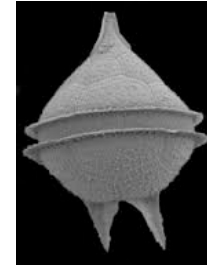
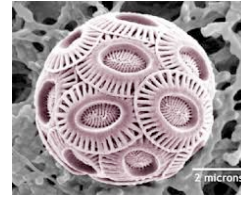
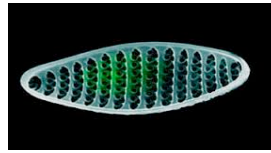
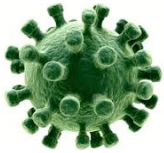


Environmental DNA as a novel approach for studying marine ecosystems in Faroese waters



Ian Salter, Havstovan

What is environmental DNA?



Environmental Forensics

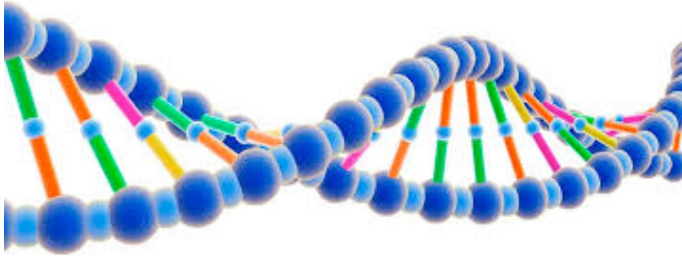
What is environmental DNA?



Criminal Forensics



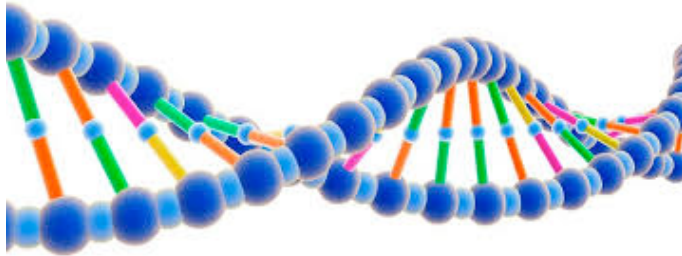
Environmental Forensics



What is environmental DNA?



Criminal Forensics



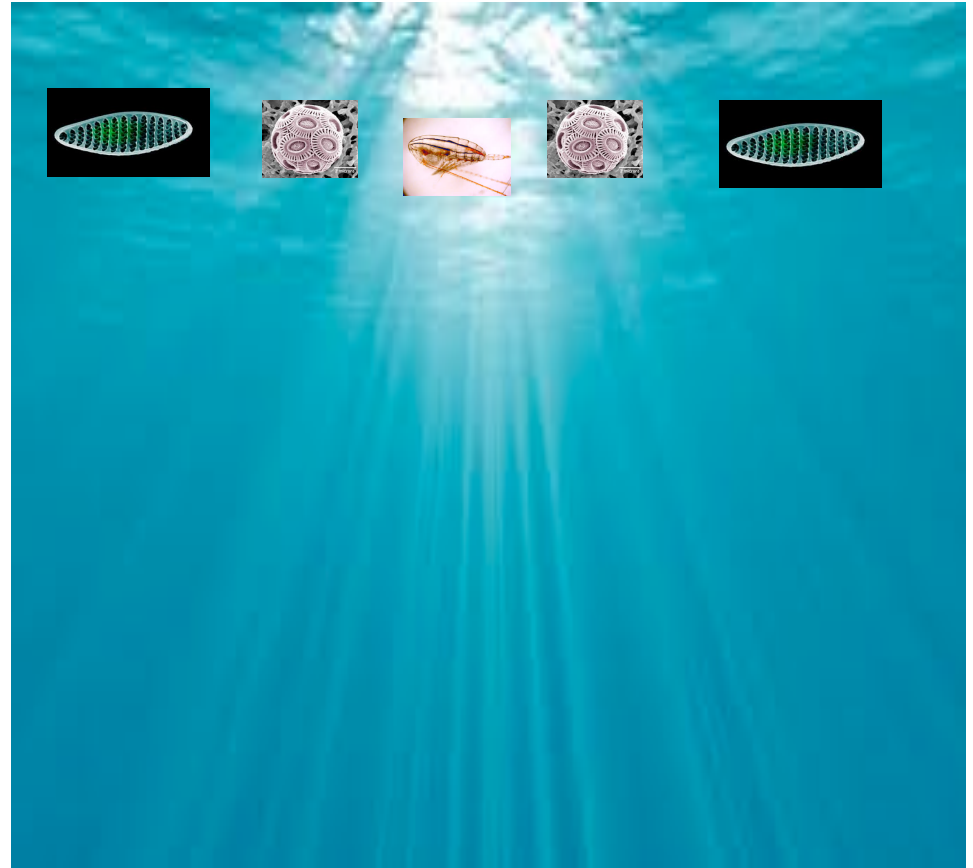
Environmental Forensics



What is environmental DNA?



