

A network diagram consisting of white dots connected by thin white lines. The dots are arranged in a roughly rectangular shape with some internal connections. Three of the dots are highlighted with concentric circles: one on the left side, one on the right side, and one at the bottom center.

ITA IN THE BAY OF BISCAY FOR CONTRASTING HABITATS

LUCA MARSAGLIA

Index

Introduction: ITA in BoB

Selection of habitats: Methodology and results

Questions and methods

Data analysis: results

Integrated Trend Analysis (ITA)

- Tool used in Integrated Ecosystem Assessment (IEA)
- Summarize changes occurred in recent decades
- Find connections between physical, ecological, anthropogenic variables
- Done for the the whole BoB

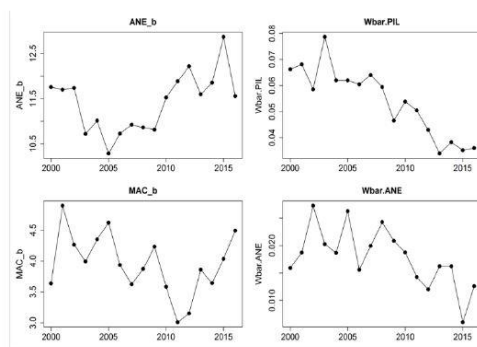
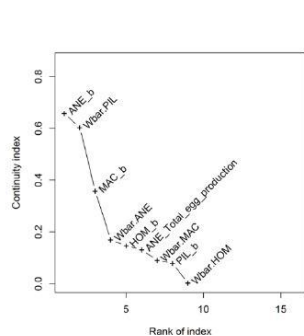


Figure 2.2.3: Results of the MAFA realized on the small pelagic fish compartment. Left: ranking of available variable according to their continuity index. Right: time-series of the most continuous variables. ANE_b is anchovy biomass, Wbar_PIL is mean weight of sardine, MAC_b is mackerel biomass, Wbar_ANE is mean weight of anchovy.

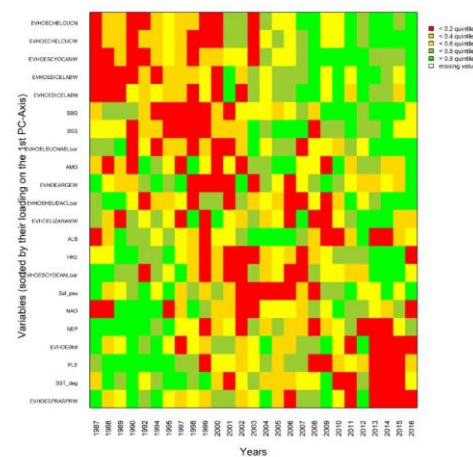


Figure 2.2.2: Traffic light plot of the temporal development of the Bay of Biscay ecosystem time-series. Variables are transformed to quantiles, colour coded (red-high values, green-low values) and sorted in numerically descending order according to their loadings of the first principle component.

Analyze contrasting areas in BoB

- **First time, an analysis for smaller / contrasting areas in BoB is performed**
- **Consider the BoB as an « averaged » ecosystem => might be problematic when we want to assess the connections with a human activity at a local scale (e.g. wind farms)**
- **The analysis could shed light on important characteristics of some areas in BoB**
- **There could be differences or shifts in trends across areas of the BoB that do not emerge from analysis over the all BoB**

ITA in BoB for contrasting habitats

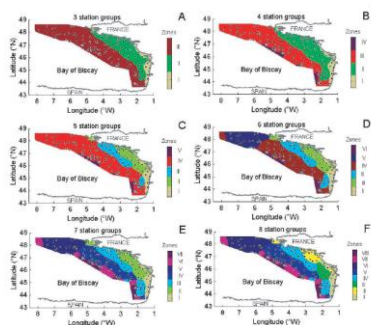


Fig. 6. Maps of the different habitat zones for the following hierarchical levels: 3(A), 4(B), 3(C), 4(D), 3(E) and 4(F). Each colour represents the median depth according to colour scale bar of Fig. 5. Dark, violet colour indicates the deeper sites (median depth > 175 m).

Defining contrasting habitats

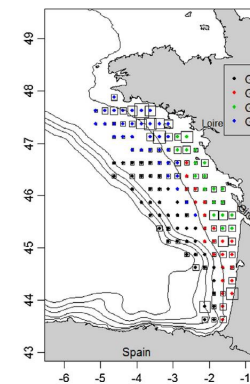
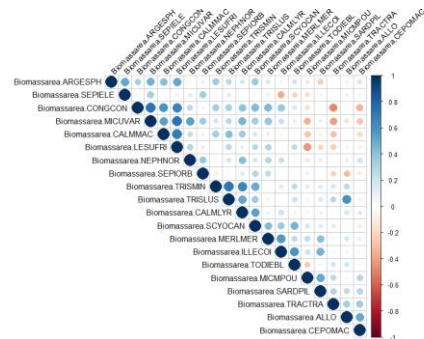
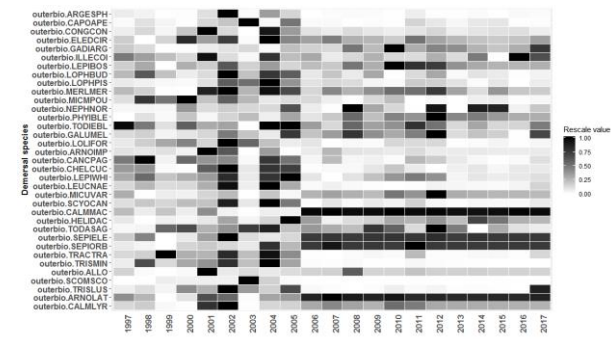


Fig. 5. Map of Bay of Biscay scapes as identified by applying MFA on all ecosystem compartments. Map of grid cell clusters and their variability in time over the years 2009-2014. Colors are that of the clusters (Fig. 3). Squares are proportional to the variability in time at each grid point (inset). The insets are 100, 200, 500, 1000, 2000 m.



