1 to 3 cm depth. Some rare individuals of *Bulimina aculeata* and *B. marginata* are found in the totally anoxic environments below 3 cm depth.

At station B (water depth 553 m), the bottom water oxygen concentration is 4.8 ml/l. The zero oxygen level is found at about 2 cm depth (Fig. 4a). Well-oxygenated conditions prevail in the first half cm. A downward diffusive zone of nitrate and nitrite extends from 0.25 to 2.25 cm depth. Fig. 4b shows that maximal densities (between 800 and 650 individuals/50 cm³) are found in the top first cm. Faunal numbers drastically drop to about 40 individuals/0.5 cm level at 1.75 cm depth. The ALD₁₀ of the total live fauna is 0.8 cm. The benthic foraminiferal fauna is strongly dominated by *Uvigerina mediterranea* (33.8%), *Uvigerina peregrina* (21.6%), *Melonis barleeanus* (9.0%), *Globobulimina affinis* (6.0%) and *Cibicidoides pachydermus* (4.7%) (Table 2b).

U. mediterranea (ALD₁₀=0.6 cm), accompanied by *C. pachydermus* (ALD₁₀=0.6 cm) and *U. peregrina* (ALD₁₀=0.8 cm), dominate the superficial sediments (Fig. 4c). Slightly deeper, between 0.5 and 1.5 cm depth, *U. peregrina* (ALD₁₀=0.8 cm) and *M. barleeanus* (ALD₁₀=0.8 cm) show maximum frequencies. *G. affinis* (ALD₁₀=2.4 cm) thrives preferentially in the deeper part of the core. It increases below 1.0 cm depth and becomes the only remaining dominant taxon below 2.25 and 6.5 cm depth.

At station A (water depth 1012 m), the oxygen concentration is 4.36 ml/l. The zero oxygen level is found at about 2.0 cm depth, whereas a nitrate + nitrite downward diffusive zone extends from 0.75 to 3.5 cm depth (Fig. 5a). Fig. 5b depicts a bimodal distribution of the benthic foraminiferal fauna. A first density maximum (180 individuals/50 cm³) is recorded around 1 cm depth, a second one (40 individuals/50 cm³) occurs deeper in the sediment, at about 6 cm depth. The ALD₁₀ of the total live fauna is 2.4 cm. *Globobulimina affinis* (28.8%), *Uvigerina peregrina* (17.9%), *Uvigerina mediterranea* (9.5%), *Hoeglundina elegans* (5.5%), *Nuttallides umboniferus* (5.3%), *Cibicidoides pachydermus* (1.9%) and *Melonis barleeanus* (1.3%) are the dominant taxa of station A (Table 2c). *N. umboniferus* (ALD₁₀=0.4 cm), *U. mediterranea* (ALD₁₀=0.6 cm), *C. pachydermus* (ALD₁₀=0.3 cm) and *H. elegans* (ALD₁₀=0.8 cm) constitute a rich superficial fauna, which is present in the well-oxygenated first cm (Fig. 5c). *U. peregrina* (ALD₁₀=1.1 cm), accompanied by *M. barleeanus* (ALD₁₀=1.7 cm), occupies slightly deeper layers and is the dominant component of the shallowest density peak. *G. affinis* (ALD₁₀=5.3 cm), accompanied by some individuals of *Glandulina ovula* (ALD₁₀=5.0 cm) and *C. oolina* (ALD₁₀=5.5 cm), dominates the rich stained fauna below 2 cm depth.

The oxygen concentration of the bottom waters at station F (water depth 1264 m) is 4.70 ml/l. Within the sediment, the zero oxygen level is reached at 6.4 cm depth (Fig. 6a). The dissolved oxygen curve shows some minor oscillations that may be the result of microenvironments caused by burrows. A nitrate diffusive gradient extends from 4.5 to 7.5 cm depth. The benthic foraminiferal

Table 2
Benthic foraminiferal census data for stations D, B, A, F and H. N.B. Numbers are not standardized for a sediment volume

	Depth															Total	ALD ₁₀	% of total fauna
	0–0.35	0.35–0.75	0.75–1	1–1.5	1.5–2	2–2.5	2.5–3	3–3.5	3.5–4	4–5	5–6	6–7	7–8	8–9	9–10			
(a) Station D 140 m																		
Perforate																		
<i>Amphicoryna scalaris</i>	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0.6	0.1
<i>Bolivina alata</i>	26	10	1	0	0	0	0	0	0	0	0	0	0	0	0	37	0.3	1.9
<i>Bolivina subaenariensis</i>	38	21	1	1	0	0	0	0	0	0	0	0	0	0	61	0.3	3.1	
<i>Bulimina aculeata</i>	1	1	0	0	2	4	2	2	2	5	3	5	4	3	34	4.7	1.7	
<i>Bulimina inflata</i>	1	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0.2	0.1	
<i>Bulimina marginata</i>	2	0	0	2	3	6	7	3	4	4	5	3	5	2	46	4.2	2.3	
<i>Canceris auriculus</i>	6	5	0	0	0	0	0	0	0	0	0	0	0	0	16	0.3	0.8	
<i>Cassidulina carinata</i>	2	3	0	1	1	0	0	0	0	0	0	0	0	0	7	0.7	0.4	
<i>Chilostomella oolina</i>	91	125	59	135	98	58	18	7	4	2	1	2	0	0	600	1.2	30.2	
<i>Coryphostoma</i> sp.	6	1	0	0	0	0	0	0	0	0	0	0	0	0	7	0.2	0.4	
<i>Dentalina</i> sp.	2	1	0	(1)	0	0	0	0	0	0	0	0	0	0	4	0.3	0.2	
<i>Dentalina ariena</i>	4	0	0	(1)	0	0	0	0	0	0	0	0	0	0	5	0.2	0.3	
<i>Elphidium advenum</i>	0	0	0	0	0	0	0	0	0	1	0	0	0	0	1	4.5	0.1	
<i>Gavelinopsis translucens</i>	0	0	0	0	0	0	0	0	1	0	0	0	0	0	1	3.8	0.1	
<i>Globobulimina affinis</i>	0	4	0	1	2	3	0	(1)	0	0	0	0	0	(1)	12	2.2	0.6	
<i>Hoeglundina elegans</i>	2	0	0	0	(1)	0	0	0	0	0	0	0	0	0	3	0.2	0.2	
<i>Hyalinea balthica</i>	14	8	1	6	0	0	0	0	0	0	0	0	0	0	53	1.6	2.7	
<i>Lenticulina peregrina</i>	1	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0.2	0.1	
<i>Nonion scaphum</i>	20	24	24	63	32	36	18	8	10	8	4	5	0	(2)	(4)	258	1.8	13.0
<i>Nonionella turgida</i>	0	0	0	1	0	1	0	(1)	0	0	0	0	0	0	3	1.8	0.2	
<i>Pseudoepionides falsobeccarii</i>	5	1	0	4	1	3	0	0	3	1	2	3	0	1	25	3.2	1.3	
<i>Rectawigerina phlegeri</i>	16	7	1	0	2	1	0	0	(1)	(1)	0	0	0	0	29	0.5	1.5	
<i>Trifarina pauperata</i>	0	0	0	0	0	1	0	0	0	0	0	0	0	0	1	2.3	0.1	
<i>Uvigerina peregrina</i>	27	7	0	0	(2)	(2)	0	0	0	(1)	0	0	0	0	39	0.4	2.0	
<i>Valvulineria bradyana</i>	196	102	25	12	1	1	2	0	0	0	0	0	0	0	339	0.4	17.0	
Total perforate	460	321	112	228	154	120	56	24	26	23	16	20	10	9	5	1584	1.3	79.6
No. species	19	17	7	14	12	12	8	8	8	8	7	8	5	5	4	25		
Porcellaneous																		
<i>Quinqueloculina seminula</i>	26	5	1	2	5	3	2	1	0	0	0	0	0	0	45	0.8	2.3	
<i>Sigmoilina</i> sp.	0	0	0	0	0	1	0	0	0	0	0	0	0	0	1	2.3	0.1	
Total porcellaneous	26	5	1	2	5	4	2	1	0	0	0	0	0	0	46	0.8	2.3	
No. species	1	1	1	1	1	2	1	1	0	0	0	0	0	0	2			

Non fossilizing agglutinated																		
<i>Clavulina cylindrica</i>	208	60	9	11	4	4	0	0	(1)	(1)	0	0	0	0	0	298	0.4	15.0
<i>Eggerella</i> sp.	2	1	3	4	0	(6)	0	0	0	(1)	0	0	0	0	0	18	0.5	0.9
<i>Haplophragmoides</i> sp.	5	2	0	1	0	0	0	0	0	0	0	0	0	0	0	8	0.4	0.4
<i>Nouria polymorphinoides</i>	11	3	1	1	0	0	0	0	0	0	0	0	0	0	0	16	0.4	0.6
<i>Reophax</i> sp.	4	2	0	0	0	0	0	0	0	0	0	0	0	0	0	6	0.3	0.3
<i>Reophax ampullacea</i>	0	0	0	1	1	0	0	0	0	0	0	0	0	0	0	2	1.5	0.1
<i>Reophax scorpiurus</i>	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0.2	0.1
Total agglutinated	231	68	13	18	5	10	0	0	1	2	0	0	0	0	0	349	0.4	17.5
No. species	10	11	7	6	4	3	0	0	0	0	0	0	0	0	0	7		
Fossilizing agglutinated																		
<i>Textularia agglutinans</i>	0	0	0	1	0	(1)	0	0	0	0	0	0	0	0	0	2	0.6	0.1
<i>Textularia sagittula</i>	1	0	0	(1)	(1)	0	0	(2)	(1)	(2)	0	0	0	0	0	8	0.0	0.4
Total agglutinated	1	0	0	2	1	1	0	2	1	2	0	0	0	0	0	10	0.7	0.5
No. species	1	0	0	2	1	1	0	1	1	1	0	0	0	0	0	2		
Total live foraminifera	710	386	125	254	160	143	63	28	32	45	28	31	29	15	21	1989	1.1	100.0
No. species	30	29	15	21	17	17	9	9	8	8	7	8	5	5	4	36		

Depth

	0–0.25	0.25–0.5	0.5–0.75	0.75–1	1–1.5	1.5–2	2–2.5	2.5–3	3–3.5	3.5–4	4–5	5–6	6–7	7–8	8–9	9–10	Total	ALD ₁₀	% of total fauna
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(b) Station B 553 m

Perforate

Indet	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0.1	0.1
<i>Amphicoryna scalaris</i>	1	0	1	0	1	1	0	0	0	0	0	0	0	0	0	0	4	0.9	0.3
<i>Bolivina alata</i>	0	0	1	2	2	1	0	0	0	0	0	0	0	0	0	0	6	1.1	0.4
<i>Bolivinita quadrilatera</i>	5	2	0	0	0	0	0	0	0	(1)	0	0	0	0	0	0	8	0.2	0.6
<i>Bulimina marginata</i>	5	4	0	1	1	3	1	4	0	0	0	0	0	0	0	0	19	1.2	1.4
<i>Chilostomella oolina</i>	0	0	0	2	0	0	0	0	1	0	0	0	0	0	0	0	3	1.7	0.2
<i>Cibicides lobatulus</i>	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0.6	0.1
<i>Cibicides wuellerstorfi</i>	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0.6	0.1
<i>Cibicoides pachydermus</i>	26	16	6	3	3	6	1	2	0	0	0	0	0	0	0	0	63	0.6	4.7
<i>Dentalina</i> sp.	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0.1	0.1
<i>Elphidium</i> sp.	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0.4	0.1
<i>Gavelinopsis translucens</i>	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0.1	0.1
<i>Glandulina ovula</i>	1	2	0	0	0	0	0	0	0	0	0	0	0	0	0	0	3	0.3	0.2
<i>Globobulimina affinis</i>	0	0	1	4	19	6	13	16	10	6	3	3	0	0	(1)	0	81	2.4	6.0
<i>Gyroidina altiformis</i>	7	0	2	0	0	(1)	0	0	0	0	0	0	0	0	0	0	10	0.2	0.7
<i>Gyroidina orbicularis</i>	1	0	0	2	0	0	0	0	0	0	0	0	0	0	0	0	3	0.6	0.2
<i>Hyalinea balthica</i>	7	2	0	0	0	0	(1)	(1)	(1)	0	0	0	0	0	0	0	12	1.1	0.9
<i>Lenticulina gibba</i>	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	1	0.9	0.1

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Table 2 (continued)