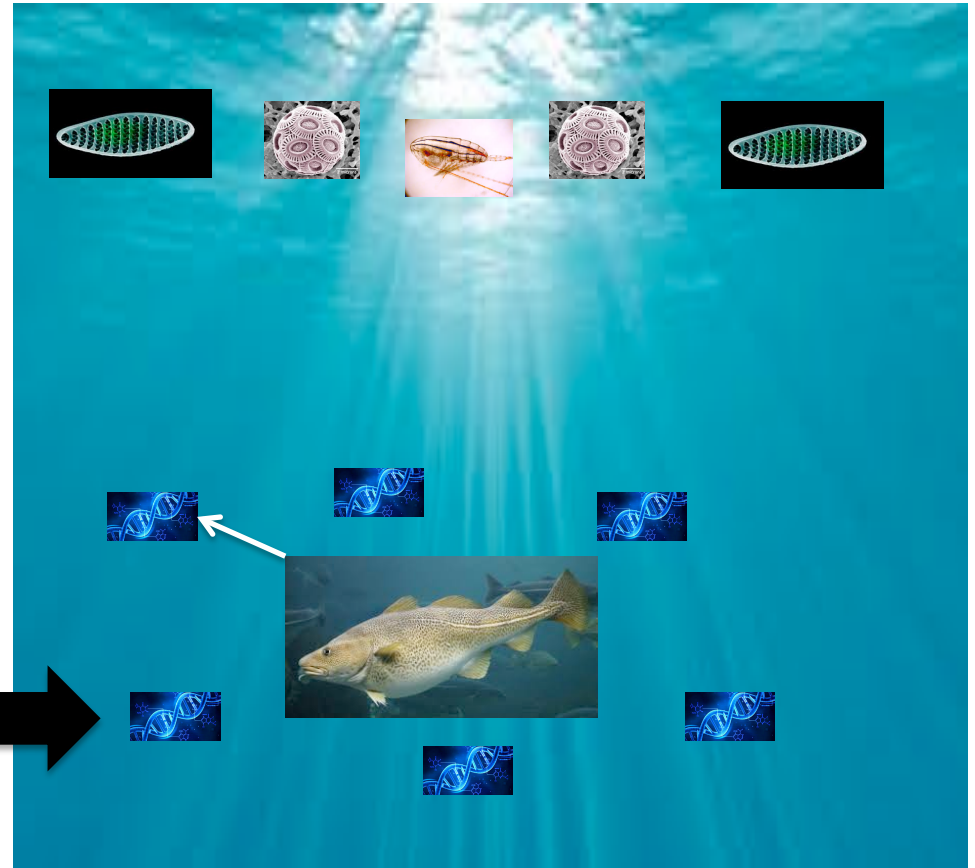
Environmental Forensics



What is environmental DNA?



Environmental Forensics



Skin cells and mucous,
sperm, eggs, secretions,
faeces, urine, blood

etc..

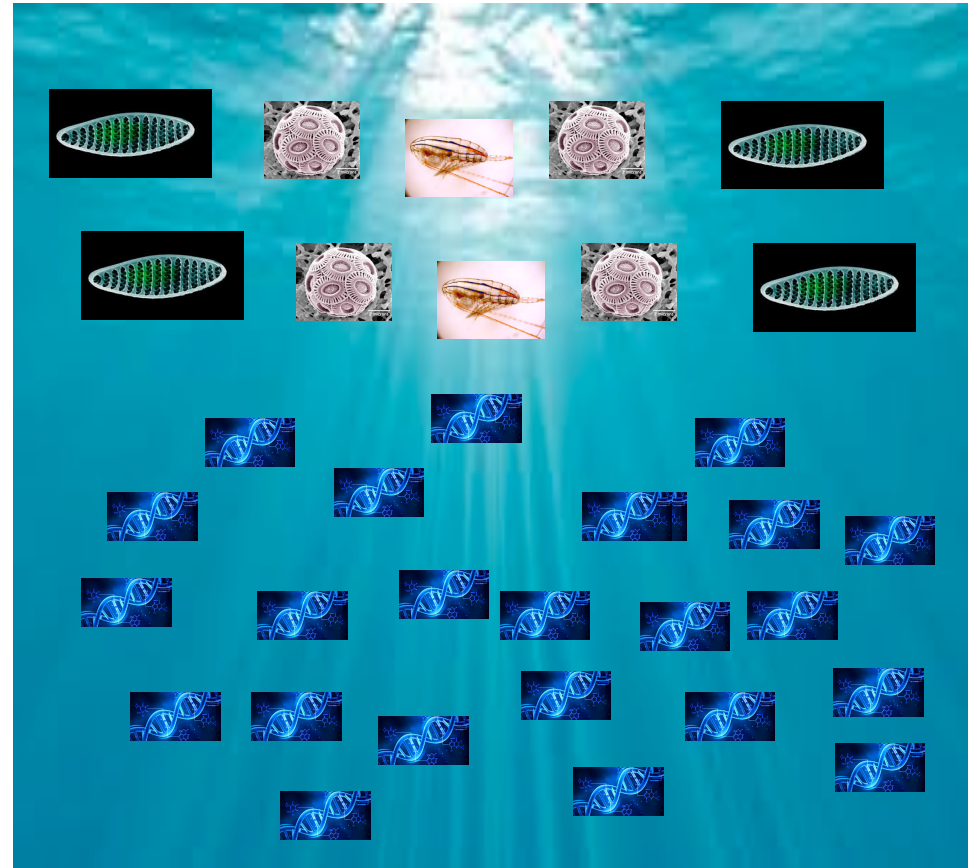
What is environmental DNA?



Environmental Forensics



Every litre of seawater
is like a "DNA-soup"



How can we use environmental DNA?



How can we use this “soup” of environmental DNA in seawater to learn something about ocean ecosystems?

How can we use environmental DNA?



We can use DNA molecules to describe species inventories

How can we use environmental DNA?



We can use DNA molecules to describe species inventories

Uses short sequences of DNA from genes present in all animals, but different depending on the species – fingerprint

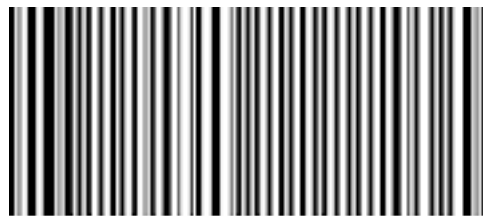
How can we use environmental DNA?



We can use DNA molecules to describe species inventories

Uses short sequences of DNA from genes present in all animals, but different depending on the species – fingerprint

Supermarket scanner uses patterns of black lines in a barcode



(00) 0 0123456 000000001 8



(00) 0 0123456 000000001 8

How can we use environmental DNA?



We can use DNA molecules to describe species inventories

Uses short sequences of DNA from genes present in all animals, but different depending on the species – fingerprint

