

Using different FlowCams to observe grazers

GlobalHAB symposium on automated in situ observations of plankton

Kristineberg, Sweden, August 2022

Terje Berge

Institute of Marine Research, Norway

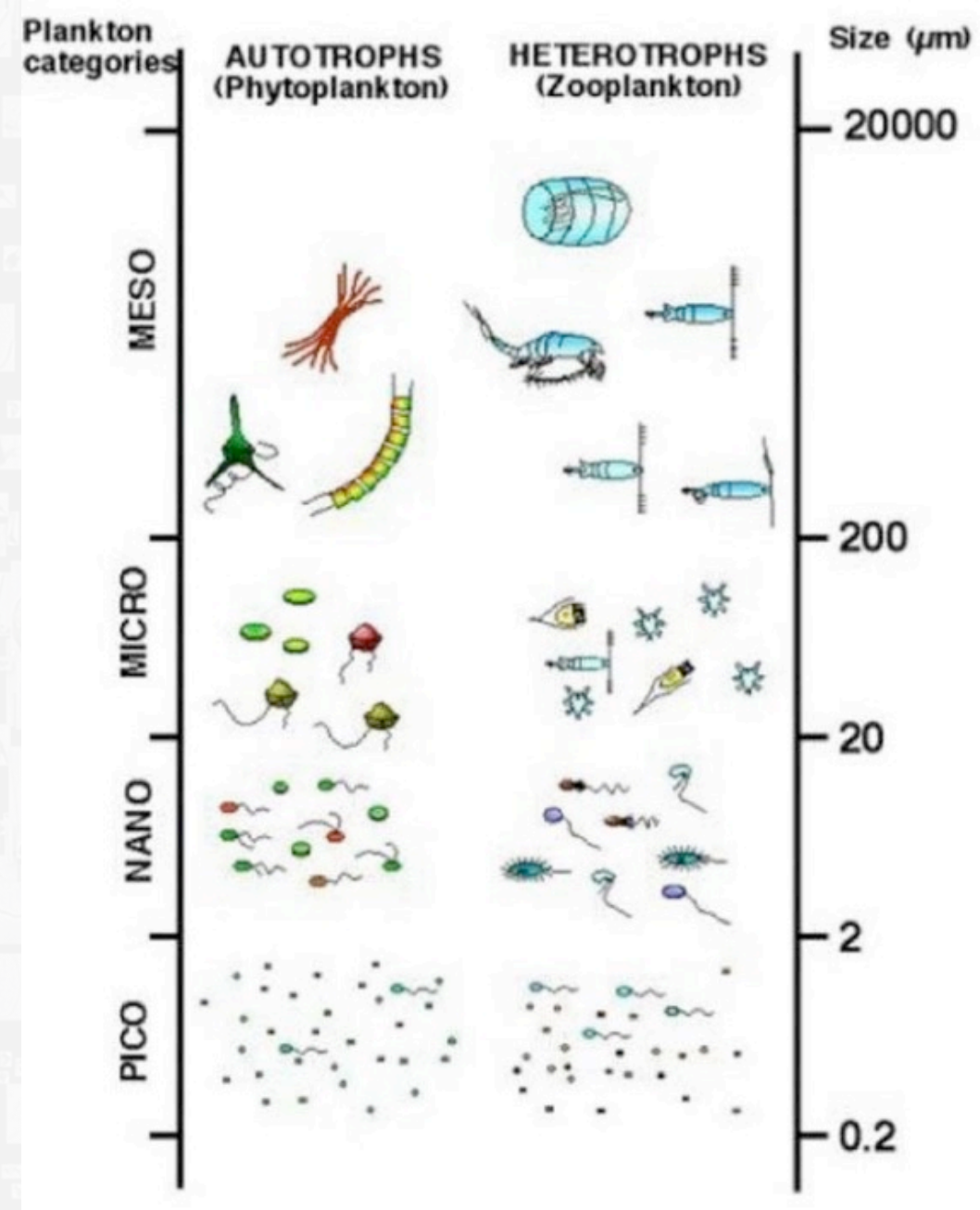


Overview of this presentation

- Objective and challenges
- How we come from FlowCam data to classified data
- Some examples of our use of premature classifiers
- Combining the data from different FlowCams and instruments

Objective

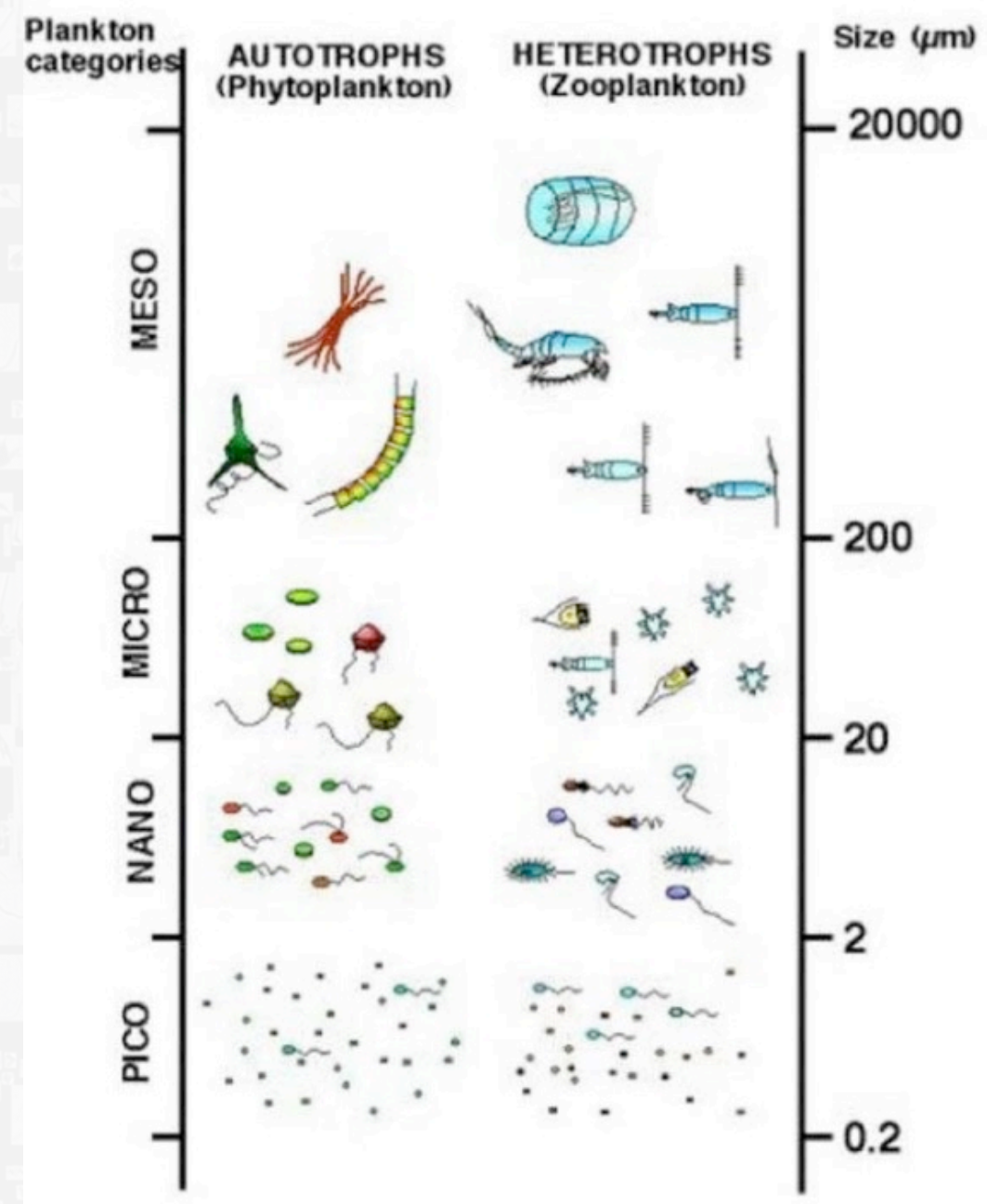
Implement FlowCam in routine monitoring of coastal «grazers» using automatic image recognition at IMR in Norway



From Alcaraz and Calbet 2003

Challenges

- Large size range (5- 8000 μm in length)
 - The need to combine different magnification settings and FlowCam instruments and make multiple training sets
- Often low abundances
 - Requires large volume of water to be imaged
- Many without pigments and fluorescence signals
 - The need to use autoimage-mode and classify large numbers of non-living particles (e.g. detritus, bubbles, background images)
 - Including diatoms
 - All image-types has to be classified



