

eDNA protocols and application to Medwave samples

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Jeannette Carlsson, Jens Carlsson, Marina Carreira, Telmo Morato, Miriam Brandt, Covadonga Orejas,
Sophie Arnaud-Haond



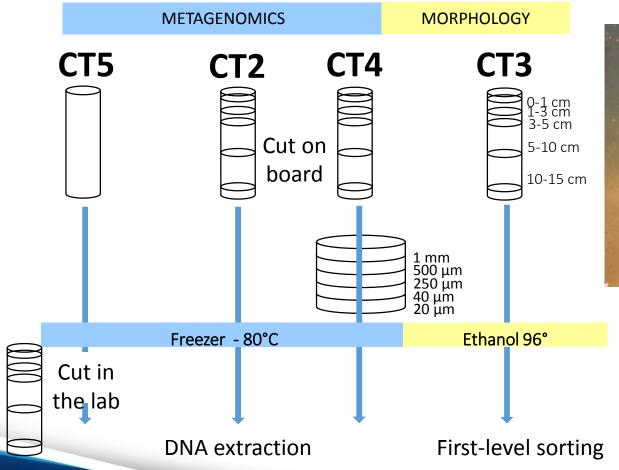
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Setting up metabarcoding protocols to unravel communities through eDNA



1) Sampling design



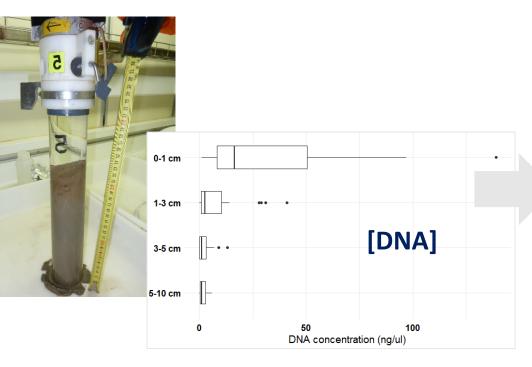


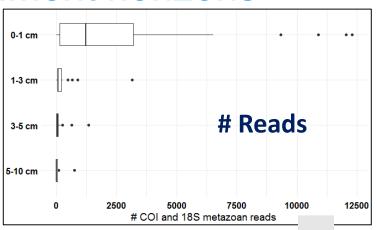
ROV-deployed cores 2400 m

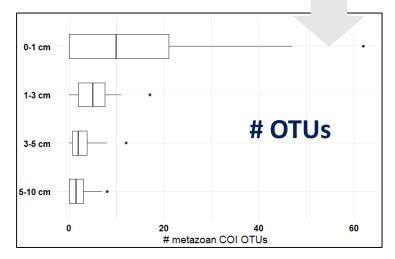




Metazoan results and sediment horizons

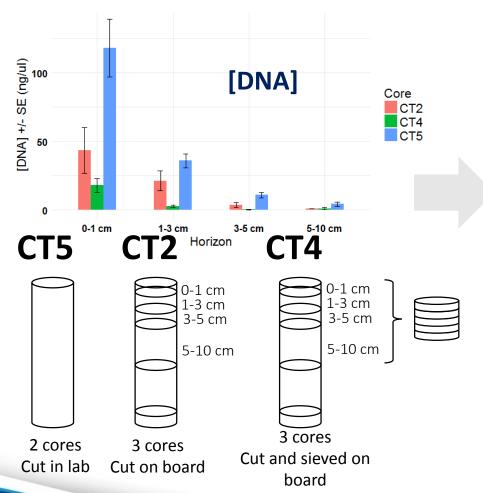


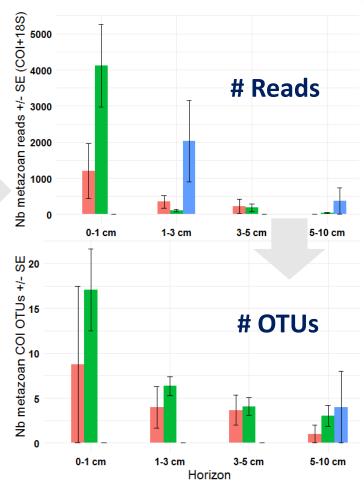




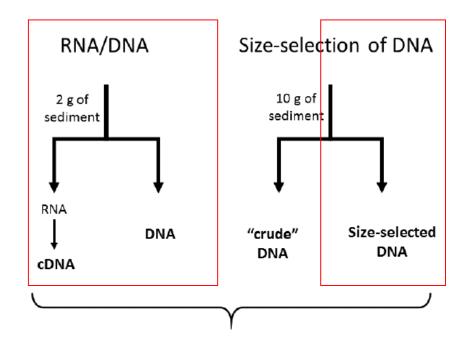


Differences between core treatments?