	Depth																Total	ALD ₁₀	% of total fauna
	0-0.25	0.25-0.5	0.5-0.75	0.75-1	1-1.5	1.5-2	2-2.5	2.5-3	3-3.5	3.5-4	4-5	5-6	6-7	7-8	8-9	9-10			
<i>Lenticulina peregrina</i>	4	3	0	0	0	0	0	0	0	0	0	0	0	0	0	0	7	0.2	0.5
<i>Lenticulina rotulata</i>	2	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	3	0.4	0.2
<i>Melonis barleeanus</i>	14	33	27	23	14	4	1	1	3	1	0	0	0	0	0	0	121	0.8	9.0
<i>Nodosaria</i> sp.	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0.6	0.1
<i>Nuttallides umboniferus</i>	2	2	1	0	0	0	0	0	0	0	0	(1)	0	0	0	0	6	0.3	0.4
<i>Pseudoeponides falsobeccarii</i>	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	1	3.3	0.1
<i>Robertinoides</i> sp.	3	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	3	0.1	0.2
<i>Siphogenerina</i> sp.	13	7	4	1	0	2	1	1	1	0	0	0	0	0	0	0	30	0.6	2.2
<i>Trifarina angulosa</i>	2	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	3	0.3	0.2
<i>Uvigerina elongatastriata</i>	0	0	5	15	31	3	3	2	0	(1)	0	(1)	0	0	0	0	62	1.2	4.6
<i>Uvigerina mediterranea</i>	119	123	108	54	20	14	4	2	0	(1)	0	(1)	0	0	0	0	454	0.6	33.8
<i>Uvigerina peregrina</i>	30	43	83	78	43	7	3	2	0	0	0	0	0	(1)	0	0	290	0.8	21.6
<i>Uvigerina proboscidea</i>	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0.1	0.1
Total perforate	246	238	243	187	134	48	29	38	19	8	3	6	0	1	1	0	1201	0.8	89.3
No. species	21	12	15	13	9	11	9	10	7	4	1	4	0	1	1	0	33		
Porcellaneous																			
<i>Cruciloculina</i> sp.	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0.1	0.1
<i>Cyclogyra</i> sp.	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0.1	0.1
<i>Pyrgo depressa</i>	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0.1	0.1
<i>Sigmoilina</i> sp.	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0.1	0.1
Total porcellaneous	4	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	4	0.1	0.3
No. species	4	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	4		
Non fossilizing agglutinated																			
<i>Ammolagena</i> sp.	1	3	1	0	0	0	0	0	0	0	0	0	0	0	0	0	5	0.4	0.4
<i>Ammoscalaria</i> sp.	0	1	1	0	0	0	0	0	0	0	0	0	0	0	0	0	2	0.5	0.1
<i>Clavulina cylindrica</i>	0	0	1	6	3	0	0	0	0	0	0	0	0	0	0	0	10	1.0	0.7
<i>Cribrostomoides subglobosum</i>	8	5	4	0	2	1	0	0	0	0	0	0	0	0	0	0	20	0.5	1.5
<i>Cyclammina cancellata</i>	5	4	2	0	0	(1)	(1)	0	0	0	0	0	0	0	0	0	13	0.3	1.0
<i>Eggerella</i> sp.	1	2	1	1	1	0	0	0	0	0	0	0	0	0	0	0	6	0.6	0.4
<i>Haplophragmoides</i> sp.	0	2	0	0	0	0	0	0	0	0	0	0	0	0	0	0	2	0.4	0.1
<i>Reophax</i> sp.	21	19	6	4	0	1	1	0	0	0	0	0	0	0	0	0	52	0.4	3.9
<i>Trochammina inflata</i>	0	0	1	1	0	0	0	0	0	0	0	0	0	0	0	0	2	0.8	0.1
Total agglutinated	36	36	17	12	6	3	2	0	0	0	0	0	0	0	0	0	112	0.5	8.3
No. species	10	14	11	7	6	4	3	0	0	0	0	0	0	0	0	0	9		

Fossilizing agglutinated																				
<i>Bigenerina nodosaria</i>	5	2	1	0	1	0	0	0	0	0	0	0	0	0	0	0	0	9	0.4	0.7
<i>Pseudoclavulina crustata</i>	2	3	0	1	1	2	1	0	0	0	0	0	0	0	0	0	0	10	0.9	0.7
<i>Siphotextularia affinis</i>	4	1	2	0	2	0	0	0	0	0	0	0	0	0	0	0	0	9	0.5	0.7
Total agglutinated	11	6	3	1	4	2	1	0	0	0	0	0	0	0	0	0	0	28	0.6	2.1
No. species	3	3	2	1	2	1	1	0	0	0	0	0	0	0	0	0	0	3		
Total live foraminifera																				
No. species	297	280	263	200	144	53	32	38	19	8	3	6	0	1	1	0	1345	0.8	100.0	
<i>Glomospira</i> spp	1	4	6	4	11	10	11	7	1	1	0	0	0	0	0	0	56	1.6		
Arborescent indet	34	30	44	61	22	12	0	0	0	0	0	0	0	0	0	0	203	0.7		
Depth																				
	0–0.25	0.25–0.5	0.5–0.75	0.75–1	1–1.5	1.5–2	2–2.5	2.5–3	3–3.5	3.5–4	4–4.5	4.5–5.5	5.5–6.5	6.5–7.5	7.5–8.5	8.5–9.5	9.5–10.5	Total	ALD ₁₀	% of total fauna
(c) Station A 1012 m																				
Perforate																				
<i>Bulimina inflata</i>	1	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	2	0.2	0.4
<i>Bulimina marginata</i>	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	1	4.8	0.2
<i>Chilostomella oolina</i>	0	0	0	0	0	0	0	0	0	0	1	1	1	1	0	0	0	4	5.5	0.8
<i>Cibicides pachydermus</i>	4	2	3	0	0	0	0	0	0	0	0	0	0	0	0	0	0	9	0.3	1.9
<i>Glandulina ovula</i>	0	0	1	1	0	0	0	0	1	2	2	0	1	1	2	0	1	12	5.0	2.5
<i>Globobulimina affinis</i>	0	0	0	0	0	0	3	13	8	10	10	17	44	27	5	0	0	137	5.3	28.8
<i>Gyroidina altiformis</i>	1	0	2	0	0	0	0	0	0	0	0	0	0	0	0	0	0	3	0.5	0.6
<i>Gyroidina orbicularis</i>	0	0	0	0	2	0	(1)	0	(1)	0	0	0	0	0	0	0	0	4	1.3	0.8
<i>Hoeglundina elegans</i>	4	4	4	6	7	1	0	0	0	0	0	0	0	0	0	0	0	26	0.8	5.5
<i>Lenticulina</i> sp.	0	0	1	0	1	0	0	0	0	0	0	0	0	0	0	0	0	2	0.9	0.4
<i>Melonis barleeanus</i>	0	0	0	0	3	1	2	0	0	0	0	0	0	0	0	0	0	6	1.7	1.3
<i>Nuttallides umboniferus</i>	8	8	6	2	1	0	0	0	0	0	0	0	0	0	0	0	0	25	0.4	5.3
<i>Pullenia</i> sp.	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	1	2.3	0.2
<i>Siphogenerina</i> sp.	0	0	0	0	0	0	0	0	0	1	0	0	1	0	0	0	0	2	4.9	0.4
<i>Rosalina</i> sp.	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0.1	0.2
<i>Uvigerina mediterranea</i>	5	12	12	12	4	0	0	0	0	0	0	0	0	0	0	0	0	45	0.6	9.5
<i>Uvigerina peregrina</i>	2	1	10	22	37	13	0	0	0	0	0	0	0	0	0	0	0	85	1.1	17.9
Total perforate	26	28	39	43	55	15	7	13	10	13	13	19	47	29	7	0	1	365	2.7	76.7
No. species	9	6	9	5	7	3	4	1	3	3	3	3	4	3	2	0	1	20		
Porcellaneous																				
<i>Biloculinella irregularis</i>	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0.9	0.2
<i>Cyclogyra</i> sp.	2	0	0	1	0	(1)	0	0	0	0	0	0	0	0	0	0	0	4	0.3	0.8
<i>Pyrgo depressa</i>	0	0	2	0	0	0	0	0	0	0	0	0	0	0	0	0	0	2	0.6	0.4

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Table 2 (continued)

	Depth																	Total	ALD ₁₀	% of total fauna
	0–0.25	0.25–0.5	0.5–0.75	0.75–1	1–1.5	1.5–2	2–2.5	2.5–3	3–3.5	3.5–4	4–4.5	4.5–5.5	5.5–6.5	6.5–7.5	7.5–8.5	8.5–9.5	9.5–10.5			
<i>Pyrgo subsphaerica</i>	0	0	1	1	2	0	0	0	0	0	0	0	0	0	0	0	0	4	1.0	0.8
<i>Pyrgoella sphaera</i>	2	1	0	0	0	0	(1)	0	0	0	0	0	0	0	0	0	0	4	0.2	0.8
<i>Quinqueloculina seminula</i>	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0.1	0.2
<i>Quinqueloculina</i> sp.	0	0	1	0	0	(2)	0	0	0	0	0	0	0	0	0	0	0	3	0.2	0.6
<i>Sigmoilopsis schlumbergeri</i>	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0.1	0.2
Total porcellaneous	6	1	4	3	2	3	1	0	0	0	0	0	0	0	0	0	0	20	0.6	4.2
No. species	4	1	3	3	2	2	1	0	0	0	0	0	0	0	0	0	0	9		
Non fossilizing agglutinated																				
<i>agglutinant</i> spA	1	1	3	1	0	0	0	0	0	0	0	0	0	0	0	0	0	6	0.5	1.3
<i>agglutinant</i> spB	0	0	4	4	4	0	0	0	0	0	0	0	0	0	0	0	0	12	0.9	2.5
<i>Ammoglobigerina</i> sp.1	1	1	5	1	1	0	4	2	0	1	1	1	0	1	0	0	0	19	2.1	4.0
<i>Ammoglobigerina</i> sp.2	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0.4	0.2
<i>Ammoscalaria</i> sp.	0	5	3	0	2	0	0	0	0	0	0	0	0	0	0	0	0	10	0.6	2.1
<i>Cystamina pauciloculata</i>	0	0	3	4	4	3	0	1	0	0	0	0	0	0	0	0	0	15	1.2	3.2
<i>Cyclamina</i> sp.	0	2	1	1	0	(1)	0	0	0	0	0	0	0	0	0	0	0	5	0.6	1.1
<i>Haplophragmoides</i> sp.	0	0	0	0	1	5	0	0	(1)	0	0	0	0	0	0	0	0	7	1.7	1.5
<i>Karrerella bradyi</i>	1	1	5	0	0	0	0	0	0	0	0	0	0	0	0	0	0	7	0.5	1.5
<i>Reophax</i> sp.	1	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	2	0.5	0.4
<i>Saccamina</i> sp.	0	2	3	1	0	0	0	0	0	0	0	0	0	0	0	0	0	6	0.6	1.3
Total agglutinated	4	13	27	13	12	9	4	3	1	1	1	1	0	1	0	0	0	90	0.6	18.9
No. species	4	7	8	7	5	3	1	2	1	1	1	1	0	1	0	0	0	11		
Fossilizing agglutinated																				
<i>Textularia</i> sp.	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0.6	0.2
Total agglutinated	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0.6	0.2
No. species	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1		
Total life foraminifera	36	42	71	59	69	27	12	16	11	14	14	20	47	30	7	0	1	476	2.4	
No. species	17	14	20	15	14	8	6	3	4	4	4	4	4	4	2	0	1	40		
<i>Glomospira</i> spp	0	0	0	0	1	1	3	1	0	2	1	1	0	0	0	0	0	10		
Arborescent indet	0	28	77	112	267	72	14	2	0	0	0	3	0	0	0	0	0	575		