From Alcaraz and Calbet 2003

Different instruments for different plankton size fractions

Flowcam macro, FCM (50-2000 μ m)

- Mesozooplankton
- WP2 and multinet plankton net trawls
- Formalin fixed

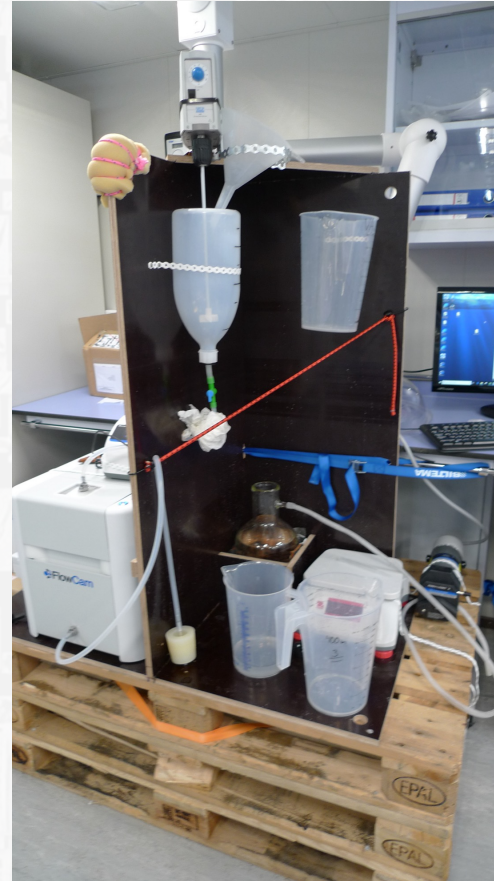
Flowcam VS-1, FCVS (35-500 μ m)

- Microplankton
- **Unfiltered** seawater samples (500 ml)
- Lugol's fixed

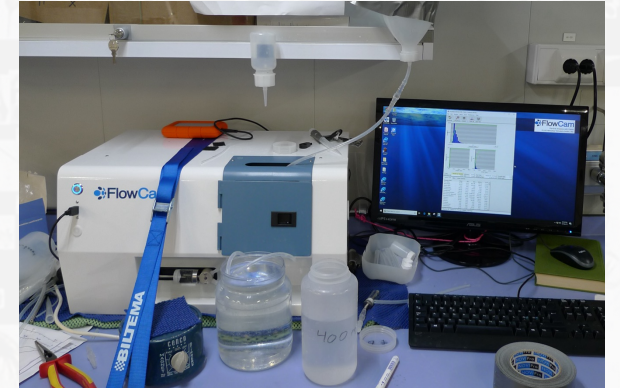
Flowcam 8400, FC8400 (5-300 μ m)

- Nano/microplankton
- Lugol fixed

Flowcam macro



Flowcam VS



Flowcam 8400



Flowcam principle in short

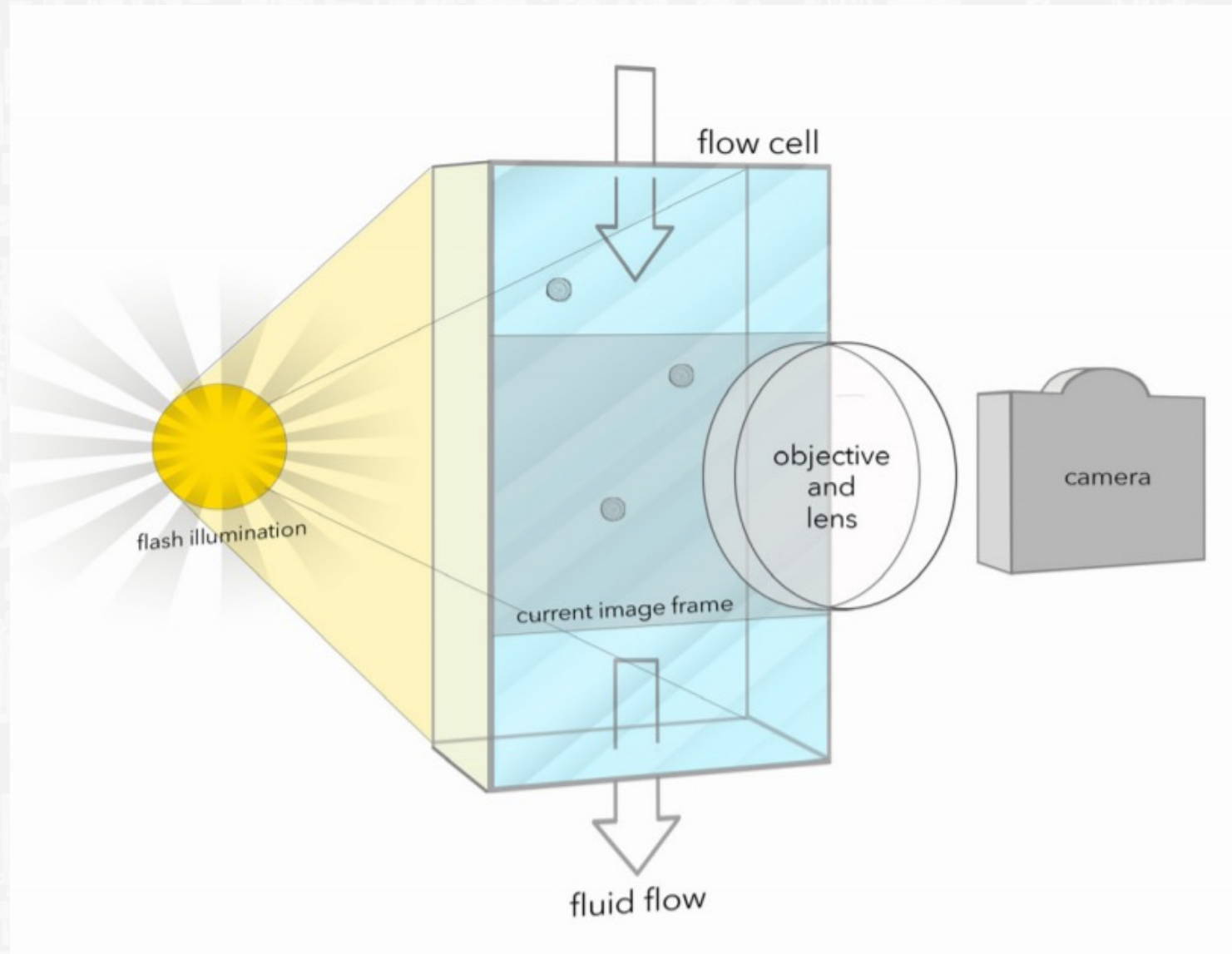
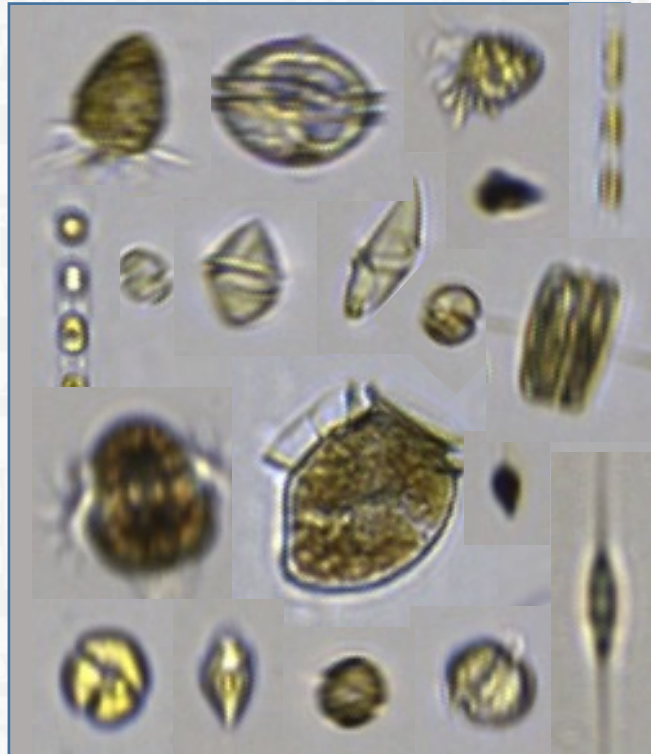


