

# Spatial distribution of crustacean demersal assemblages in trawl grounds off Galician and Cantabrian shelf

**Esther Abad<sup>1</sup>, Julio Valeiras<sup>1</sup>, Eva Velasco<sup>2</sup>, Alberto Serrano<sup>3</sup>, Antonio Punzón<sup>3</sup>, Francisco Velasco<sup>3</sup>**

<sup>1</sup>Centro Oceanográfico de Vigo, Subida a Radio Faro, 50-52, 36290, Vigo, Spain [esther.abad@vi.ieo.es](mailto:esther.abad@vi.ieo.es)

<sup>2</sup>Centro Oceanográfico de Vigo, Casiada a Radio Paro, 30-32, 36290, Vigo, Spain [COSTER.vg](#)

<sup>3</sup>Centro Oceanográfico de Santander, Promontorio San Martín s/n, 39004, Santander, Spain



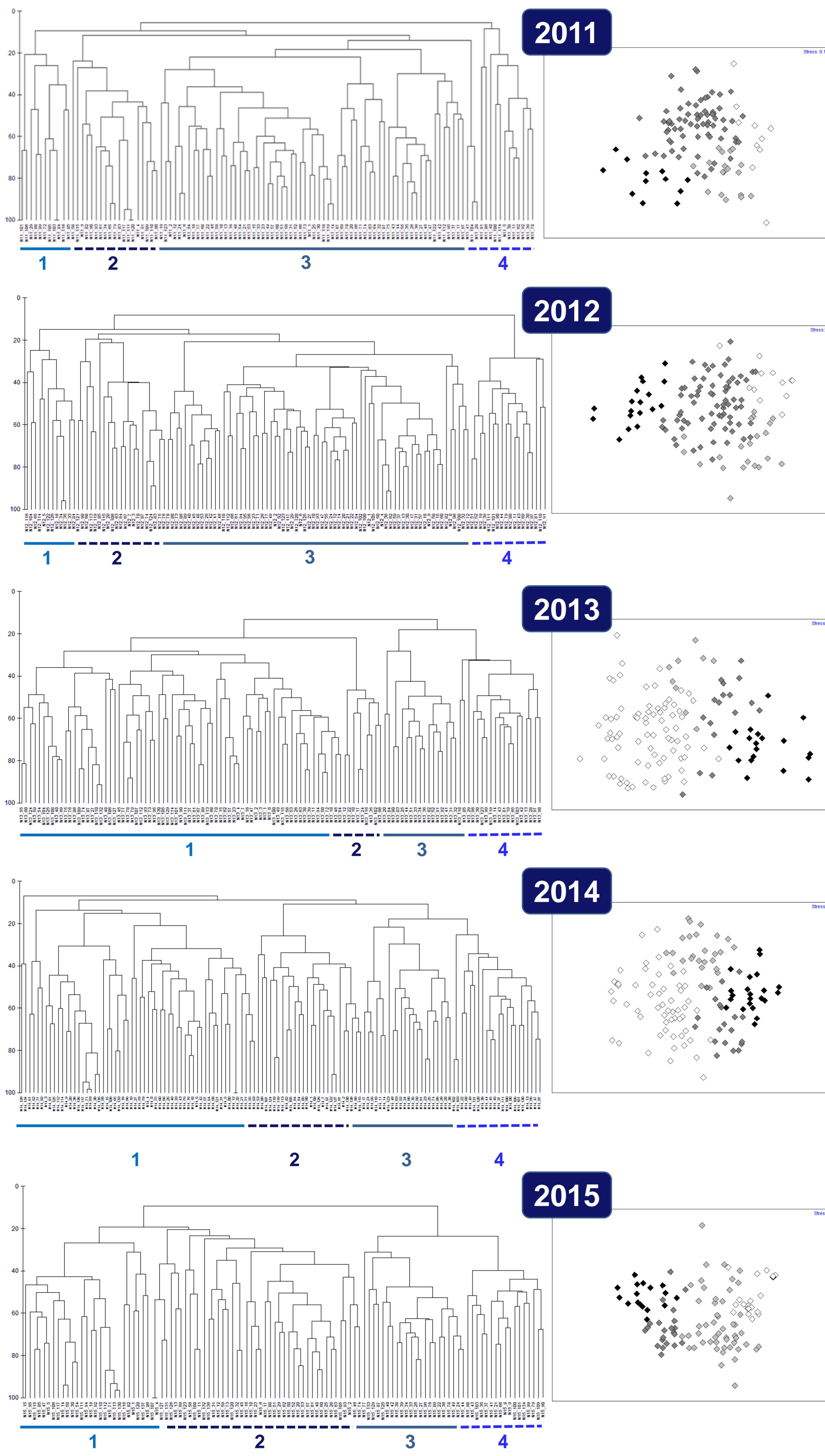
# SIEBM

# DO★MAR



# Introduction

In spite of crustaceans are the second group in abundance after fish in the Galician and Cantabrian continental shelf, studies on the spatial distribution patterns in the area are scarce. In the framework of the Ecosystem approach to fisheries management, it is necessary to study catches and discards and to increase the knowledge on the ecological aspects of these species at a regional scale.



## Figure 2. Cluster results for analyzed period (2011-2015) and MDS plots.

**Table 2. SIMPER results: species contributing most to similarity in the 4 cluster groups.**

Group 1		Group 2		Group 3		Group 4																	
Average similarity:	25.71	Average similarity:	24.01	Average similarity:	27.69	Average similarity:	38.92																