Data analysis within areas



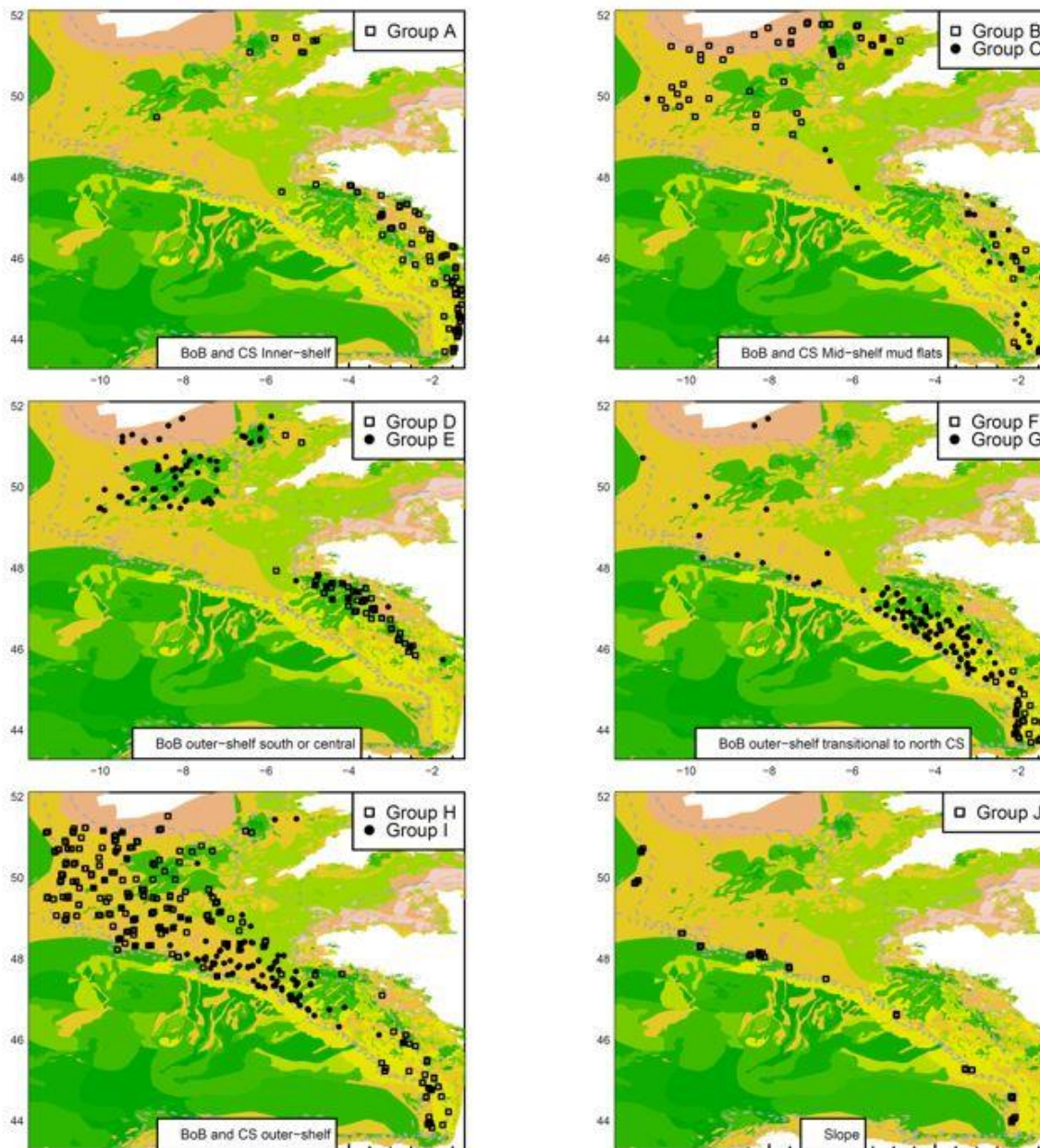


Table 2 - Summary of habitats and benthic communities (A to J) as defined from clustering process for combined sampled years for the Bay of Biscay (BoB) and the Celtic Sea (CS), indicate mean species richness for all years, total number of observed stations.

Habitat Name (groups)	Main location	Median depth	Habitat summary
Inner Shelf (A)	BoB and CS	A:38m	Various shallow coastal habitats in the vicinity of semi-enclosed bays or under estuarine influence
Inner/Mid Shelf (B,C)	BoB and CS	B:78m C:81m	Various soft bottom habitats in shallow coastal areas
Mid-shelf Mud flats (D,E)	North BoB and Central CS	D:103m, E:107m	Mid-shelf muddy habitats
Outer shelf south (F)	Southern BoB	106m	Sandy dominated habitat in the intermediate area surrounding the shelf break
Outer shelf central (G)	BoB	138m	patchy muddy-sands to sandy habitats, transitional between the slope and the mid-shelf mud flats
Outer shelf transitional (I)	CS and BoB "transition"	146m	Sandy dominated habitat in the intermediate area surrounding the shelf break
Outer shelf north (H)	BoB south and north CS	143m	Sandy dominated habitat in the intermediate area surrounding the shelf break
Slope (J)	CS and BoB	489m	Muddy grounds on the upper slope of the shelf break

CS: Celtic Sea - BoB: Bay of Biscay

Comprehensive geo-referenced data over large sea areas and spatio-temporal sampling scale. These data series, we applied a generic procedure to define ecosystem spatial units in the Bay of Biscay. The multi-table analysis method known as MFA, thus separating in the analysis the spatial patterns across ecosystem components. The MFA results are consistent over the years together with a map in agreement with sub-regional production system assessments. Also, it highlights the importance of ecosystem description, assessment and man-

Figure 3 - Assemblages, The Inner/mid shelf habitats (groups A, B and C), Mid-shelf mud flats (groups D and E), Outer shelf habitats (F : south BoB, G: central, I: Transitional, H: north CS mainly and south BoB)

Se

Souissi et al

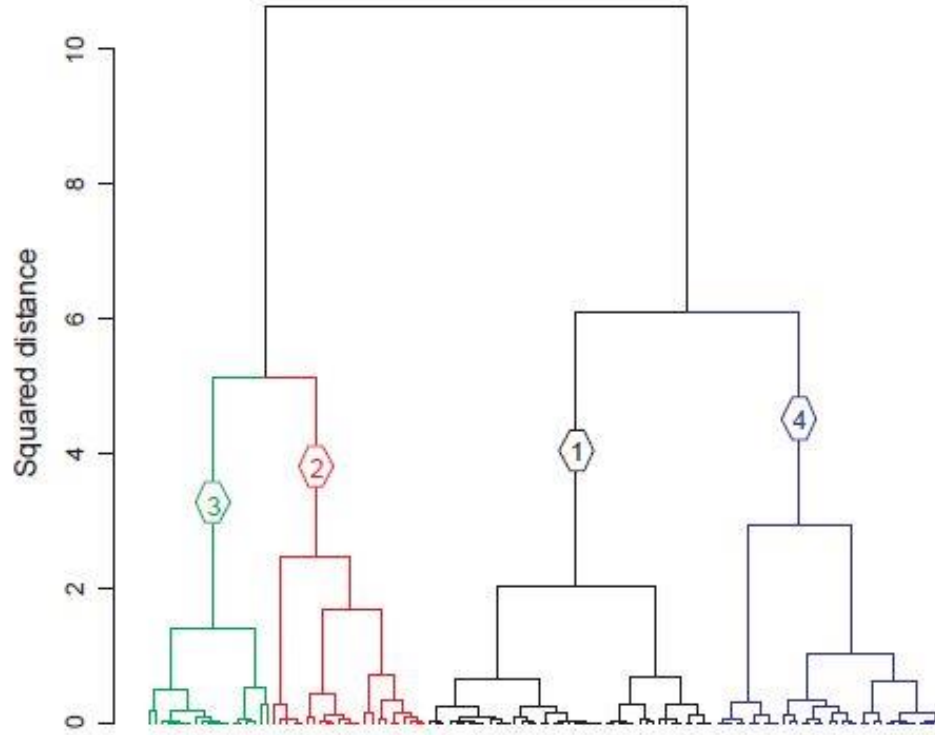
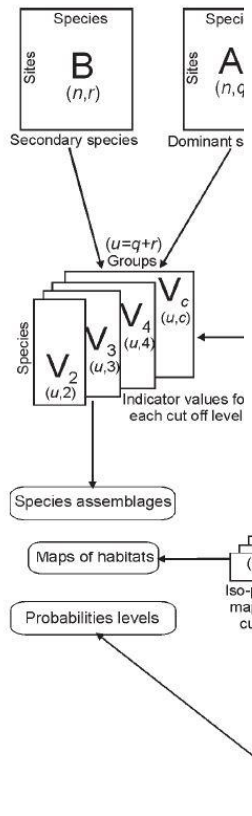