Image examples from FC8400, FCVS and FCM

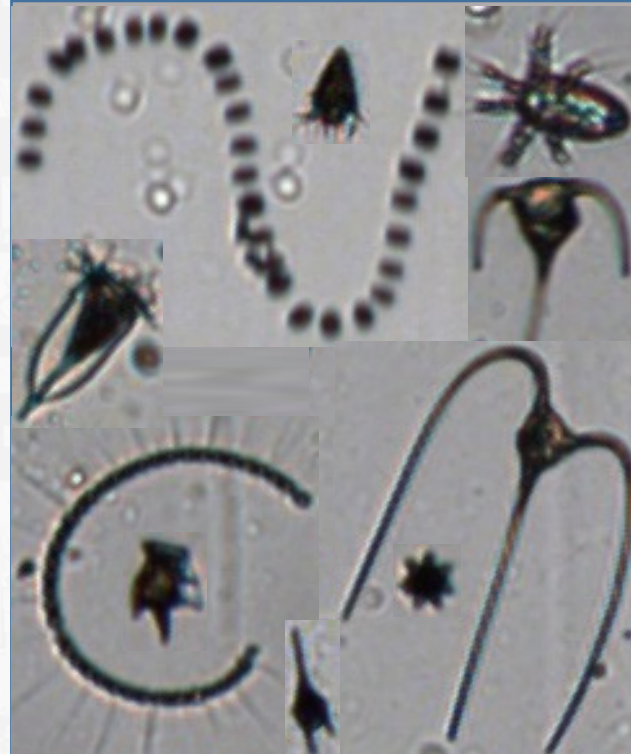
10x objective, 100 μm flowcell



FC8400

Nano- microplankton

2x objective, 800 μm flowcell



FCVS

Microplankton

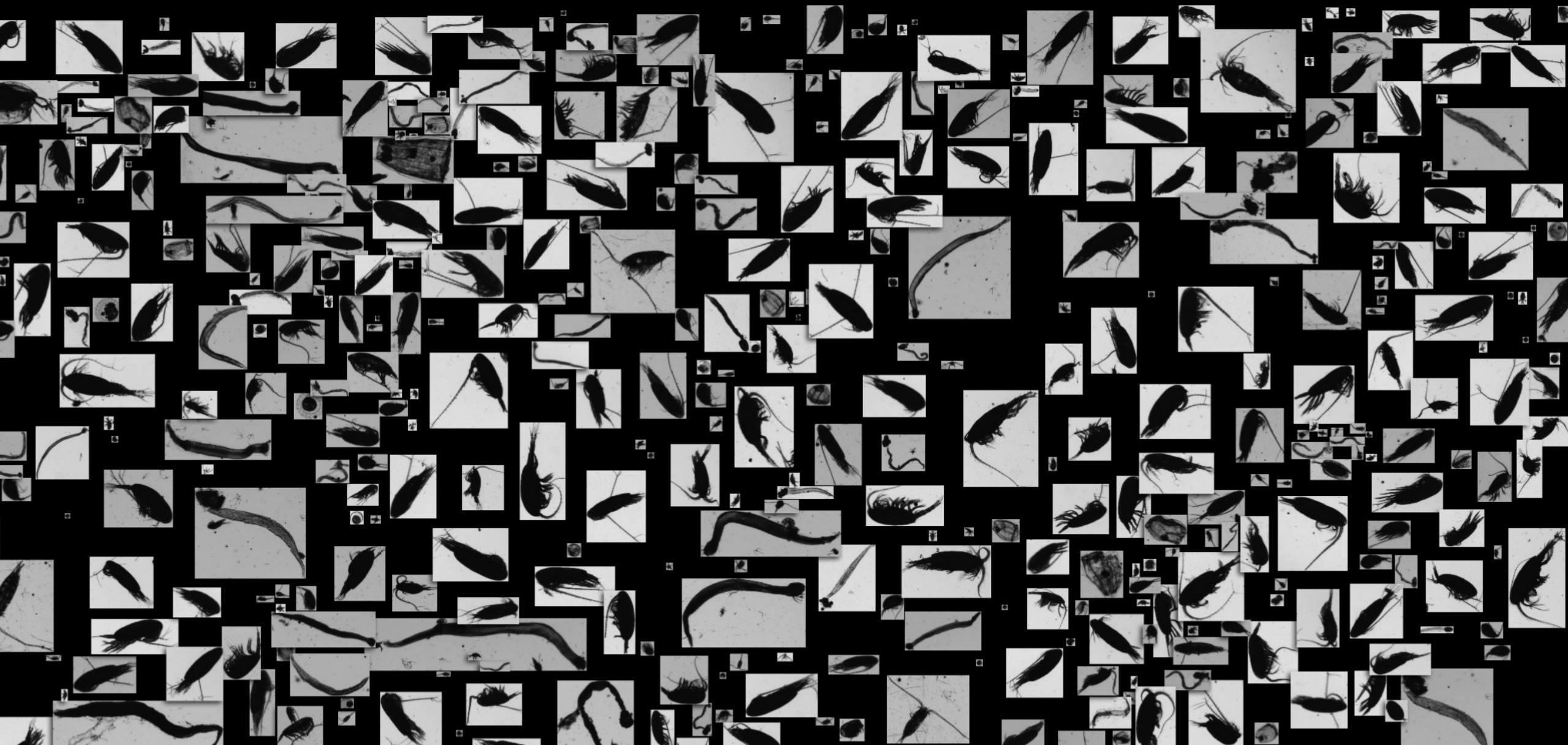
0.5x objective, 800 μm flowcell



FCM

Mesozooplankton

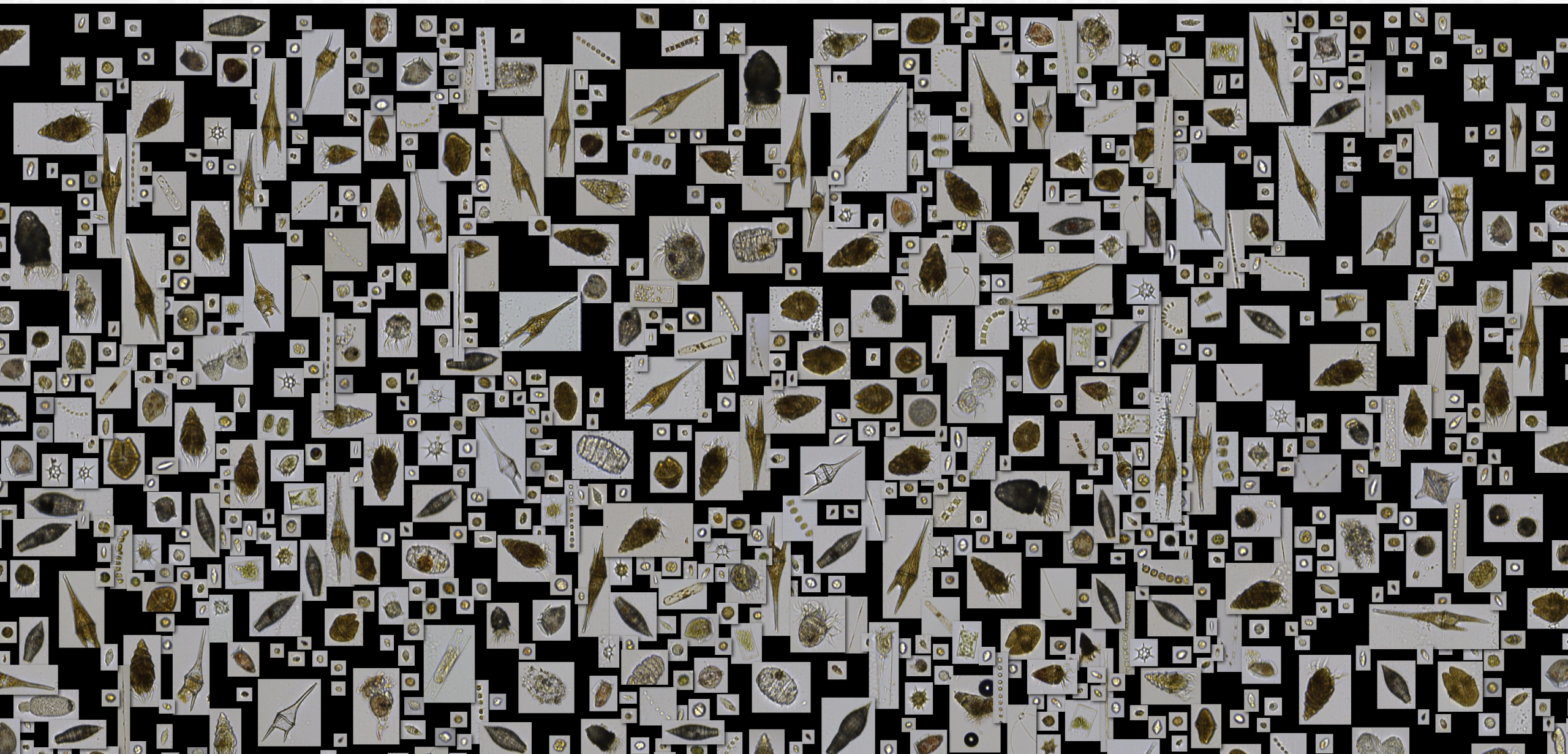
Mesozooplankton – FCM – 12.5 x magnification – 0.5x objective