	Depth															Total	ALD ₁₀	% of total fauna	
	0-0.25	0.25-0.5	0.5-0.75	0.75-1	1-1.5	1.5-2	2-2.5	2.5-3	3-3.5	3.5-4	4-5	5-6	6-7	7-8	8-9				9-10
(d) Station F 1264 m																			
Perforate																			
<i>Bulimina inflata</i>	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0.1	0.8
<i>Cibicidoides pachydermus</i>	0	0	0	1	0	3	0	0	0	0	0	0	0	0	0	0	4	1.5	3.3
<i>Cibicidoides robertsonianus</i>	1	1	0	3	1	0	0	1	0	1	0	0	0	0	0	0	8	1.4	6.6
<i>Gavelinopsis translucens</i>	2	0	0	0	0	0	(1)	0	0	0	0	0	0	0	0	0	3	0.1	2.5
<i>Glandulina ovula</i>	0	0	1	0	0	0	0	0	0	0	0	0	0	1	2	0	4	8.2	3.3
<i>Globbulimina affinis</i>	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	1	7.5	0.8
<i>Gyroidina umbonata</i>	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	1	1.8	0.8
<i>Hoeglundina elegans</i>	4	0	6	2	0	0	0	0	0	0	0	0	0	0	0	0	12	0.5	9.8
<i>Lenticulina peregrina</i>	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0.4	0.8
<i>Melonis barleeanus</i>	0	0	1	1	3	0	0	0	0	0	0	0	0	0	0	0	7	1.4	5.7
<i>Oridorsalis umbonatus</i>	1	0	2	0	0	0	0	0	0	0	0	0	0	0	0	0	3	0.5	2.5
<i>Pullenia bulloides</i>	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	1	5.5	0.8
<i>Robertinoides</i> sp.	0	0	2	0	1	0	0	0	0	0	0	0	0	0	0	0	3	0.8	2.5
<i>Uvigerina mediterranea</i>	2	1	0	0	0	0	(2)	0	0	0	0	0	0	0	0	0	5	0.2	4.1
<i>Uvigerina peregrina</i>	11	2	12	5	4	0	6	0	1	0	0	0	0	0	0	0	41	0.9	33.6
Total perforate	22	5	24	12	9	4	11	1	1	1	0	1	0	2	2	0	95	1.2	77.9
No. species	7	4	6	5	4	2	4	1	1	1	0	1	0	2	1	0	15		
Porcellaneous																			
<i>Quinqueloculina seminula</i>	3	4	0	0	0	0	0	0	0	0	0	0	0	0	0	0	7	0.3	5.7
<i>Quinqueloculina</i> sp.	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	1	2.3	0.8
<i>Pyrgo depressa</i>	0	1	1	0	0	0	0	0	0	0	0	0	0	0	0	0	2	0.5	1.6
Total porcellaneous	3	5	1	0	0	0	1	0	0	0	0	0	0	0	0	0	10	0.5	8.2
No. species	1	2	1	0	0	0	1	0	0	0	0	0	0	0	0	0	3		
Non fossilizing agglutinated																			
<i>Ammoglobigerina</i> sp.	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	1	1.3	0.8
<i>Ammoscalaria</i> sp.	0	0	0	0	0	0	2	0	0	0	0	0	0	0	0	0	2	2.3	1.6
<i>Cyclammina</i> sp.	0	0	0	0	3	0	0	0	0	0	0	0	0	0	0	0	3	1.3	2.5
<i>Cribrostomoides subglobosum</i>	0	0	0	0	0	1	3	0	0	0	0	0	0	0	0	0	4	2.1	3.3
<i>Karrerella bradyi</i>	0	0	2	0	0	0	1	0	0	0	0	0	0	0	0	0	3	1.2	2.5
<i>Reophax</i> sp.	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	1	2.3	0.8

(continued on next page)

Table 2 (continued)

	Depth																Total	ALD ₁₀	% of
	0–0.25	0.25–0.5	0.5–0.75	0.75–1	1–1.5	1.5–2	2–2.5	2.5–3	3–3.5	3.5–4	4–5	5–6	6–7	7–8	8–9	9–10			
<i>Saccamina</i> sp.	2	0	0	0	0	0	(1)	0	0	0	0	0	0	0	0	0	3	0.1	2.5
Total agglutinated	2	0	2	0	4	1	8	0	0	0	0	0	0	0	0	0	17	1.5	13.9
No. species	1	0	1	0	2	1	5	0	0	0	0	0	0	0	0	0	7		
Total live foraminifera	27	10	27	12	13	5	20	1	1	1	0	1	0	2	2	0	122	1.2	100.0
No. species	9	6	8	5	6	3	10	1	1	1	0	1	0	2	1	0	25		
Arborescent indet	27	10	27	12	13	5	20	1	1	1	0	1	0	2	2	0	122		

	Depth																Total	ALD ₁₀	% of total fauna
	0–0.25	0.25–0.5	0.5–0.75	0.75–1	1–1.5	1.5–2	2–2.5	2.5–3	3–3.5	3.5–4	4–5	5–6	6–7	7–8	8–9	9–10			
(e) Station H 1993 m																			
Perforate																			
<i>Bulimina inflata</i>	0	2	1	0	0	0	0	0	0	0	0	0	0	0	0	0	3	0.5	1.7
<i>Cibicidoides pachydermus</i>	0	0	0	2	0	3	0	0	0	0	0	0	0	0	0	0	5	1.4	2.8
<i>Chilostomella oolina</i>	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	1	6.5	0.6
<i>Fissurina</i> sp.	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	1	1.8	0.6
<i>Gavelinopsis translucens</i>	0	1	1	0	2	0	0	0	0	0	0	0	0	0	0	0	4	0.9	2.2
<i>Globbulimina affinis</i>	0	0	0	0	0	0	0	0	0	1	0	4	0	0	0	0	5	6.1	2.8
<i>Gyroidina umbonata</i>	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	1	3.8	0.6
<i>Gyroidina orbicularis</i>	0	0	0	4	5	5	0	0	1	2	1	0	0	0	0	0	18	1.9	10.1
<i>Hoeglundina elegans</i>	12	16	18	5	5	2	0	0	0	0	0	0	0	0	0	0	58	0.6	32.4
<i>Lenticulina peregrina</i>	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	1	0.9	0.6
<i>Pullenia</i> sp.	0	0	0	3	0	0	0	0	0	0	0	0	0	0	0	0	3	0.9	1.7
<i>Uvigerina peregrina</i>	0	27	3	2	11	1	0	0	0	(1)	0	0	0	0	0	0	45	0.7	25.1
Total perforate	12	46	23	17	23	12	0	0	1	4	2	0	5	0	0	0	145	1.0	81.0
No. species	1	4	4	6	4	5	0	0	1	3	2	0	5	0	0	0	12		
Porcellaneous																			
<i>Biloculinella irregularis</i>	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	1	1.3	0.6
<i>Quinqueloculina seminula</i>	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	1	0.9	0.6
<i>Pyrgo depressa</i>	0	0	1	1	1	0	0	0	0	0	0	0	0	0	0	0	3	0.9	1.7
<i>Pyrgo subsphaerica</i>	0	0	1	0	1	1	0	1	0	0	0	0	0	0	0	0	4	1.6	2.2
<i>Scutularis</i> sp.	0	2	0	0	0	0	0	0	0	0	0	0	0	0	0	0	2	0.4	1.1
<i>Sigmoilopsis schlumbergeri</i>	1	2	0	0	0	0	0	0	0	0	0	0	0	0	0	0	3	0.3	1.7

Total porcellaneous	1	4	2	2	3	1	0	1	0	0	0	0	0	0	0	0	14	0.9	7.8
No. species	1	2	2	2	3	1	0	1	0	0	0	0	0	0	0	0	6		
Non fossilizing agglutinated																			
<i>Ammoscalaria</i> sp.	2	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	3	0.3	1.7
<i>Cribrostomoides subglobosum</i>	0	0	2	0	2	0	0	0	0	0	0	0	0	0	0	0	4	0.9	2.2
<i>Cyclamina</i> sp.	0	0	2	0	0	0	0	0	0	0	0	0	0	0	0	0	2	0.6	1.1
<i>Cystamina pauciloculata</i>	0	0	2	0	0	0	0	0	0	0	0	0	0	0	0	0	2	0.6	1.1
<i>Haplophragmoides</i> sp.	0	1	0	1	0	0	0	0	0	0	0	0	0	0	0	0	2	0.6	1.1
<i>Karrerella bradyi</i>	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0.4	0.6
<i>Reophax ampullacea</i>	1	2	0	0	0	0	0	0	0	0	0	0	0	0	0	0	3	0.3	1.7
<i>Reophax scorpiurus</i>	2	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	2	0.1	1.1
<i>Reophax</i> sp.	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0.6	0.6
Total agglutinated	5	4	8	1	2	0	0	0	0	0	0	0	0	0	0	0	20	1.0	11.2
No. species	1	3	5	1	2	1	0	0	0	0	0	0	0	0	0	0	9		
Total live foraminifera	18	54	33	20	28	13	0	1	1	4	2	0	5	0	0	0	179	1.0	100.0
No. species	3	9	11	9	9	7	0	1	1	3	2	0	5	0	0	0	27		
Arborescent indet.	0	4	4	12	6	20	4	3	0	0	1	0	0	0	0	0	54		

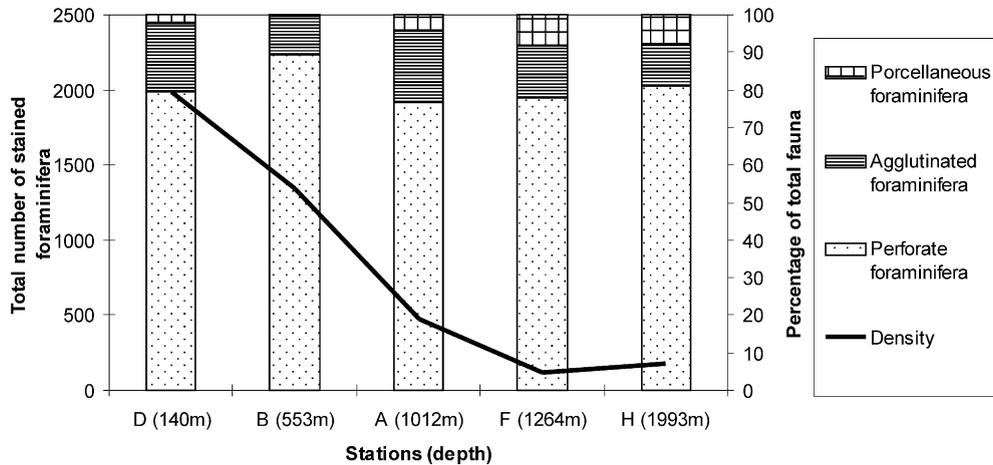


Fig. 2. Abundance of the three main foraminiferal groups, and the density of foraminiferal faunas (total number of stained foraminifera per core (720 cm³)) along the bathymetrical transect.

fauna (Fig. 6b) is largely limited to the upper 3 cm (ALD₁₀=1.2 cm); maximum values are found in the top first cm. *Uvigerina peregrina* (33.6%), *Hoeglundina elegans* (9.8%), *Cibicidoides robertsonianus* (6.6%), *Melonis barleeanus* (5.7%), *Quinqueloculina seminula* (5.7%), *Uvigerina mediterranea* (4.1%), and *Cibicidoides pachydermus* (3.3%) are the main taxa encountered at station F (Table 2d). Although the relatively low numbers of live individuals do not allow very precise observations of the vertical distribution, *Q. seminula* (ALD₁₀=0.3 cm) appears to be limited to the topmost 0.5 cm, whereas *U. peregrina* (ALD₁₀=0.9 cm), and *H. elegans* (ALD₁₀=0.5 cm) can be found slightly deeper (Fig. 6c). Just as at stations B and A, *M. barleeanus* (ALD₁₀=1.4 cm) shows a subsurface maximum at about 1 cm depth.

At station H (water depth 1993 m), the oxygen concentration is 5.85 ml/l. The zero oxygen level is positioned at about 6 cm depth, whereas a weak nitrate + nitrite downward diffusive zone seems to be present between 2.5 and 6.5 cm depth (Fig. 7a). The live fauna is almost completely limited to the uppermost 2 cm of sediment; a maximum density (about 100 individuals/50 cm³) is found at the 0.0–0.5 cm level (Fig. 7b). The ALD₁₀ of the total fauna is 1.0 cm. *Hoeglundina elegans* (32.4%),

Uvigerina peregrina (25.1%), *Gyroidina orbicularis* (10.1%), *Cibicidoides pachydermus* (2.8%), and *Globobulimina affinis* (2.8%) are the main taxa encountered at station H (Table 2d, Fig. 7c). *H. elegans* (ALD₁₀=0.6 cm) and *U. peregrina* (ALD₁₀=0.7 cm) dominate the faunas in the first half cm. *G. orbicularis* (ALD₁₀=1.9 cm) and *C. pachydermus* (ALD₁₀=1.4 cm) appear mainly between 0.5 and 2 cm depth. Some rare individuals of *G. affinis* live in the deepest part of sediment, around the zero oxygen level at about 6 cm depth (Table 3).