Supermarket scanner uses patterns of black lines in a barcode



MIKLAGARÐUR



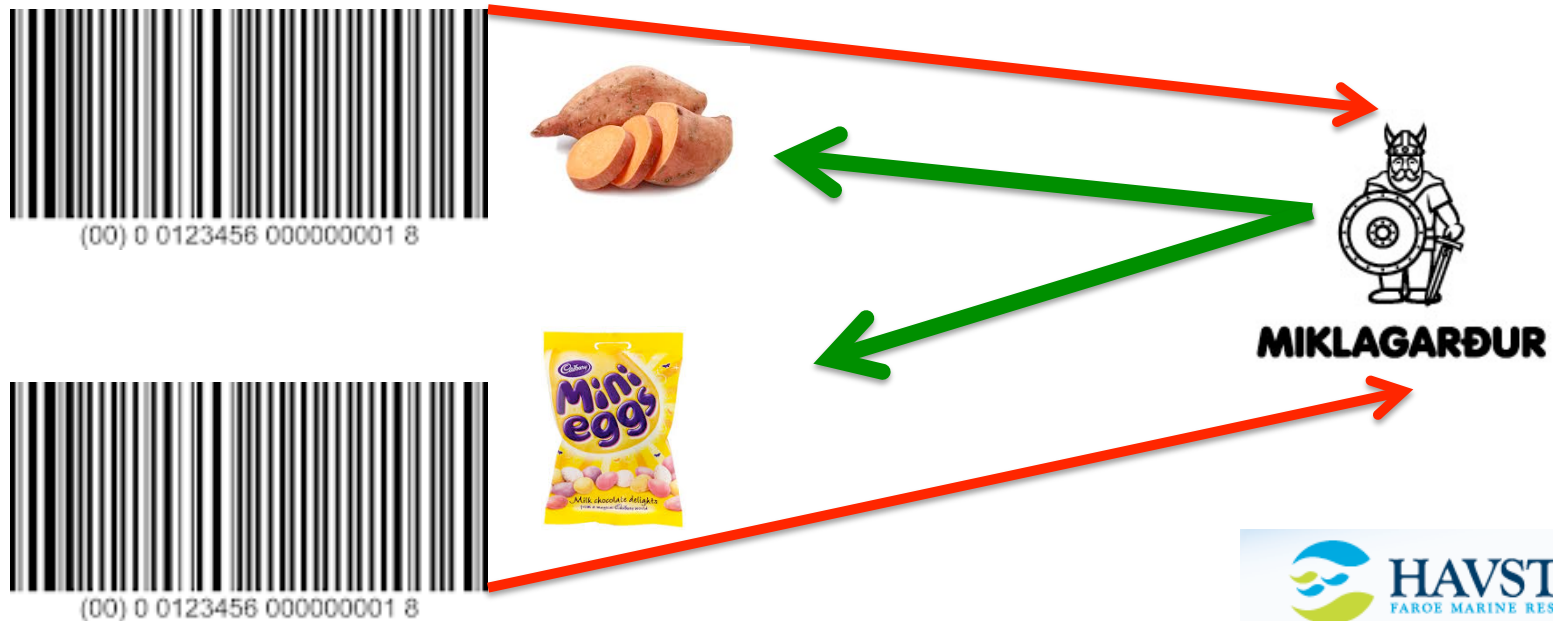
How can we use environmental DNA?



We can use DNA molecules to describe species inventories

Uses short sequences of DNA from genes present in all animals, but different depending on the species – fingerprint

Supermarket scanner uses patterns of black lines in a barcode



How can we use environmental DNA?

We can use DNA molecules to describe species inventories

Uses short sequences of DNA from genes present in all animals, but different depending on the species – fingerprint

Supermarket scanner uses patterns of black lines in a barcode

Metabarcoding

AGCCCGATTGCCAGT
ACGATTACGATATAA
ATCGAGGACATATA

AGCCCGATTGCCAGT
ACGATTACGATATAA
ATCGAGGACATATA



Computer
Database

How can we use environmental DNA?



BOLD SYSTEMS DATABASES IDENTIFICATION TAXONOMY WORKBENCH RESOURCES LOGIN

TAXONOMY

Kingdoms of Life Being Barcoded

 SEARCH TAXONOMY

7,826,119

Specimen Records

5,803,721

Specimens with Barcodes

268,055

Species with Barcodes

Animals:

- Acanthocephala [588]
- Annelida [58603]
- Arthropoda [6282571]
- Brachiopoda [226]
- Bryozoa [2776]
- Chaetognatha [589]
- Chordata [551571]
- Cnidaria [18451]
- Cycliophora [326]
- Echinodermata [40678]
- Gnathostomulida [18]
- Hemichordata [163]
- Mollusca [139953]
- Nematoda [14173]
- Nemertea [2178]
- Onychophora [690]
- Platyhelminthes [16723]
- Porifera [3840]
- Priapulida [44]
- Rotifera [9743]
- Sipuncula [757]
- Tardigrada [1985]
- Xenoturbellida [7]

Plants:

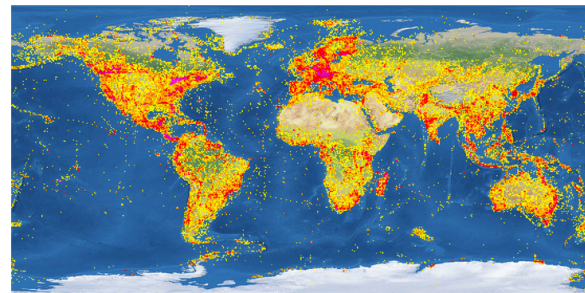
- Bryophyta [11214]
- Chlorophyta [14677]
- Lycopodiophyta [1129]
- Magnoliophyta [338188]
- Pinophyta [6870]
- Pteridophyta [10475]
- Rhodophyta [49214]

Fungi:

- Ascomycota [83592]
- Basidiomycota [54951]
- Chytridiomycota [277]
- Glomeromycota [3516]
- Myxomycota [226]
- Zygomycota [3137]

Protists:

- Chlorarachniophyta [67]
- Ciliophora [706]
- Heterokontophyta [6195]
- Pyrrophytophyta [2281]



How can we use environmental DNA?

We can use DNA molecules to describe species inventories

Uses short sequences of DNA from genes present in all animals, but different depending on the species – fingerprint