Fig. 3. Cluster dendrogram of grid cells obtained by their hierarchical clustering in the MFA space. The distance considered is the squared Euclidian distance between average grid cell points in the MFA plane made of the first two principal components. Four clusters were retained, which are shown in different colors.

time series was from 2009
ms of variable are defined

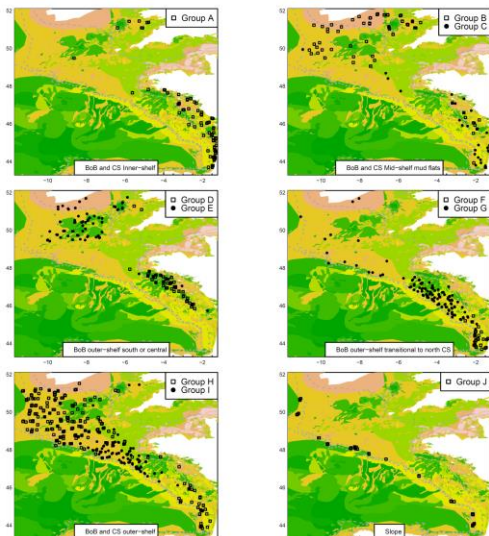
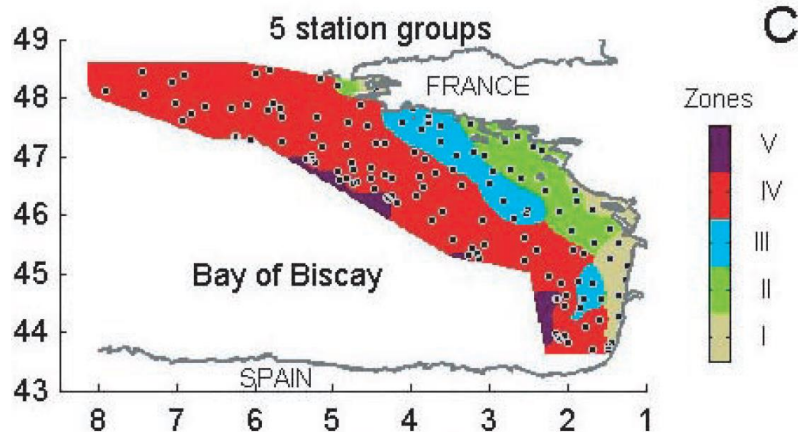
Selection
No
Yes
Yes
No
Yes
No
Yes
Yes
No
No
No
Yes
No
No
Yes
Yes
Yes
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No
No
Yes
No
Yes
Yes

ent to
id average
g to map
ecosystem structure

Fig. 2. Diagram of the analysis steps for the multivariate mapping method. The different steps explaining the connections between the input matrices A (n; sites, q; dominant species) and B (n; sites, r; secondary species) and the final resulting map of fish habitats and indicator species for each hierarchical level. The different steps of the diagram are detailed in Materials and Methods section.

Selection of habitats – Studies used

Souissi et al. 2001



Laffargue – EVHOE
(Unpublished)

Table 2 - Summary of habitats and benthic communities (A to J) as defined from clustering process for combined sampled years for the Bay of Biscay (BoB) and the Celtic Sea (CS), indicate mean species richness for all years, total number of observed stations.

Habitat Name (group)	Main location	Median depth	Habitat summary
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Mud flats (D, E)	North BoB and Central CS	D: 103m, E: 107m	Mid-shelf muddy habitats
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CS: Celtic Sea - BoB: Bay of Biscay

Petitgas et al. 2017

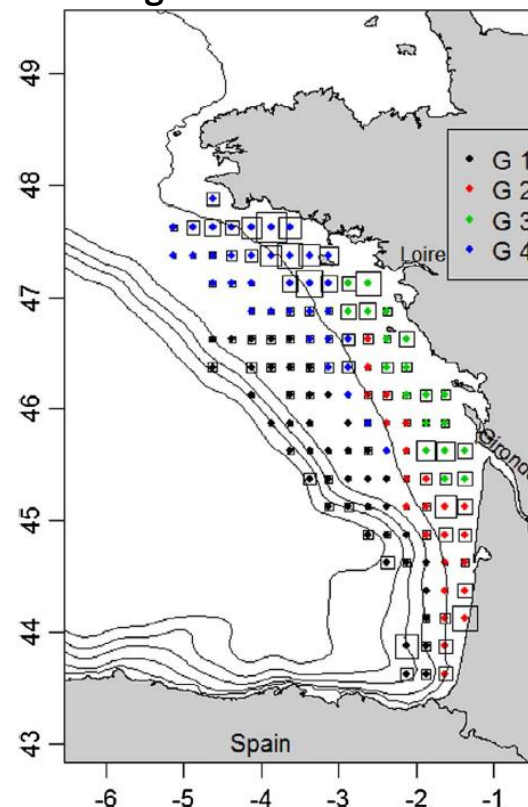


Fig. 5. Map of Bay of Biscay seascapes as identified by applying MFA on all ecosystem compartments. Map of grid cell clusters and their variability in time over the years 2009–2014. Colors are that of the clusters (Fig. 3). Squares are proportional to the variability in time at each grid point (inertia). The isobaths are 100, 200, 500, 1000, 2000 m.

There will be...

**Areas in which benthic and pelagic communities
are well structured and maintain their structure
throughout the whole year**

