



**FOLLOWING THE MEDITERRANEAN
OUTFLOW WATER:**

**A CHARACTERIZATION OF
COMMUNITY CONNECTIVITY DRIVEN
BY WATER MASSES FROM THE
ALBORAN SEA TO THE AZORES**



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2nd April 2019**

Mediterranean-Atlantic transition zone in ABYSS

1/ Water

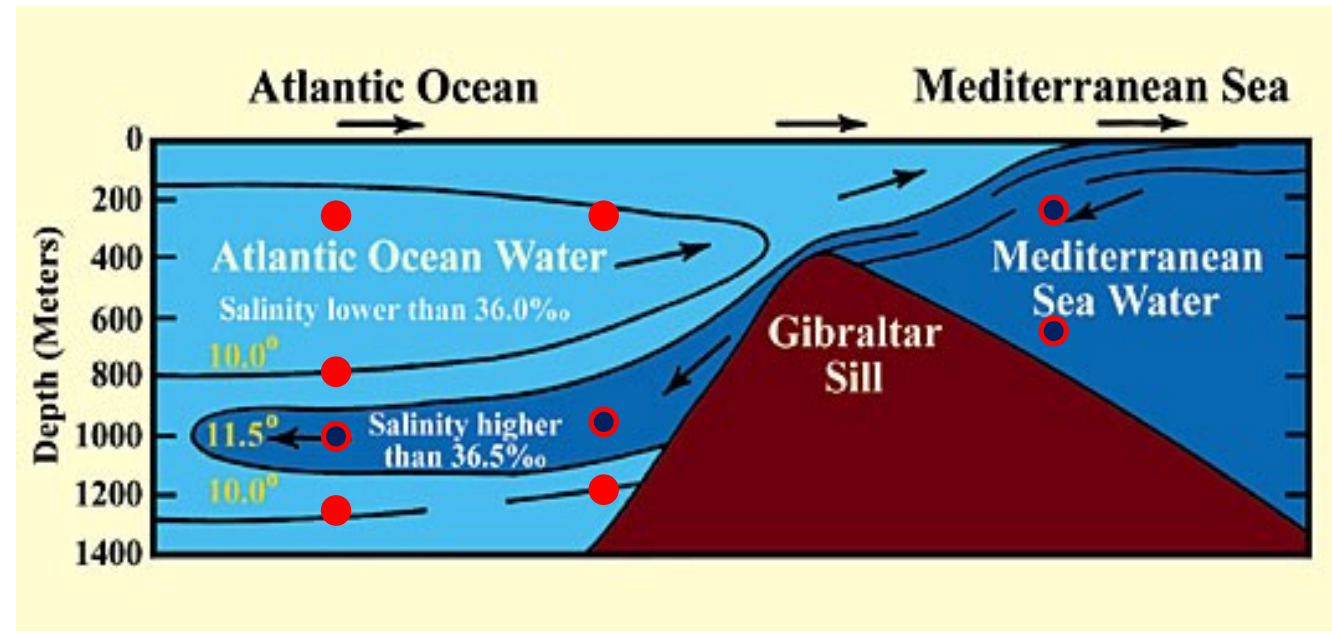
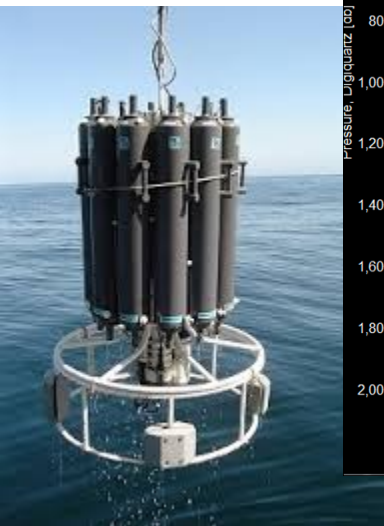
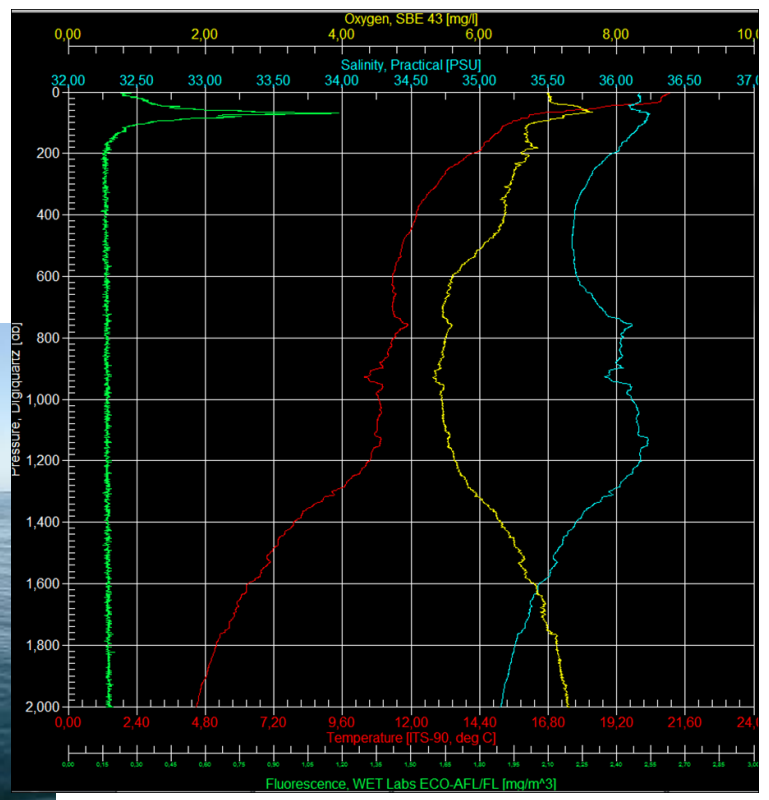
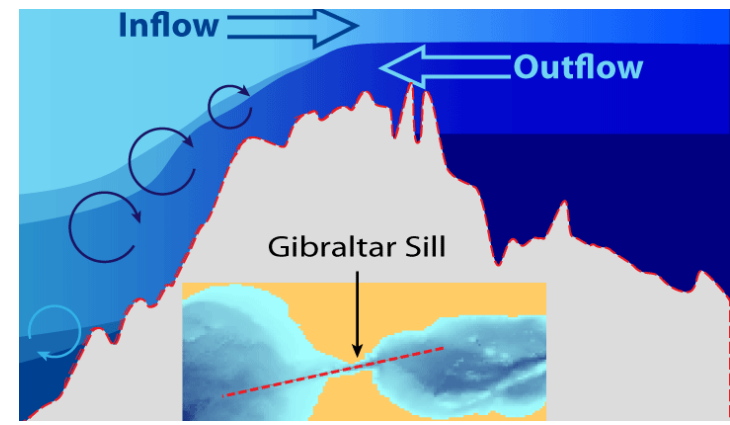
- Influence of water masses on biotic compartment
- Exploratory! (opportunistic sampling)
- Do Mediterranean water masses harbour different biotic communities than Atlantic waters?
- Does the MOW keep a stable biotic signature throughout its journey to the Azores?

2/ Sediment

- Evaluation of benthic community changes across the Mediterranean-Atlantic transition zone
- Are changes primarily explained by depth zone or ocean basin ?
- Are these changes homogenous among sediment layers, or are deeper layers differently affected than upper layers?

Water pilot study in ABYSS

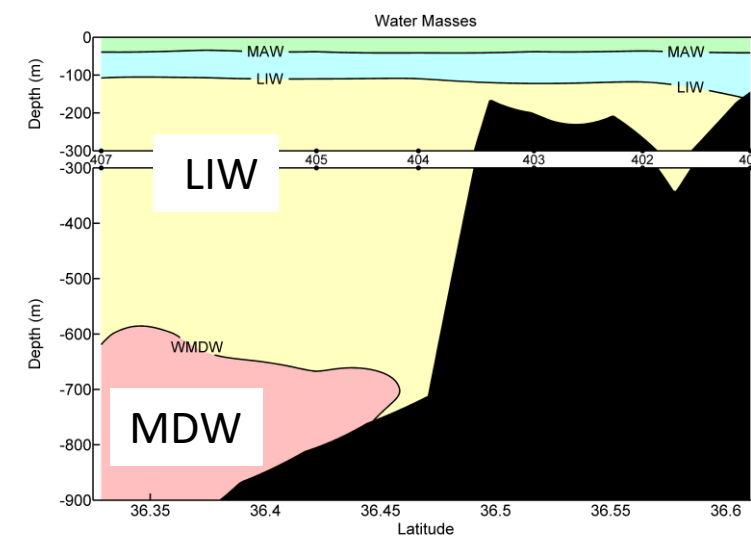
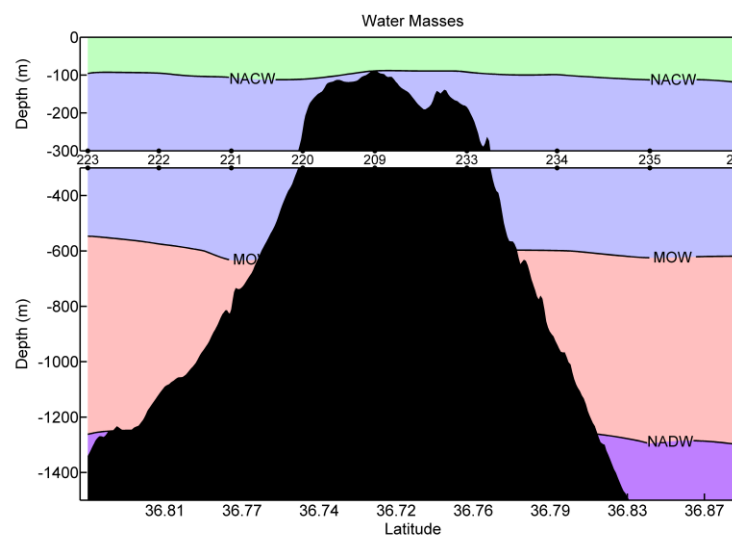
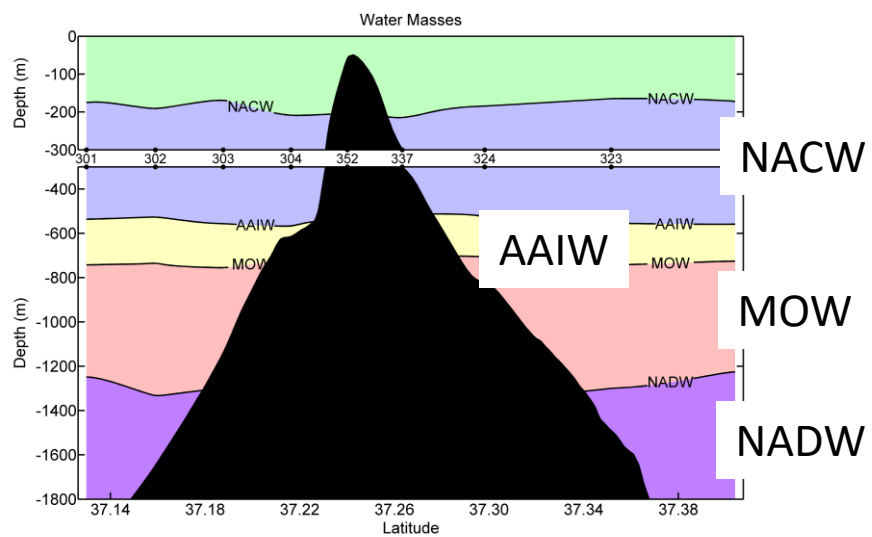
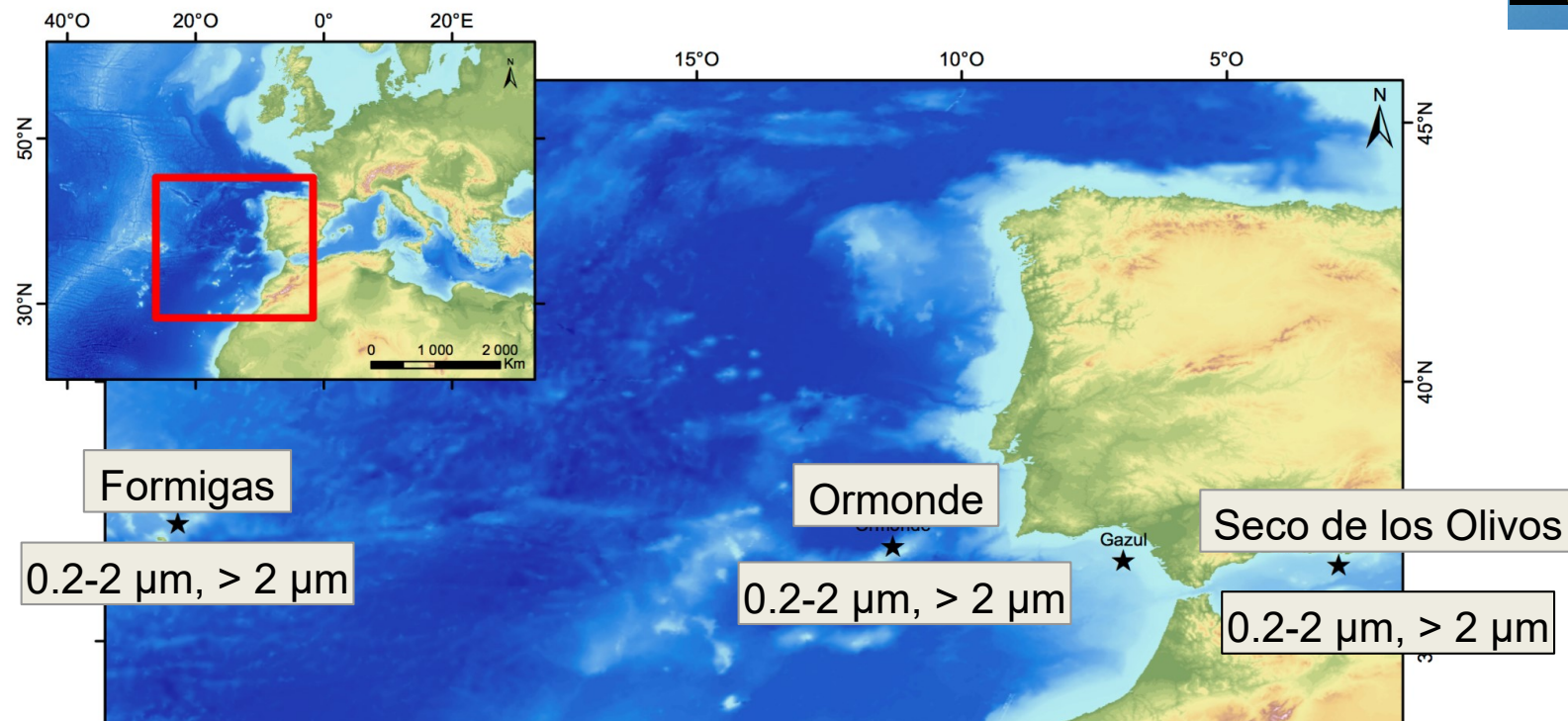
- Following the Mediterranean Outflow Water (MOW) along an East-West gradient in the Mediterranean-Atlantic transition zone