Micro- and mesozooplankton – FCVS – 20 x magnification – 2x objective



Nano- and mesozooplankton – FCM – 100 x magnification – 10x objective



FlowCams and sample volume

- **Mesozooplankton and FlowCam Macro**

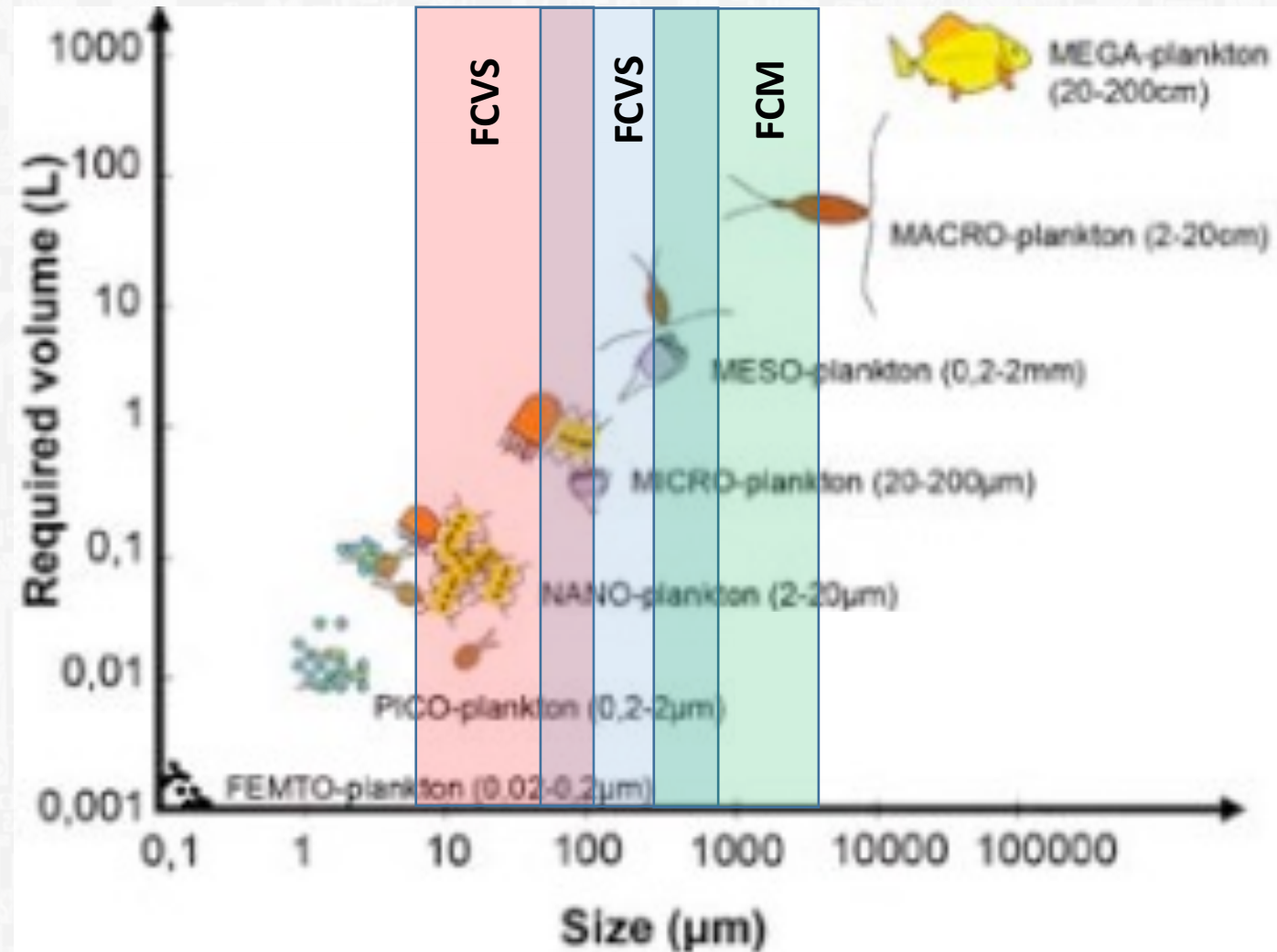
Samples are highly concentrated ($25-1000\text{m}^3$) and need splitting and dilution for imaging with FCM

- **Microzooplankton, HNF`s and mixoplankton**

Samples analysed using two flowcams:

1 - FlowCam VS - 2x objective, $800\mu\text{m}$ flowcell and flowrate of 25 ml min^{-1} . 500ml samples analyses in ca. 20 mins

2 – FlowCam 8400 – 10x objective, $100\mu\text{m}$ flowcell and a flowrate of 0.6ml min^{-1} . 40 ml^{-1} imaged in 25 mins



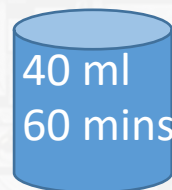
Sampling for coastal microplankton and mesozooplankton

Nano/Microplankton

Whole seawater sample
from 5m Niskin bottle



Filtration through
100 μm meshsize

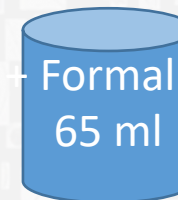


460 ml FCVS with 10x
and 100 μm flowcell

FC8400 with 10x
and 100 μm flowcell

Mesozooplankton

180 μm WP2 and Multinet tows – concentrate of
up to hundreds of m^3



Mesozooplankton < 2000
 μm is worked up manually
at home and with the FCM

Macroplankton >
2000 μm is worked
up manually at sea



Training sets and classifiers

Training sets and classifiers

- We use the r-package «**Zooimage**» (Grosjean et al. 2012) to make training sets and classifiers
- We use the r-package «**fc2zi**» by Eva Álvarez (2011, 2012 and 2014) to extract images and additional particle features from the FlowCam raw collage images in batch (up to hundreds of samples)
- In addition to naming the files according to Zooimage requirements, **fc2zi** performs an instrument artifact (e.g. background images, repeated images etc) filtration procedure and prepares the images for «Zooimage» in batch (hundreds of sample)
- The output from the process (individual images and feature table) can be used for training or classified if samples

Working with training sets

Very easy to share with specialists for annotation of images in any file or image explorer

The diagram illustrates the process of sharing training sets through a file explorer. It shows a directory structure starting with 'This PC > System (C:)' and listing various biological groups like Ciliophora, Cryptophyta, Euglenozoa, Haptophyta, Incertae sedis, Myzozoa, nonliving, and Ochrophyta. A blue arrow points to a second file explorer window showing a hierarchical folder structure: 'Dino...' > Amphidinales > Dinophyceae spp > Dinophysiales > Gonyaulacales > Gymnodinales > Peridinales > Prorocentrales. Another blue arrow points to a third file explorer window showing a grid of image thumbnails with file names like 'FBN029.2021-01-25.001.A1+A1_4' through 'FBN030.2021-01-25.001.A1+A1_66'. A final blue arrow points to a fourth file explorer window showing a grid of image thumbnails with file names like 'FBN011.2019-11-10.001.A1+A1_17' through 'FBN011.2019-11-10.001.A1+A1_154'. The background features a faint world map.