Selection of habitats – Overlap of studies strata

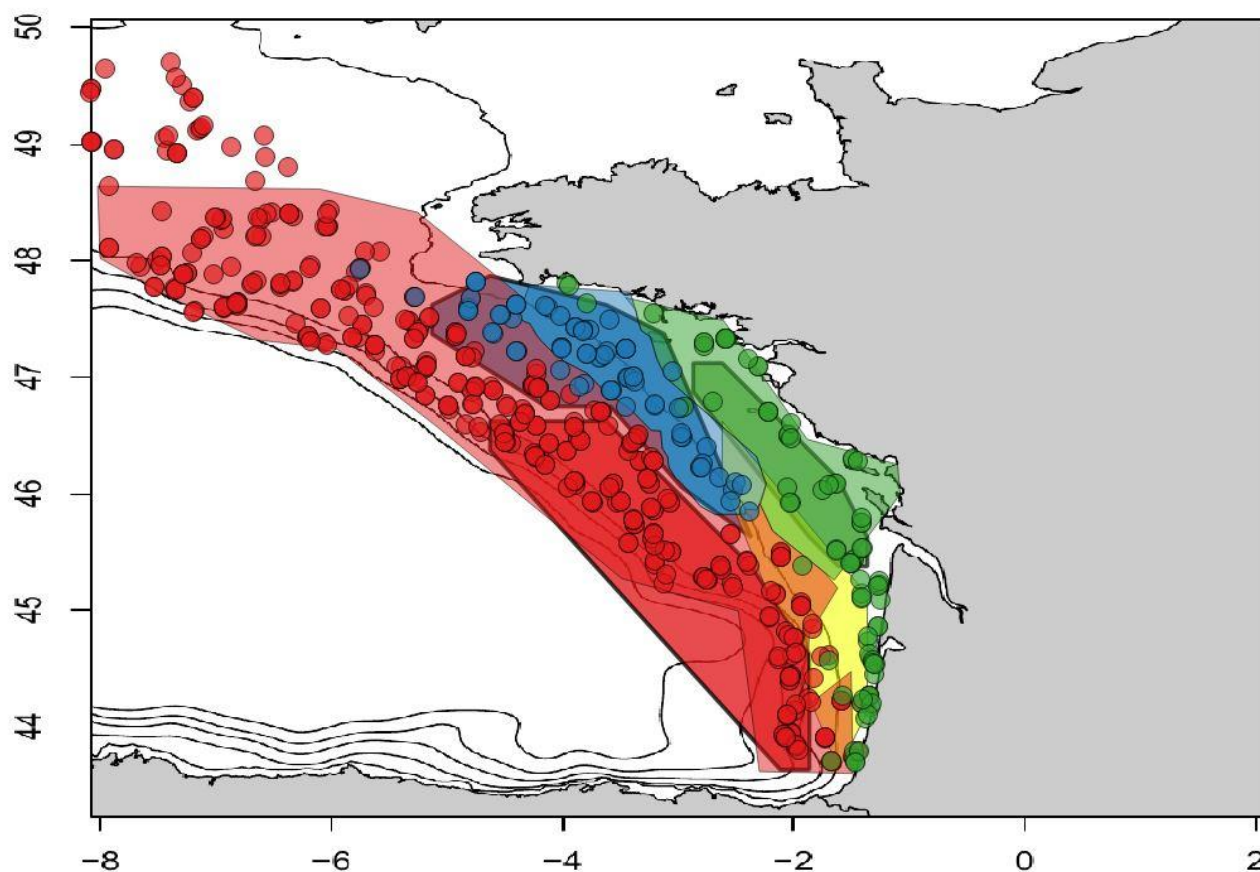


Figure 1 shows the three different strata from the selected studies Souissi et al 2001 as the non-bordered polygons, Petitgas 2017 as the bordered areas and Pascal 8 EVHOE as the points. Green represents coastal riverine – influenced areas, blue La Grande Vasière, red the outer shelf and tiled-yellow the southern area of the bay.

Selection of habitats - Overlap of studies strata

- **Overlap:**
 1. Coastal (Green)
 2. La Grande Vasière (Blue)
 3. Outer shelf (Red)

- Time of year + species depth range to cause mismatch in La Grande Vasière?

- Severe mixing in southern BoB area?

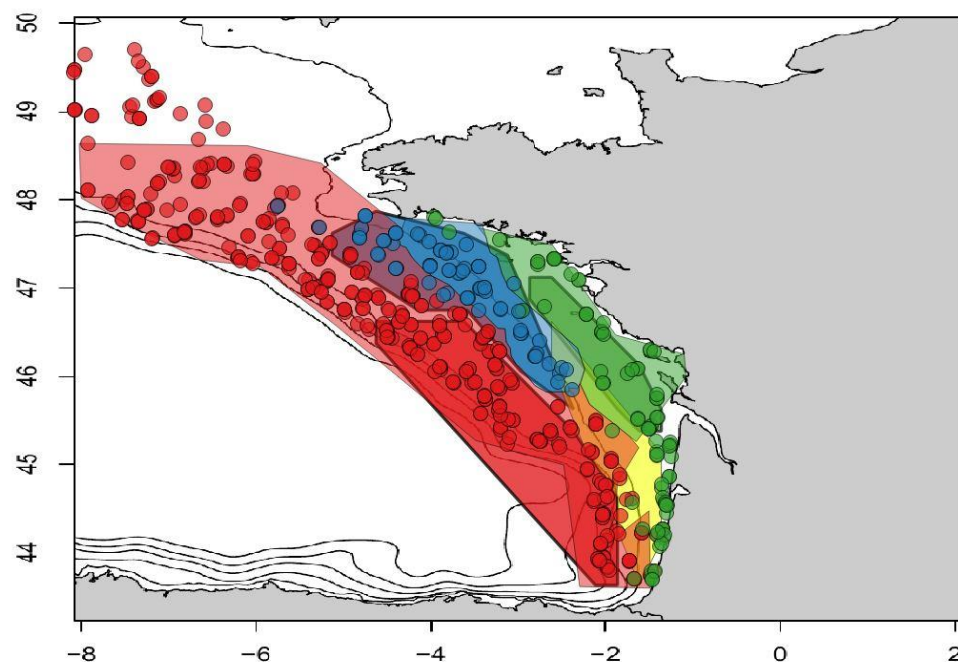
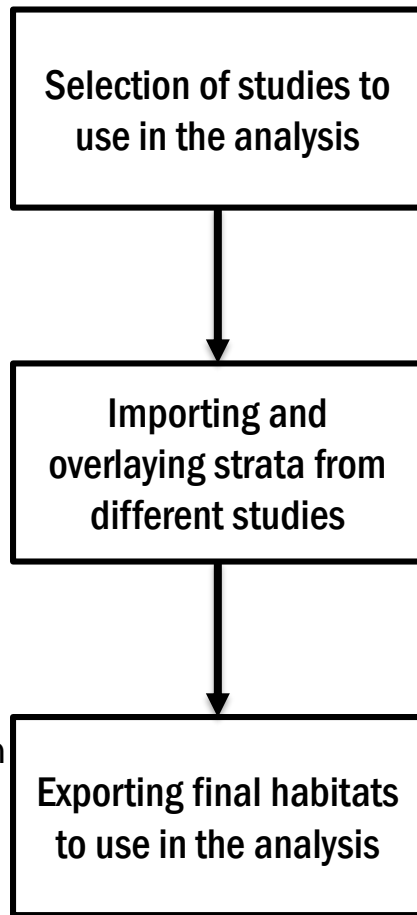


Figure 1 shows the three different strata from the selected studies Souissi et al 2001 as the background map, Petigas 2017 as the bordered areas and Pascal 8 EVHOE as the points. Green represents coastal riverine – influenced areas, blue La Grande Vasière, red the outer shelf and tiled-yellow the southern area of the bay.

Selection of habitats - Methodology

- 20 studies on BoB demersal and pelagic distribution reviewed
 - Preference to studies presenting spatial units
-
- Qgis csv transformation into points or polygons and georeferencing
 - Different layers overlaid each other
-
- Clip Function to export only areas in common between polygons



A new multivariate mapping method for studying species assemblages and their habitats: example using bottom trawl surveys in the Bay of Biscay (France)

Sarah Socolar¹, Frédéric Buisson, Rodolphe Deshayes, Jean-Benoît Audebert, Anne Claire Cabelguen, Fabien Blanchard & Jean-Christophe Poulard