4. Discussion

4.1. Benthic ecosystem redox conditions

Bottom water oxygen concentrations for our five stations vary between 4.36 and 5.85 ml/l (Table 1). These values agree very well with the data which have been published for the various water masses in the Bay of Biscay (Ogawa and Tauzin, 1973). The minimum values encountered at station A are typical of the Mediterranean outflow waters. Measurements of bottom water oxygenation during 10 successive sampling campaigns show only minor seasonal and interannual changes for our

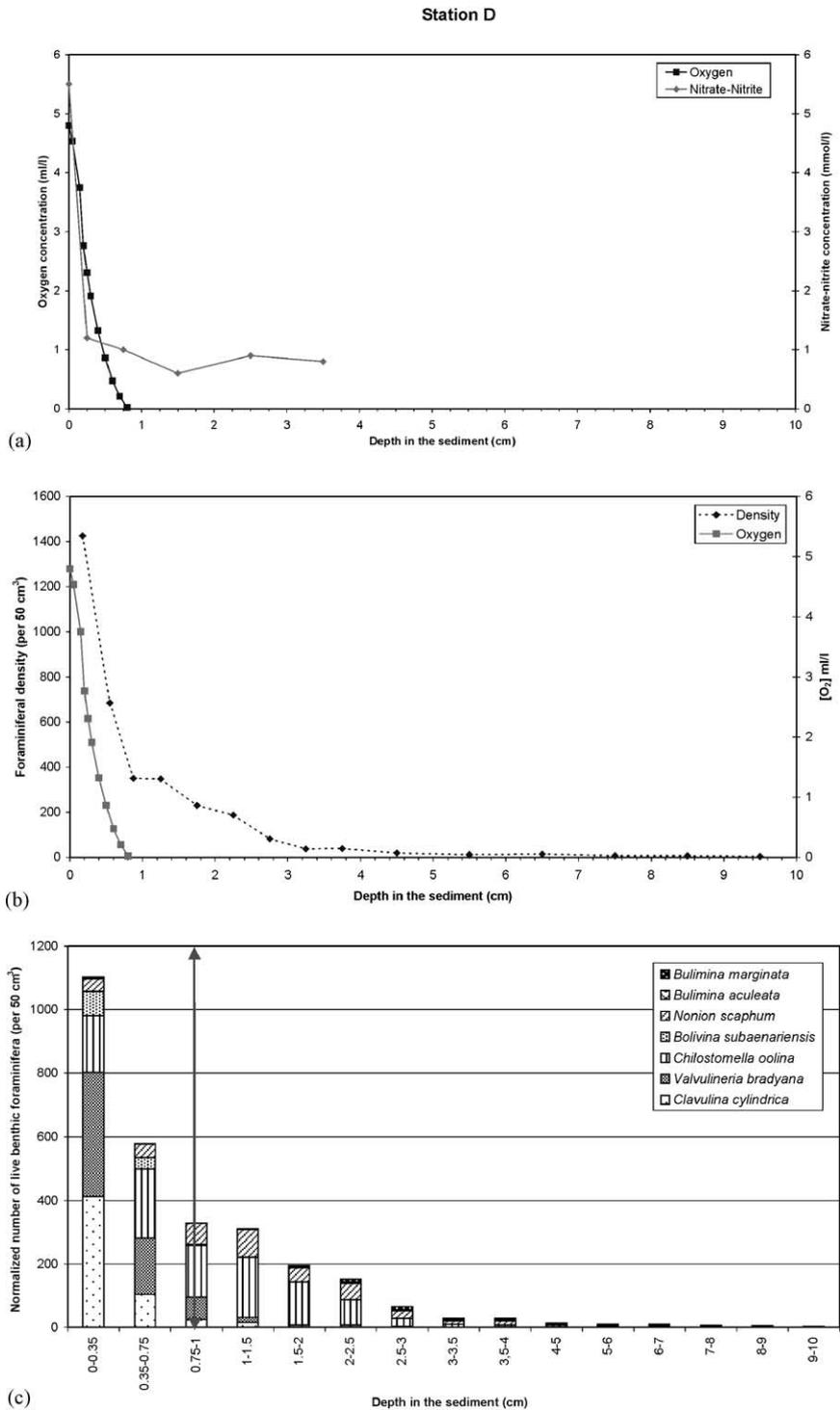


Fig. 3. Station D; (a): dissolved oxygen, and concentration of nitrate + nitrite; (b) dissolved oxygen profile and foraminiferal density (standardised for a 50 cm³ sediment volume); (c) foraminiferal distribution (number of individuals found in each level, standardised for a 50 cm³ sediment volume); double arrow represents zero oxygen boundary.

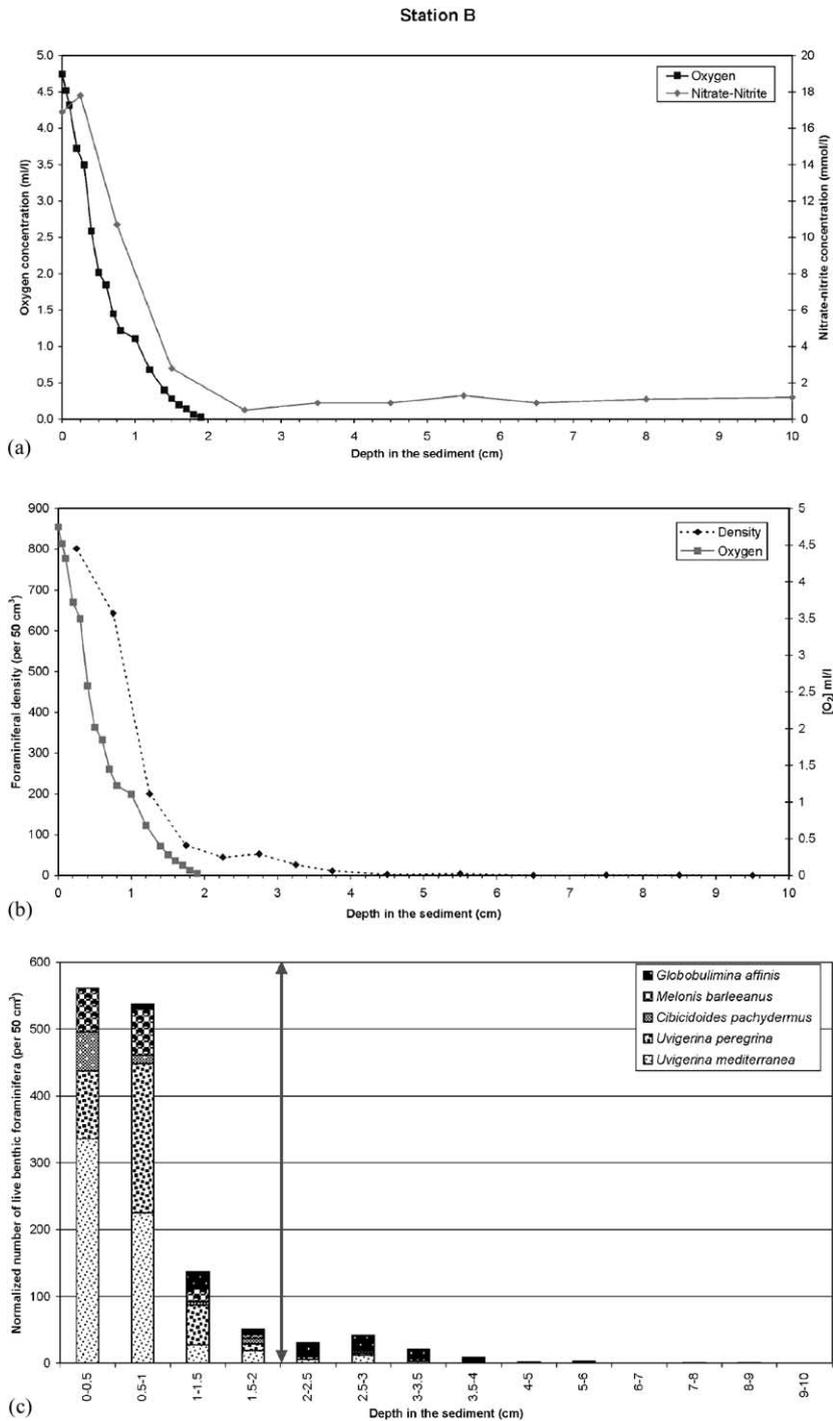


Fig. 4. Station B; (a) dissolved oxygen, and concentration of nitrate + nitrite; (b) dissolved oxygen profile and foraminiferal density (standardised for a 50 cm³ sediment volume); (c) foraminiferal distribution (number of individuals found in each level, standardised for a 50 cm³ sediment volume); double arrow represents zero oxygen boundary.

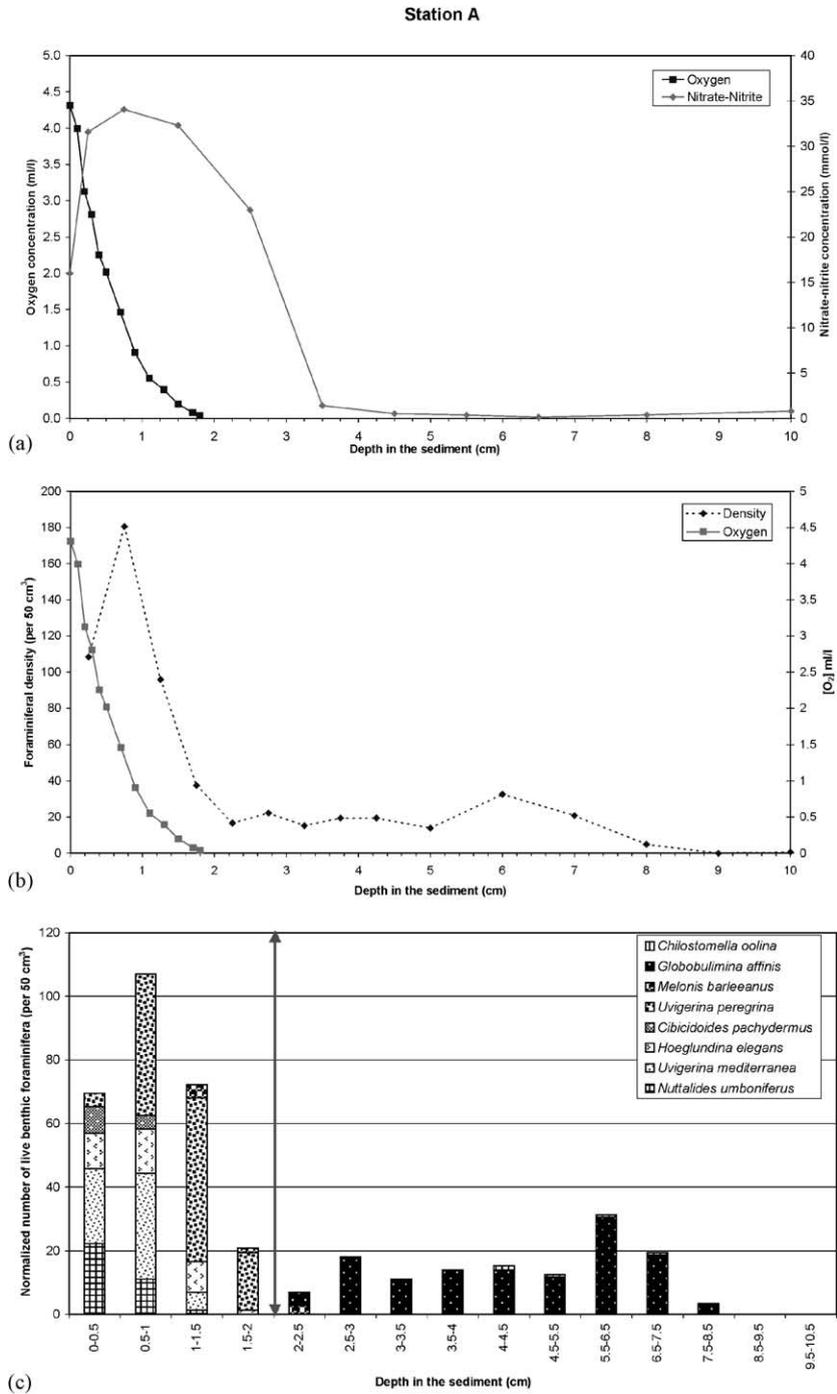


Fig. 5. Station A; (a) dissolved oxygen, and concentration of nitrate + nitrite; (b) dissolved oxygen profile and foraminiferal density (standardised for a 50 cm³ sediment volume); (c) foraminiferal distribution (number of individuals found in each level, standardised for a 50 cm³ sediment volume); double arrow represents zero oxygen boundary.