Sampling problem: only three 5L Niskin bottles at each site...
 We had to pool the filters to allow sufficient yield for DNA extraction

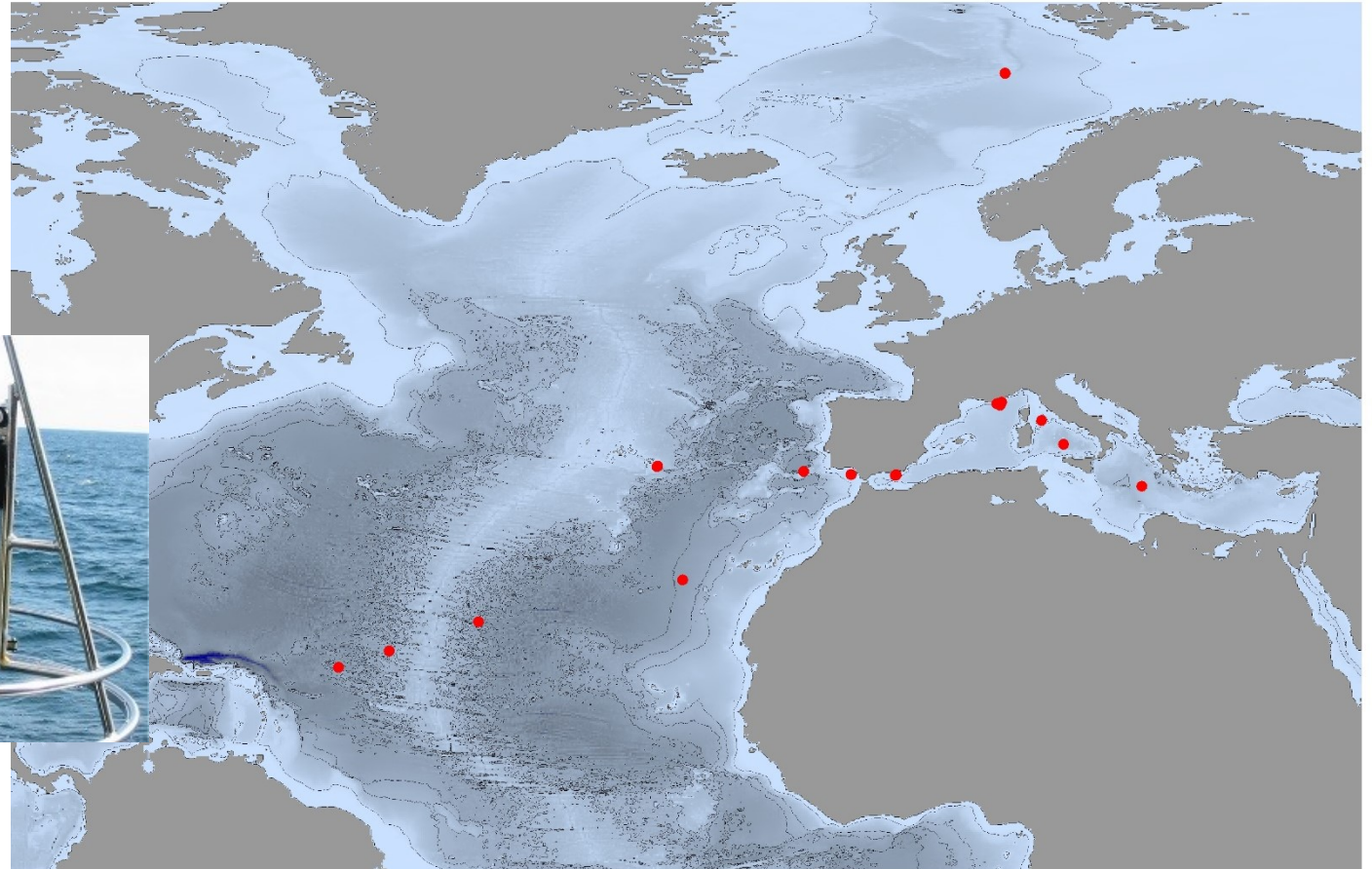
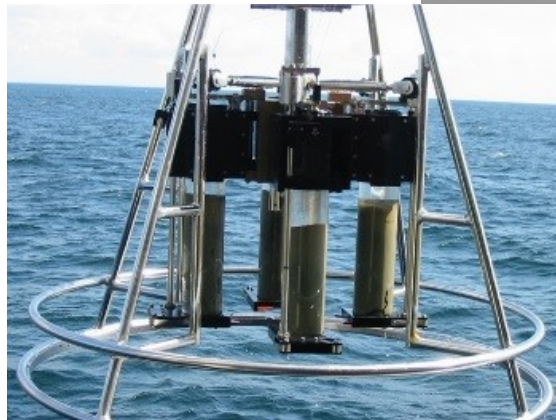
Water pilot study

- Two size fractions targeted at each site



Sediment study

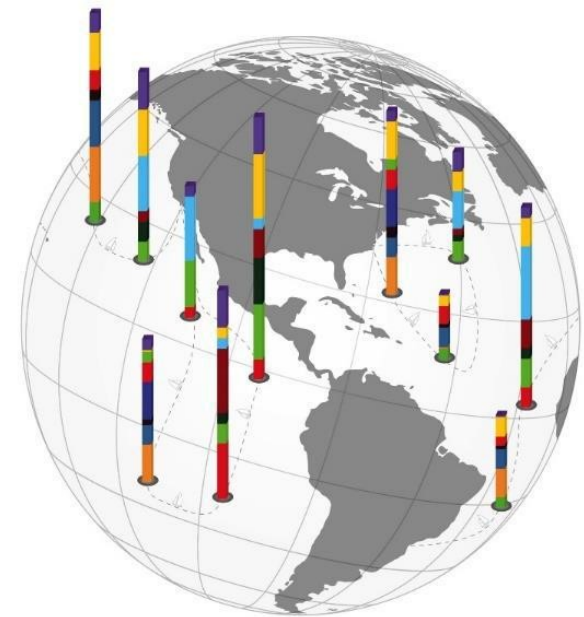
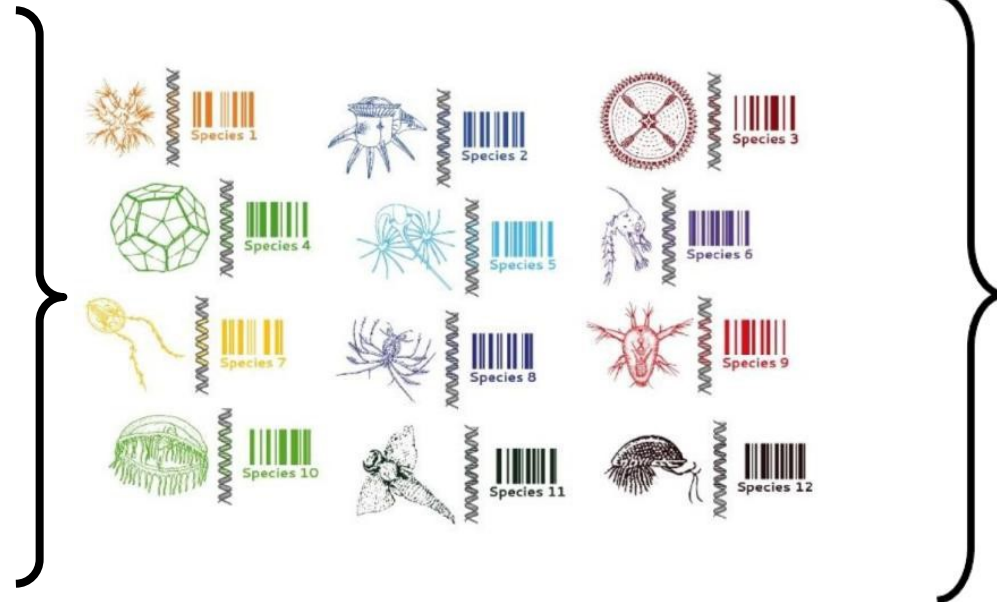
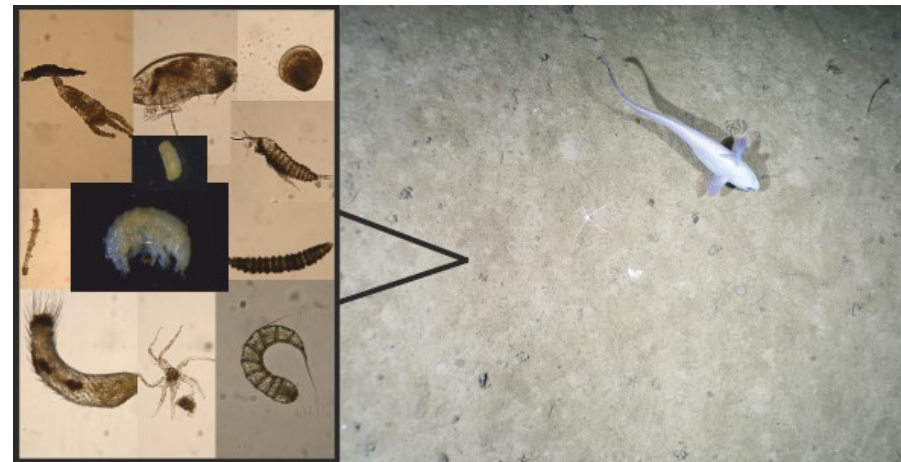
- 245 sediment samples covering the Atlantic-Mediterranean transition zone
- Triplicate sediment cores at each sampling site