SARSIA

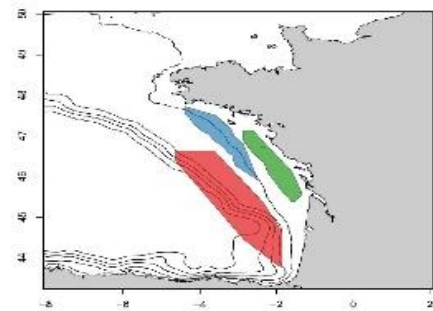
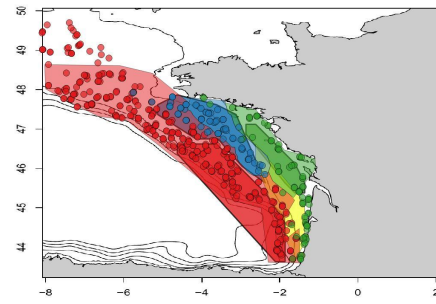
Abstract: Bottom trawl surveys in the Bay of Biscay (France) have provided a wealth of data on the distribution of demersal and pelagic species. This data has been used to study the spatial structure of species assemblages and their habitats. The aim of this study was to develop a new multivariate mapping method for studying species assemblages and their habitats. This method consists in combining the results of two different multivariate analyses: a correspondence analysis (CA) and a principal component analysis (PCA). The CA is used to study the spatial structure of species assemblages and the PCA is used to study the spatial structure of their habitats. The results of the two analyses are then combined to produce a new multivariate mapping method. This method allows to study the spatial structure of species assemblages and their habitats in a more comprehensive way than the traditional methods. The results of this study show that the new method is more powerful than the traditional methods. It allows to identify the spatial structure of species assemblages and their habitats in a more comprehensive way than the traditional methods. The results of this study show that the new method is more powerful than the traditional methods. It allows to identify the spatial structure of species assemblages and their habitats in a more comprehensive way than the traditional methods.

Ecosystem spatial structure revealed by integrative survey data

Patricio Pardo¹, María Teresa², Cristina Domínguez³, Antonio Aguilera⁴, Mónica Muñoz⁵, Julián Rodríguez⁶, Miquel Dolégal⁷

1. Universidad de Cantabria, 2. Universidad de Cantabria, 3. Universidad de Cantabria, 4. Universidad de Cantabria, 5. Universidad de Cantabria, 6. Universidad de Cantabria, 7. Universidad de Cantabria

Abstract: The spatial structure of ecosystems is a key factor to understand their functioning and evolution. This structure is determined by the spatial distribution of species and their interactions. The aim of this study was to reveal the spatial structure of ecosystems using integrative survey data. This data includes information on species distribution and their interactions. The results of this study show that the spatial structure of ecosystems is determined by the spatial distribution of species and their interactions. The results of this study show that the spatial structure of ecosystems is determined by the spatial distribution of species and their interactions.



Selection of habitats – Final habitats

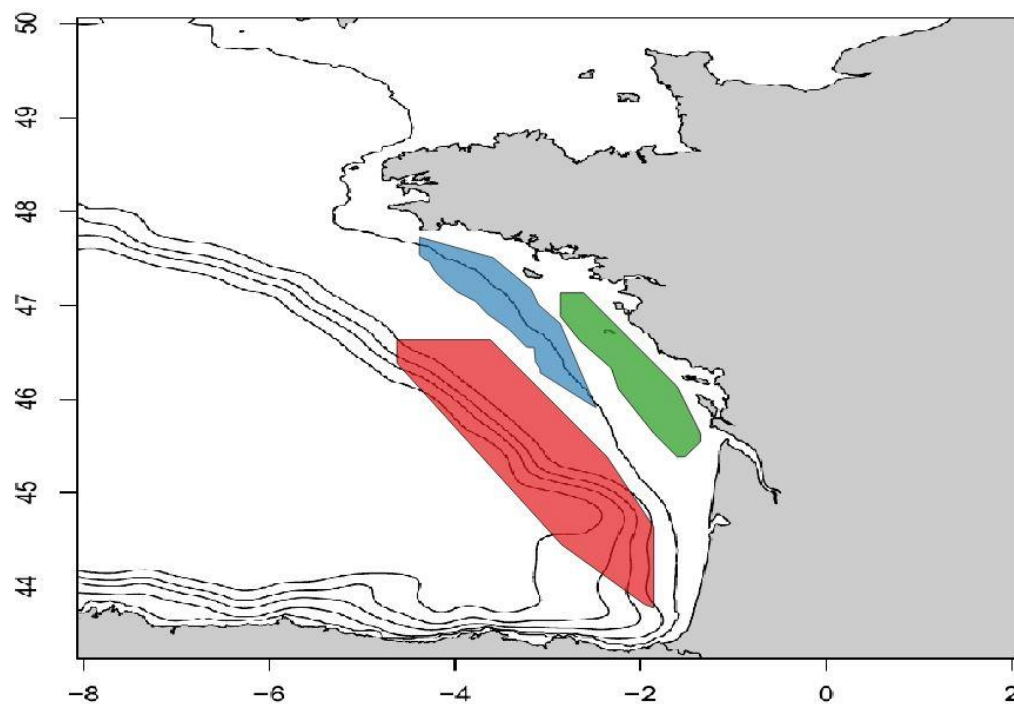


Figure 2 shows the three final core habitat for the Bay of Biscay. Green coastal riverine – influenced areas, blue La Grand Vasiere, and red the outer shelf.

Specific questions and methods

Questions	Method
<p>How to characterize the different areas in regards to community composition?</p>	<p>β diversity analysis Species Biomass Histogram</p>
<p>Do species show different trends in different areas in recent decades?</p>	<p>Time series</p>

Data used: EVHOE (Demersal species) + PELGAS (Pelagic species)

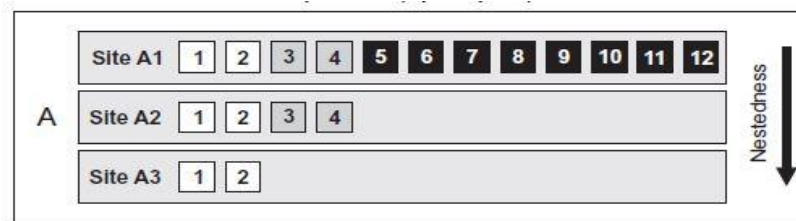
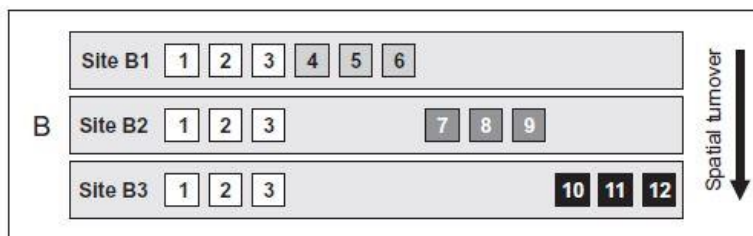
β diversity analysis – Baselga 2010

Overall β Diversity

$$\frac{\left[\sum_{i < j} \min(b_{ij}, b_{ji}) \right] + \left[\sum_{i < j} \max(b_{ij}, b_{ji}) \right]}{2 \left[\sum_i S_i - S_T \right] + \left[\sum_{i < j} \min(b_{ij}, b_{ji}) \right] + \left[\sum_{i < j} \max(b_{ij}, b_{ji}) \right]}$$

Species Turnover

Species Nestedness



$$\frac{\left[\sum_{i < j} \min(b_{ij}, b_{ji}) \right]}{\left[\sum_i S_i - S_T \right] + \left[\sum_{i < j} \min(b_{ij}, b_{ji}) \right]}$$

$$\frac{\left[\sum_{i < j} \max(b_{ij}, b_{ji}) \right] - \left[\sum_{i < j} \min(b_{ij}, b_{ji}) \right]}{2 \left[\sum_i S_i - S_T \right] + \left[\sum_{i < j} \min(b_{ij}, b_{ji}) \right] + \left[\sum_{i < j} \max(b_{ij}, b_{ji}) \right]} \times \frac{\sum_i S_i - S_T}{\left[\sum_i S_i - S_T \right] + \left[\sum_{i < j} \min(b_{ij}, b_{ji}) \right]}$$

where S_i is the total number of species in site i and S_T is the total number of species in all sites. B_{ij} and B_{ji} represent the number of species present only in site i and j respectively.