Working with training sets



Taxonomical training set – images are sorted and annotated to highest possible taxonomic level from phyla to species. Images put in annotated folder structure according to the phylogeny found in **WoRMS**

Functional training set

- functionally similar taxa can be grouped together before training

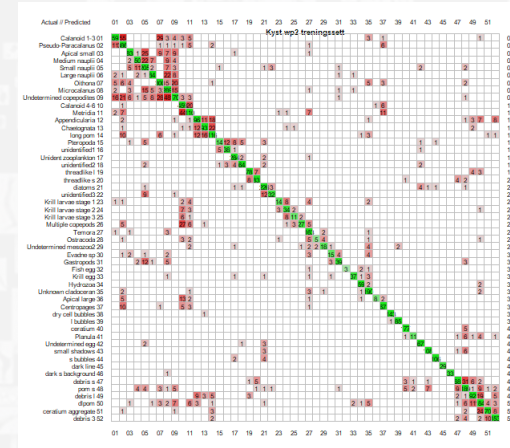
E.g. particles may be classified into **autotrophs**, **mixotrophs** and **heterotrophs**

We have been using **Random Forest** classifiers

Evaluating our classifiers

We have been using **Random Forest** classifiers

Ten fold cross validation



Visual check – images automatically sorted according to the classifier into the folders of the training data

Dictyochales	CBR001.202	CBR001.202	CBR001.202	CBR001.202
	2-04-01.001	2-04-01.001	2-04-01.001	2-04-01.001
	.T1+F1_150	.T1+F1_156	.T1+F1_263	.T1+F1_264
CBR001.202	CBR001.202	CBR001.202	CBR001.202	CBR001.202
2-04-01.001	2-04-01.001	2-04-01.001	2-04-01.001	2-04-01.001
.T1+F1_406	.T1+F1_983	.T1+F1_108	.T1+F1_127	.T1+F1_358

10 items | 1 item selected



Improving the training set

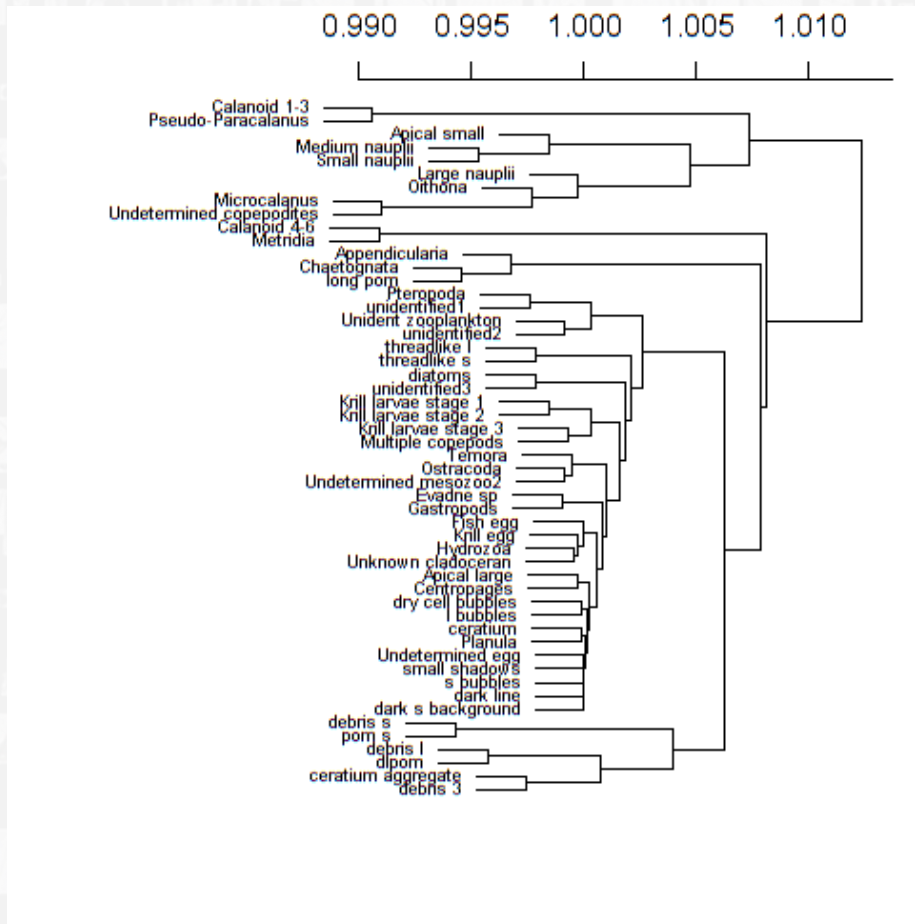
The initial and premature classifiers are used to automatically place unannotated images into the folder structure of the training set and visually checked

Problematic groups are improved by moving representative images to the correct folders, before retraining the classifier. We repeat this MANY times until we find the automatic sorting of images into the groups of the training set folder structure acceptable

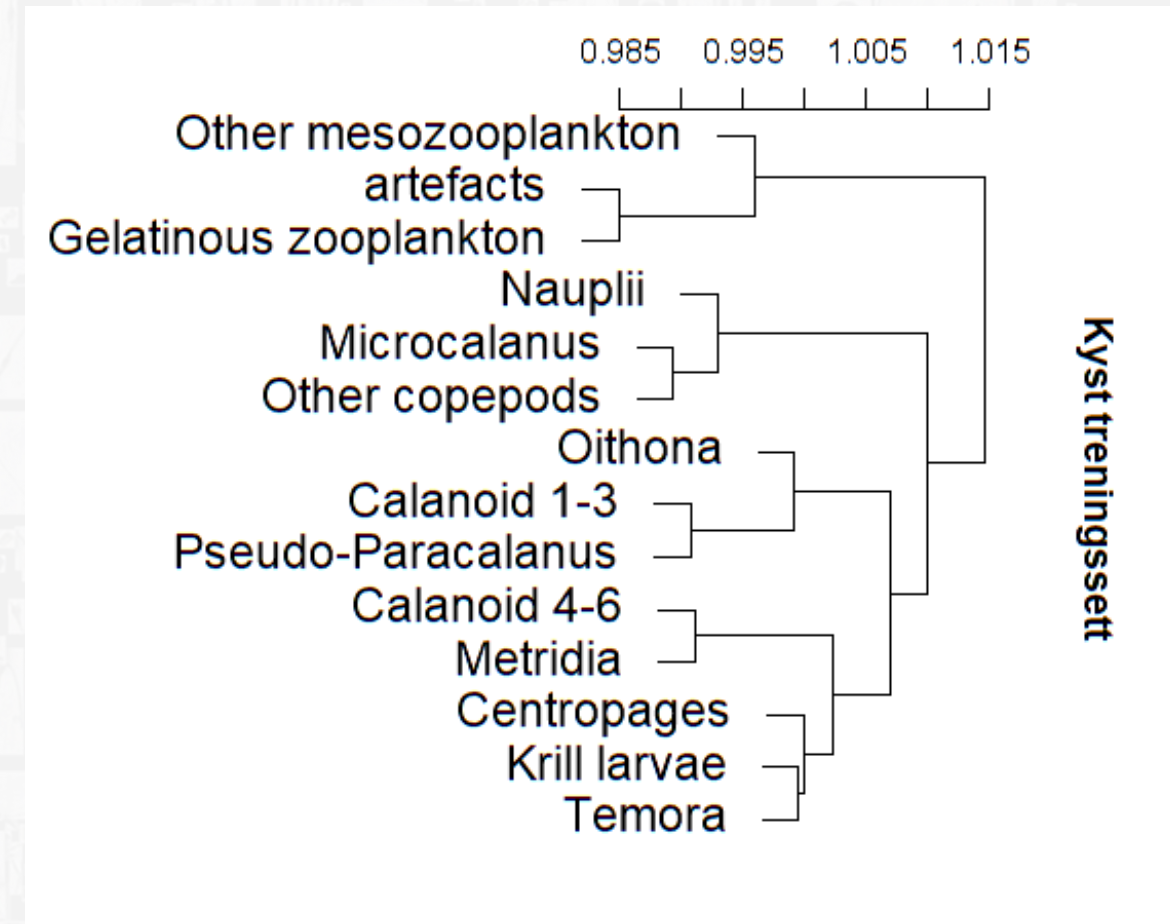
Not for HABs: The classifier will perform better at lower taxonomical detail. If you can't wait (like me) you can group the classes at a **lower level of taxonomical detail** or in a **functional** way.