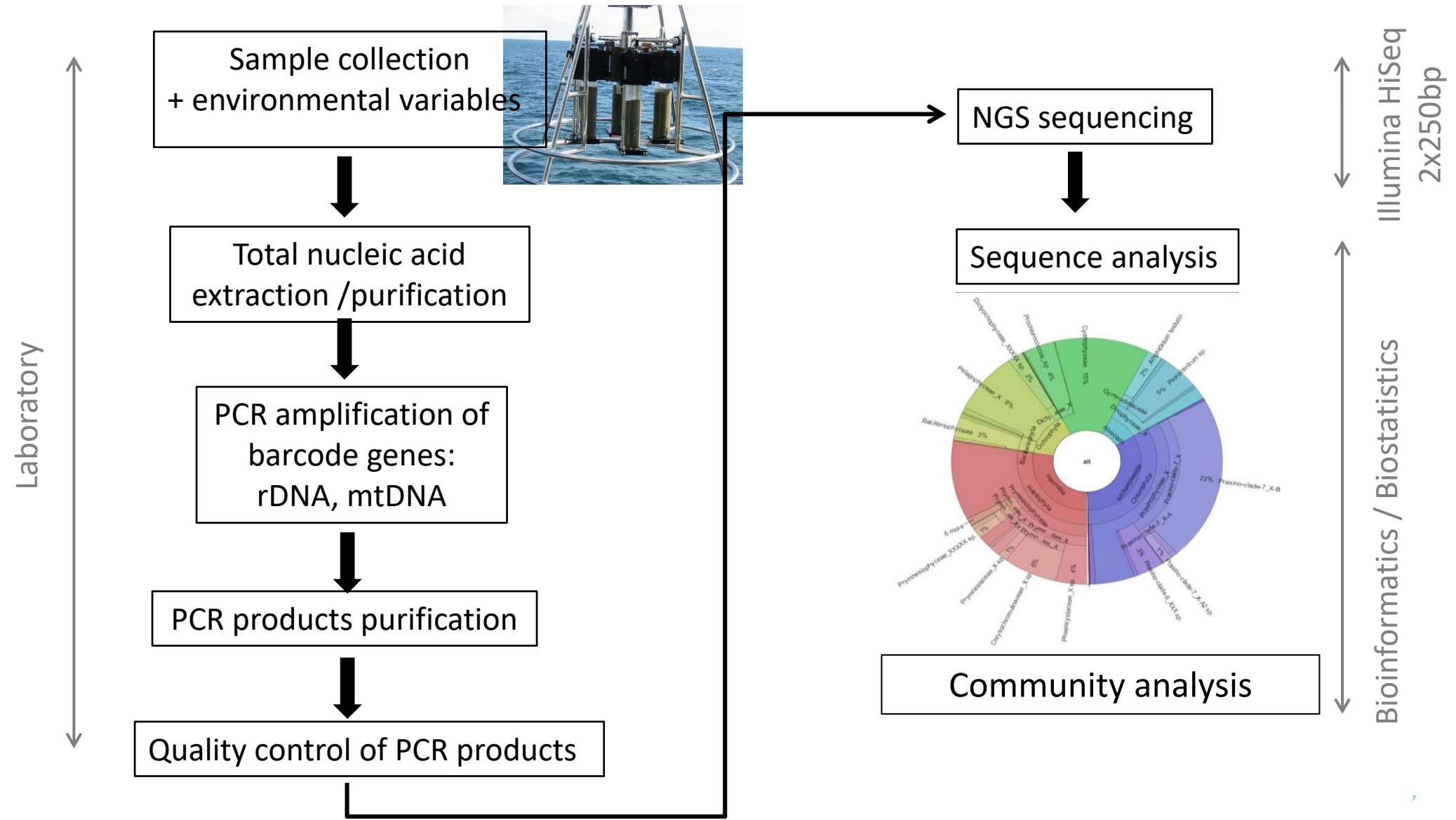
- DNA extraction from 10g of sediment using PowerMaxSoil kit

Environmental DNA metabarcoding

- based on the evolutionary species concept
- description of biodiversity in a natural sample (community complex)



General eDNA workflow



METAZOANS

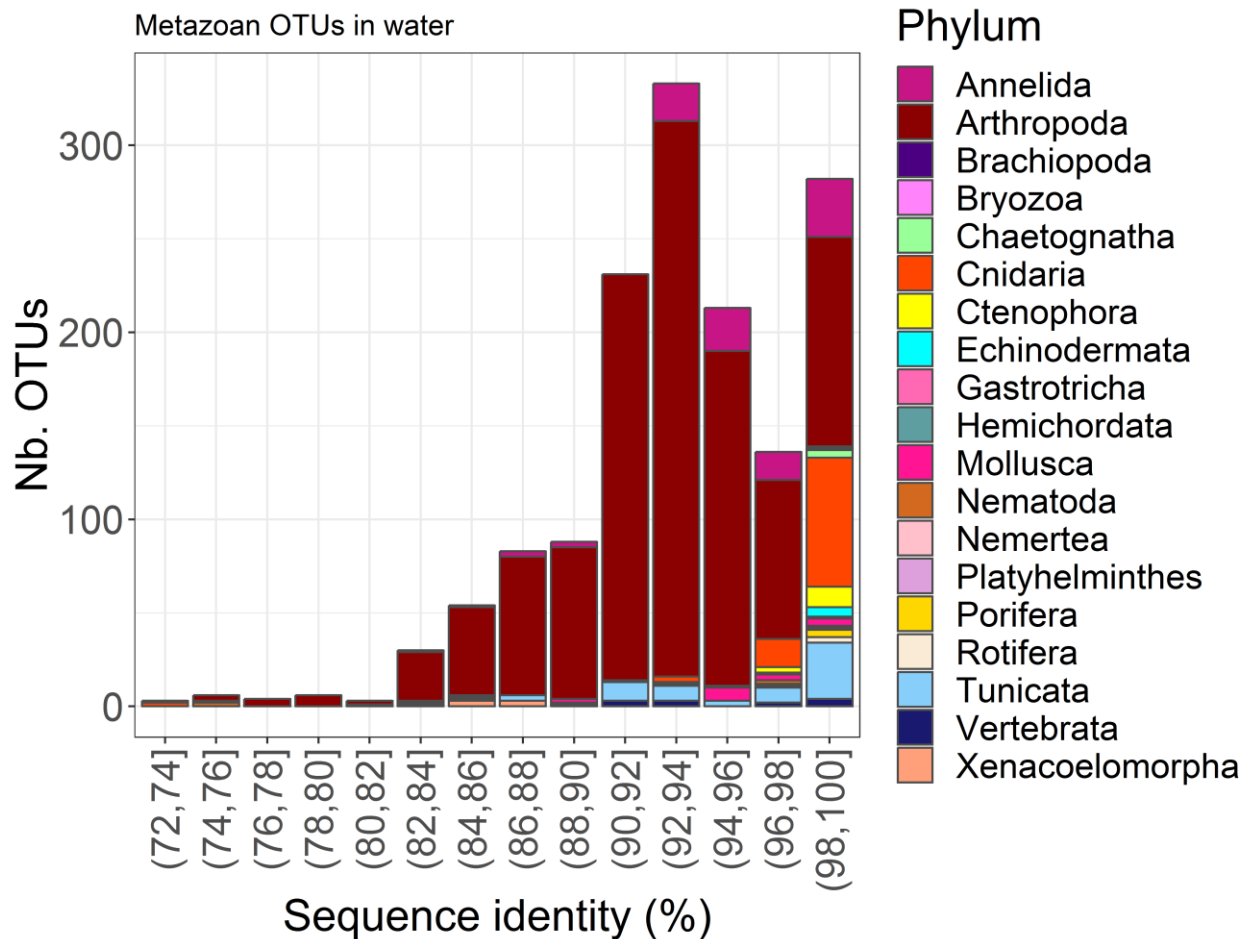
PRIMERS:

18S V1-V2

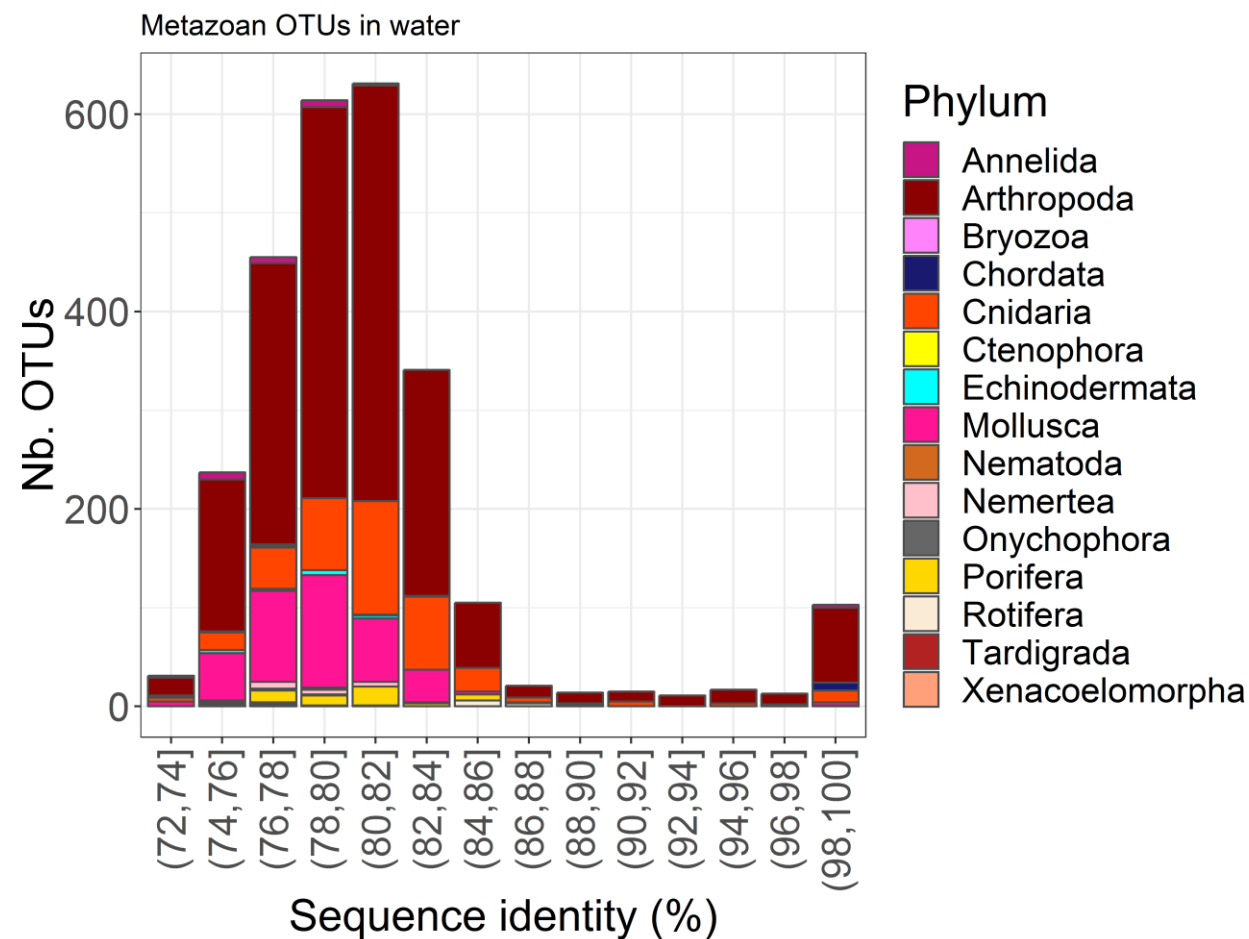
COI

Data overview for metazoans in the water column

• 18S-V1 rDNA



• Mitochondrial COI



Taxonomic compositions, 18S-V1

Phylum

- Cnidaria
- Tunicata
- Chaetognatha
- Arthropoda
- Nematoda
- Gastrotricha
- Mollusca
- Nemertea
- Hemichordata
- Annelida
- Porifera
- Platyhelminthes
- Ctenophora
- Vertebrata
- Xenacoelomorpha
- Echinodermata