β diversity analysis

Demersal species (EVHOE)

Pelagic species (PELGAS)

Species
Turnover

β_{sim}	Coast	Gran Vasière
Gran Vasière	0.24	
Outer Shelf	0.35	0.16

β_{sim}	Coast	Gran Vasière
Gran Vasière	0	
Outer Shelf	0	0.083

Species
Nestedness

β_{sne}	Coast	Gran Vasière
Gran Vasière	0.024	
Outer Shelf	0.13	0.20

β_{sne}	Coast	Gran Vasière
Gran Vasière	0.043	
Outer Shelf	0.043	0

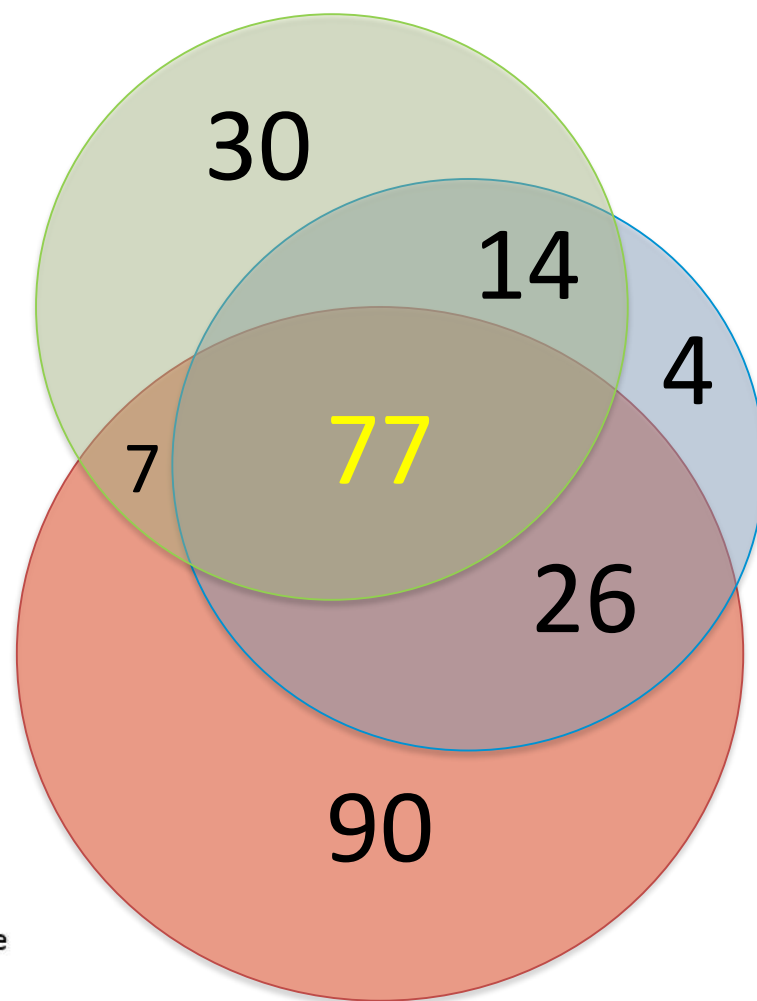
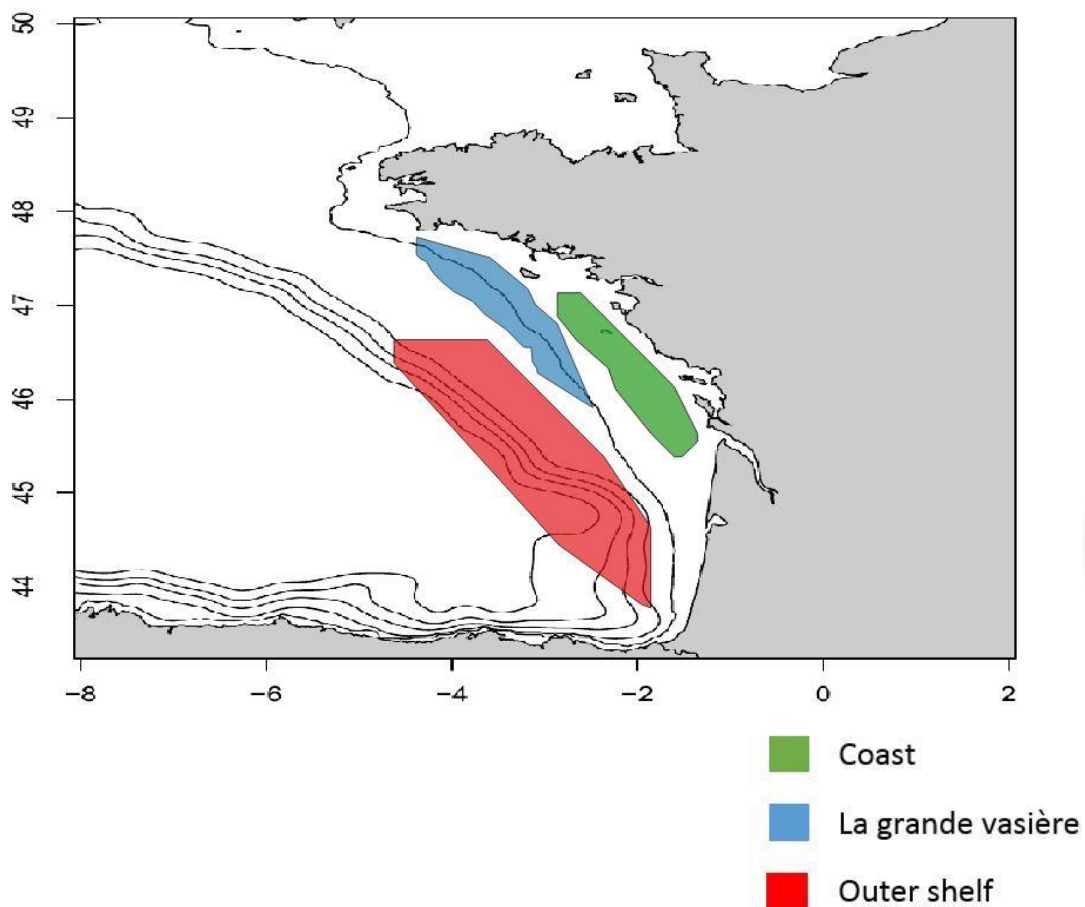
Overall β
Diversity

