

Amphipods and benthic biocoenosis on the Coasts of Alboraya-Albuixech (Spain, Gulf of Valencia, Western Mediterranean)

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The studied zone, north of Valencia city, corresponds to a typical sandy coast ecosystem of 5 km length. This area is suffering an important anthropic pressure (residual waters flows, fisheries, and urban, industrial, touristic and agricultural activities) which originates a general regression. Harbours and breakwaters, are the hard substrates that can be found there.

GINER (1989) studied the infralittoral biocoenosis distribution based on the molluscan fauna, and MARTI (1989) using information from amphipod fauna contributes to a better characterization of these biocoenosis in the area.

Different methods have been used to take samples of the different types of biocoenosis such as scoop net, Aberdeen double side anchor dredge, Agassiz trawl and scraped surfaces 25x25 cm, from 21 stations (15 from soft substrates and 6 from hard ones). The location of these stations were chosen on purpose to define the whole conditions of the studied area.

RESULTS.

SUPRALITTORAL ZONE.

* **LDL biocoenosis:**

It can be found along the shoreline. The sediment is formed by a mixture of pebbles and fine sand, on which masses of several types of organic debris and *Posidonia oceanica* rhizome fibrils are located. This biocoenosis is characterized by the existence of high density populations of *Orchestia platensis* together with sporadic specimens of *Talorchestia deshayesii*.

MEDIOLITTORAL ZONE.

* **AP biocoenosis:**

Species mentioned by LEDOYER (1968), BELLAN-SANTINI & LEDOYER (1973) from shallow algae populations and from high polluted areas as *Jassa marino-rata*, *Corophium acutum*, *C. insidiosum*, *Caprella aquilibra* and *Elasmopus rapax* have been located on artificial rocky substrates at the study area. Among all of them, the last species characterizes the *Mytilus galloprovincialis* and *Corallina elongata* facies, and it is only found in those facies at the studied area.

INFRALITTORAL ZONE.

* **SFHN biocoenosis:**

It has a very slow specific richness, with only dispersed individuals from nearby biocoenosis, *Corophium saxtonae*, *Harpinia pectinata*, *Siphonocoetes sabatieri* and *Urothoe poseidonis* can be found. It may be due to the sensitiveness of this group to highly polluted waters, pointed out by DAVIN (1981), and to the artificial structures settled --harbours, breakwaters, urban effluents-- which also alter the hidrological and sedimentary factors.

* **SFBC biocoenosis:**

It reaches 9-10 m depth where the upper limit of the *P. oceanica* meadows is settled. The existence of *Pericouloides longimanus*, *Pariambus typicus*, *Amphelisca brevicornis*, *Leucothoe incisa*, *Microprotopus maculatus* and *Urothoe poseidonis* helps to the precise characterization of this biocoenosis.

High densities of *Siphonocoetes sabatieri* and *Gammarus crinicornis*, typical species from low salinity environments, are found in some sectors of this biocoenosis under the influence of fresh-waters flows. Fluctuations in the populations of these species can be observed, and there is a substitution from *S. sabatieri* to *G. crinicornis* in winter probably due to the variability of the hidrological and sedimentary factors and to the opportunist nature of them.

* **Posidonia oceanica meadows:**

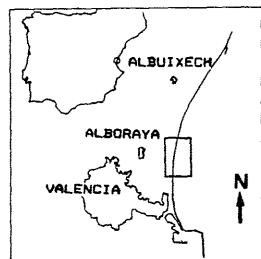
It shows a high regression degree (1 or 2 shoots/m² to -10 m) buried by sandy sediment where dense *Caulerpa prolifera* patches are developed. This biocoenosis is characterized in the sampled area by *Erichthonius punctatus*, *Maera inaequipes*, *Orchomene humilis* and by *Leucothoe richiardii* which are found in the meadow and in the enclaves of biological sciaphilic concreted algae and porifera. All these species have been already mentioned in the rhizome terraces of *P. oceanica* (CHEVREUX, 1910; HARMELIN, 1964; LEDOYER, 1962; LEDOYER, 1968).

* **SGCF biocoenosis:**

It is found in big pot-holes and channels in the *P. oceanica* meadows. The existence of *Monocouloides carinatus*, *Pontocrates arenarius*, *Ceradocus semiserratus*, *Gonnes coalita* y *Socarnes erythrophthalma*, characterizes this biocoenosis perfectly.

* **Enclaves of circalittoral biological concreted masses:**

They are developed on the dead rhizome terraces of *P. oceanica*. These enclaves are identified by the presence of *Iphimedia serripes*, *Lysianassa pilicornis* and *Pseudoprotella phasma*, typical species of circalittoral zone and coralligenous bottoms, and also by the exclusive localization of an unidentified species of *Maera*.