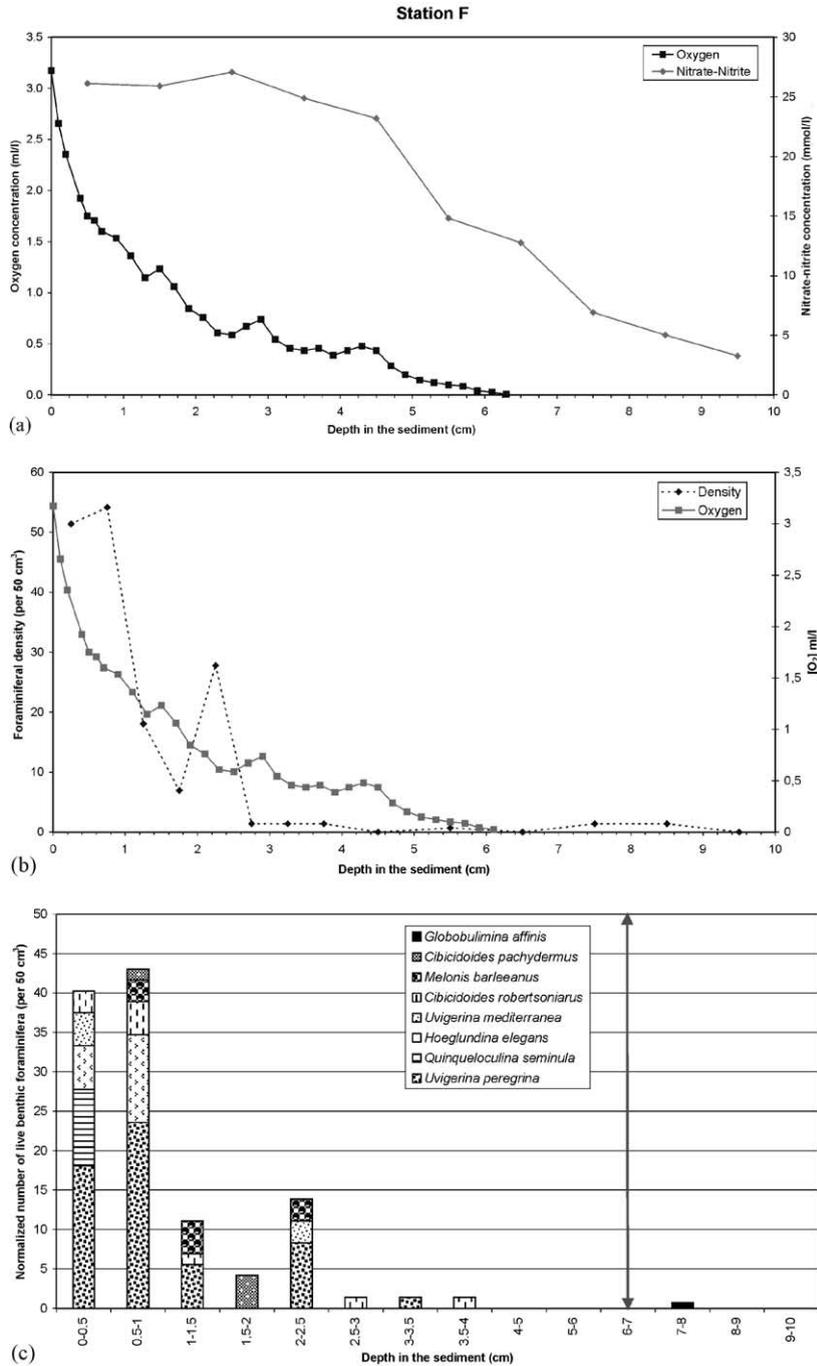


Fig. 6. Station F; (a) dissolved oxygen, and concentration of nitrate + nitrite; (b) dissolved oxygen profile and foraminiferal density (standardised for a 50 cm³ sediment volume); (c) foraminiferal distribution (number of individuals found in each level, standardised for a 50 cm³ sediment volume); double arrow represents zero oxygen boundary.

Table 3

Average living depth (ALD₁₀) of foraminiferal species and (between parentheses) the number of individuals on which the calculation is based. Only occurrences of ≥ 5 individuals are shown. The numbers in italic represent dominant taxa with a relative proportion $\geq 5\%$ in at least one of the stations. Microhabitat patterns are summarised as shallow infaunal (SI), intermediate infaunal (II) or deep infaunal taxa (DI)

Species (> 5 specimen/core)	Stations, ALD ₁₀ (number of specimen)					Total ALD ₁₀	Microhabitat
	D	B	A	F	H		
<i>Bolivina alata</i>	0.3 (37)	1.1 (6)				0.41	SI
<i>Bolivina subaenariensis</i>	0.3 (61)					0.33	SI
<i>Bolivinita quadrilatera</i>		0.2 (7)				0.20	SI
<i>Bulimina aculeata</i>	4.9 (32)					4.97	DI
<i>Bulimina marginata</i>	4.2 (44)	1.2 (19)				3.30	DI
<i>Cancris auriculus</i>	0.3 (11)					0.35	SI
<i>Cassidulina carinata</i>	0.7 (7)					0.71	SI
<i>Chilostomella oolina</i>	1.2 (600)					1.20	II
<i>Cibicidoides pachydermus</i>		0.6 (63)	0.3 (9)		1.4 (5)	0.62	SI
<i>Cibicidoides robertsonianus</i>				1.4 (8)		1.40	II
<i>Coryphostoma</i> sp.	0.2 (7)					0.23	SI
<i>Glandulina ovula</i>			5.0 (12)			5.00	DI
<i>Globobulimina affinis</i>	2.2 (10)	2.4 (80)	5.3 (137)		6.1 (5)	4.24	DI
<i>Gyroidina altiformis</i>		0.2 (9)				0.24	SI
<i>Gyroidina orbicularis</i>					1.9 (18)	1.90	II
<i>Hoeghundina elegans</i>			0.8 (26)	0.5 (12)	0.6 (58)	0.62	SI
<i>Hyalinea balthica</i>	1.6 (53)	1.1 (9)				1.52	II
<i>Lenticulina peregrina</i>		0.2 (7)				0.24	SI
<i>Melonis barleeanus</i>		0.8 (121)	1.7 (6)	1.4 (7)		0.85	SI/II
<i>Nonion scaphum</i>	1.8 (252)					1.81	II
<i>Nuttallides umboniferus</i>		0.3 (5)	0.4 (25)			0.41	SI
<i>Pseudoeponides falsobeccarii</i>	3.2 (25)					3.20	DI
<i>Rectuvigerina phlegeri</i>	0.5 (29)					0.49	SI
<i>Siphogenerina</i> sp.		0.6 (30)				0.65	SI
<i>Uvigerina peregrina</i>	0.4 (38)	0.8 (289)	1.1 (85)	0.9 (41)	0.7 (44)	0.79	SI/II
<i>Uvigerina elongatastriata</i>		1.2 (60)				1.25	II
<i>Uvigerina mediterranea</i>		0.6 (453)	0.6 (45)			0.59	SI
<i>Valvulineria bradyana</i>	0.4 (339)					0.40	SI
<i>Quinqueloculina seminula</i>	0.8 (45)			0.3 (7)		0.71	SI
<i>Ammoglobigerina</i> sp.1			0.6 (9)			0.64	SI
<i>Ammolagena</i> sp.		0.4 (5)				0.38	SI
<i>Ammoscalaria</i> sp.			0.6 (10)			0.63	SI
<i>Clavulina cylindrica</i>	0.4 (296)	1 (10)				0.38	SI
<i>Cribrostomoides subglobosum</i>		0.5 (20)				0.48	SI
<i>Cyclammina cancellata</i>		0.3 (11)				0.26	SI
<i>Cystammina pauciloculata</i>			1.1 (14)			1.12	II
<i>Eggerella</i> sp.	0.5 (10)	0.6 (6)				0.52	SI
<i>Haplophragmoides</i> sp.	0.4 (8)		1.7 (6)			0.94	II
<i>Karreriella bradyi</i>			0.5 (7)			0.52	SI
<i>Nouria polymorphinoides</i>	0.4 (16)					0.36	SI
<i>Reophax</i> sp.	0.3 (6)	0.4 (52)				0.40	SI
<i>Saccammina</i> sp.			0.6 (6)			0.58	SI
<i>Bigenerina nodosaria</i>		0.4 (9)				0.36	SI
<i>Pseudoclavulina crustata</i>		0.9 (10)				0.93	II
<i>Siphotextularia affinis</i>		0.5 (9)				0.51	SI

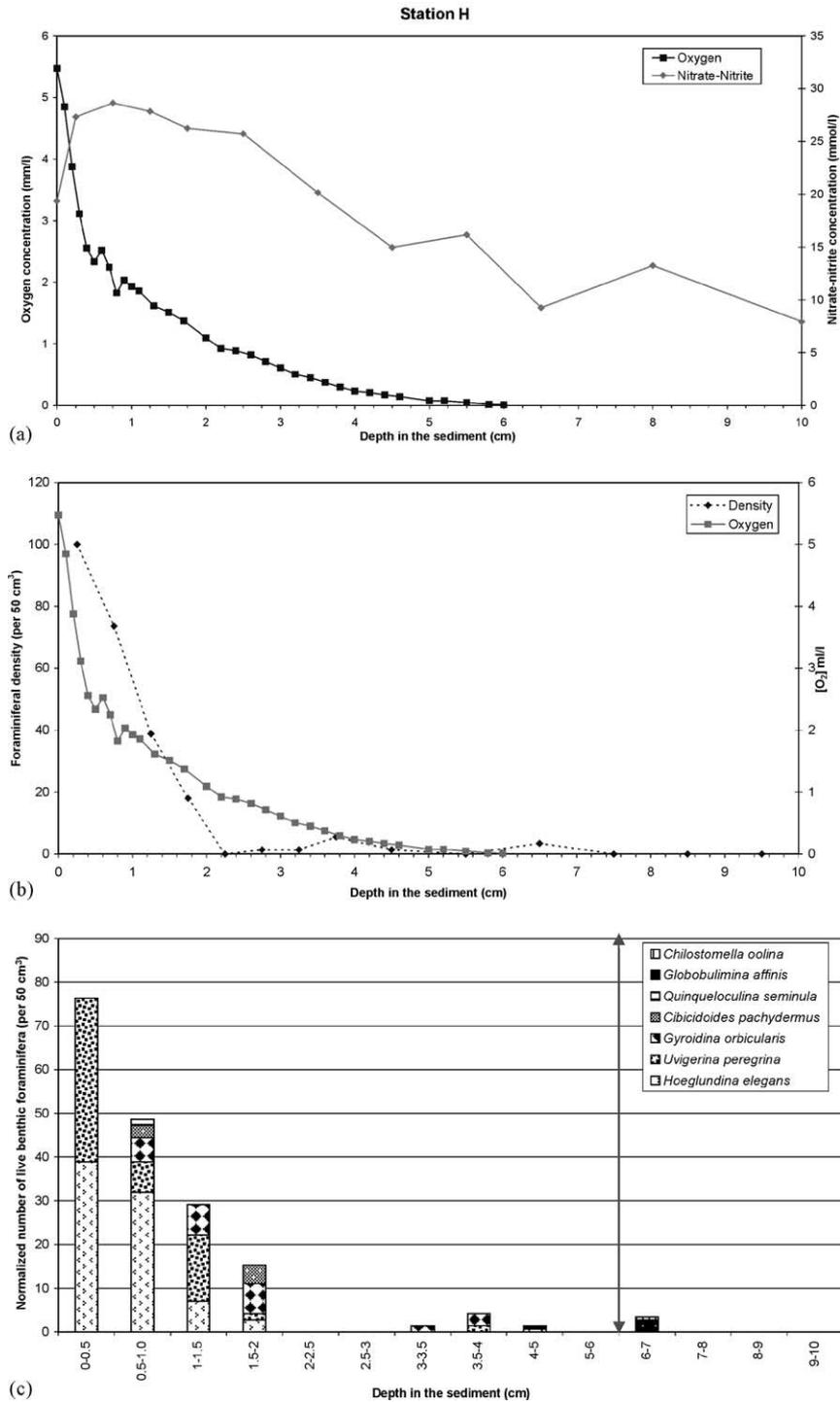


Fig. 7. Station H; (a) dissolved oxygen, and concentration of nitrate + nitrite; (b) dissolved oxygen profile and foraminiferal density (standardised for a 50 cm³ sediment volume); (c): foraminiferal distribution (number of individuals found in each level, standardised for a 50 cm³ sediment volume); double arrow represents zero oxygen boundary.

five stations (Anschutz et al., 1999; Hyacinthe et al., 2001), suggesting that the oxygen concentration of the bottom waters is only very weakly influenced by the variability of the flux of organic matter to the ocean floor. The penetration of free oxygen into the sediment varies from only 8 mm at the shallow station D to more than 6 cm at the deep stations F and H (Table 1). These values strongly suggest that the oxygen gradient in the interstitial waters is only weakly dependent on bottom water oxygenation, but is strongly influenced by the rate of oxygen consumption within the sediment. This rate depends on the oxic degradation of organic matter and oxidation of upward diffusing reduced components. But the oxygen penetration depth is relatively stable through the year (Anschutz et al., 1999; Hyacinthe et al., 2001). Our flux calculations, based on primary production and water depth, suggest a labile organic carbon flux to the ocean floor that is about 14 times higher in station D (where oxygen penetration is minimal) than at station H (where oxygen penetration is maximal).