Species	Av. Abund	Av. Sim	Sim/SD	Contrib%	Cum. %	Species	Av. Abund	Av. Sim	Sim/SD	Contrib%	Cum. %	Species	Av. Abund	Av. Sim	Sim/SD	Contrib%	Cum. %	Species	Av. Abund	Av. Sim	Sim/SD	Contrib%	Cum. %
<i>Pagurus excavatus</i>	3.7	12.84	0.78	49.93	49.93	<i>Pagurus prideaux</i>	14.6	10.54	0.69	43.91	43.91	<i>Plesionika heterocarpus</i>	193.7	7.55	0.74	27.28	27.28	<i>Pasiphaea sivado</i>	58.4	8.55	1.59	21.98	21.98
<i>Macropodia longipes</i>	2.3	4.01	0.45	15.59	65.53	<i>Macropodia longipes</i>	2.5	5.52	0.54	22.98	66.89	<i>Chlorotocus crassicornis</i>	8.9	4.77	0.62	17.25	44.53	<i>Pagurus alatus</i>	12.2	5.77	1.21	14.83	36.81
<i>Chlorotocus crassicornis</i>	1.8	2.84	0.38	11.06	76.58	<i>Plesionika heterocarpus</i>	171.8	3.04	0.38	12.66	79.55	<i>Munida sarsi</i>	3410.3	4.61	0.45	16.66	61.19	<i>Polycheles typhlops</i>	7.8	4.78	1.17	12.27	49.08
<i>Pagurus prideaux</i>	8.8	1.66	0.28	6.44	83.02	<i>Chlorotocus crassicornis</i>	1.2	1.05	0.23	4.36	83.91	<i>Macropodia longipes</i>	1.7	2.57	0.45	9.29	70.47	<i>Plesionika martia</i>	19.6	4.46	0.76	11.46	60.54
<i>Plesionika heterocarpus</i>	2.7	1.31	0.28	5.1	88.12	<i>Pagurus excavatus</i>	0.6	0.8	0.21	3.34	87.25	<i>Pontophilus spinosus</i>	3.0	1.42	0.46	5.13	75.61	<i>Dichelopandalus bonnieri</i>	10.3	2.74	0.66	7.04	67.58
<i>Liocarcinus depurator</i>	0.4	0.52	0.14	2.02	90.14	<i>Munida intermedia</i>	1.2	0.5	0.18	2.08	89.33	<i>Pagurus excavatus</i>	1.4	1.16	0.29	4.19	79.8	<i>Philocheras echinulatus</i>	7.3	2.48	0.66	6.38	73.96