Supermarket scanner uses patterns of black lines in a barcode

Metabarcoding

AGCCCGATTGCCAGT
ACGATTACGATATAA
ATCGAGGACATATA



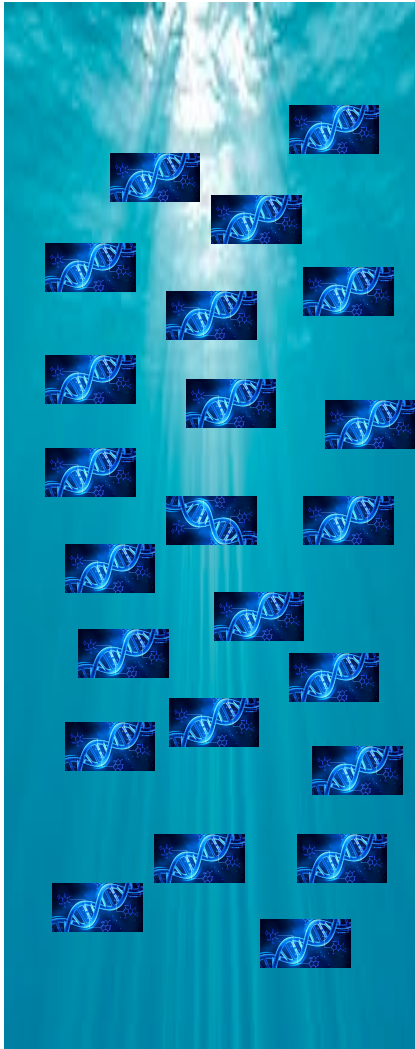
AGCCCGATTGCCAGT
ACGATTACGATATAA
ATCGAGGACATATA



Computer
Database

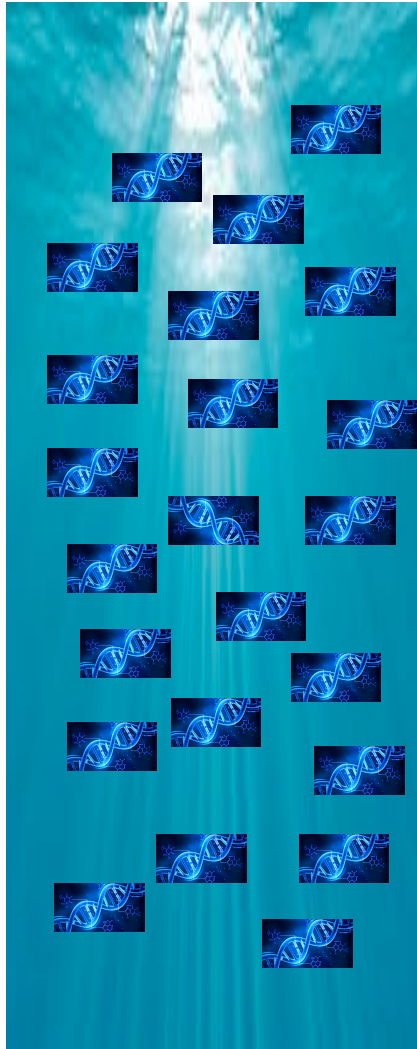
How can we use environmental DNA?

DNA soup

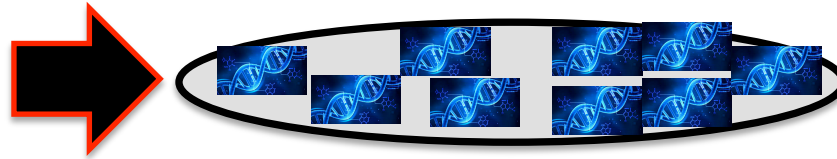


How can we use environmental DNA?

DNA soup

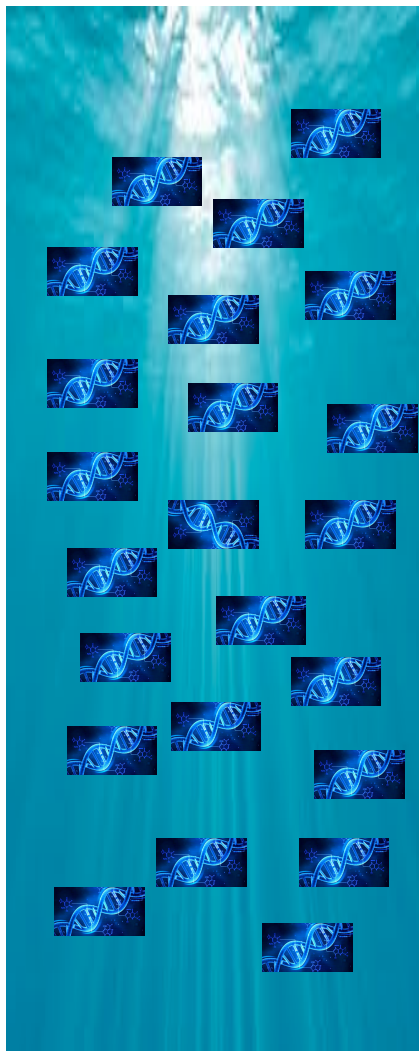


Collect and extract DNA

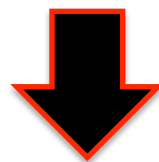
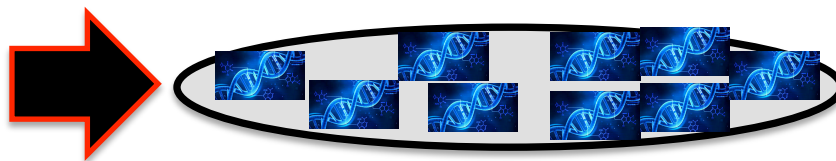


How can we use environmental DNA?

DNA soup



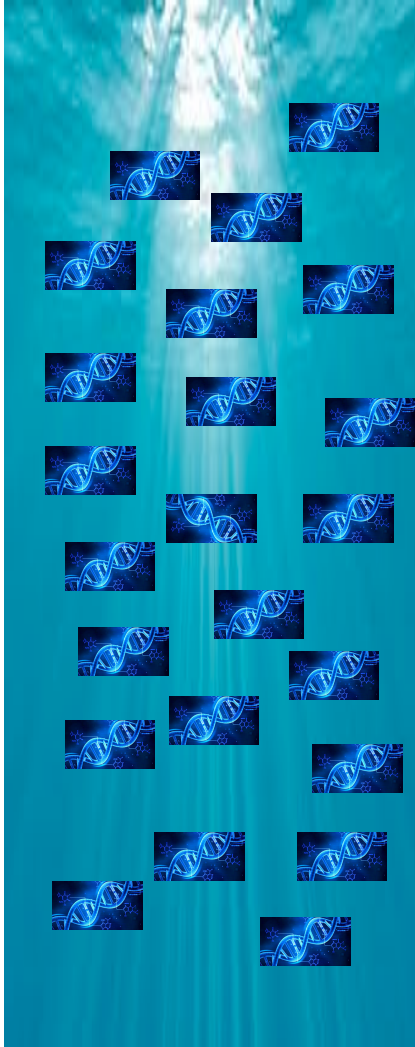
Collect and extract DNA



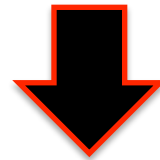
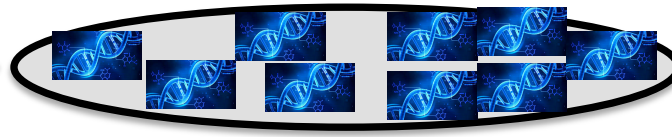
DNA sequencer

How can we use environmental DNA?