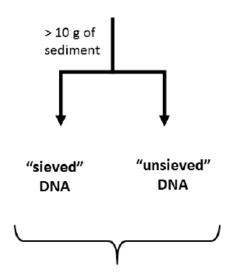




Tests in course with ABYSS project, including Medwaves samples



Sieving of sediment



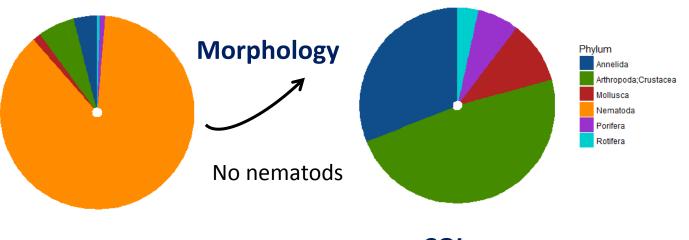
- EssNaut: 1 site, CT2 of PL06, PL07, PL11
- MDW: 4 sites, 1 core from each (CT2)

 EssNaut: 1 site, CT2 (unsieved) and CT4 (sieved) of PL06, PL07, PL11



First results with DNA/Morphology





Taxa

Others

Chordata

Cnidaria

Mollusca

Nemertea

Placozoa

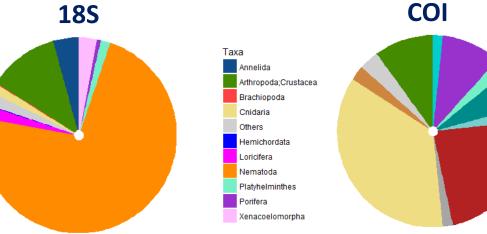
Porifera

Rotifera

Platyhelminthes

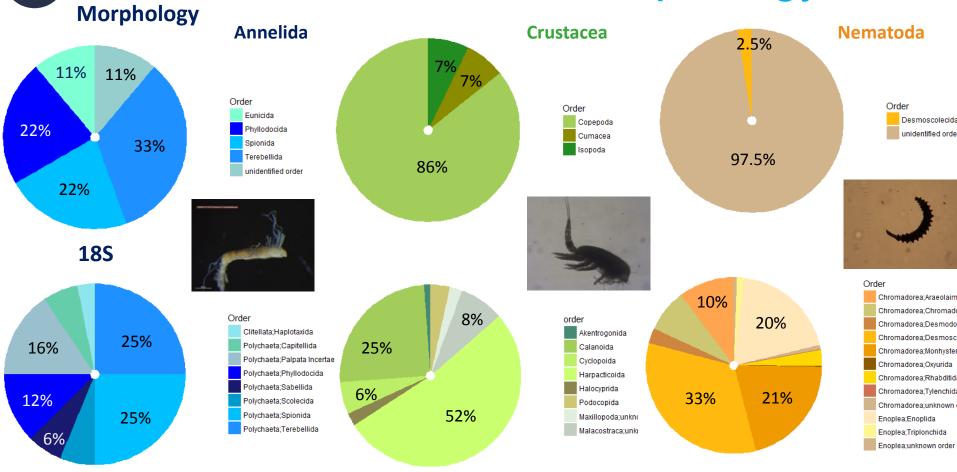
Arthropoda:Crustacea

Metazoa;Multi-affiliation





First results with DNA/Morphology





In course

- Remaining tests on RNA and size selected DNA
- Other comparisons including Medwaves samples (IEO)
- Selection of bioinformatic suites finalized in the framework of Ifremer ABYSS project, that will be used for Atlas analysis as well

Thank You!



Presenter details:

Name, details go here...

WP3: eDNA protocols set up at Ifremer & UCD In collaboration with IEO and IMAR

Project Contact Details:

Coordination: Professor Murray Roberts

murray.roberts@ed.ac.uk

Project Management: Dr. Katherine Simpson

katherine.simpson@ed.ac.uk

Communication & Press: Dr. Claudia Junge

claudia@aquatt.ie

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