The figure displays a map of the North Sea region, specifically focusing on the Skagerrak and Kattegat areas. The map shows several ship trajectories represented by lines connecting diamond-shaped markers. These markers are color-coded according to four cluster groups defined in the legend:

- Cluster Group 1: White diamonds
- Cluster Group 2: Brown diamonds
- Cluster Group 3: Black diamonds with a white center
- Cluster Group 4: Solid black diamonds

Two inset images at the bottom center show examples of ships from Cluster Groups 1 and 3. The left inset shows a white ship (Cluster 1), and the right inset shows a dark-colored ship (Cluster 3). The background of the map is a satellite-style image of the sea surface.

## **Figure 1. Study area and spatial distribution of hauls classified by cluster groups**

## Material and methods

The study area includes the Galician and Cantabrian continental shelf and slope. Data come from IEO surveys carried out on the oceanographic vessels “Cornide de Saavedra” and “Miguel Oliver” from 2011 to 2015 in autumn. A total of 614 trawling hauls were sampled using an otter trawl “Baca” type. The sampling unit was a 30-minute haul during daytime at a speed of 3 knots in a randomly stratified scheme by depth. Crustacean abundance data were fourth-root transformed and subjected to cluster analysis using the Bray-Curtis similarity index to characterize crustacean assemblages. SIMPER analysis was used to identify species typifying each group and a non-metric multidimensional scaling analysis was carried out.

**Table 1. Abundance (Numbers by haul  $\pm$  SD) of most abundant species (more than 50 individuals) by clusters groups.**