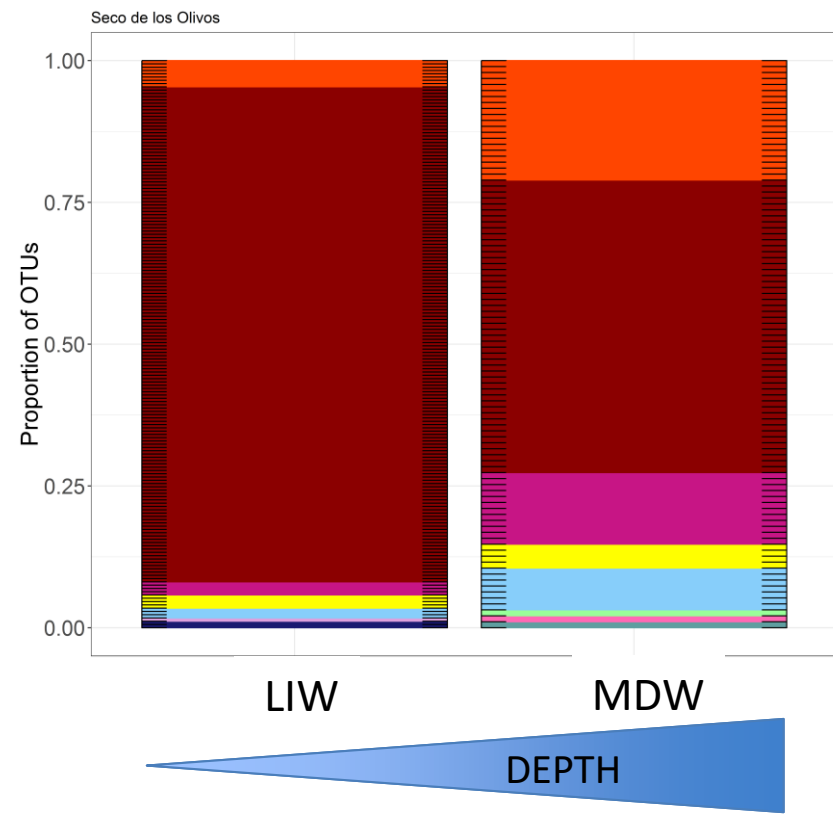
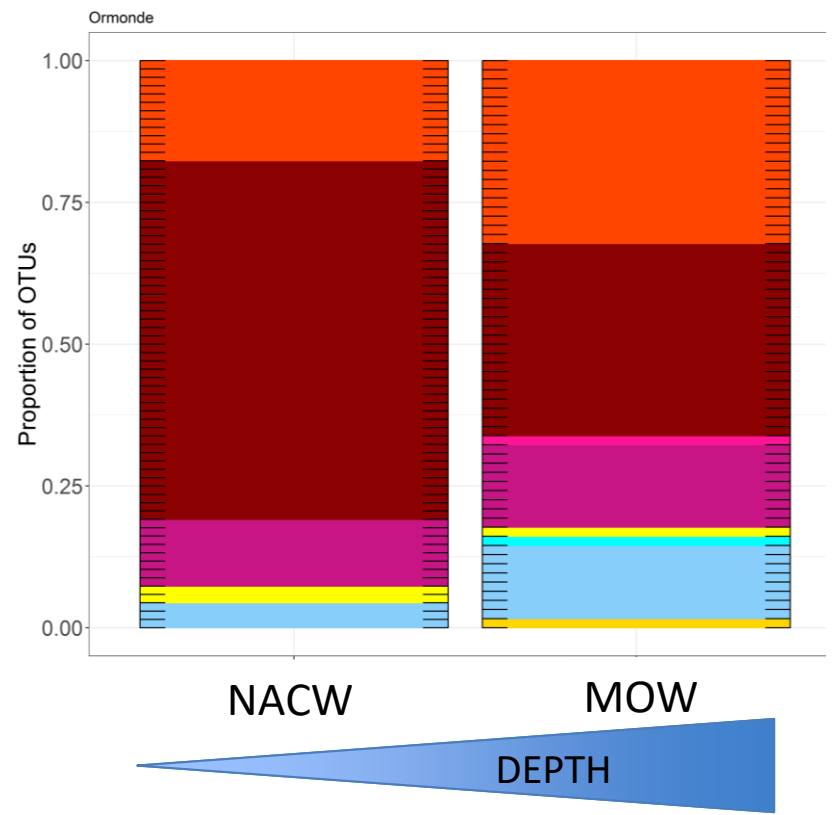
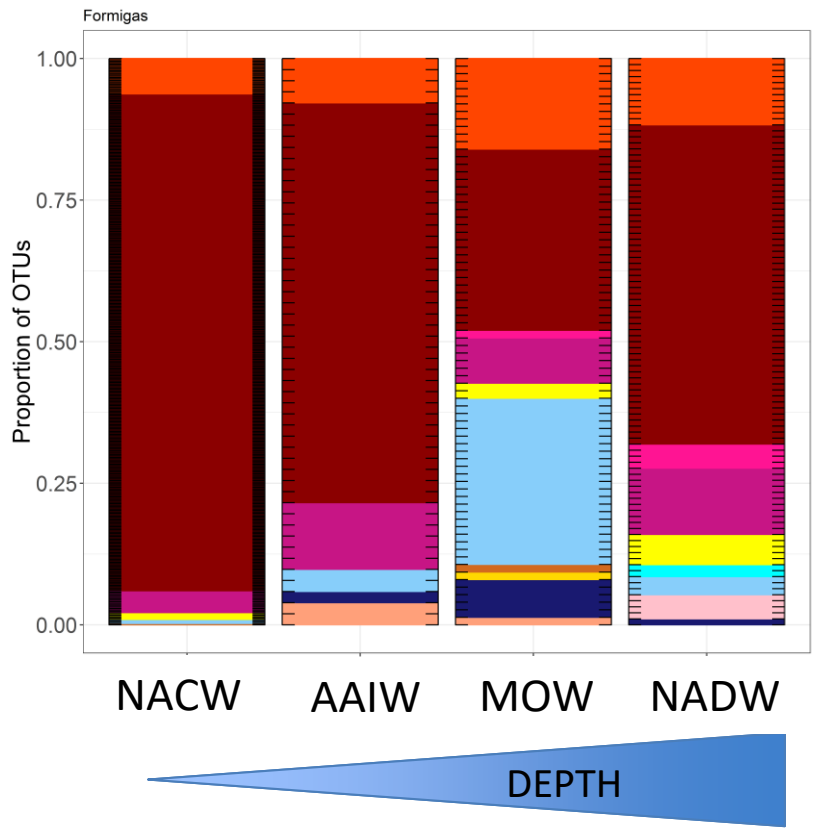
WEST

EAST

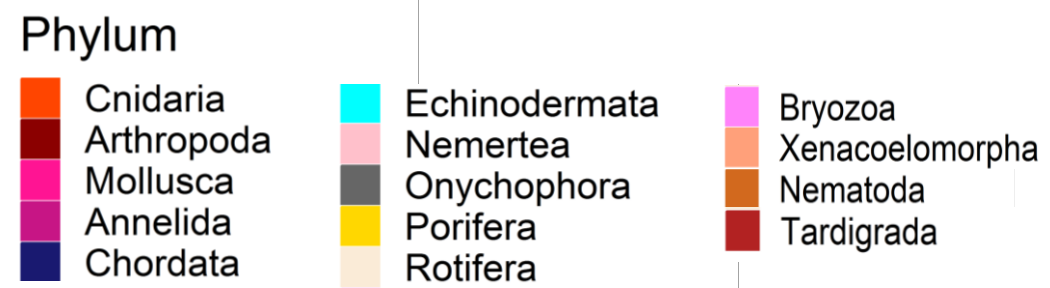
Azores

Southwest Portugal

Alboran Sea



Taxonomic compositions, COI



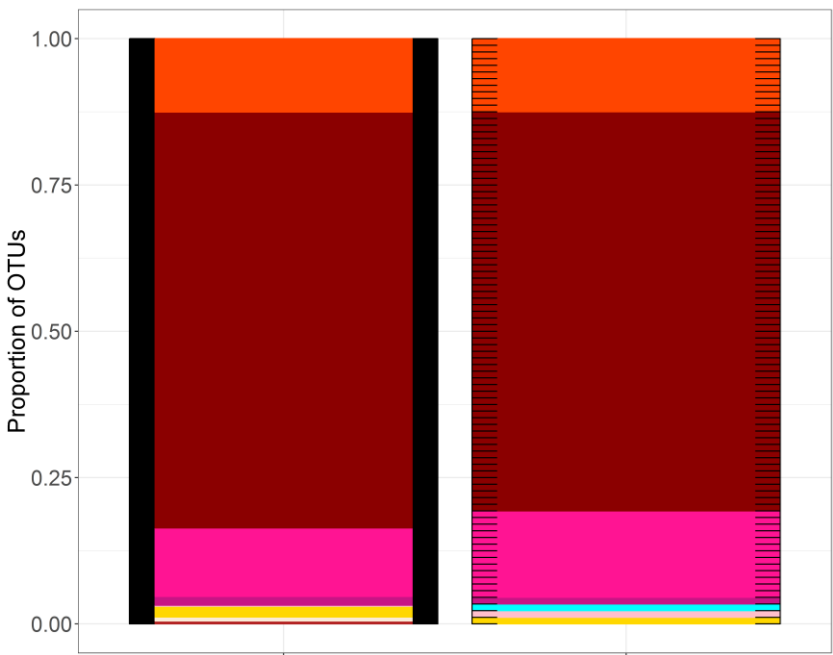
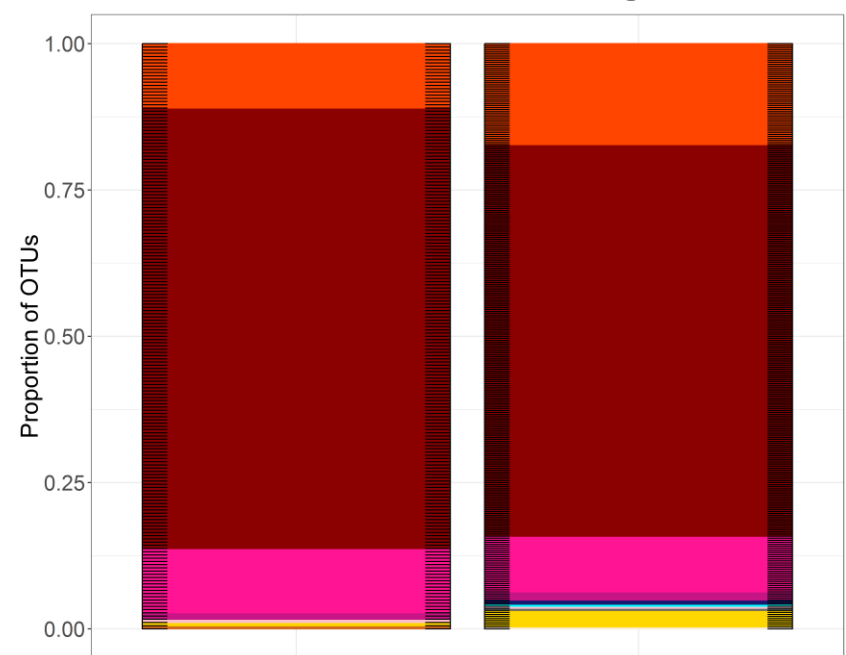
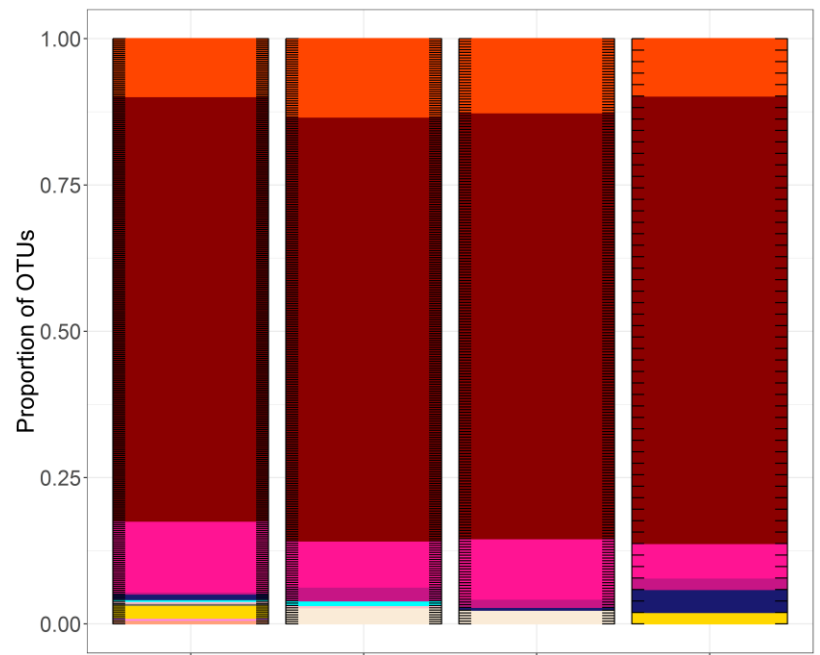
WEST

EAST

Azores

Southwest Portugal

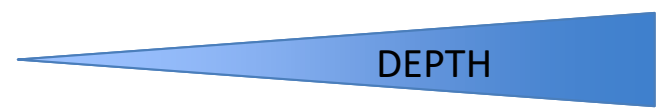
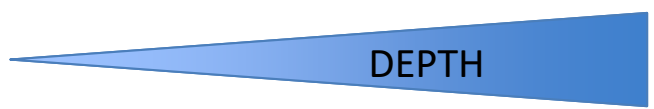
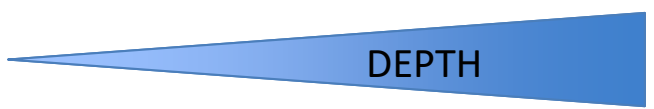
Alboran Sea