Setting the level of taxonomic or functional detail before classifying samples

Classifier trained on all groups in the training set



Classifier trained on merged groups in the training set





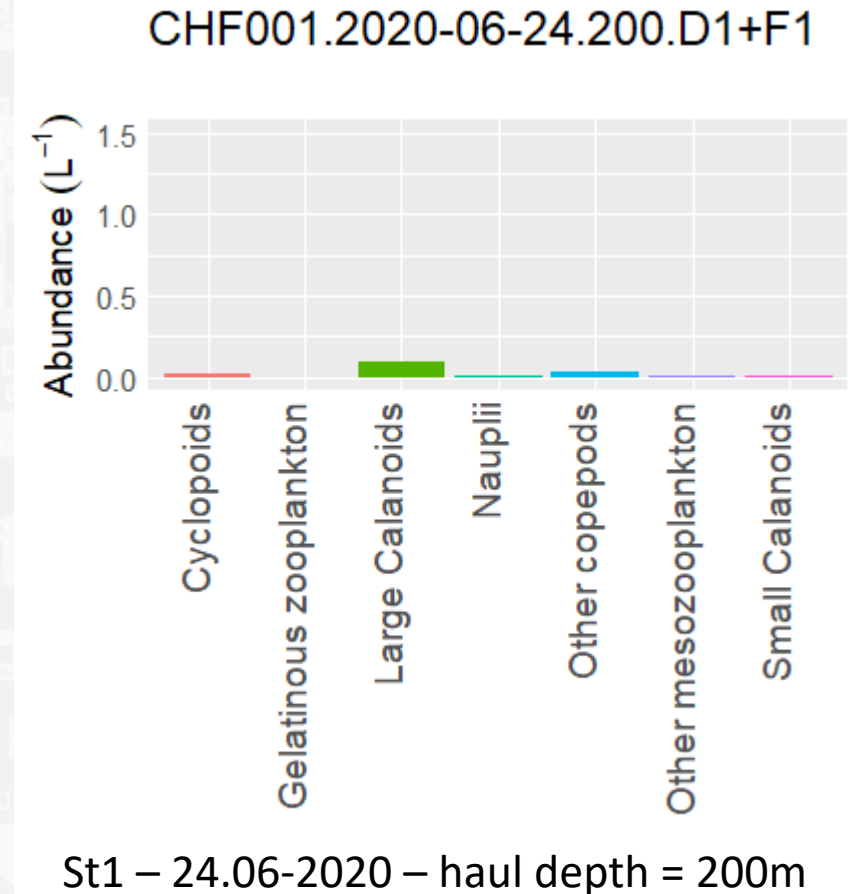
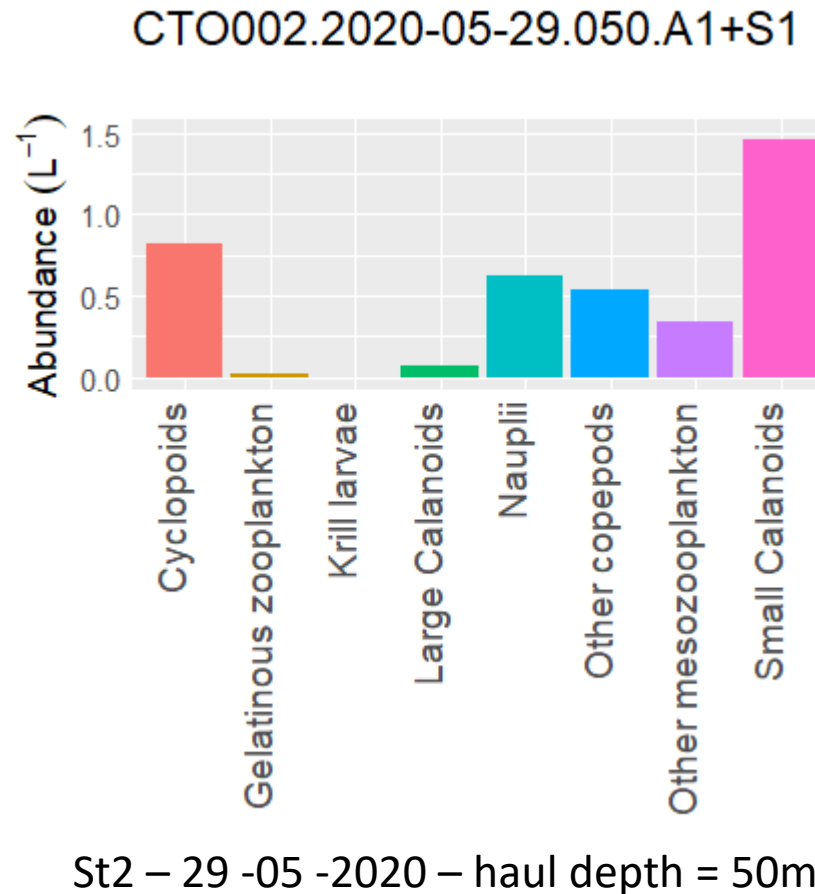
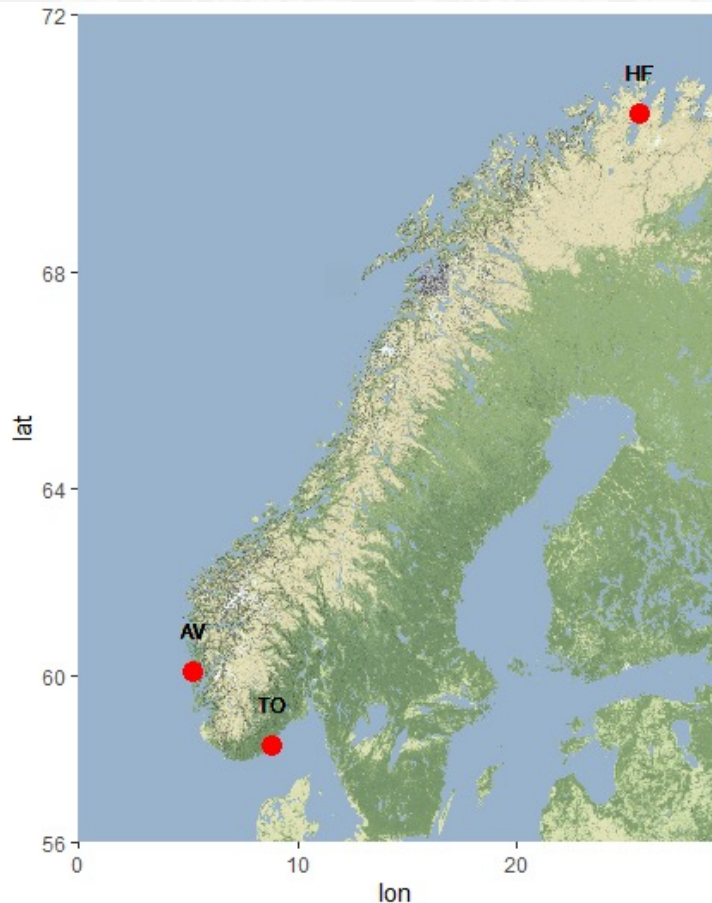
Data output from classification Taxa