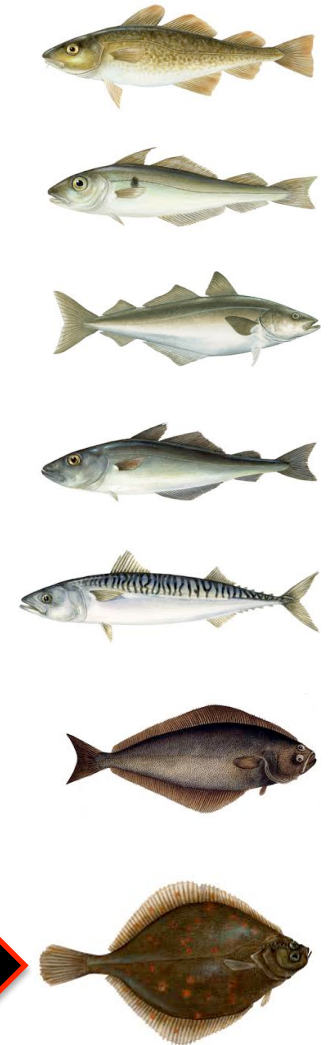
DNA soup



Collect and extract DNA



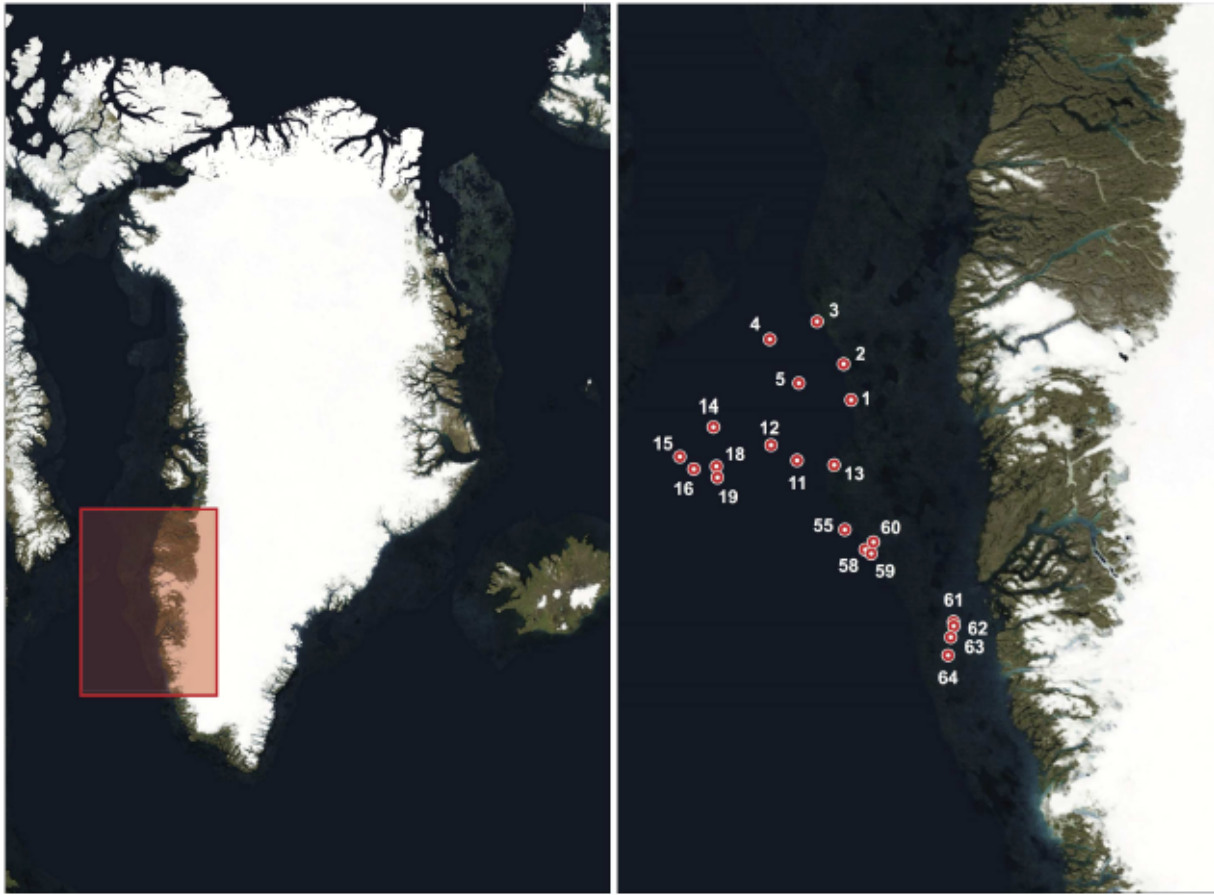
DNA sequencer



Species inventory

Passive Monitoring of bottom fish communities

How well does eDNA monitoring perform in reality?



Bottom trawl samples

eDNA Monitoring of bottom fish communities

How well does eDNA monitoring perform in reality?