As soon as all free oxygen has been consumed for the degradation of reactive organic matter, other oxidants are used to continue the remineralisation of labile organic compounds (Froelich et al., 1979; Fenchel and Finlay, 1995). A first step of anaerobic degradation is the reduction of nitrate and nitrite in dinitrogen. This redox reaction takes place mostly below the zero oxygen level, and causes an upward diffusion of newly formed ammonia from the anoxic to oxic layers. In hypoxic sediments, ammonia is oxidised by nitrifying bacteria in nitrate and nitrite, which results in a downward diffusion of nitrate (and nitrite) from oxic to anoxic layers. At the shallowest station D, where the organic flux is maximal, the zone of nitrate reduction is probably situated in the uppermost cm of the sediment (Fig. 3a). Towards the deeper, more oligotrophic stations, the zone of downward diffusion of nitrate + nitrite gradually deepens, from 0.25–2.25 cm at station B to 0.75–3.5 cm at station A, to about 2.5–7.5 cm at stations F and H. Sulphate reduction may be another important mechanism involved in the anaerobic degradation of organic matter; this phenomenon typical of eutrophic ecosystems was

observed only in station D. Important bacterial consortia are supposed to induce these anaerobic redox reactions, which ultimately result in almost complete degradation of both labile and refractory organic matter (Hargrave, 1970; Carney, 1989). The bacterial consortia may constitute an important additional source of labile organic matter below the oxic zone, and could trigger an important recycling of organic carbon under anaerobic conditions (Fenchel and Finlay, 1995).

4.2. Faunal characteristics

It is generally accepted that organic matter influences the composition of the foraminiferal fauna both qualitatively and quantitatively (Thiel, 1983; Berger and Diester Haas, 1988; Altenbach and Sarnthein, 1989; Herguera and Berger, 1991; Altenbach, 1988; Gooday, 1993; Jorissen et al., 1998). In our study area, the faunas of outer shelf station D contain about 15 times more stained foraminifera than those of the deeper stations F and H (Fig. 2). This difference is probably induced by the increase of the vertical labile organic matter flux to the ocean floor towards shallow water. Also the number of species varies between the stations; by far the highest number (49) of taxa is found at station B. This elevated number can probably be explained by the fact that the fauna contains both outer shelf and upper slope elements. The deepest stations F and H have by far the lowest number of taxa, but this is probably at least partially due to the fact that the samples contain about 15 times fewer individuals here.

At all stations, the live faunas are strongly concentrated in the first cm of the sediment, suggesting a dependence on the flux of labile, easily consumable organic matter. Despite the strong faunal concentration in the first cm, the ALD₁₀ of the total faunas (Fig. 8a and b) is not uniform: between 0.8 and 1.2 cm for the relatively shallow stations D and B, as well as for the much deeper stations F and H, but about 2.4 cm for station A (1012 m). The deepening of the ALD₁₀ for the total fauna at station A is mainly the result of an important increase of the ALD₁₀ of the infaunal species *G. affinis* and *C. oolina*, which are dominant faunal elements here (with a cumulative

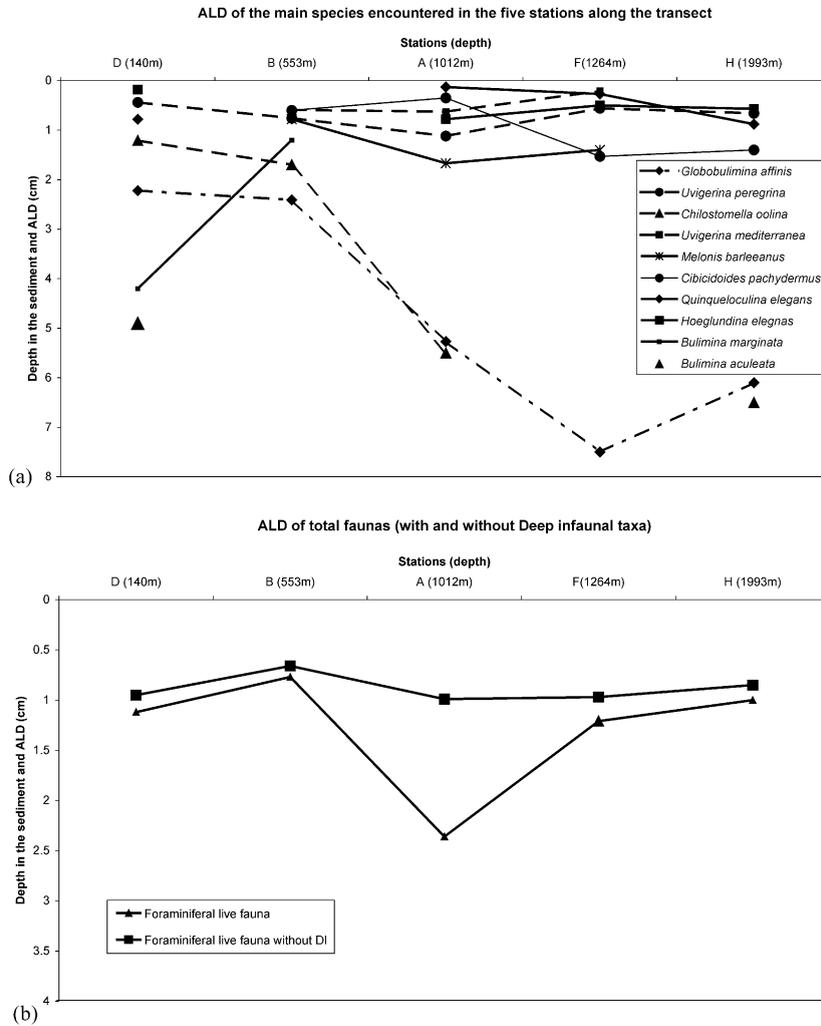


Fig. 8. (a) ALD₁₀ of the main taxa along the bathymetric transect; (b) ALD₁₀ of the total faunas (with and without deep infaunal taxa (DI)).

percentage of 30%) (Fig. 8a). Although their living depth further deepens at the deeper stations F and H, their relative proportion (and thus also their influence on the total faunal ALD₁₀) decreases strongly there. Thus, the very shallow ALD₁₀ of the latter 2 stations is caused mainly by the scarcity of deep infaunal elements. When we consider the ALD₁₀ of the total faunas without these two potentially deep infaunal species (Fig. 8b), the result is a very stable ALD₁₀ of all superficially living taxa (between 0.5 and 0.8 cm).

The shallower station D contains about 2000 individuals that are largely restricted to shallow and intermediate infaunal positions. Apparently this fauna reflects a high labile organic flux to the ocean floor. *Valvulineria bradyana* and *Clavulina cylindrica* dominate the first half cm. They can be considered as two shallow infaunal species, tolerant of low oxygen values, which respond to a high food level at the sediment-water interface. The first species has been described in low oxygen sediments from the centre of the Adriatic Sea mud belt

(Jorissen, 1987, 1988; Van der Zwaan and Jorissen, 1991). The dominant species at this station, *Chilostomella oolina*, has been described as an intermediate or deep infaunal species well adapted to suboxic conditions (Corliss, 1985; Van der Zwaan and Jorissen, 1991; Sen Gupta and Machain-Castillo, 1993; Bernhard and Sen Gupta, 1999). It settles together with *Nonion scaphum* in strongly suboxic and anoxic sediments down to 3-cm depth. In these stressed environments, these two species appear to have a competitive advantage over more superficially living taxa. They may proliferate because of the input of large quantities of organic matter into the deeper sediment layers, remineralised by anaerobic pathways, and the near-absence of less resistant competing taxa (Rathburn and Corliss, 1994). Next, an association consisting of *Bulimina aculeata*, *Bulimina marginata*, and *Pseudoepionides falsobecarii* is found deep in the sediment, under completely anoxic conditions. Several authors (e.g. Lutze and Coulbourn, 1984; Jorissen, 1987; Hermelin and Shimmield, 1990; Verhallen, 1987; Bernhard and Alve, 1996) have described *Bulimina* species as typical elements

in extremely eutrophic and dysoxic settings. In view of their consistent distribution with a surface as well as a deep maximum (Jorissen, 1999, Figs. 10.6d and 10.7f), and the systematic association with sea urchins in our material, we think that the deep occurrences of this association can be explained by their colonisation of macrofaunal burrows. These environments could be attractive because of the strongly increased bacterial activity in the burrow walls (e.g. Fenchel and Jørgensen, 1977).

A major faunal change takes place between stations D (140 m) and B (553 m) (Fig. 9). The latter station is strongly dominated by Uvigerinids (*U. mediterranea* and *U. peregrina*). These shallow infaunal species have been described in a wide variety of eutrophic settings, needing an exported labile organic flux of at least 2.5 g C/m²/yr (e.g. Lutze and Coulbourn, 1984; Corliss, 1985; Corliss and Emerson, 1990; Corliss, 1991; De Rijk et al., 2000; Morigi et al., in press). They can be considered as feeding on labile organic matter in rather well oxygenated shallow infaunal microhabitats. It is important to notice that *U. peregrina*

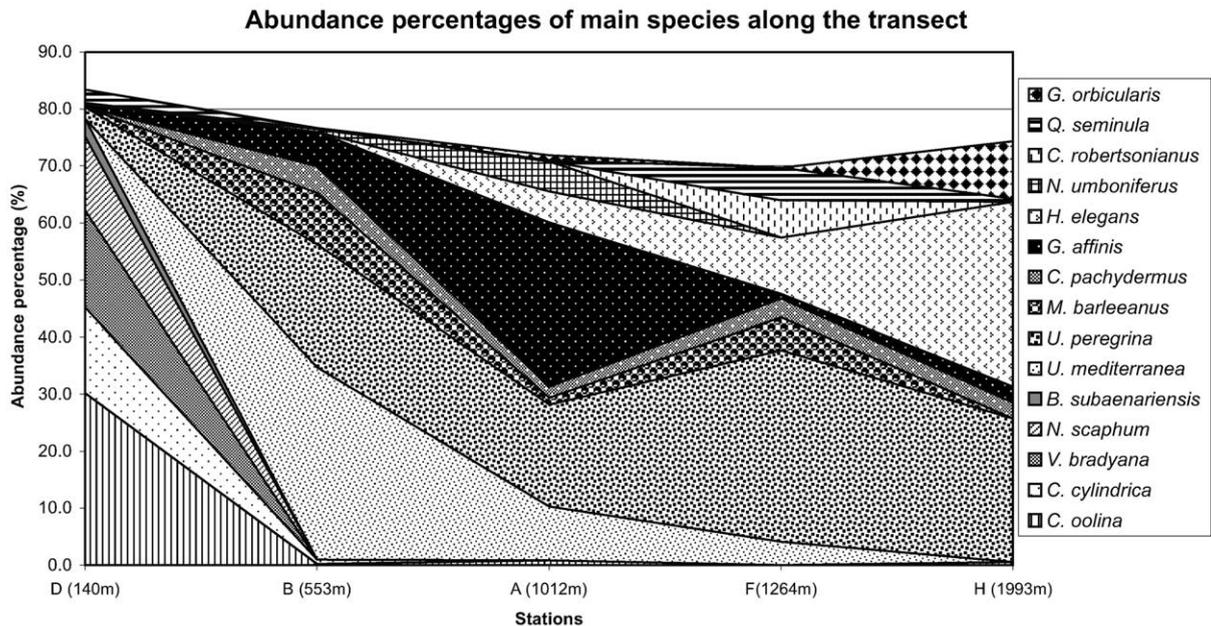


Fig. 9. Composition of the benthic live foraminiferal faunas (in % of the total fauna).

is consistently found deeper than *U. mediterranea*. This could be caused by a slightly higher tolerance for dysoxic conditions. A study of the seasonal variation at station B, however (Fontanier et al., in preparation), shows rather important changes of the microhabitat depth of both taxa through the year, in a context where the oxygen profiles show a very limited variability. Both taxa show maximum microhabitat depths in the richest samples (taken in the most productive periods), which strongly suggests a dependency of the living depth on the quality of the organic matter, high quality organic matter being more abundant in deeper sediment layers during the most eutrophic periods. The deeper microhabitat of *U. peregrina* would then show a slightly higher tolerance for low quality organic matter. A consequence of our observations is that the $\delta^{13}\text{C}$ of *U. peregrina* can hardly be considered as typical of bottom waters.