Example of data output from single FlowCam runs

Mesozooplankton sample – WP2 net 180 μ m

Southern Norway – Torungen St2

Northern Norway - Holmfjord

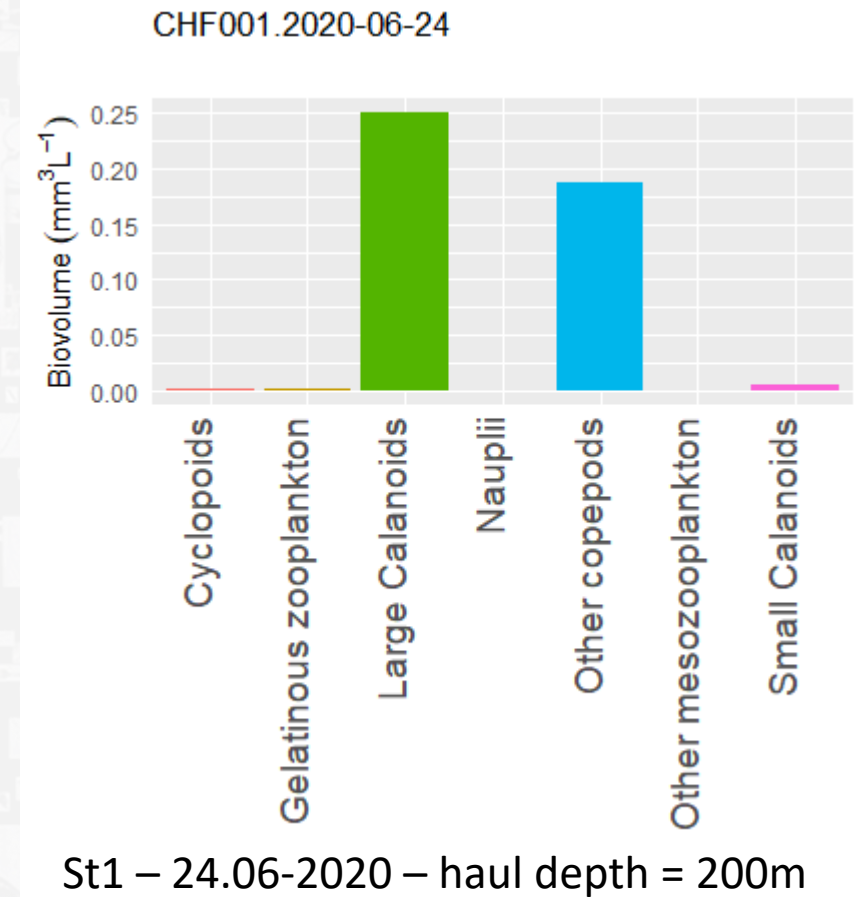
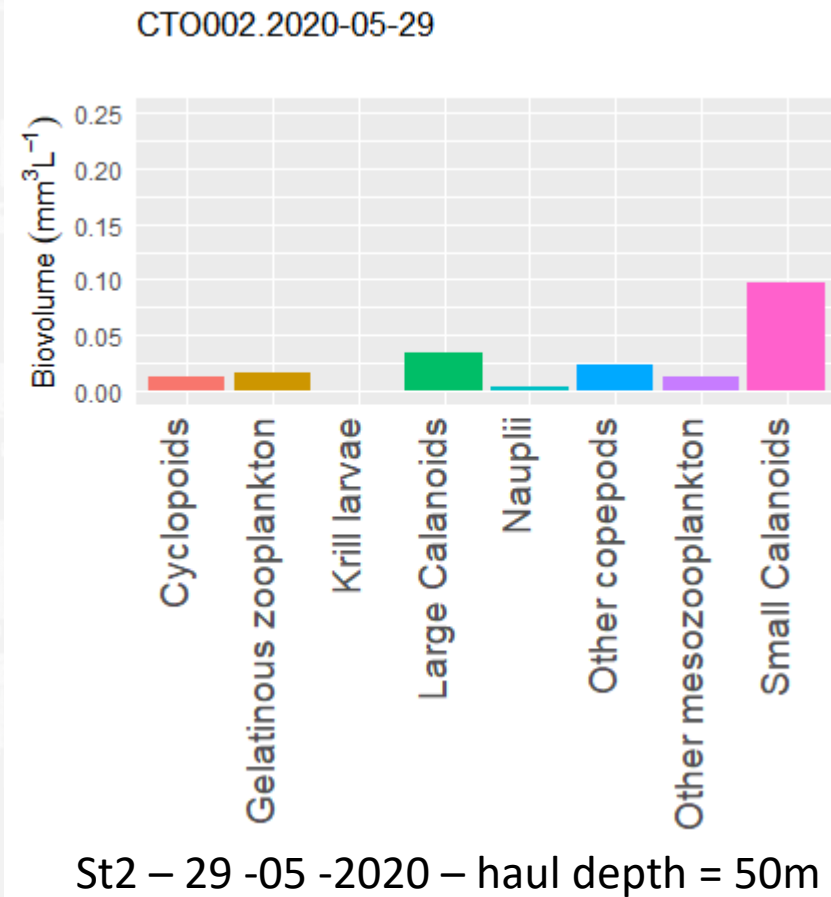
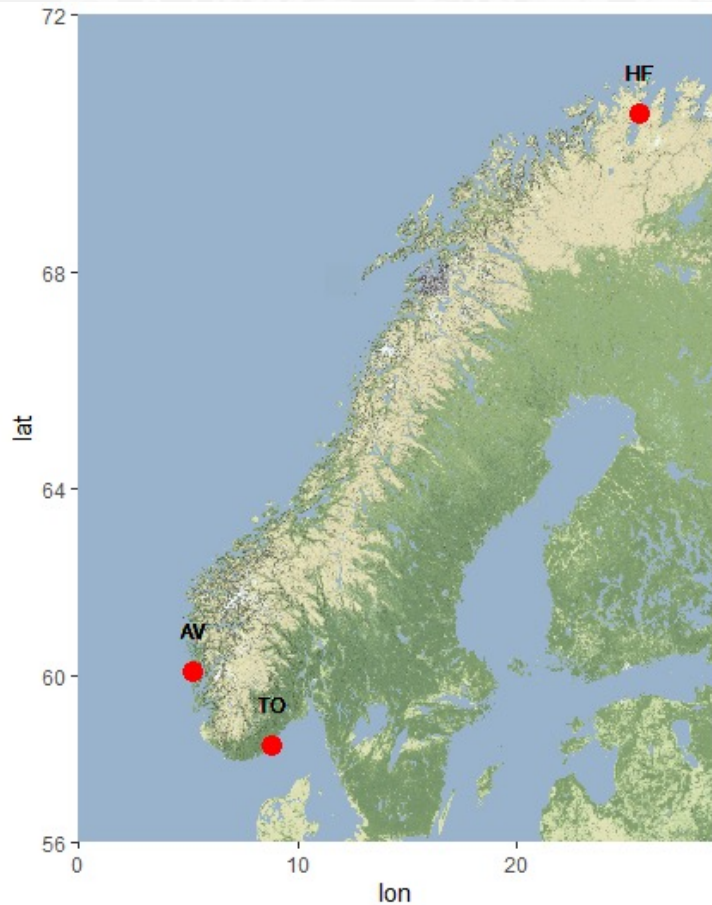


Example of data output from single FlowCam runs

Mesozooplankton sample – WP2 net 180 μ m

Southern Norway – Torungen St2

Northern Norway - Holmfjord

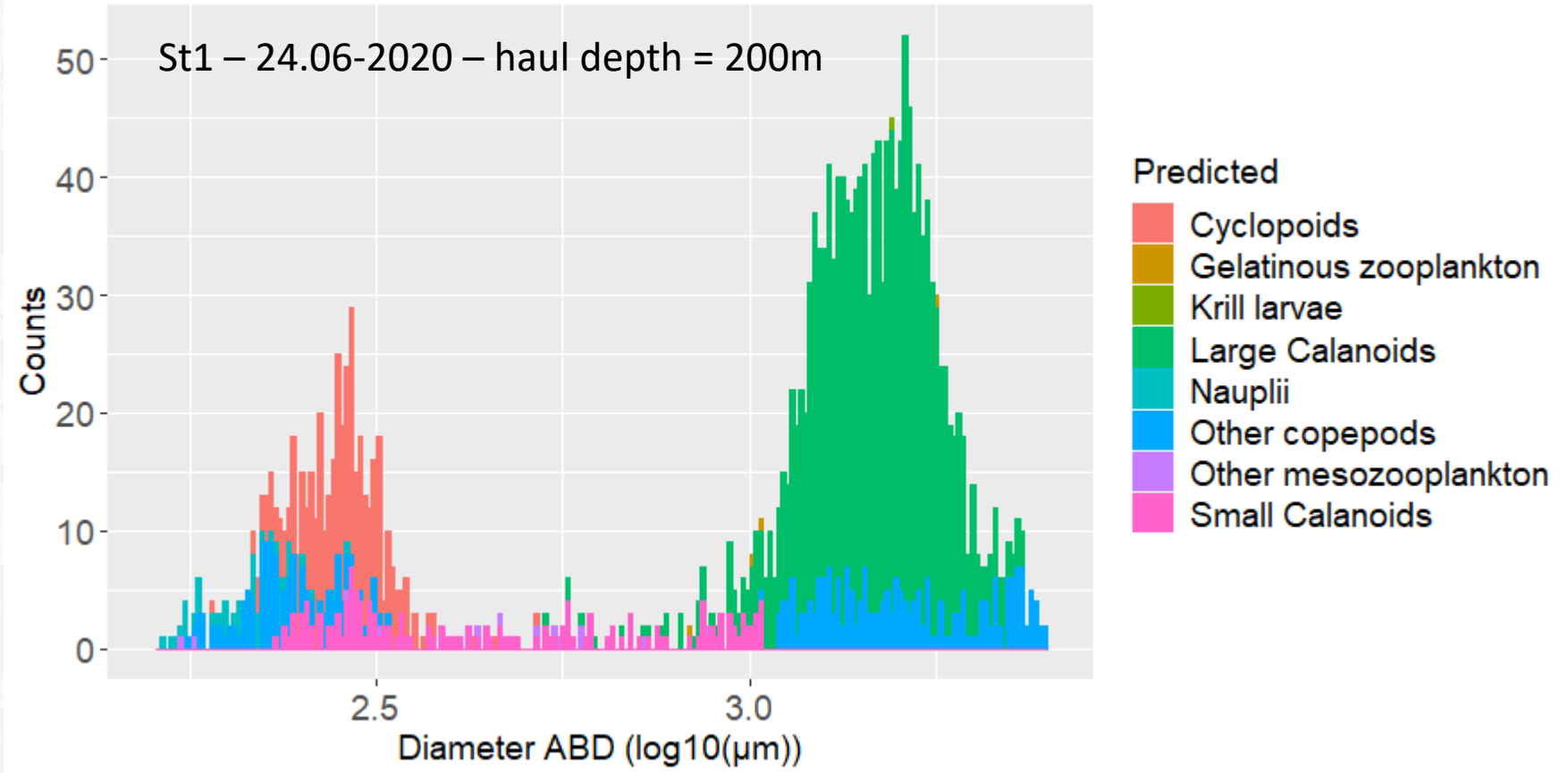
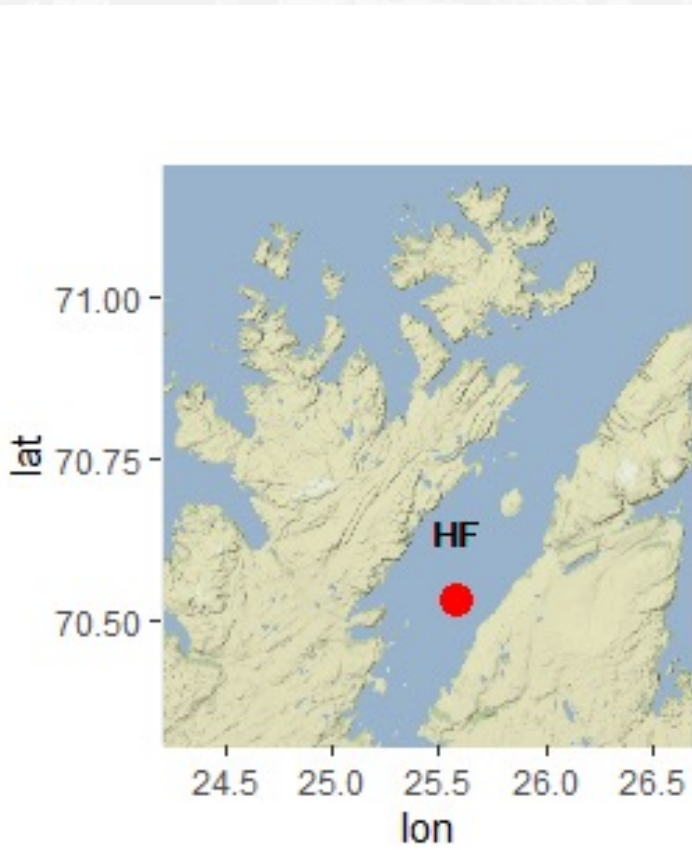