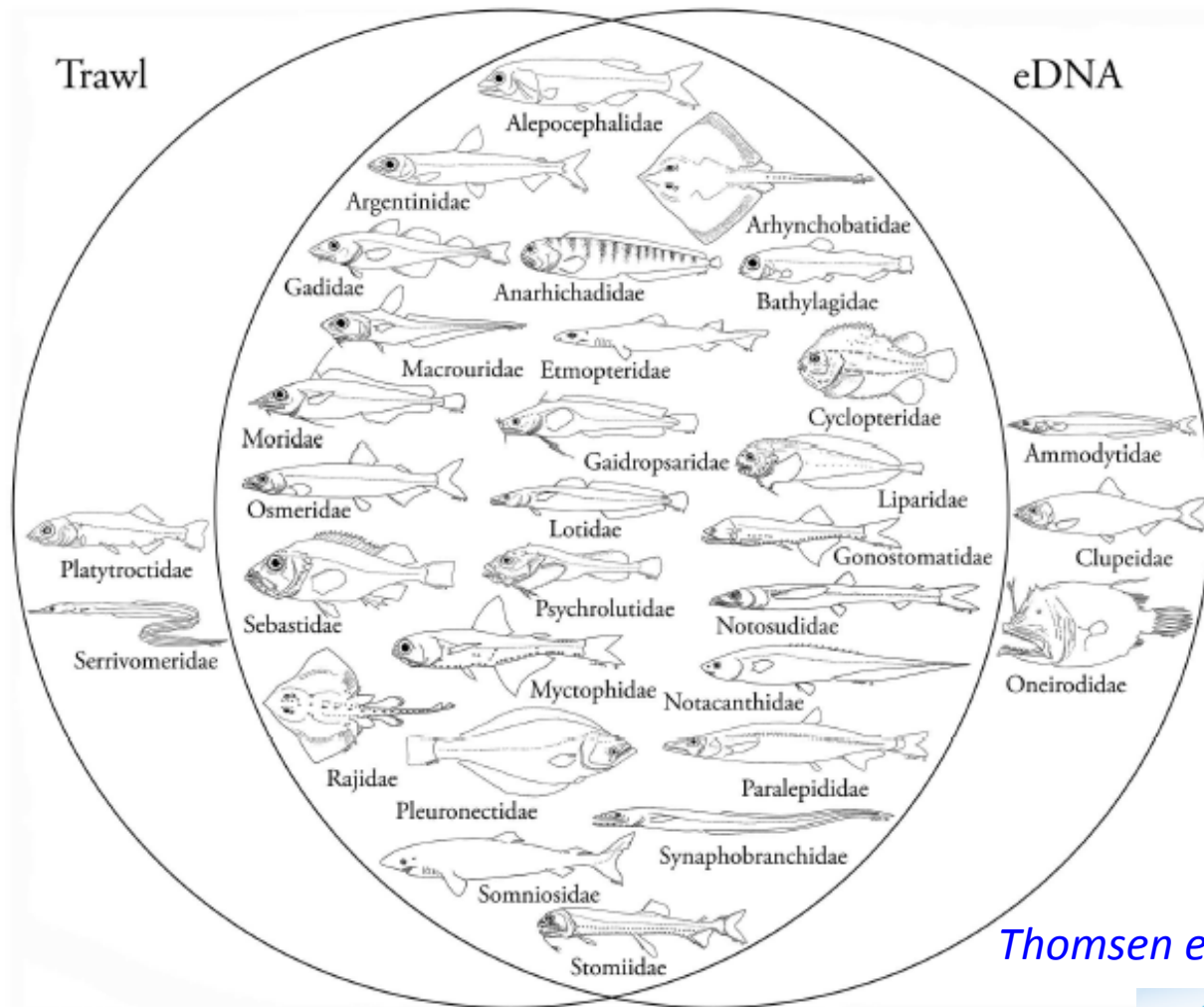
2 litre



Thomsen et al. 2016 PLoS ONE

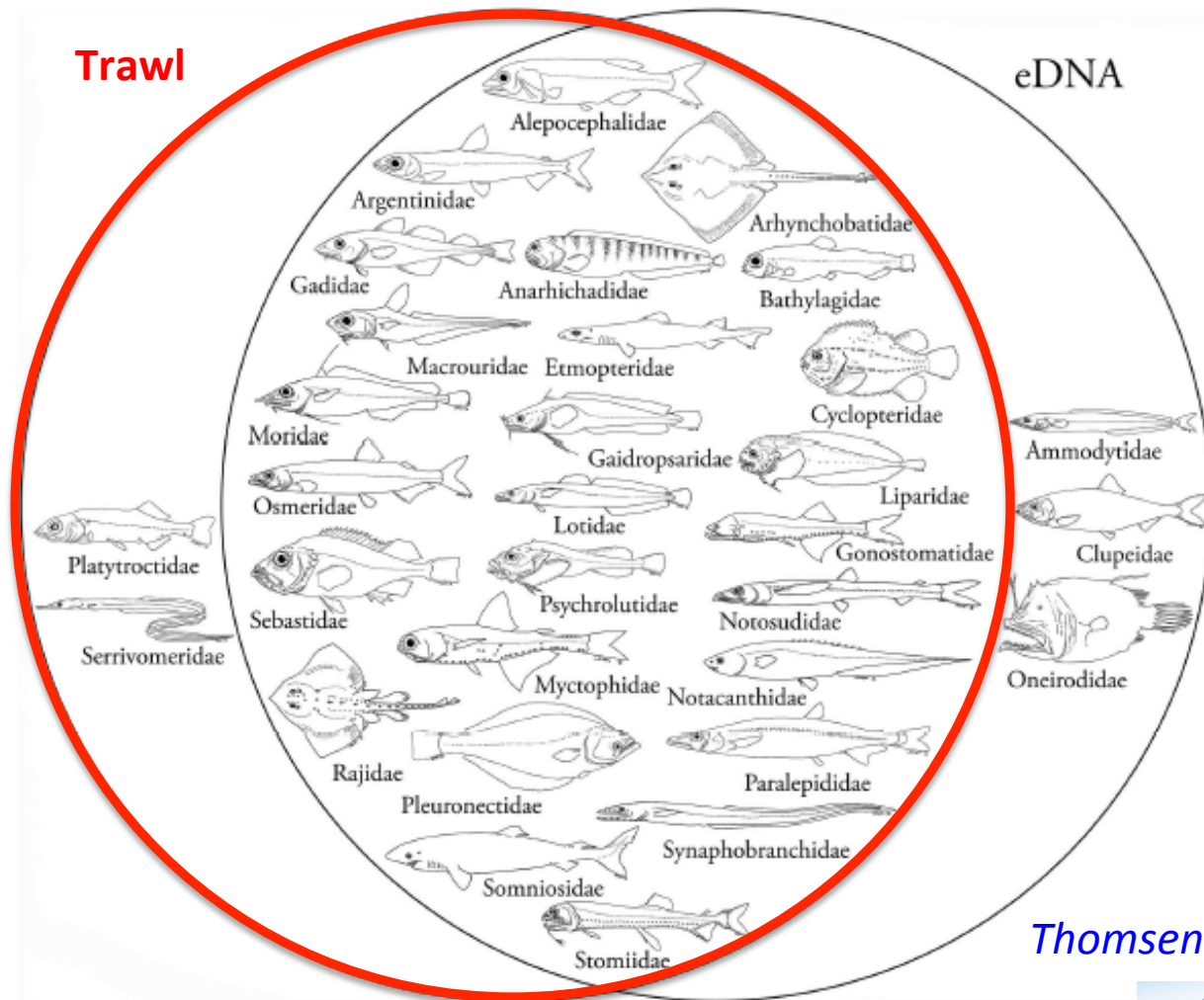
1 hour bottom trawl

eDNA Monitoring of bottom fish communities



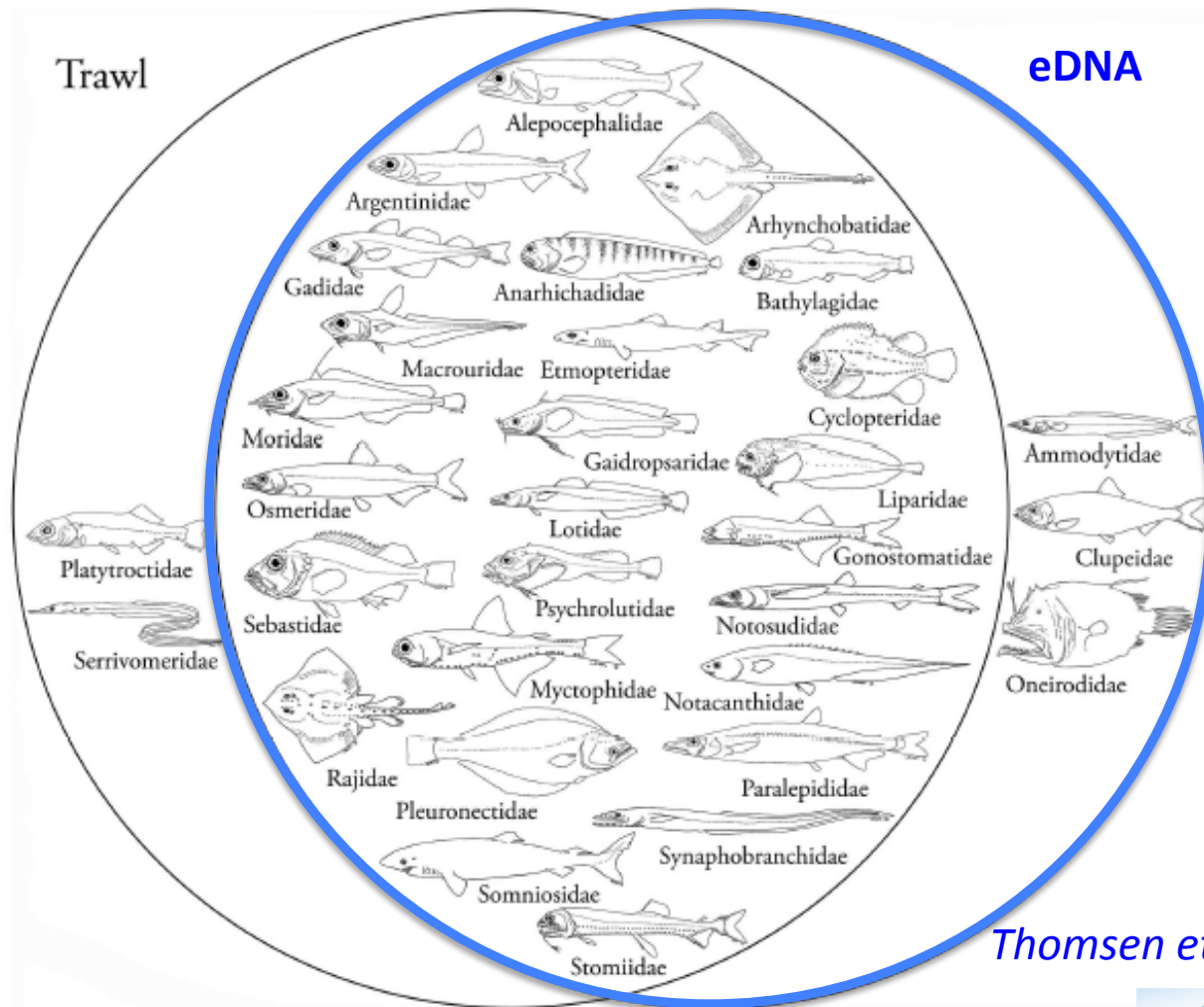
Thomsen et al. 2016 PLoS ONE

eDNA Monitoring of bottom fish communities



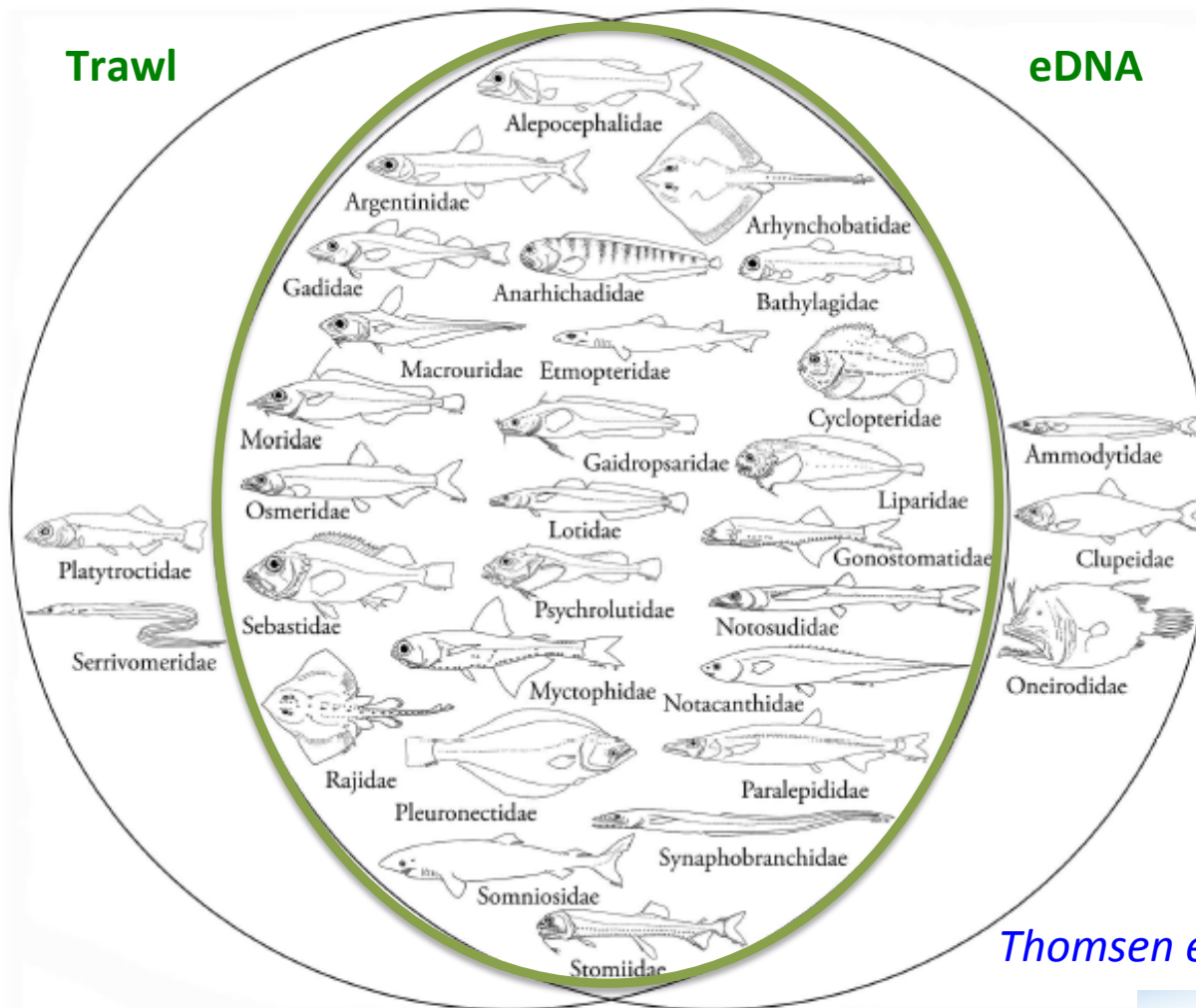
Thomsen et al. 2016 PLoS ONE

eDNA Monitoring of bottom fish communities



Thomsen et al. 2016 PLoS ONE

eDNA Monitoring of bottom fish communities



26 of 28

>93%
Detection
rate

Thomsen et al. 2016 PLoS ONE

eDNA Monitoring of bottom fish communities

eDNA monitoring works very well for describing presence of different species

Incredibly useful for invasive species monitoring, pathogens, species distribution modeling etc.

eDNA Monitoring of bottom fish communities



eDNA monitoring works very well for describing presence of different species

Incredibly useful for invasive species monitoring, pathogens, species distribution modeling etc.

