

# Habitat diversity and postlarval benthic decapod crustacean assemblages in shallow waters of the Ría de A Coruña (NW Spain)

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**INTRODUCTION.** Distribution and abundance of postlarval phases of coastal benthic decapod crustaceans is related to multiple processes affecting both planktonic (larval supply and differential settlement) and benthic (differential mortality, habitat selection and density-dependence) stages. The high diversity of decapod assemblages in coastal ecosystems results from the interaction between the above processes and the existing high environmental variability and habitat diversity. This study is an attempt to characterize patterns of habitat use and distribution of juvenile and adult stages of benthic postlarval decapod species in shallow bottoms, identifying the main environmental variables affecting these patterns. The comparative capacity of different habitats to support juvenile and adult diversity was also characterized.

**ANALYSIS.** Generalized linear models (GLM) were employed to determine the weighted effect of the environmental conditions over the spatial patterns of the postlarval community. The best subsets procedure using the Akaike Information Criterion (AIC) was used to select the best model, and a normal log model was assumed. Null models were employed to extract common diversity patterns among variable combinations. Individual-based rarefaction curves (Gotelli and Graves, 1996), which generate comparative estimates of species number independently of the differences between sample size of the groups compared, were employed as a richness index. The probability of an interspecific encounter PIE (Hurlbert, 1971) was used as an evenness measure. The abundance levels for simulation were fixed according to the sample with lowest abundance to allow the comparison among samples. Both evenness and rarefaction were estimated using EcoSim software (Gotelli and Entsminger, 2001), which uses a Monte Carlo procedure, and 1000 replicate simulations were performed for each estimate.

**RESULTS.** Shallow bottoms of the Ría de A Coruña show a wide diversity of habitats and support a highly abundant and diverse community of decapods. Our results support the hypothesis that these areas constitute essential nursery habitats for benthic species.

A lower specific diversity was found for adults than for juveniles in nearly all cases (Fig. 1). This difference was more pronounced in Canabal (the most external sampling site). Evenness values for both stages were high (Average PIE > 0.65) in most sampling locations (Table 1). The most internal site (Portocobo) presented a higher homogeneity in richness and evenness values for adults than for juveniles.

The most abundant species occurred in higher density for juvenile over adult stages in most cases (Fig. 2). All adults belonged to species also found for juveniles. Three main patterns of juvenile and adult distribution were identified: species without adult presence in the sampled area, such as *Galathea strigosa*; species that remain in the same area, but with observable ontogenic habitat shifts, which is the case of *Eualus occultus* and species that keep similar habitat use patterns along the life cycle, e.g. *Pisidia longicornis*.

Although all the tested variables showed an important effect on the distribution of decapod species, variability in spatial patterns is mainly due to habitat type, which is a micro-scale operating factor. Meso-scale variability related to sites along the Ría, as well as other micro-scale factors, i.e. depth and wave exposure, have a lower effect on spatial distribution of post-recruits.

A similar pattern in the composition of parameters included in the most parsimonious models (GLM) was found between juveniles and adults on rocky bottoms (Table 2). In both cases habitat type was the most frequent variable in models for the most abundant species (87.5% for juveniles; 64.3% for adults), while wave exposure shows the lowest percentage of occurrence in models for both juveniles (56.2%) and adults (50%).

The GLM results for kelp holdfast samples showed a lower effect of all the factors on juvenile and adult patterns, with no observable differences among variables. A high proportion of the models obtained for this group of samples were not significant. This suggests that the influence of environmental conditions could be attenuated inside kelp holdfasts. These results, altogether with the ontogenic distribution patterns of the species found in holdfasts as well as the significant differences observed in diversity between them and rocky habitats, suggest that holdfasts are used as a temporary shelter rather than a nursery habitat.