NACW AAIW MOW NADW

NACW MOW

LIW MDW



Alpha diversity

Location

- Azores
- Southwest Portugal
- Alboran Sea

• 18S-V1

• COI

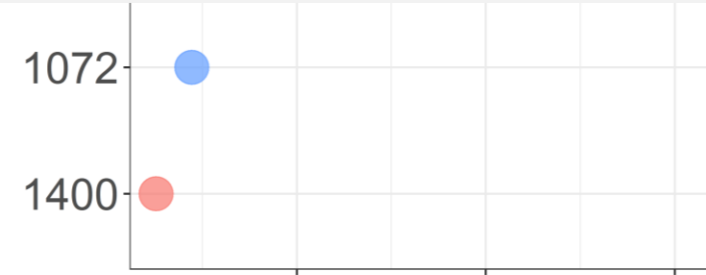
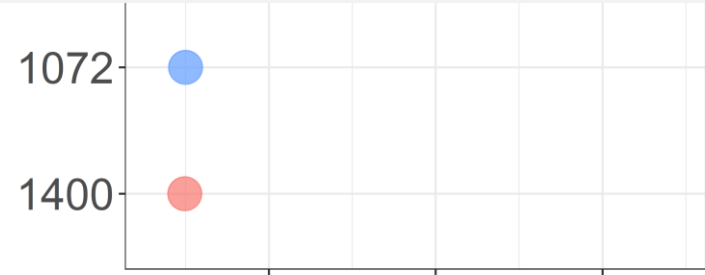
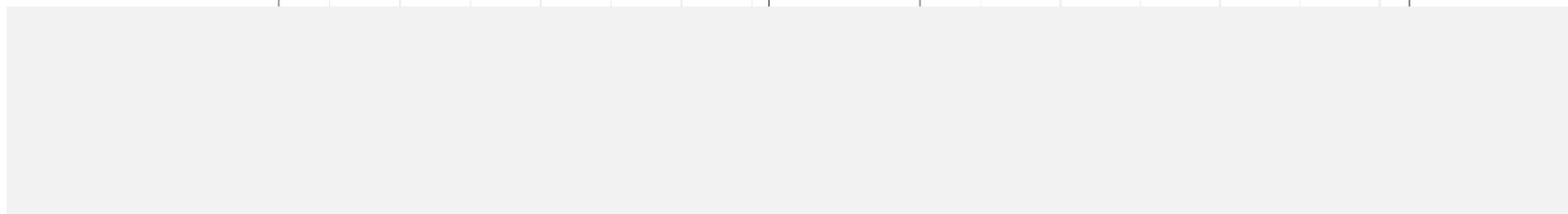
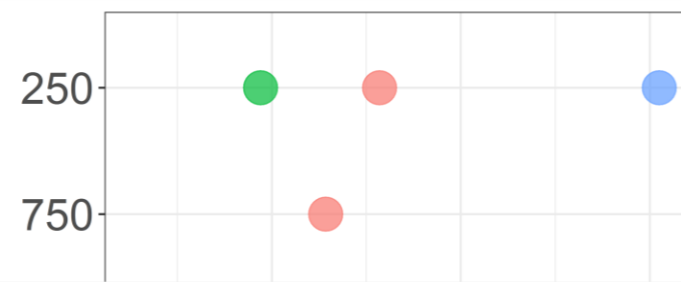
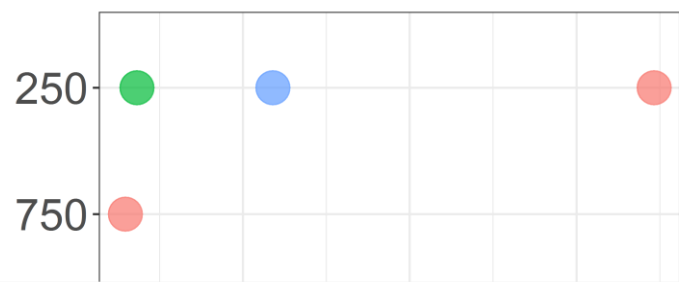
Upper waters (NACW, LIW)

Antarctic waters (AAIW)

MOW

MDW

Atlantic Deep waters (NADW)



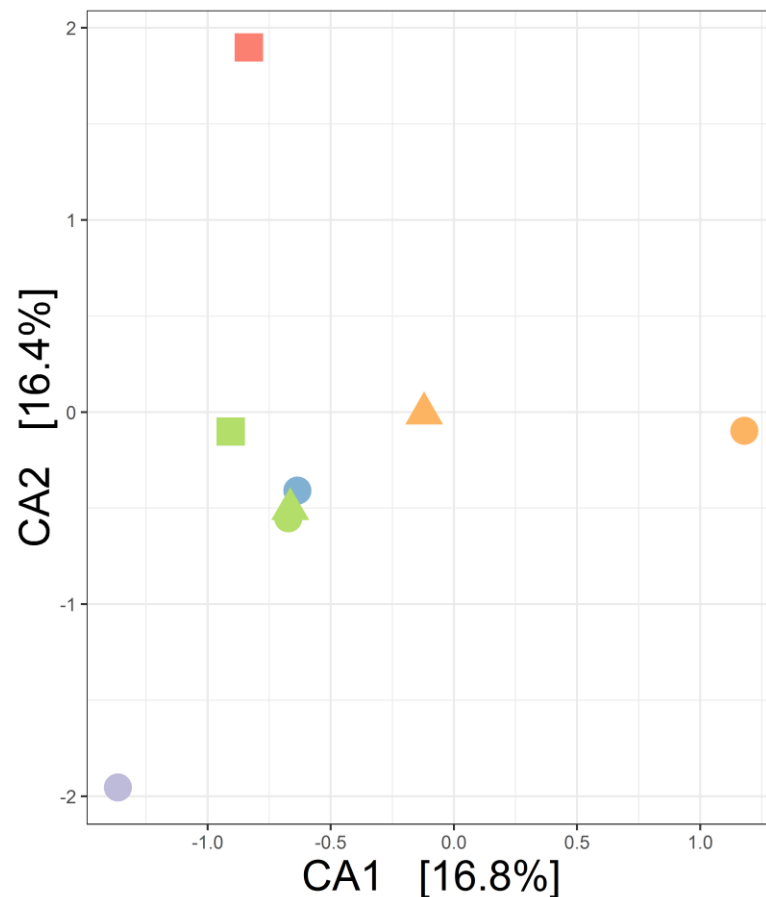
Nb. metazoan OTUs

Nb. metazoan OTUs

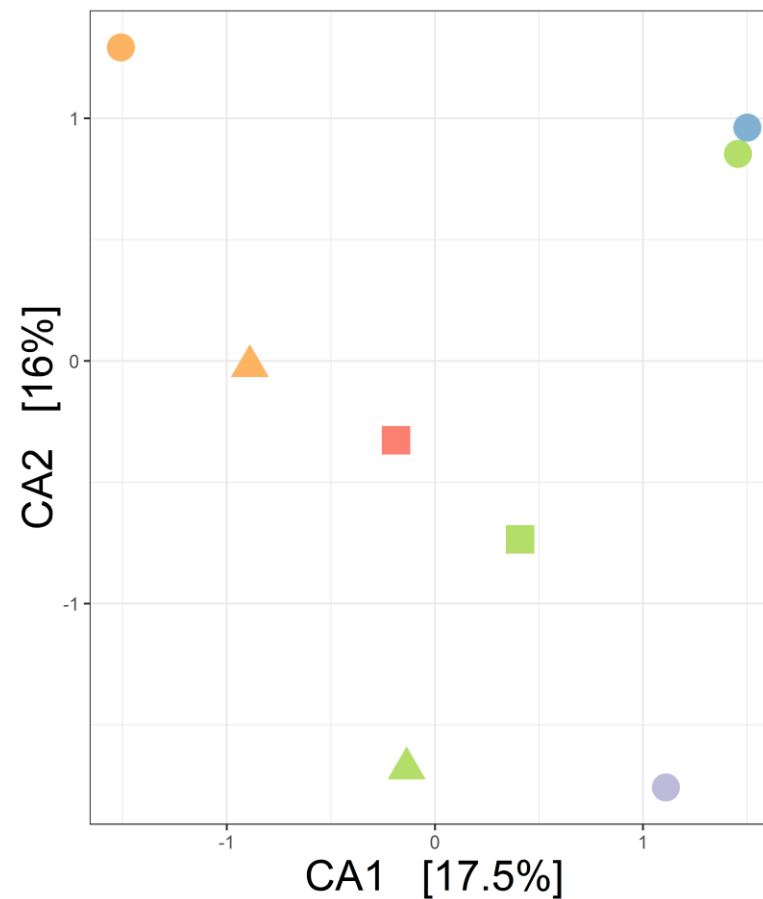
Beta diversity: influence of water masses on metazoan communities

- ~60% of variation in data explained by water mass
- Strong segregation surface vs deep
- Communities strongly vary with location

• 18S-V1



• COI



Location

- Azores
- ▲ Southwest Portugal
- Alboran Sea

water.mass

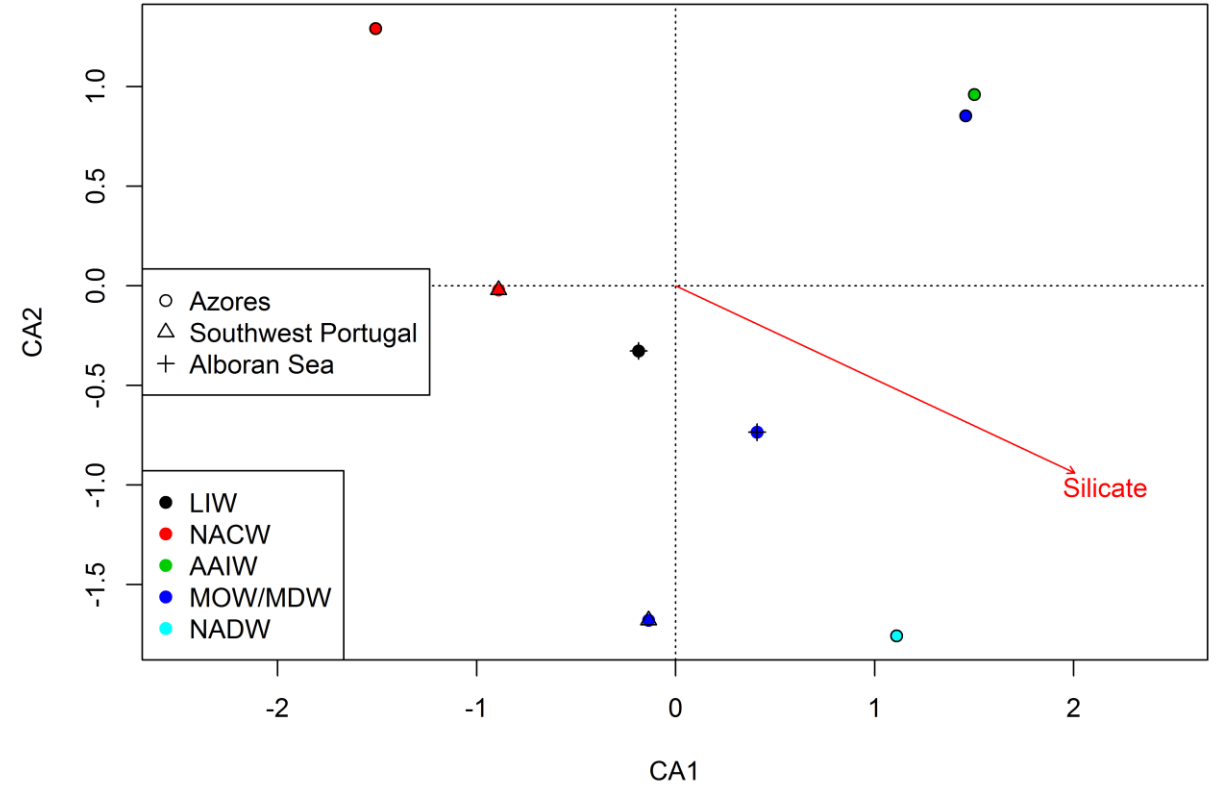
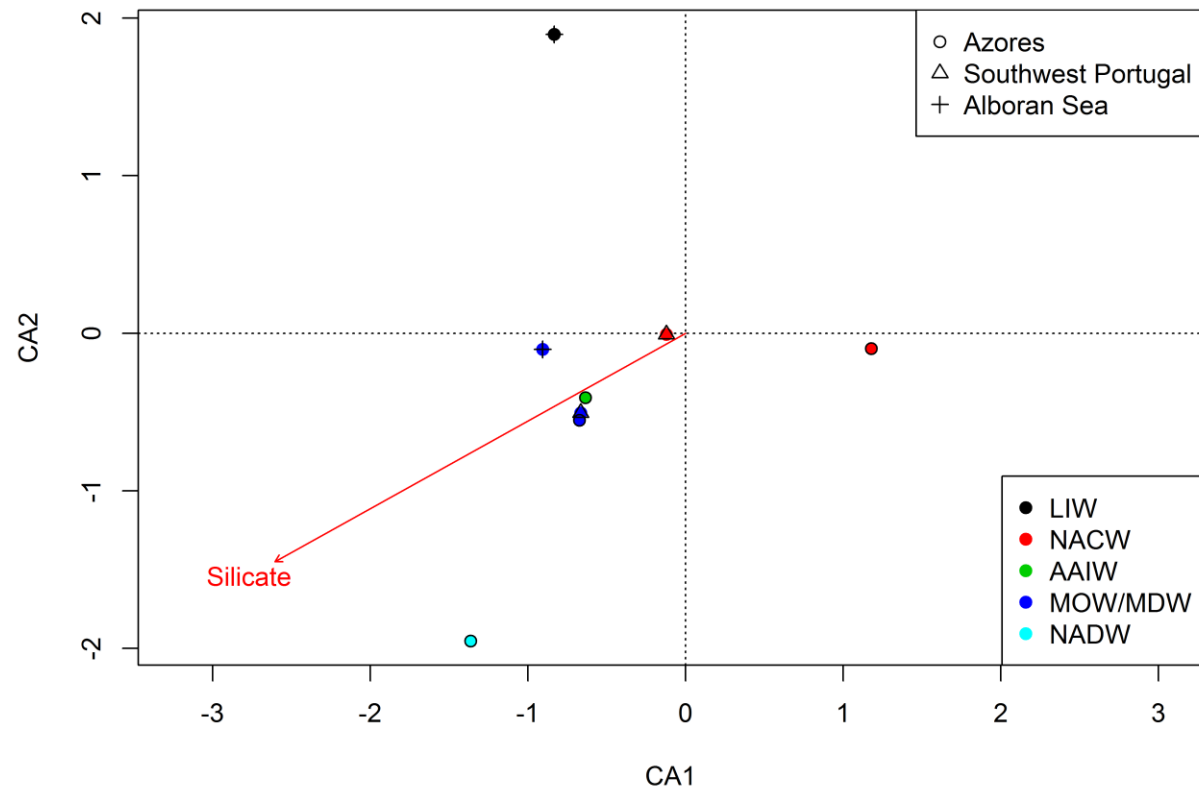
- LIW
- NACW
- AAIW
- MOW/MDW
- NADW