Map of the studied zone

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The food web of *Posidonia oceanica* beds around the Island of Ischia (Gulf of Naples -Italy) : a new trophic index

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The *Posidonia oceanica* system, characterized by high productivity and biomass partly exported to other coastal systems, supports a v. complex food web (Chessa et al., 1982). Many trophic studies can need to define the energy flow through the system: gut content analysis, calorific measurements, weight data, trophic groups analysis etc (ref. Kikuchi et Peres, 1977). The trophic behaviour of macrobenthic species sampled by a bottom trawl in different prairies analyzed and a new index is proposed as a feeding behaviour descriptor. The use of such an index allows for a precise description of feeding habits of a species and is useful in direct spat representations and multivariate analysis techniques.

Twelve samples were obtained in different *Posidonia* prairies around island of Ischia: the first 6 in winter, the others in summer collected at depths ranging from 15 to 25 meters with a bottom trawl with a 2 cm mesh, deep frozen and then fixed in 70% alcohol. The contents of each species in each sample were identified, quantified means of an arbitrary code ranging from 0 to 4 (0=absent; 4=v. abundant) and recorded in a matrix "species/food items". A multivariate analysis was performed on such data to define the principal component of the trophic model in the studied prairies (fig.1). It is a fact that generally, in such analysis, the observation points are ordered mainly on the first two axes, one of which can be mainly related to prey size while on the second the food items are ordered according to type -pl or animal- of prey. If the principal components of such trophic model are these, it is possible to redescribe the species on the basis these factors. The species were coded on the basis of a two digit parameter calculated as follows:

- the first digit represents the feeding habit -vegetarian carnivorous- of the species and is calculated by the formula:

$$\text{first digit} = (\sum V - \sum C) / \sum M \quad \text{where:}$$

V=abundance of vegetal items; C= abundance of carnivorous item M=abundance of each considered item. In such a way we can distinguish omnivorous organisms (first digit close to 0), pure carnivores (first digit close to -1), pure herbivores (first digit close to 1).

- The second digit represents the size scale of the prey and calculated by the formula:

$$\text{second digit} = \ln (\sum (PS_i \times N_i) / \sum M) \quad \text{where:}$$

PS_i= mean prey size (size measured in mm or in mg) of prey "i"; N_i= abundance; M= as defined previously. The use of logarithmic scale allows to discriminate microphagous organisms, eating prey items larger than 1 mm, (second digit negative) from macrophagous ones (second digit with positive values).

A total of 76 species were collected, while 26 food items were identified in the gut contents. A matrix was then compiled on the basis of the trophic code and it is possible to see that, plotting different species on a x-y system using the two digits of the code descriptors, a model very close to the multivariate one is obtained. In particular we observe that the principal clusters showed by a P.C. are reproduced in the new code based trophic model (fig. 1 and 2). Species can be then grouped into trophic categories as shown in fig. 1 and analyzed by common mathematical methods and such groups are far better descriptors than those generally defined only on an empirical basis. As one can observe, the two representations -using the 1 items and trophic index plotted- directly provide a single model of prairie, in which the importance of vegetarian and detritivorous organisms is assessed. Predators are represented only by a few species of fishes plotted in the second quadrant. The model obtained is used to classify the 96 considered species into calculated trophic categories as far as the trophic index calculated for each species be considered relatively constant of any ecosystem, while the position in a multivariate analysis model depends on the contribution of other observations considered.

The trophic index described can be considered a valid solution to compare data resulting from different investigations and may be used to define the differences observed in the feeding behaviour of a species studied on different temporal or spatial scales. The use of this index allows to get over the first ordination model, given that food items are generally ordered on the basis of size and qualitative components. Our research can thus be directed in the definition of measurable comparable trophic groups and food web models.

P.C.A.

Classification by two digits code

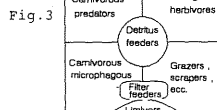
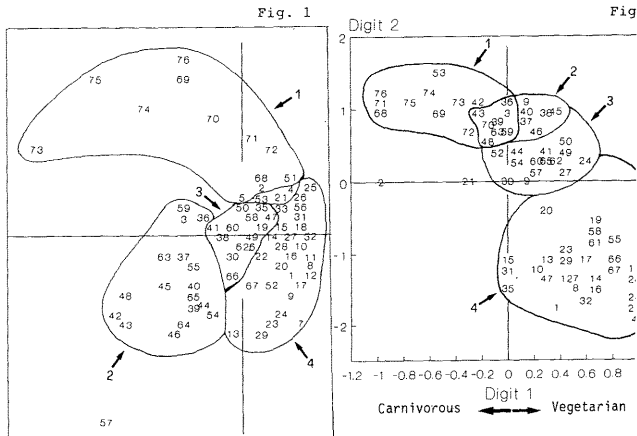


Fig. 3
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