How about as a quantitative approach?

Does the concentration of cod DNA in seawater reflect the biomass of cod in a given location?

eDNA monitoring of bottom fish communities

COD-e-DNA (Fisheries Ministry)
October 2017 - October 2019

Relationship between cod biomass and cod DNA copies on the Faroese Bank

Compare cod biomass from trawls with cod
DNA molecules in seawater samples



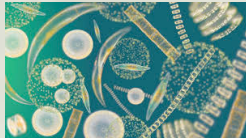
eDNA monitoring of entire ecosystems

1-2 Litres
Seawater

Bacteria



Phytoplankton



Zooplankton



Fish



Marine mammals



Marker Genes

16S Ribosomal

18S Ribosomal

COI Mitochondria

12S Mitochondria

COI Mitochondria

Ecosystem dynamics

Species distribution

Biogeography

Diversity

Food-web structure

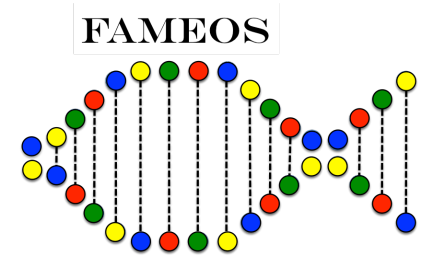
Invasive species

eDNA monitoring of entire ecosystems

FAMEOS (Faroese Research Council)

January 2018 - December 2019

Faroese Marine Ecosystem Observing Study



Faroese Marine Ecosystem Observing Study

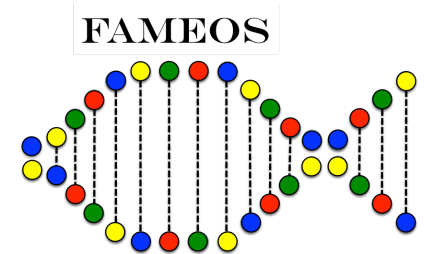
eDNA monitoring of entire ecosystems



FAMEOS (Faroese Research Council)

January 2018 - December 2019

Faroese Marine Ecosystem Observing Study



Faroese Marine Ecosystem Observing Study

Integrating environmental-DNA based estimates of diversity with essential ocean variables

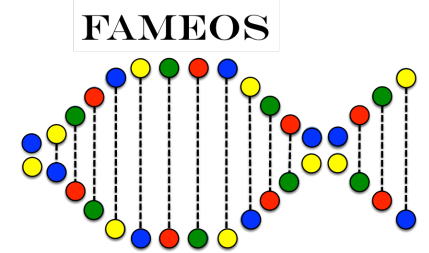
eDNA monitoring of entire ecosystems



FAMEOS (Faroese Research Council)

January 2018 - December 2019

Faroese Marine Ecosystem Observing Study



Faroese Marine Ecosystem Observing Study

Integrating environmental-DNA based estimates of diversity with essential ocean variables

Objectives

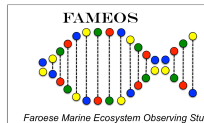
Time-series monitoring program of marine biodiversity and ecosystem dynamics in Faroese waters

Establish a robust baseline to understand the effects of climate variability and anthropogenic pressures

FAMEOS

Faroese Marine Ecosystem Observing Study

Ian Salter (ians@hav.fo) (Havstovan); Sólva Jacobsen (Havstovan); Durita Sørensen (Havstovan); Eilíf Gaard (Havstovan); Katja Metfies (AWI, Germany); Halina Tegetmeyer (Uni. Bielefeld, Germany).



Rationale

The biology of the oceans is critical for our survival. It provides half of the oxygen we breathe, absorbs our CO₂ emissions and forms the basis of a marine food-web that provides 1/6th of the protein we consume globally. Natural and anthropogenic pressures are changing the physical and chemical properties of the oceans in a way that is likely to impact ecosystem structure. This is especially relevant in the Faroe Islands that has such a strong cultural and economic dependence on it's marine environment. It is now considered critical to develop observing systems that integrate physical, chemical and biological components to characterize baselines of marine ecosystem health. Sustained measurements of biodiversity within structured observation frameworks are currently lacking, limiting our capacity to report on changes in their status. FAMEOS aims to develop an observing system in Faroese waters that integrates DNA-based estimates of ecosystem structure and biodiversity with essential ocean variables.

Essential Ocean Variables

Essential Ocean Variables (EOVs) are observation parameters that are able to describe the changing physical and chemical state of our oceans

They are selected by panels of experts that represent supranational organizations such as UNESCO and UNEP and form part of a common Global Ocean Observing System (GOOS)



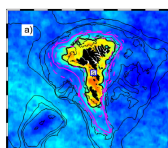
Biodiversity

Biodiversity underpins almost every aspect of human development. It helps produce food, clean water, regulates climate and controls disease transmission.

Biodiversity is being rapidly degraded.

Keeping track of and protecting our planets biodiversity is one of the United Nations Sustainable Development Goals (SDGs).

Phase 1 – Coastal Observations



Weekly monitoring at station Skopun (S)

Representative of shelf ecosystem due to strong tidal mixing

Image courtesy of Sólva Eliassen

Samples will be collected for following parameters:

| | |
|----------------------------|--|
| <u>Existing parameters</u> | <u>New parameters</u> |
| Temperature | ¹³ C and ¹⁵ N |
| Salinity | CaCO ₃ and SiO ₂ |
| Nutrients | Cytometry |
| Chlorophyll | Microbial DNA |
| Plankton | Macrobial DNA |

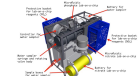
Phase 2 - Technology Corner



FerryBox: Underway monitoring systems on ships of opportunity (RV Norrøna)



FlowCam: Semi-automated image recognition systems for phytoplankton



Autonomous Samplers: Takes and preserves water samples for later analysis

Future Perspectives

FAMEOS will run for three years from January 2018 to December 2020. It is intended to form the basis for launching a sustained marine biodiversity observation program. Future initiatives will extend FAMEOS to include novel sampling platforms like ships of opportunity and autonomous sensor and sampler platforms.

Thank you for your attention



Fiskimálaráðið



RESEARCH PARK
iNOVA

- A new way to do science