Surface
Antarctic
Deep

Beta diversity: influence of environmental variables

• 18S-V1

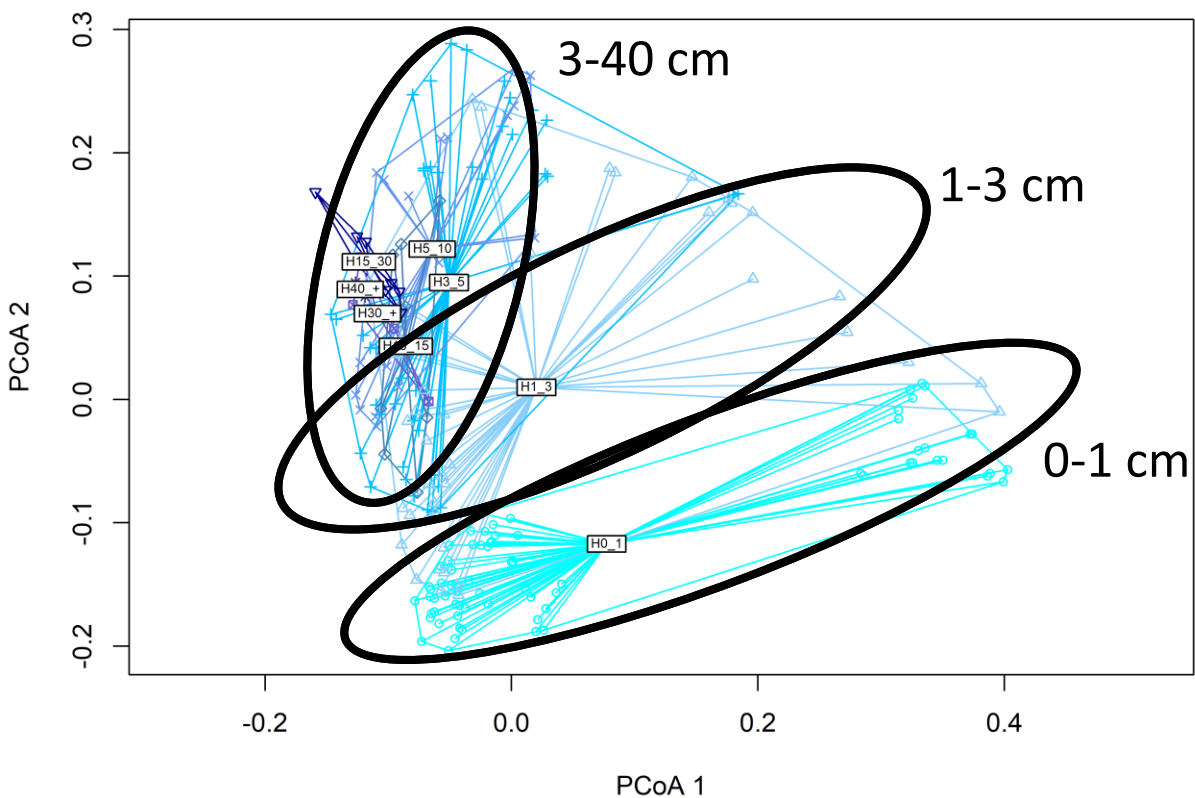
• COI



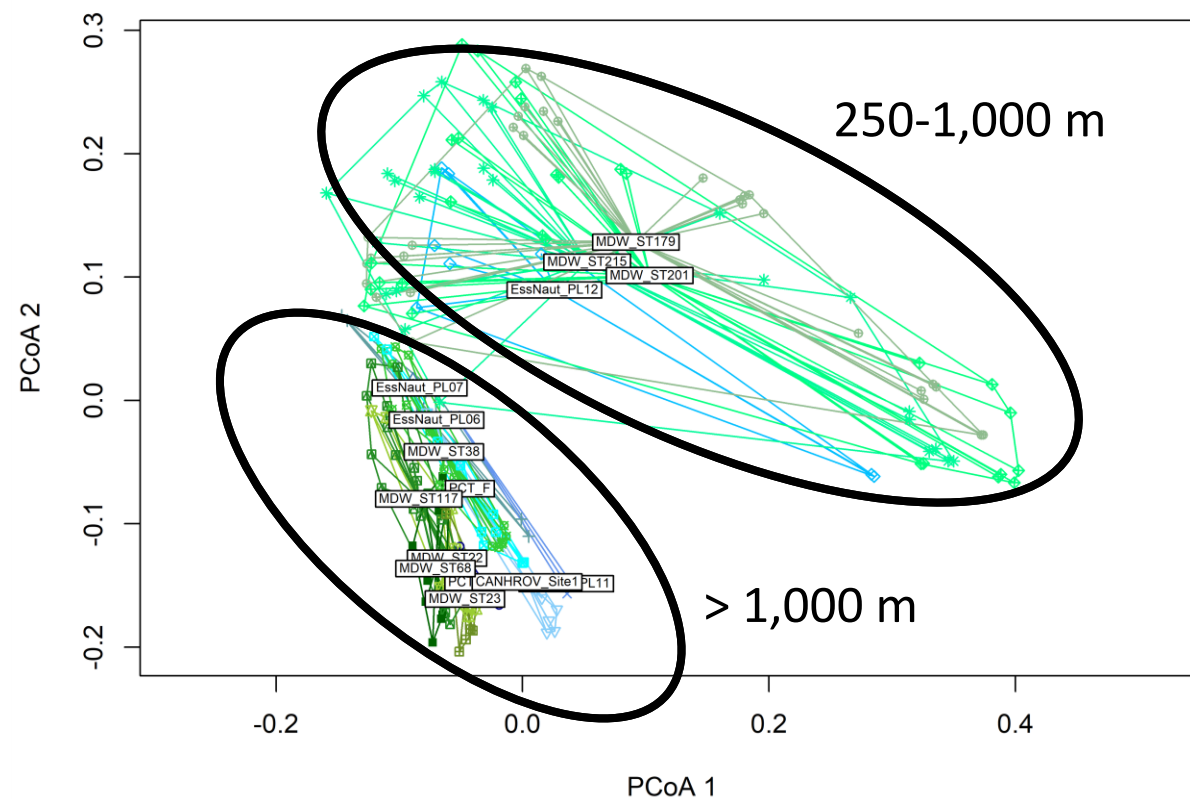
- Silicate concentration significantly correlated to community composition changes
- Silicate concentrations may explain the depth segregation

Metazoan communities: comparison of benthic communities (18S)

- segregation according to depth layer in sediment



- segregation according to depth region, not ocean basin

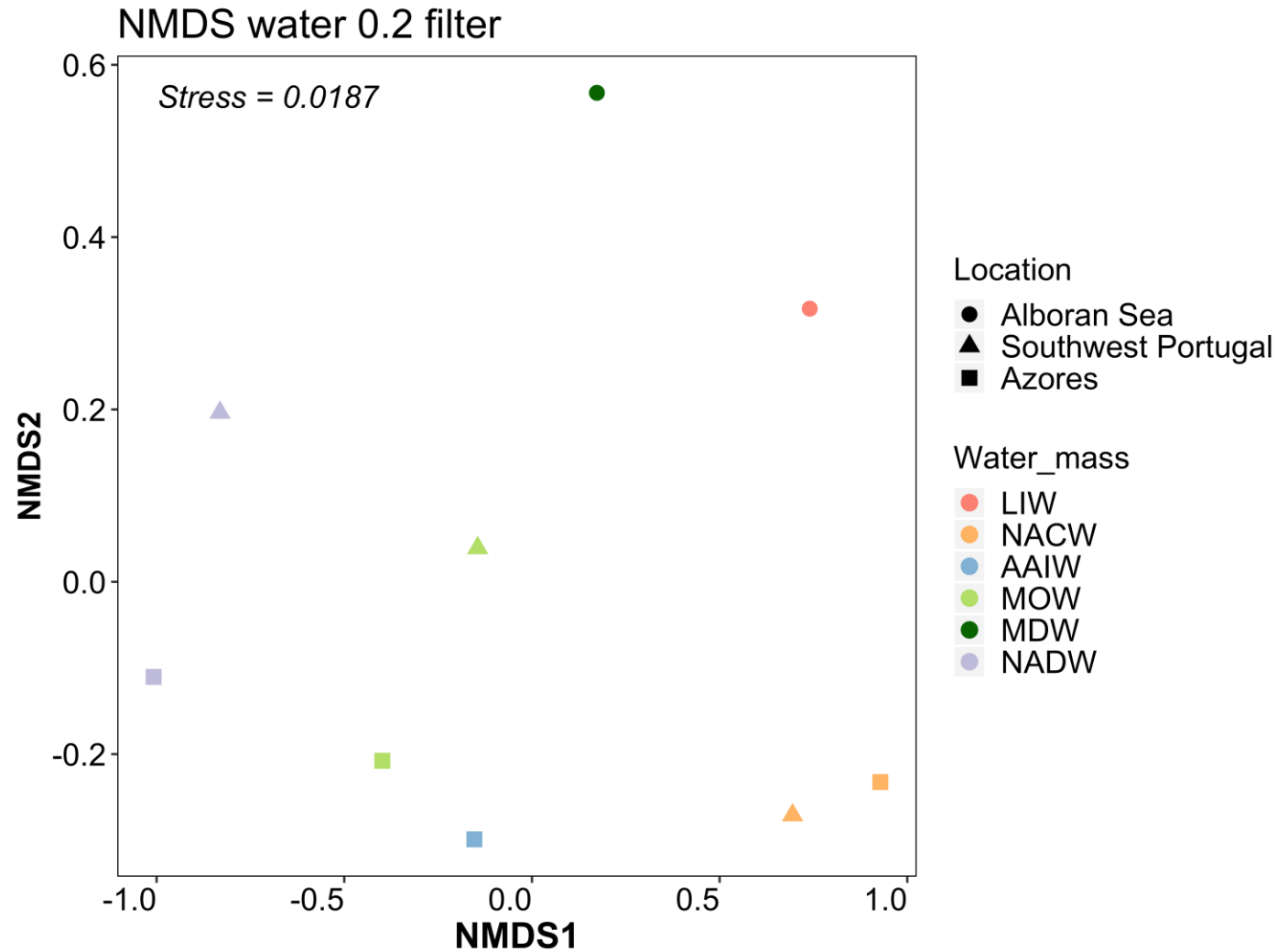


PROKARYOTES

PRIMERS:

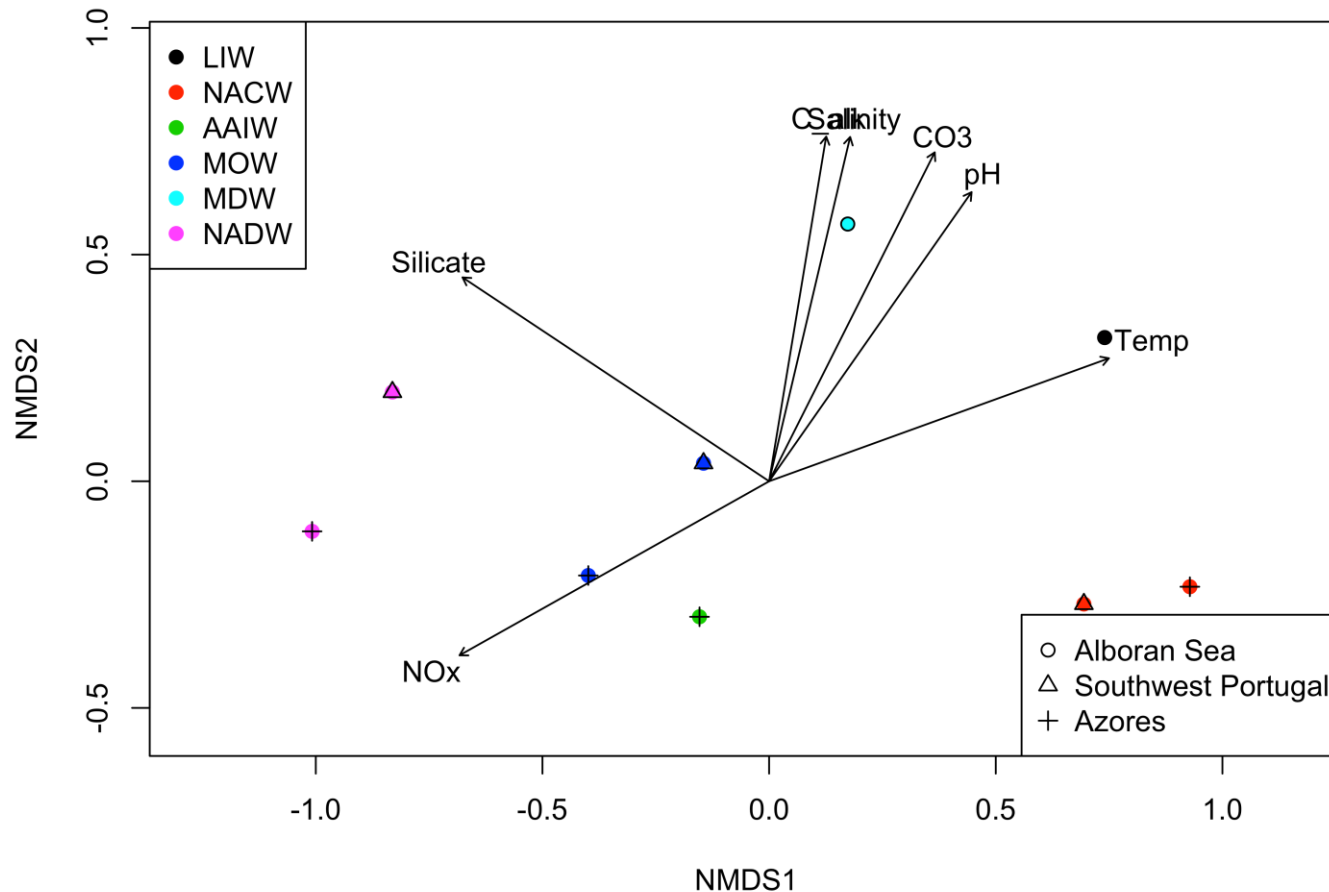
16S V4V5

Prokaryotic communities: comparison between water masses



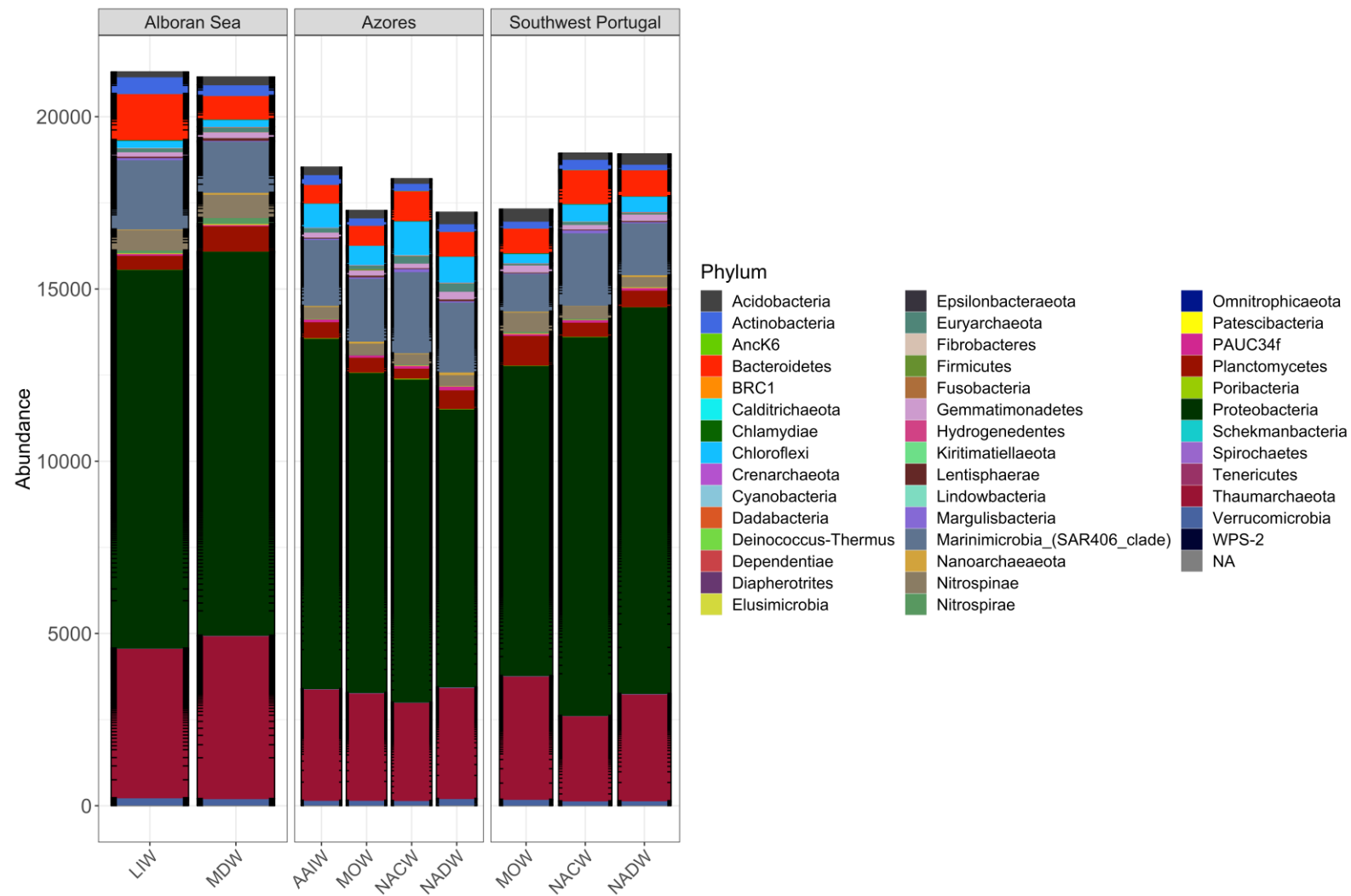
- Segregation of Mediterranean vs. Atlantic samples
- Segregation by depth and sampling location

Prokaryotic communities: comparison between water masses



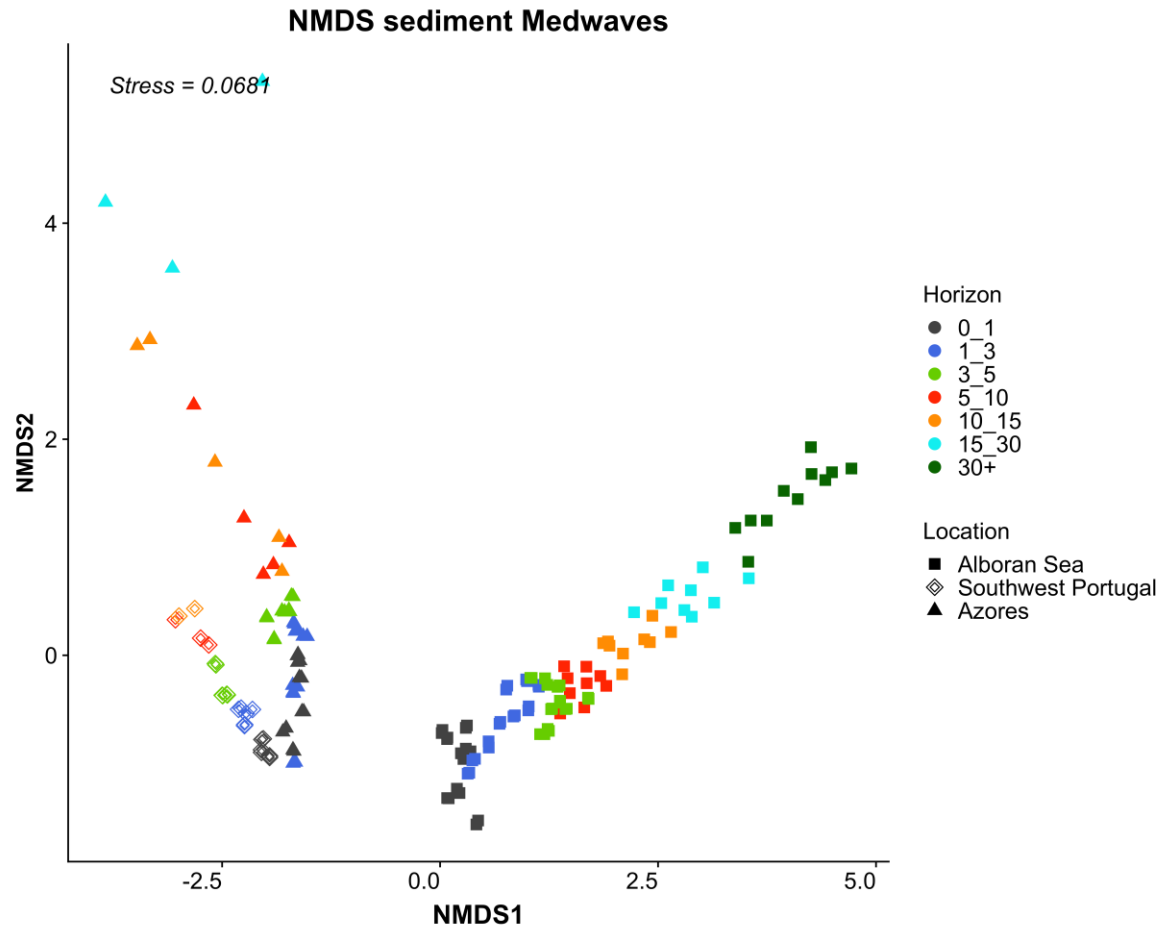
- Many physical-chemical parameters explain community structure
- Nitrates and silicates may best explain the depth segregation