	ROCKY BOTTOMS				KELP HOLDFASTS			
	Wave	Depth	Exposure	Species	Wave	Depth	Exposure	Species
<i>Pisidia longicornis</i>	1	1	1	1	1	1	1	1
<i>Athanas nitescens</i>	1	1	1	1	1	1	1	1
<i>Thorulus cranchii</i>	1	1	1	1	1	1	1	1
<i>Hippolyte varians</i>	1	1	1	1	1	1	1	1
<i>Plumulus spp.</i>	1	1	1	1	1	1	1	1
<i>Processa edulis</i>	1	1	1	1	1	1	1	1
<i>Xantho spp.</i>	1	1	1	1	1	1	1	1
<i>Pinnixa denticulata</i>	1	1	1	1	1	1	1	1
<i>Porcellana platycheles</i>	1	1	1	1	1	1	1	1
<i>Palaeomon spp.</i>	1	1	1	1	1	1	1	1
<i>Galathea strigosa</i>	1	1	1	1	1	1	1	1
<i>Eualus occultus</i>	1	1	1	1	1	1	1	1
Percentage	44.3	57.1	56.2	56.2	52.4	54.3	48.1	50.0

Table 2. GLM models for rocky bottoms and kelp holdfasts.



**SAMPLING.** A stratified random survey was carried out along the coast of the Ría de A Coruña (Galicia, NW Spain) during the months of July and August 1998, using a suction sampler operated by divers. Three sampling sites along the main axis of the Ría were selected to identify the meso-scale variability and an array of sampling locations were defined within each site according to the combination of three micro-scale variables: wave exposure, depth and habitat type. Four replicates were taken in each location.

WAVE EXPOSURE
Exposed
Sheltered

DEPTH
0: Intertidal
1: 3-5 m
2: 8-12 m

HABITAT TYPE
Sandy bottoms
Rocky bottoms
Flat rock surfaces covered with algae
Cobble and boulder pools
Kelp ( <i>Saccorhiza polyschides</i> ) holdfasts

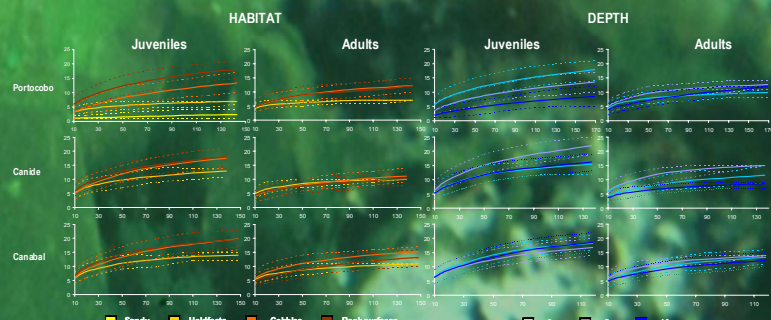


Fig. 1. Rarefaction diversity curves. The Y axis represents the expected richness for a given sample size (no. of individuals).

	Abund. level	Portocobo			Canabal			Canabal			
		Juveniles	Adults	PIE	Juveniles	Adults	PIE	Juveniles	Adults	PIE	
Sandy	143	0.021	0.00026	0.715	0	0	0	0.525	0.00137	0.762	0.00026
		0.586	0.00046	0.715	0	0	0	0.839	0.00018	0.689	0.00044
		0.451	0.00184	0.839	0	0	0	0.206	0.00155	0.693	0
Cobbles	138	0.739	0.00094	0.667	0.0106	0	0	0.861	0.0002	0.859	0
		0.856	0.00024	0.741	0.00048	0	0	0.795	0.0007	0.690	0.00086
		0.830	0.00008	0.844	0	0	0	0.700	0.00009	0.854	0.00036
Rocky	148	0.881	0.00026	0.851	0.00096	0	0	0.886	0.00017	0.816	0.00023
		0.902	0.00009	0.815	0.0001	0	0	0.870	0.00021	0.805	0.00021

Table 1. Evenness values.

Sandy bottom coverage was significant only in sheltered areas of one sampling site and therefore GLM was not fitted. The diversity and evenness found in this habitat were very low due to the dominance of the Paguridae. Early juveniles of this group showed an average density of 427 Ind/m<sup>2</sup> (Fig. 3), reaching 1227 Ind/m<sup>2</sup> at 10 m deep. Data found support the hypothesis that these bottoms are essential recruitment habitats for this group.

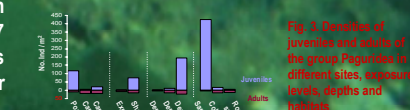


Fig. 2. Density of juveniles and adults of the group Paguridae in different sites, exposure levels, depths and habitats.

## LITERATURE CITED.

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