β_{sor}	Coast	Gran Vasière
Gran Vasière	0.25	
Outer Shelf	0.49	0.36

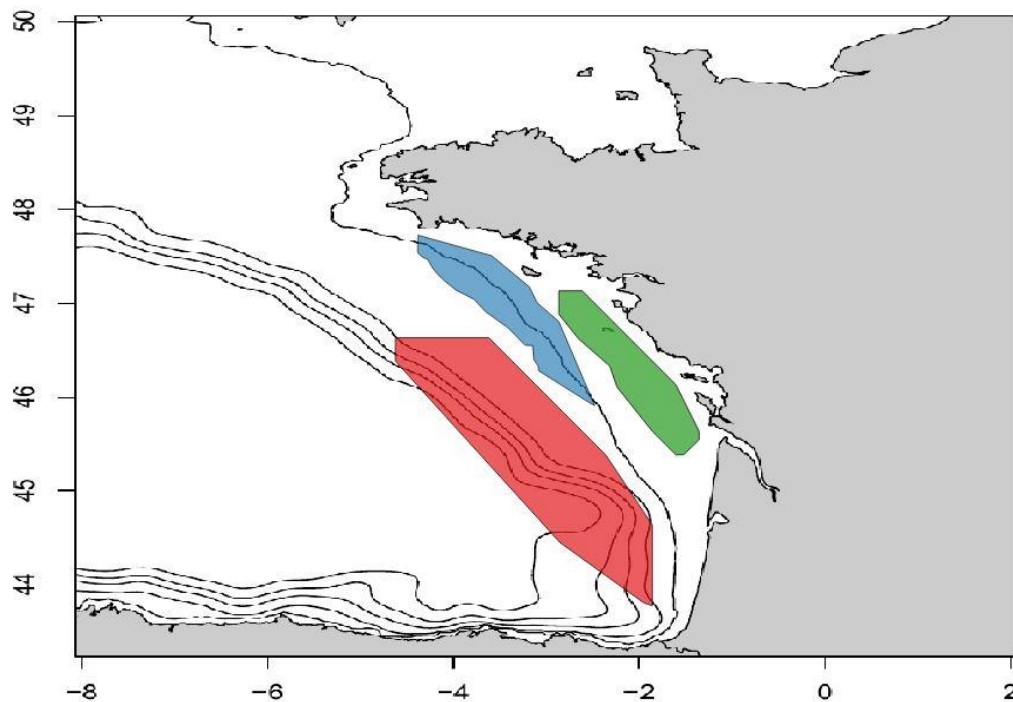
β_{sor}	Coast	Gran Vasière
Gran Vasière	0.043	
Outer Shelf	0.043	0.083

- The coast has a high species turnover with both La Gran Vasiere and the Outer Shelf. Coast has species that are not present in neither Outer and Gran Vasiere, and Vice Versa.
- The diversity between Gran Vasiere and Outer is a quite even mix of species turnover and nestedness.