Prokaryotic communities: comparison between water masses



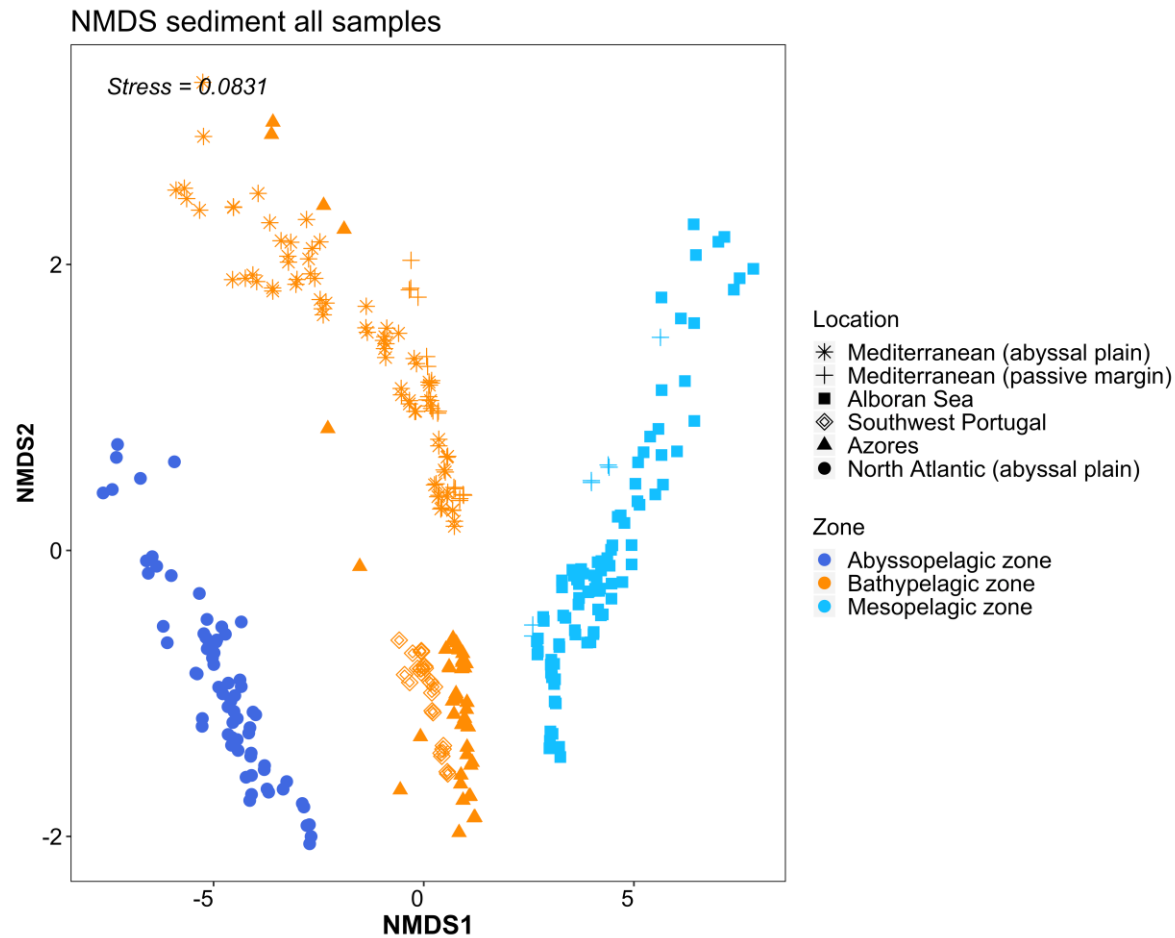
- No striking change in pattern at this scale

Prokaryotic communities: comparison of benthic communities



- Separation between Mediterranean and Atlantic communities
- Clear segregation according to depth layer in sediment

Prokaryotic communities: comparison of benthic communities



- Clear segregation by depth region

Perspectives

➤ Metazoans

- Benthic communities in the Mediterranean-Atlantic transition zone
- Coupling between pelagic and benthic communities?

➤ Prokaryotes

- Archaeal data
- Oligotyping : high resolution taxonomic analysis of most abundant members of the communities

Thank you for listening!



Acknowledgements

- Pourquoi Pas les Abysses? Team & Ifremer Bioinformatics group
- Ifremer
- Génoscope (Julie Poulain and Patrick Wincker)
- Researchers and crew of the MEDWAVES cruise (ATLAS H2020 project)

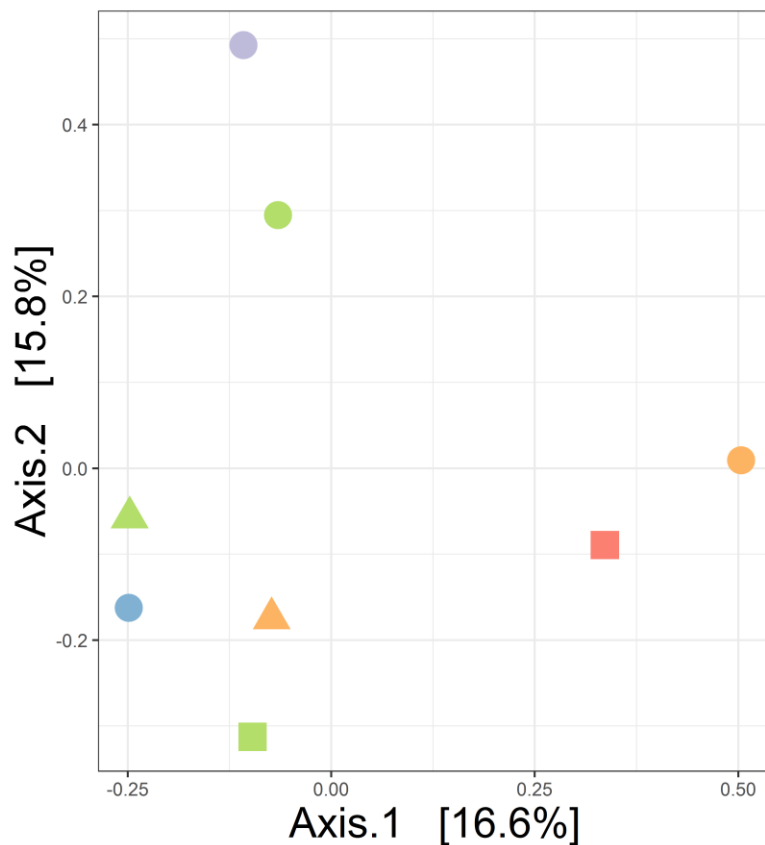


Beta diversity: PCoA on Jaccard distances

- Significant effect of water mass on community structure
- Strong segregation surface vs deep
- MOW always intermediate, but strong variations with location

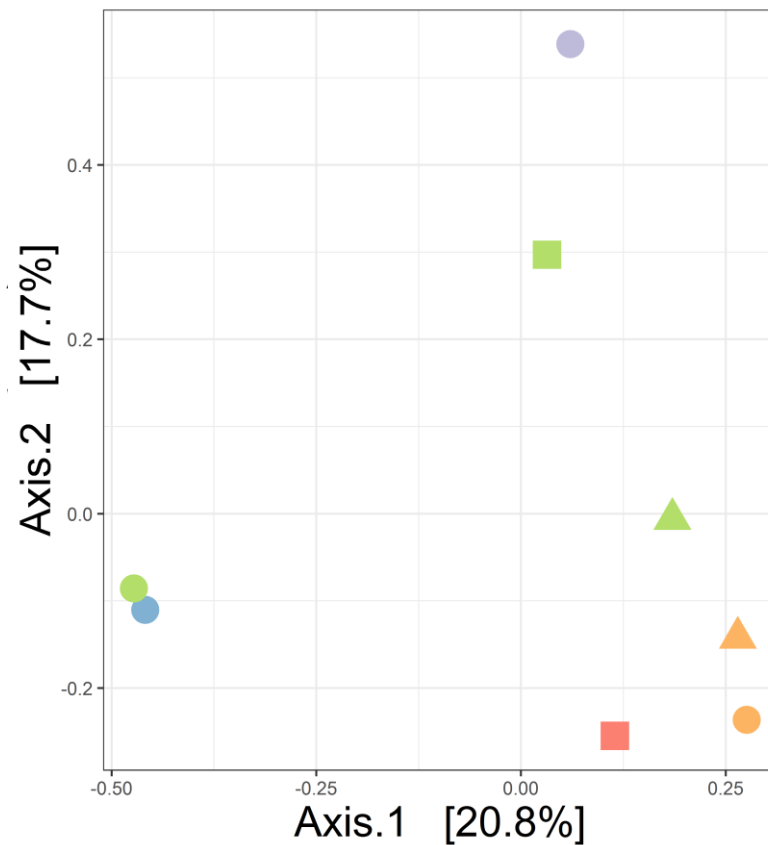
• 18S-V1

Jaccard PCoA



• COI

Jaccard PCoA



Location

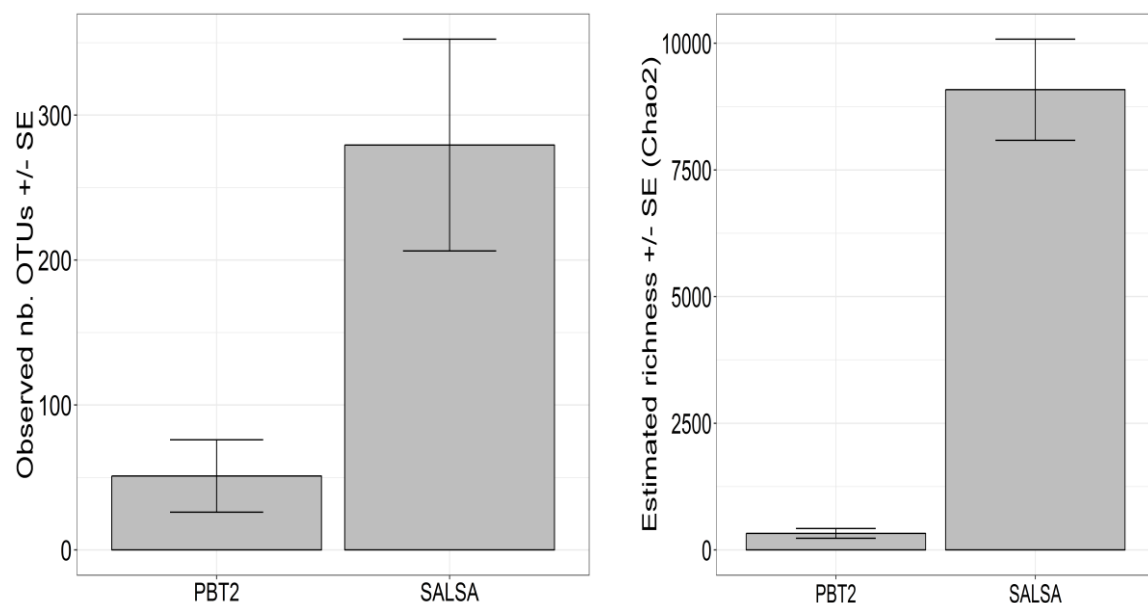
- Azores
- ▲ Southwest Portugal
- Alboran Sea

water.mass

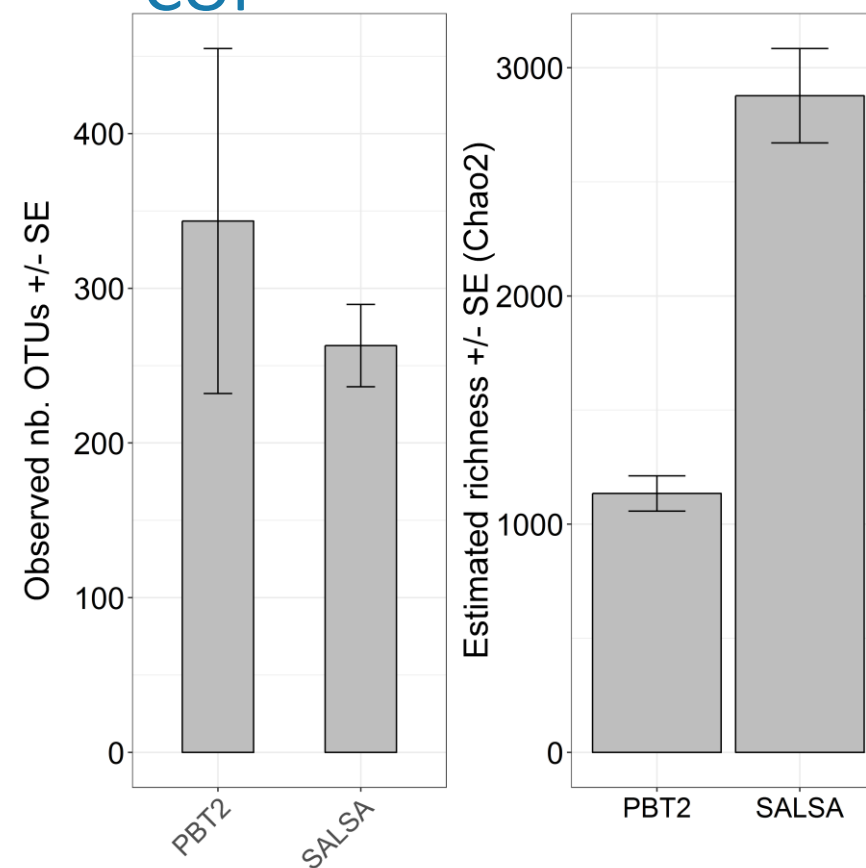
- LIW
 - NACW
 - AAIW
 - MOW/MDW
 - NADW
- } Surface
 } Antarctic
 } Deep

Water pilot study: sampling methods in EssNaut

• 18S-V1



• COI



- 12 phyla detected in PBT2, 21 phyla in SALSA
- Similar taxonomic composition for main phyla

- 12 phyla detected in PBT2, 13 phyla in SALSA
- Similar taxonomic composition