Deeper in the sediment, the infaunal niches are occupied by *M. barleeanus* and (still deeper) *G. affinis*. This combination of species has been described in numerous mesotrophic-eutrophic oceanic ecosystems (e.g. Harloff and Mackensen, 1997). Jorissen et al. (1995) observe that *M. barleeanus* occurs systematically in the lower part of the oxic zone, where nitrate production (by nitrifying bacteria) occurs, whereas *G. affinis* is consequently found in the upper part of the totally anoxic zone, where nitrate reduction occurs. Our results confirm this pattern. They suggest that both species are dependent on aerobic and anaerobic bacterial stocks degrading more or less refractory organic matter. These bacterial stocks should be concentrated directly below the major redox fronts, where maximum foraminiferal abundances have been described (e.g. Mullins et al., 1985; Linke and Lutze, 1993; Thomsen and Altenbach, 1993; our data). Both species would either prey directly on the bacterial stocks (Lee, 1979; Thomsen and Altenbach, 1993; Kitazato, 1994), or feed themselves with the break-off products. Our data confirm these ideas. We assume that most of the labile components will be consumed in the well-oxygenated topmost cm of the sediment, and will not arrive at the depth where *M. barleeanus* and *G. affinis* live. In the much more eutrophic station D, on the other hand, the near-

absence of both species may be explained by the availability of important amounts of labile organic matter in the sub-surface dysoxic and anoxic levels, which are colonised by *C. oolina* and *N. scaphum*, species that appear to be less able to subsist on low quality organic matter. The total dominance of *C. oolina* in the levels just below Mediterranean sapropel S1 (Jorissen et al., in preparation), which were deposited in a strongly dysoxic setting with an important labile organic matter influx, would be another argument in favour of this interpretation.

Dominant taxa found at station A are essentially the same as those found in station B, showing that the change from North Atlantic Central Waters to MW has no visible impact on the benthic faunas. The almost perfect separation between the group of superficial infaunal taxa (dominated by *Nuttallides umboniferus*, *U. peregrina* and *U. mediterranea*) from a deep infaunal assemblages (dominated by *Globobulimina affinis*) is striking. The appearance of more oligotrophic shallow infaunal taxa such as *H. elegans*, *N. umboniferus* and *C. pachydermus* (compare Corliss, 1985; Corliss and Emerson, 1990; Corliss, 1991), shows the influence of the gradually decreasing organic flux towards deeper areas. Also rather oligotrophic arborescent agglutinated taxa (Jones and Charnock, 1985), which have not been included in our counting results, are very abundant at this station (575 fragments). The high quantities of *G. affinis* in the anoxic sediments from 2 to 8 cm depth can be explained in several ways. They could be (1) dead, but still staining, partially decomposed individuals, (2) live, but inactive animals with a lowered metabolism, or (3) active, facultative aerobic individuals, which dwell on an important stock of food made available by bacterial activity. The source of the originally (before bacterial conversion/partial break-off) refractory organic matter could be a former refractory organic matter input by slope failure processes.

The poor faunas (in terms of faunal density) of stations F (1264 m) and H (1993 m) show faunal changes characterising the further decrease of the trophic level. *U. mediterranea* and *G. affinis* show a strong relative frequency decrease, and *U. pere-*

grina (station F) and *H. elegans* become dominant. Furthermore, the faunas become largely restricted to the well-oxygenated top 2 cm of the sediment. The slightly deeper zone of nitrate reduction is almost devoid of benthic foraminifera. In the top of the anoxic zone, only very few individuals of *G. affinis* are found. This faunal evolution translates the trend towards oligotrophic ecosystems, where metabolisable organic matter is limited to the uppermost sediment layers. Apparently bacterial conversion of refractory organic matter in deeper, dysoxic/anoxic sediment layers is a minor process here.

4.3. Benthic foraminiferal microhabitats

The dependence of the benthic foraminiferal microhabitat on the availability of food and the oxygenation of the benthic ecosystem has been schematised in a conceptual model by Jorissen et al. (1995), as shown in Fig. 10. The so-called TROX-model explains that in very oligotrophic environments, shallow infaunal species, which are adapted to a low trophic level, will thrive close to the sediment-water interface in a well oxygenated setting. The scarcity of organic matter introduced into the sediment (because of weak bioturbation and almost complete consumption of organic compounds in the first mms of the sediment)

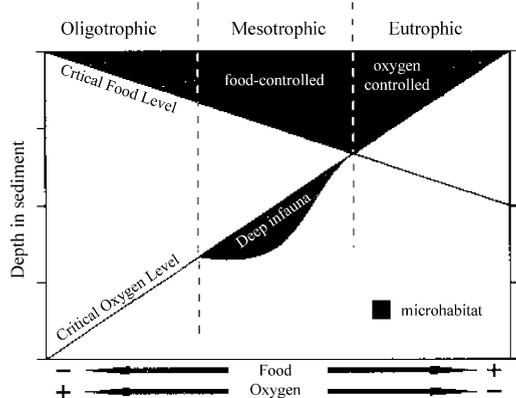


Fig. 10. TROX-model (Trophic condition and Oxygen concentration) explaining the vertical distribution of foraminifera in the top cm of the sediment (after Jorissen et al., 1995); see text for full explanation.

prevents colonisation of the deeper sediment layers by infaunal taxa. But also in much more eutrophic conditions, where the principal redox front is positioned close to the sediment surface, shallow infaunal taxa are limited to the first mm of the sediment; in this case, they have only a limited tolerance for low oxygen conditions. According to the TROX-model, faunal penetration will be maximal in mesotrophic settings, where oxygen penetration is relatively deep, and more or less labile food particles are introduced at depth in the sediment by bioturbating macrofauna. In such mesotrophic environments, grazing on anaerobic bacterial stocks (or their break-up products) has been proposed as an explanation for the presence of rather deep-living intermediate and deep infaunal species. It is evident that the foraminiferal taxa have a dynamic behaviour, allowing them to adapt rapidly to changing conditions. Inherent to the TROX-model is the notion that the control of the foraminiferal microhabitat is dual. Whereas oxygenation is considered as the main controlling parameter in eutrophic ecosystems, food availability is supposed to control the vertical faunal distribution in more oligotrophic environments.

Our data fully confirm the main distributional trends predicted by the TROX-model. At stations D and B, which represent the eutrophic extreme in our study, the faunas are concentrated in the first 2 cm. In the “mesotrophic” station A, faunal penetration is deepest, and an important population of deep infaunal *G. affinis* is found at considerable depth in the sediment. Stations F and H, finally, represent the most oligotrophic situation in our study, and faunas are once again limited to the first centimetres of the sediment.

Despite the good fit to our data, it is evident that the TROX-model oversimplifies the foraminiferal response to the controlling environmental parameters for a number of reasons:

- (1) The TROX-model does not insist enough on the dependency of pore water oxygenation on the organic flux. Since the supply of metabolisable organic matter controls the oxygen consumption in the sediment and the localisation of the successive redox fronts, it is evident

that the organic flux is the main parameter controlling the foraminiferal distribution in the sediment.

- (2) Recent studies (e.g. Alve and Bernhard, 1995; Fenchel and Finlay, 1995; Moodley et al., 1997; Jannink et al., 1998) show that anoxic conditions do not have a direct lethal effect for the majority of species. If severe dysoxia in bottom and pore waters cause the disappearance of certain taxa, this is possibly so, because reproduction is inhibited in such environments. Therefore, benthic ecosystem oxygenation appears to be a less important factor than was suggested by Jorissen et al. (1995), and in consequence, the quantity and quality of food particles appear to be by far the most important parameters.
- (3) The TROX-model depicts trends for the total fauna, but individual taxa may show important differences in their tolerance levels with respect to the main controlling parameters.

- (4) It is evident that competition between taxa for high quality food particles is an important factor causing microhabitat differences (Van der Zwaan et al., 1999). The deep microhabitat of several infaunal taxa may be caused by their low competitiveness in the more attractive sediment surface niches (which are much richer in easily metabolisable organic matter).
- (5) Benthic ecosystems may know important seasonal or interannual variability. The foraminiferal faunas will respond to rapid changes in the organic flux or bottom water oxygenation by a shift of their microhabitat structure (Barmawidjaja et al., 1992; Ohga and Kitazato, 1997).

In Fig. 11, we show the microhabitat occupation at a species level in the range of meso-oligotrophic environments represented by our stations. Labile, particulate organic matter will be concentrated at the sediment surface. In extremely oligotrophic

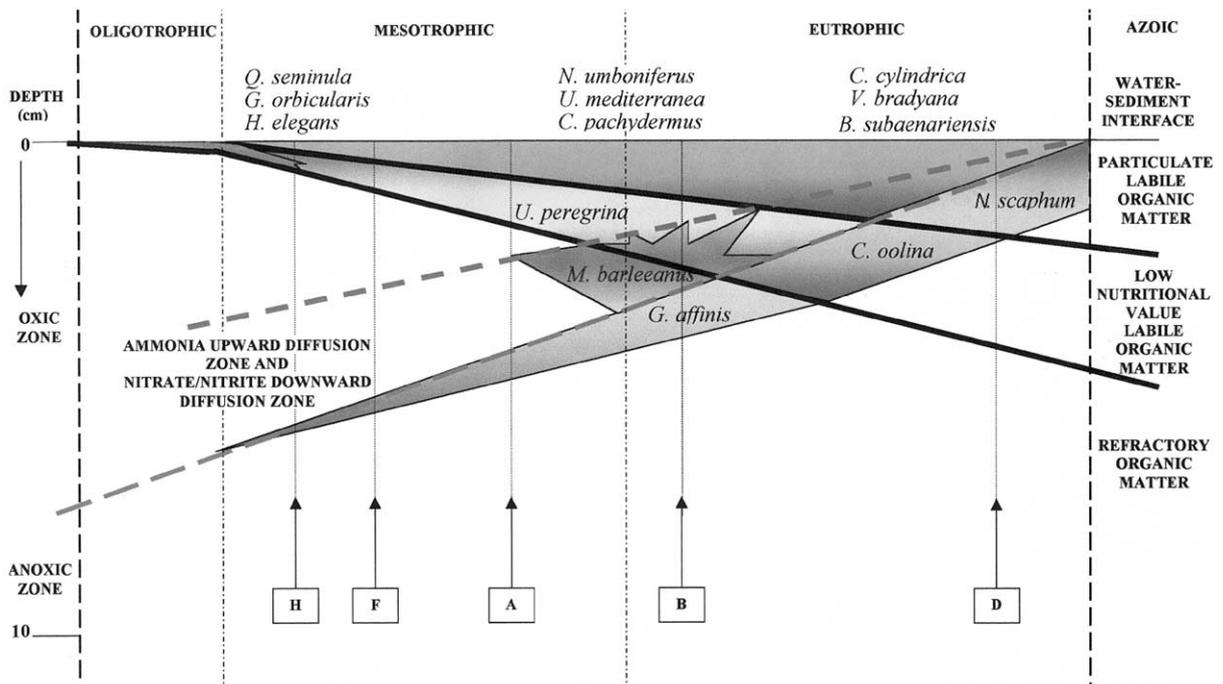


Fig. 11. Microhabitat distribution and specific foraminiferal composition along the bathymetric transect in the Bay of Biscay; the approximate position of our five stations are indicated; see text for full explanation.

ecosystems such easily metabolisable material will be very scarce, or even be absent, except for rare phytodetritus deposit events (Gooday, 1993). Towards slightly less oligotrophic sites (our stations H and F), part of this material will be introduced in the top mm of the sediment. The result is the creation of a niche inhabited by very competitive, shallow infaunal taxa. In rather oligotrophic settings such faunas are dominated by *H. elegans*, *Q. seminula*, and *C. pachydermus*, in more mesotrophic settings (stations A, B) by *U. mediterranea*, *N. umboniferus* and *G. orbicularis*, and in more eutrophic environments (station D) by *C. cylindrica*, *V. bradyana* and *B. subaenariensis* (Fig. 11). All these taxa seem to combine a preference for high quality resources (or a low tolerance for low quality food), a high competitiveness and perhaps a limited tolerance for anoxic conditions. When in eutrophic ecosystems, labile, easily metabolisable matter is introduced into anoxic parts of the sediment column (because of a shallow zero oxygen level), other, more low-oxygen resistant species, such as *C. oolina* and *N. scaphum*, will take over this superficial niche. Immediately below this surface zone with easily metabolisable organic matter, an environment is found where organic matter is less reactive (for instance in faecal pellets). *U. peregrina* seems to be one of the most tolerant species for this lower quality organic matter. In oligotrophic ecosystems, at the basal part of this zone, reactive organic matter becomes very scarce, or even absent within the sediment, and, consequently, benthic foraminifera are no longer found. Still deeper, concentrations of reactive organic matter will be associated with bacterial activity, and will logically be positioned around important redox boundaries (Lee, 1979; Mullins et al., 1985; Linke and Lutze, 1993; Kitazato, 1994; Jorissen, 1999). One of the first of these fronts may be that of nitrifying bacteria. Between this front and the zero oxygen level, *M. barleeanus* seems to be the best-adapted taxon. The main locus of bacterial activity appears to be concentrated in the topmost part of the anoxic layer. Here, abundant benthic foraminiferal faunas dominated by *G. affinis* may be present. In the more oligotrophic environments (stations H and F), the burial of organic compounds becomes

insignificant and can no longer sustain important anaerobic bacterial stocks and associated deep infaunal foraminifera. When reading Fig. 11, one should, of course, realise that the limits between the various types of resources are not sharp, and consequently, large overlaps between the various taxa occur. Furthermore, ecosystems are always dynamic. The trophic level may show important short-term fluctuations, and the benthic faunas will respond accordingly. In general, more opportunistic taxa will profit from such ecosystem instability.