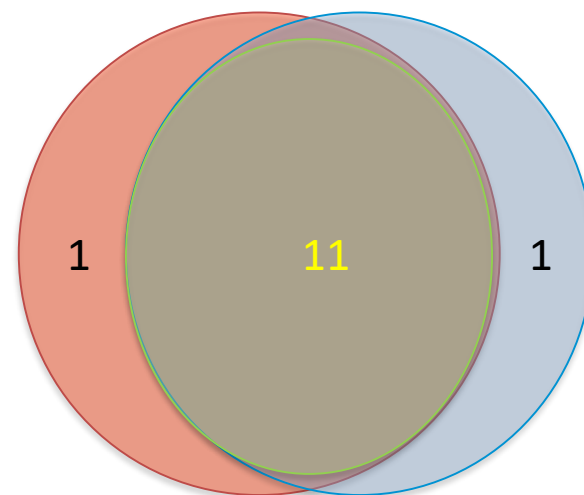
β diversity analysis - EVHOE



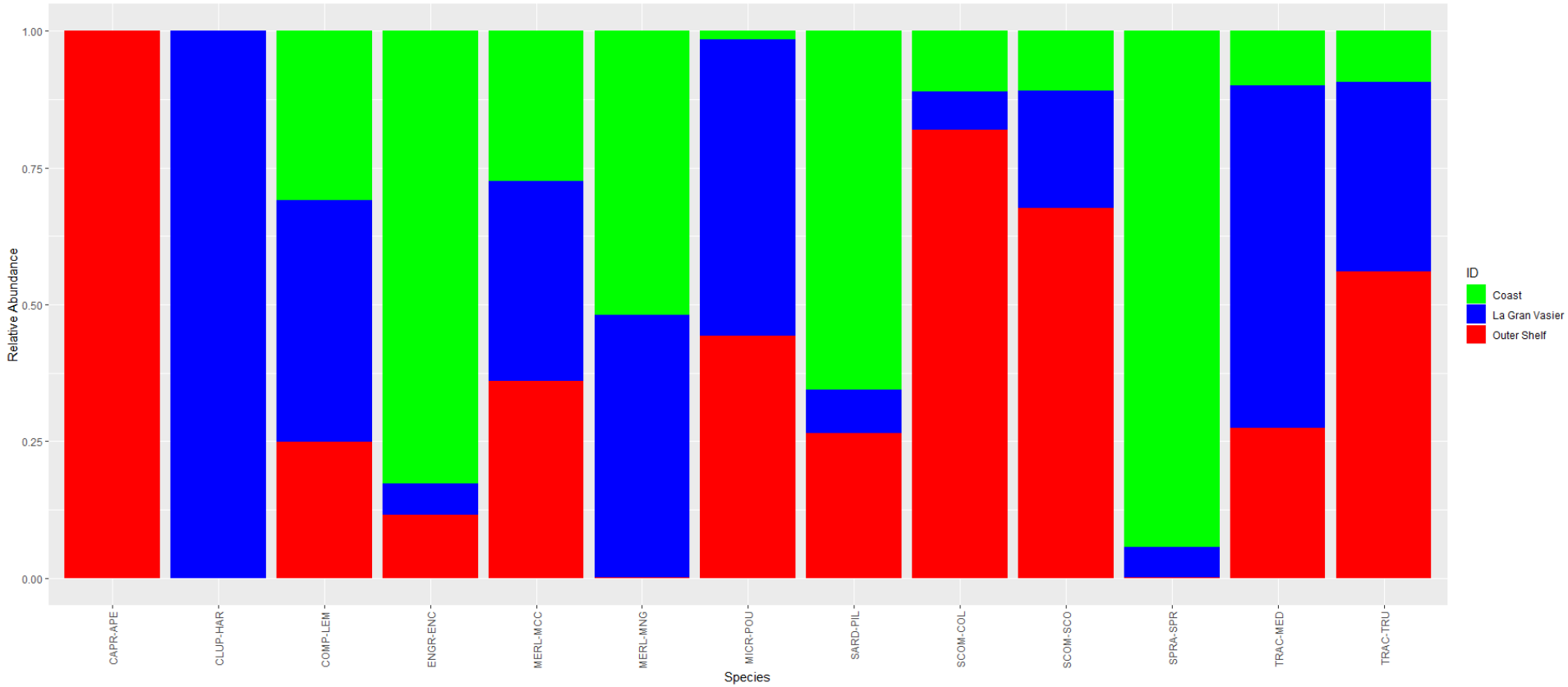
β diversity analysis - PELGAS



- Coast
- La grande vasière
- Outer shelf



Species biomass histogram- PELGAS

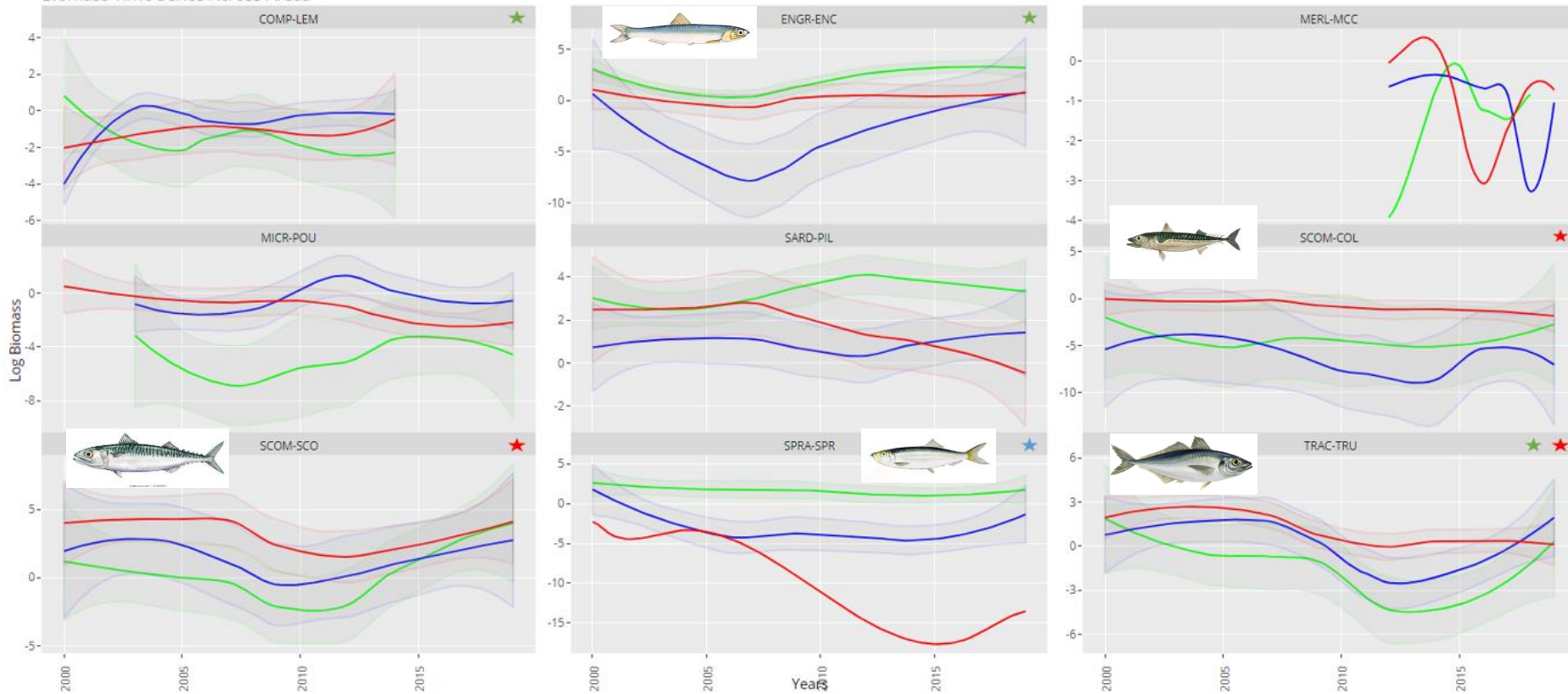


Questions and methods

Questions	Method
<p>How to characterize the different areas in regards to community composition?</p>	<p>Species biomass x area histogram B diversity analysis</p>
<p>Do species show different trends in different areas in recent decades?</p>	<p>Time series</p>

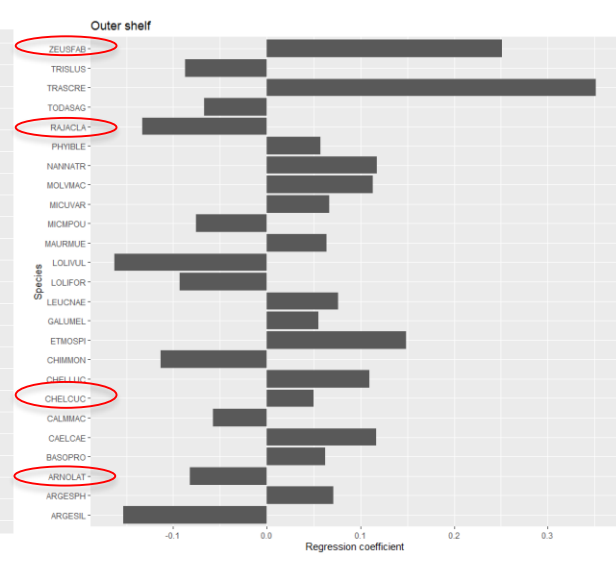
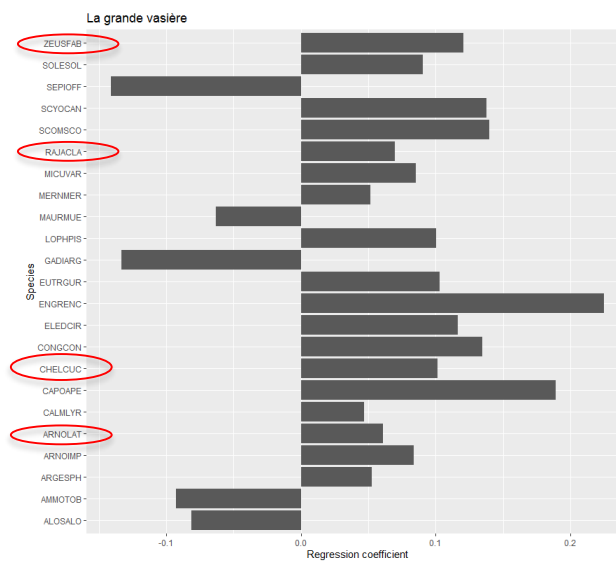
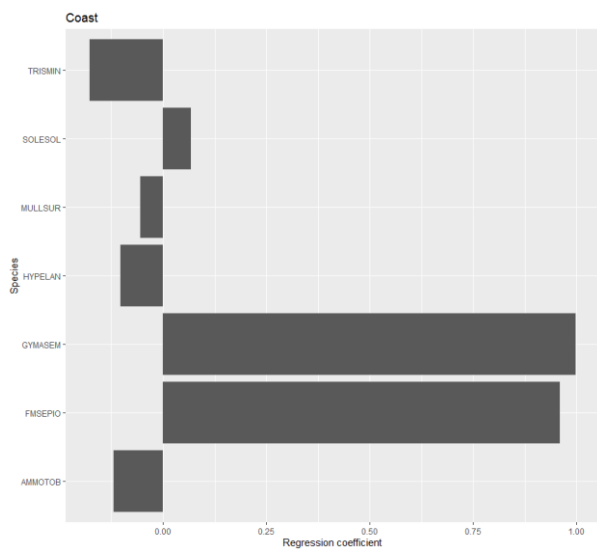
Time series - PELGAS

Biomass Time Series Across Areas



- Coast
- La grande vasière
- Outer shelf

Regression coefficients - EVHOE



Conclusion

- **Match between demersal- defined habitat and pelagic-defined habitat**
- **Demersal fishes show different community composition in different habitats**
- **The three areas seem different according to β diversity**
- **Some trends of demersal fishes seem to differ across areas**
- **Pelagic fishes composition and trends are more homogenous across habitats compared to demersal**

Perspective

- **Looking at finer scale (e.g. cohorts instead of entire population)**
- **Looking at mean weight**
- **Including other informations (abiotic, anthropogenic) if spatial resolution allows it**



QUESTIONS?

MERCI POUR VOTRE ATTENTION