Species	Cluster groups			
	1	2	3	4
<i>Alpheus glaber</i>	0.28 ± 0.01	0.10 ± 0.00	0.18 ± 0.00	0.13 ± 0.00
<i>Anapagurus laevis</i>	0.08 ± 0.00	0.09 ± 0.00	0.18 ± 0.01	0.11 ± 0.01
<i>Aristeomorpha foliacea</i>	0.00 ± 0.00	0.00 ± 0.00	0.00 ± 0.00	0.65 ± 0.03
<i>Chlorotocus crassicornis</i>	1.70 ± 0.02	1.22 ± 0.02	8.93 ± 0.08	0.16 ± 0.01
<i>Dichelopandalus bonnieri</i>	0.02 ± 0.00	0.10 ± 0.00	11.05 ± 0.16	10.34 ± 0.43
<i>Ephyrina figureirai</i>	0.00 ± 0.00	0.00 ± 0.00	0.04 ± 0.00	1.02 ± 0.08
<i>Galathea sp.</i>	0.29 ± 0.01	0.18 ± 0.00	0.21 ± 0.01	0.03 ± 0.00
<i>Geryon trispinosus</i>	0.00 ± 0.00	0.00 ± 0.00	0.00 ± 0.00	3.23 ± 0.13
<i>Goneplax rhomboides</i>	0.88 ± 0.03	0.18 ± 0.00	0.29 ± 0.01	0.86 ± 0.02
<i>Inachus dorsettensis</i>	0.36 ± 0.01	0.20 ± 0.01	0.13 ± 0.00	0.00 ± 0.00
<i>Inachus leptochirus</i>	0.03 ± 0.00	0.37 ± 0.02	0.00 ± 0.00	0.01 ± 0.00
<i>Liocarcinus depurator</i>	0.33 ± 0.01	0.39 ± 0.02	3.16 ± 0.10	0.02 ± 0.00
<i>Macropipus tuberculatus</i>	0.39 ± 0.01	0.50 ± 0.02	2.51 ± 0.04	0.14 ± 0.01
<i>Macropodia longipes</i>	2.17 ± 0.03	2.43 ± 0.04	1.70 ± 0.02	0.73 ± 0.02
<i>Munida intermedia</i>	0.49 ± 0.01	1.16 ± 0.06	11.02 ± 0.50	0.14 ± 0.01
<i>Munida iris</i>	0.11 ± 0.00	0.93 ± 0.07	27.79 ± 1.54	0.02 ± 0.00
<i>Munida sarsi</i>	0.25 ± 0.01	1.17 ± 0.06	3410.32 ± 64.49	0.91 ± 0.03
<i>Nephrops norvegicus</i>	0.08 ± 0.00	0.05 ± 0.00	1.19 ± 0.02	4.23 ± 0.12
<i>Pagurus alatus</i>	0.22 ± 0.01	0.97 ± 0.03	1.95 ± 0.04	12.20 ± 0.23
<i>Pagurus bernhardus</i>	0.78 ± 0.03	0.00 ± 0.00	0.00 ± 0.00	0.00 ± 0.00
<i>Pagurus excavatus</i>	3.39 ± 0.03	0.56 ± 0.01	1.36 ± 0.02	0.05 ± 0.00
<i>Pagurus prideaux</i>	8.13 ± 0.17	14.48 ± 0.40	11.36 ± 0.24	0.56 ± 0.03
<i>Parapagurus pilosimanus</i>	0.00 ± 0.00	0.00 ± 0.00	0.00 ± 0.00	0.61 ± 0.03
<i>Pasiphaea multidentata</i>	0.00 ± 0.00	0.00 ± 0.00	0.00 ± 0.00	0.70 ± 0.03
<i>Pasiphaea sivado</i>	0.09 ± 0.00	0.16 ± 0.00	10.16 ± 0.26	58.35 ± 1.56
<i>Philoceras echinulatus</i>	0.09 ± 0.00	0.13 ± 0.01	1.65 ± 0.03	7.32 ± 0.17
<i>Plesionika heterocarpus</i>	2.48 ± 0.06	170.43 ± 6.24	193.70 ± 3.08	0.11 ± 0.01
<i>Plesionika martia</i>	0.00 ± 0.00	0.00 ± 0.00	0.05 ± 0.00	19.58 ± 0.28
<i>Polycheles typhlops</i>	0.01 ± 0.00	0.01 ± 0.00	0.14 ± 0.01	7.80 ± 0.17
<i>Pontophilus norvegicus</i>	0.00 ± 0.00	0.00 ± 0.00	0.09 ± 0.00	2.01 ± 0.06
<i>Pontophilus spinosus</i>	0.41 ± 0.01	0.23 ± 0.01	3.00 ± 0.04	4.03 ± 0.16
<i>Processa canaliculata</i>	0.10 ± 0.00	0.31 ± 0.01	0.38 ± 0.00	0.64 ± 0.02
<i>Psathyrocaris infima</i>	0.01 ± 0.00	0.01 ± 0.00	0.00 ± 0.00	0.73 ± 0.03
<i>Sergia robusta</i>	0.00 ± 0.00	0.00 ± 0.00	0.00 ± 0.00	1.15 ± 0.04
<i>Solenocera membranacea</i>	0.22 ± 0.01	0.28 ± 0.01	1.05 ± 0.01	5.23 ± 0.14
<i>Systellaspis debilis</i>	0.04 ± 0.00	0.02 ± 0.00	0.07 ± 0.00	3.97 ± 0.18
Total	23.42 ± 0.00	196.65 ± 0.00	3703.70 ± 0.01	147.76 ± 0.09

## Results

## 1 Shallow shelf assemblage with a medium depth of 135 m typical

2. Shelf assemblage with a medium depth of 164 m typified by the pagurid *Pagurus prideaux*.
  3. Upper slope assemblage with a medium depth of 230 m typified by the pandalid *Plesionika heterocarpus*.
  4. Middle slope assemblage with a medium depth of 536 m typified by the pasiphaeid *Pasiphaea sivado*.

**Depth appears to be the most important factor in structuring assemblages, just like it has been described in the study area for other associations.**