



# Biogeographical patterns in the deep ocean: environmental, biological, and historical drivers in the North Atlantic

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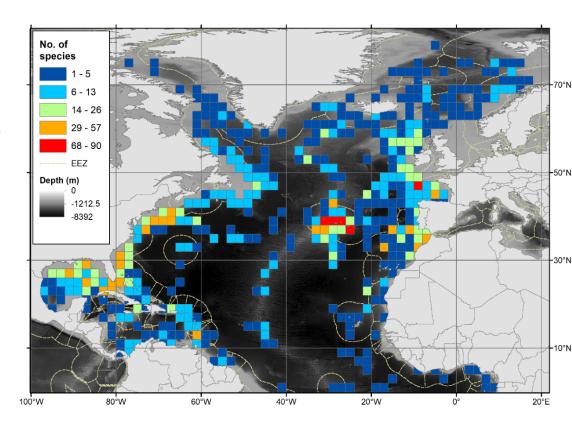


#### **RATIONALE**

Effective management of VMEs should be based on the full understanding of ecological processes and the assessment of the different scales structuring VMEs species diversity and communities.

#### **Research Questions:**

- 1. Are existing biogeographic classifications adequate to represent deep-sea VME biogeography?
- 2. Are current patterns of distribution in the North Atlantic a result of larval dispersal or environmental adaptation mechanisms?

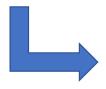


### RQ1

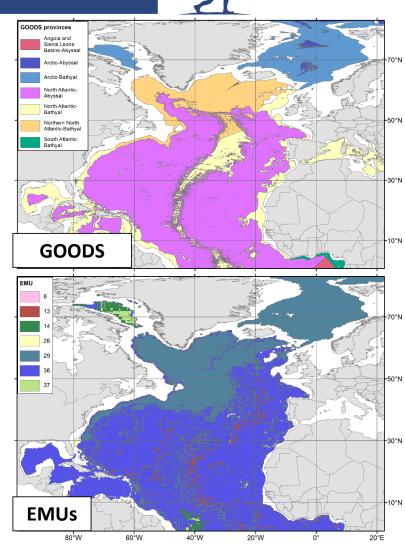
# Are existing biogeographic classifications adequate for VME taxa?

#### **Background**

- The lack of biogeographic data in the deep sea has pushed for approaches based on physiognomic proxies (i.e. bathymetry, oceanographic variables) not validated with species data.
- GOODS and EMUs have implemented this approach with expert knowledge and statistical modelling, respectively.



- nMDS ordination and ANOSIM to test for significant differences in VME assemblages among GOODS and EMUS provinces.
- Exploration of the effect of longitude on dispersal.



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#### **RESULTS**

- Significant spatial structure in assemblage composition:
  - **GOODS**: † R Global value = **depth has strong effect**
  - EMUs:  $\int R$  global value = no pattern in the nMDS plot
- A longitudinal gradient was evident in GOODS.

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).25 -	<b>A</b>	•	• •	•	•		<ul><li>Northern bathyal</li><li>Arctic bathyal</li><li>Abyssal</li></ul>	0.25 -	•			•	•		<ul><li>29</li><li>36</li><li>37</li></ul>
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0.25 -	<b>A</b>	<b>A</b>	<b>*</b> *				<ul><li>Caribbean</li><li>▲ Central</li><li>◆ Eastern</li><li>• Western</li></ul>	-0.25 -	•			•	•	•	<ul><li>Caribb</li><li>Centra</li><li>Easter</li></ul>
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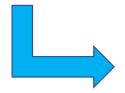
		Global R	P-value	
(A) ALL TAXA	EMUs GOODS	0.196 0.440	0.0005* 0.0005*	
(B) EMUs	Scleractinia Octocorallia	0.168 0.130	0.014* 0.039*	
	Porifera	0.162	0.005*	
(C) GOODS	Scleractinia	0.048	0.335	
	Octocorallia	0.177	0.054	
	Porifera	0.262	0.006*	



#### **CONCLUSIONS**



- Expert driven classification (GOODS) performed better than purely statistical approaches (EMUs).
- Important effect of depth → Bathymetry co-varies with many factors that influence deep-sea species distribution patterns.
- Some evidence of an eastern and western differentiation in assemblage composition was observed in the nMDS ordination of the GOODS provinces only.
- Longitudinal patterns were not observed in the cluster analysis that included species from upper bathyal depths  $(200 800 \text{ m}) \rightarrow$  Topographic effect?



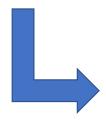
Refined GOODS could be implemented! (east and west Atlantic separation)

#### Larval dispersal vs. environmental adaptation



#### **Background**

• Biotic and abiotic interactions control community structure at varying spatial and temporal scales, and generate spatial patterns that need to be assessed to disentangle the ecological processes structuring these communities.

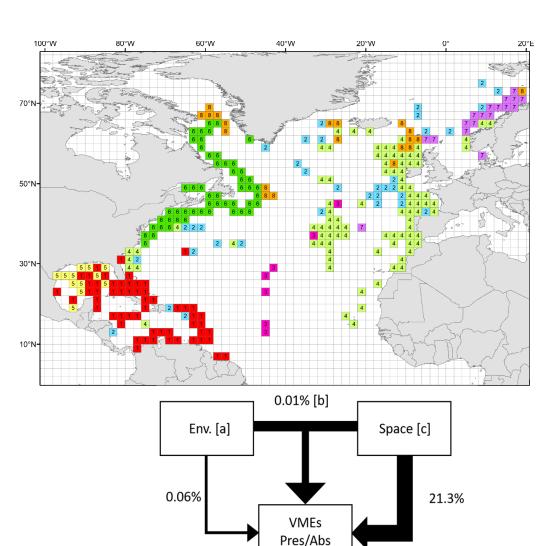


- We aimed to describe the distribution of VMEs and relate these to environmental factors at basin scale.
- Unravel the relative importance of environmental versus larval dispersal mechanisms in the biogeographical structure of VMEs.
- Spatial variation in multivariate data through distance-based Moran's eigenvector mapping (dbMEM) and redundancy analysis (RDA).

#### **RESULTS**

- 8 biogeographic clusters representing all VME taxa.
- dbMEM analysis provided vectors representing broad-scale patterns.
- Full spatial and environmental model explained 21.3 % of the variation of the data:

VME Presence/absence ~ T + Aragonite Saturation state + Calcite Saturation state + SD Oxygen + SD Si + EPC + pH + Currents Speed + Salinity + spatial eigenfunctions (broad-scale)



78%

Unexplained [d]



#### **CONCLUSIONS**



- Change in resolution of environmental variables to match species data masks their potential effect.
- Role of space indicates that present broad-scale patterns of deep-sea VME distribution are likely a result of topography, distance-decay relationships or historical events.
- Biogeographical clusters were driven by the oceanographic conditions characterising the water masses present in each geographic area.
- Larval dispersal mechanisms, primarily, and environmental processes (spatially structured environmental variables) not fully captured at the resolution of our study, potentially have determined the present-day distribution of complex habitats formed by VMEs in the North Atlantic.

Implications for spatial management measures

#### **Acknowledgements**









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#### **Thank You**

## **Patlas**

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