5. Conclusions

1. The dissolved oxygen concentration of the bottom waters is not seriously influenced by the exported organic flux. The interstitial waters, on the contrary, show a clear linkage: at the shallowest site (D), high and relatively constant organic matter inputs result in a redox front close to the sediment-water interface. At the deeper sites, the zero oxygen level is found much deeper in the sediment, as a direct result of the decreasing organic flux.
2. The flux of organic matter to the ocean floor is the main parameter that controls the density and the composition of benthic foraminiferal faunas along the bathymetrical transect.
3. The oxygenation of the pore waters does not have a direct control on the vertical distribution of most foraminiferal taxa. Most species are able to adapt to severe dysoxia or even anoxia, provided that sufficient high quality food particles are available.
4. The TROX-model (Jorissen et al., 1995) is confirmed by the present data. Faunal penetration is indeed maximal in mesotrophic conditions, where the advantageous conditions of high-oxygen concentration and ample high quantity food availability may be found relatively deep in the sediment. However, the variability of the total faunal ALD₁₀ (Average Living Depth) is mainly determined by deep infaunal taxa. Most shallow infaunal taxa show only a minor variability in their microhabitats between the various stations.

5. Certain taxa, such as *N. scaphum* and *C. oolina* appear to be typical for settings where labile organic matter is introduced into strongly dysoxic or anoxic environments. In case of an organic matter of lower quality, they are replaced by *M. barleeanus* and *G. affinis*.
6. Rather consistently, *U. peregrina* is found slightly below the most superficially living taxa, confirming that it is incorrect to consider the stable isotopic composition of its shell as representative for the bottom waters.

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Appendix A

Differentiating between *Uvigerina peregrina* and *Uvigerina mediterranea* may be difficult sometimes. This is especially the case when *U. peregrina* types lose the spinose aspect of their ornamentation. A good overview of various *U. peregrina* morphotypes is given in Van der Zwaan et al. (1986), dealing with *Uvigerina* taxonomy.

In general, differences between the 2 species are as follows:

- (1) *U. mediterranea* has a rather bulky test, with a low length/width ratio, the opposite is true for

U. peregrina. Adult *U. mediterranea* are much bigger than adult *U. peregrina*.

- (2) *U. mediterranea* has a rather large first chamber, much bigger than *U. peregrina*. Exception: microspheres of *U. mediterranea*, which may cause problems.
- (3) *U. mediterranea* has rather “gentle” costae, which are widely spaced, and continuous. *U. peregrina* has very sharp costae, less distance between them. The costae are very often crenulated, because from an evolutionary point of view (*U. peregrina* is very far from *U. mediterranea*); the costae have formed out of a series of spines. This difference can particularly well be seen on the basis of the test (first chamber); this is completely smooth in *U. mediterranea*, and uneven, covered with very small spines in *U. peregrina*.
- (4) *U. peregrina* may have spinose ornamentation (the last chamber, or the surface between the costae), this can almost always be detected on the first chamber (see before); *U. mediterranea* has never such spinose ornamentation.
- (5) Although not a 100% characteristic, the neck of *U. mediterranea* is better developed, and very often there is a prominent lip.
- (6) Again not a 100% characteristic: the neck of *U. mediterranea* is very often placed in a kind of depression; the aperture of *U. peregrina* is always 100% terminal, on the highest point of the test.

Characteristic species from the outer-shelf and slope environments of the Bay of Biscay were identified by using designation and references to plates and figures in literature on Atlantic and Mediterranean foraminifera (see Table 4).

Table 4

Species	References
<i>Amphicoryna scalaris</i> (Batsch), 1791	Jones (1994), pl. 63, Figs. 28–31
<i>Bigenerina nodosaria</i> d’Orbigny, 1826	Jones (1994), pl. 44, Figs. 19–24
<i>Biloculinella irregularis</i> (d’Orbigny) 1839	d’Orbigny (1839), pl. 8, Figs. 20 and 21
<i>Bolivina alata</i> (Seguenza), 1862	Schiebel (1992), pl. 1, Fig. 2

Table 4 (continued)

Species	References
<i>Bolivina spathulata</i> (Williamson), 1858	Jorissen (1987), pl. 1, Fig. 5
<i>Bolivina subaenariensis</i> Cushman, 1922	Phleger et al. (1953), pl. 7, Figs. 24 and 25
<i>Bolivinita quadrilatera</i> (Schwager), 1866	Jones (1994), pl. 42, Figs. 8–12
<i>Bulimina costata</i> d'Orbigny, 1826	Van Leeuwen (1989), pl. 8, Figs. 2 and 3
<i>Bulimina inflata</i> Seguenza, 1862	Van Leeuwen (1989), pl. 8, Fig. 4
<i>Cancris auriculus</i> (Fichtel & Moll), 1942	Jones (1994), pl. 106, Fig. 4
<i>Cassidulina carinata</i> Silvestri, 1896	Phleger et al. (1953), pl. 9, Figs. 32–37
<i>Chilostomella oolina</i> Schwager, 1878	Jones (1994), pl. 55, Figs. 12–14
<i>Cibicides lobatulus</i> Walker and Jacob, 1798	Jones (1994), pl. 92, Fig. 10
<i>Cibicides wuellerstorfi</i> (Schwager), 1866	Van Leeuwen (1989), pl. 10, Figs. 1–9
<i>Cibicidoides pachydermus</i> (Rzehac), 1886	Jones (1994), pl. 94, Fig. 9
<i>Cibicidoides robertsonianus</i> (Brady), 1881	Van Leeuwen (1989), pl. 9, Figs. 1–3
<i>Clavulina cylindrica</i> d'Orbigny, 1952	Hofker (1932), Figs. 18 and 19
<i>Cribrostomoides subglobosum</i> (M. Sars), 1868	Jones (1994), pl. 34, Figs. 8–10
<i>Cyclammina cancellata</i> Brady, 1879	Jones (1994), pl. 37, Figs. 8–16
<i>Cystammina pauciloculata</i> (Brady), 1879	Jones (1994), pl. 41, Fig. 1
<i>Dentalina ariena</i> Patterson and Pettis, 1986	Jones (1994), pl. 62, Figs. 27–31
<i>Elphidium advenum</i> Cushman, 1922	Phleger et al. (1953), pl. 6, Fig. 15
<i>Gavelinopsis translucens</i> (Phleger & Parker), 1951	Schiebel (1992), pl. 4, Fig. 5
<i>Glandulina ovula</i> d'Orbigny, 1846	Jones (1994), pl. 61, Figs. 17–22
<i>Globobulimina affinis</i> d'Orbigny, 1826	Phleger et al. (1953), pl. 6, Fig. 32
<i>Glomospira charoides</i> Jones & Parker, 1860	Phleger et al. (1953), pl. 5, Fig. 1
<i>Glomospira gordialis</i> Jones & Parker, 1860	Phleger et al. (1953), pl. 5, Fig. 2
<i>Gyroidina altiformis</i> Stewart & Stewart, 1930	Jorissen (1987), pl. 1, Fig. 11
<i>Gyroidina umbonata</i> (Silvestri), 1898	Parker (1958), pl. 3, Figs. 19 and 20
<i>Hanzawaia boueana</i> (d'Orbigny), 1846	Jorissen (1987), pl. 3, Fig. 10
<i>Hoeglundina elegans</i> (d'Orbigny), 1826	Phleger et al. (1953), pl. 9, Figs. 24 and 25
<i>Hyalinea balthica</i> (Schroeter), 1783	Jones (1994), pl. 112, Fig. 1 and 2
<i>Karriella bradyi</i> (Cushman), 1911	Jones (1994), pl. 41, Figs. 1–4
<i>Lenticulina gibba</i> d'Orbigny, 1839	Jones (1994), pl. 69, Figs. 8 and 9
<i>Lenticulina peregrina</i> (Schwager), 1866	Cushman and McCulloch (1950), pl. 39, Fig. 5
<i>Melonis barleanus</i> (Williamson), 1858	Van Leeuwen (1989), pl. 13, Figs. 1 and 2
<i>Nonion scaphum</i> (Fichtel and Moll), 1798	Jones (1994), pl. 109, Fig. 12
<i>Nonionella turgida</i> (Williamson), 1858	Jones (1994), pl. 109, Figs. 17–19
<i>Nouria polymorphinoides</i> Heron–Allen & Earland, 1914	Loeblich and Tappan (1988a, b), pl. 123, Figs. 11 and 12
<i>Nuttallides umboniferus</i> (Cushman), 1933	Van Leeuwen (1989), pl. 15, Figs. 11–13; pl. 16, Figs. 1–7
<i>Oridorsalis umbonatus</i> Reuss, 1851	Van Leeuwen (1989), pl. 17, Figs. 1–13
<i>Pseuclavulina crustata</i> Cushman, 1936	Jorissen (1987), pl. 1, Fig. 1
<i>Pseudoepionides falsobeccarii</i> Rouvillois, 1974	Jorissen (1987), pl. 4, Fig. 3
<i>Pullenia bulloides</i> (d'Orbigny), 1826	Phleger et al. (1953), pl. 10, Fig. 19
<i>Pyrgo depressa</i> (d'Orbigny), 1826	Jones (1994), pl. 2, Figs. 12, 16 and 17
<i>Pyrgo subsphaerica</i> d'Orbigny, 1839	Cushman (1929), pl. 18, Figs. 1 and 2
<i>Pyrgoella sphaera</i> (d'Orbigny), 1839	Jones (1994), pl. 2, Fig. 4
<i>Quinqueloculina seminula</i> (Linné), 1758	Jones (1994), pl. 5, Fig. 6
<i>Rectuwigerina phlegeri</i> Le Calvez, 1959	Le Calvez (1959), pl. 1, Fig. 11
<i>Reophax ampullacea</i> Brady, 1881	Jones (1994), pl. 30, Fig. 6
<i>Reophax scorpiurus</i> Montfort, 1808	Loeblich and Tappan (1988a, b), pl. 44, Figs. 1–3
<i>Sigmoilopsis schlumbergeri</i> Silvestri, 1904	Jones (1994), pl. 8, Figs. 1–4
<i>Siphotextularia affinis</i> Fornasini, 1883	Kohl (1985), pl. 2, Fig. 5
<i>Textularia agglutinans</i> d'Orbigny, 1839	Jones (1994), pl. 43, Figs. 1–3
<i>Textularia sagittula</i> Defrance, 1824	Jorissen (1987), pl. 3, Fig. 12
<i>Trifarina angulosa</i> (Williamson), 1858	Jones (1994), pl. 74, Figs. 17 and 18
<i>Trifarina pauperata</i> (Heron–Allen & Earland), 1932	Timm (1992), pl. 6, Fig. 4

Table 4 (continued)

Species	References
<i>Trochammina inflata</i> (Montagu), 1808	Jones (1994), pl. 41, Fig. 4
<i>Uvigerina elongatastriata</i> (Colom), 1952	Van der Zwaan et al. (1986), pl. 6, Figs. 1–8
<i>Uvigerina mediterranea</i> Hofker, 1932	Van der Zwaan et al. (1986), pl. 5, Figs. 1–7
<i>Uvigerina peregrina</i> Cushman, 1923	Van der Zwaan et al. (1986), pl. 1, Figs. 1–6
<i>Uvigerina proboscidea</i> Schawger, 1866	Van der Zwaan et al. (1986), pl. 12, Figs. 1–4
<i>Valvulineria bradyana</i> (Fornasini), 1900	Jorissen (1987), pl. 4, Figs 1 and 2

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