Data output from classification
Size structure

Example of data output from single FlowCam runs

Mesozooplankton sample – WP2 net 180 μ m

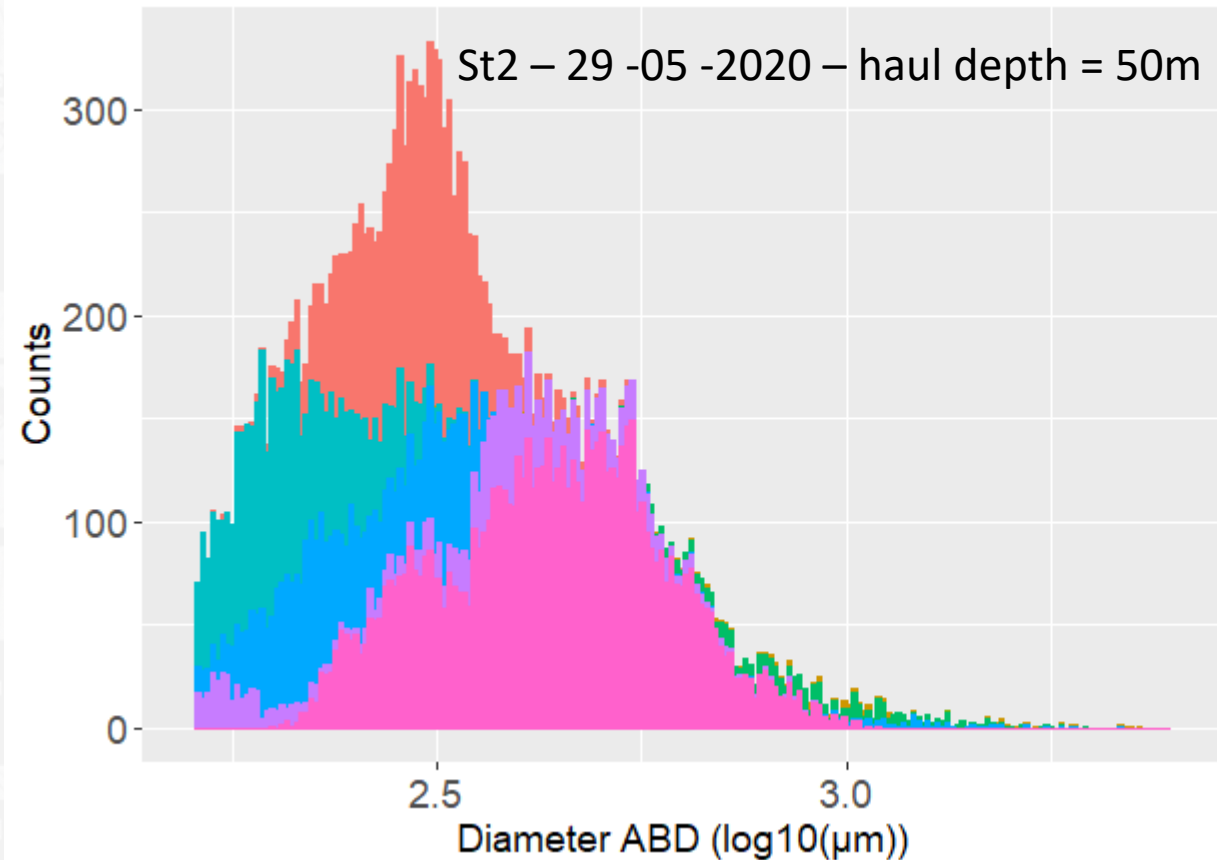
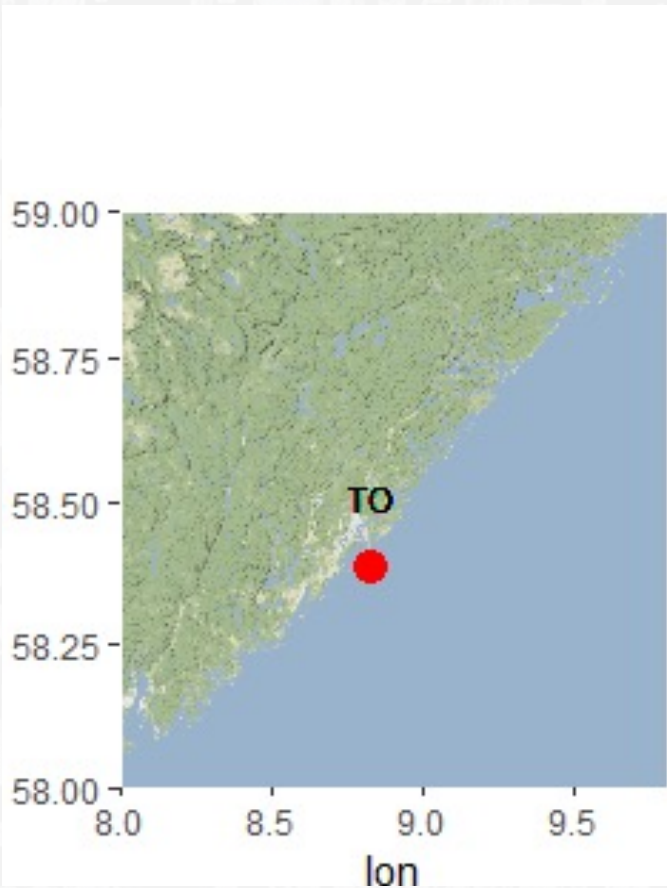
Mesozooplankton - Northern Norway - Holmfjord



Example of data output from single FlowCam runs

Mesozooplankton sample – WP2 net 180 μ m

Mesozooplankton - Southern Norway – Torungen St2

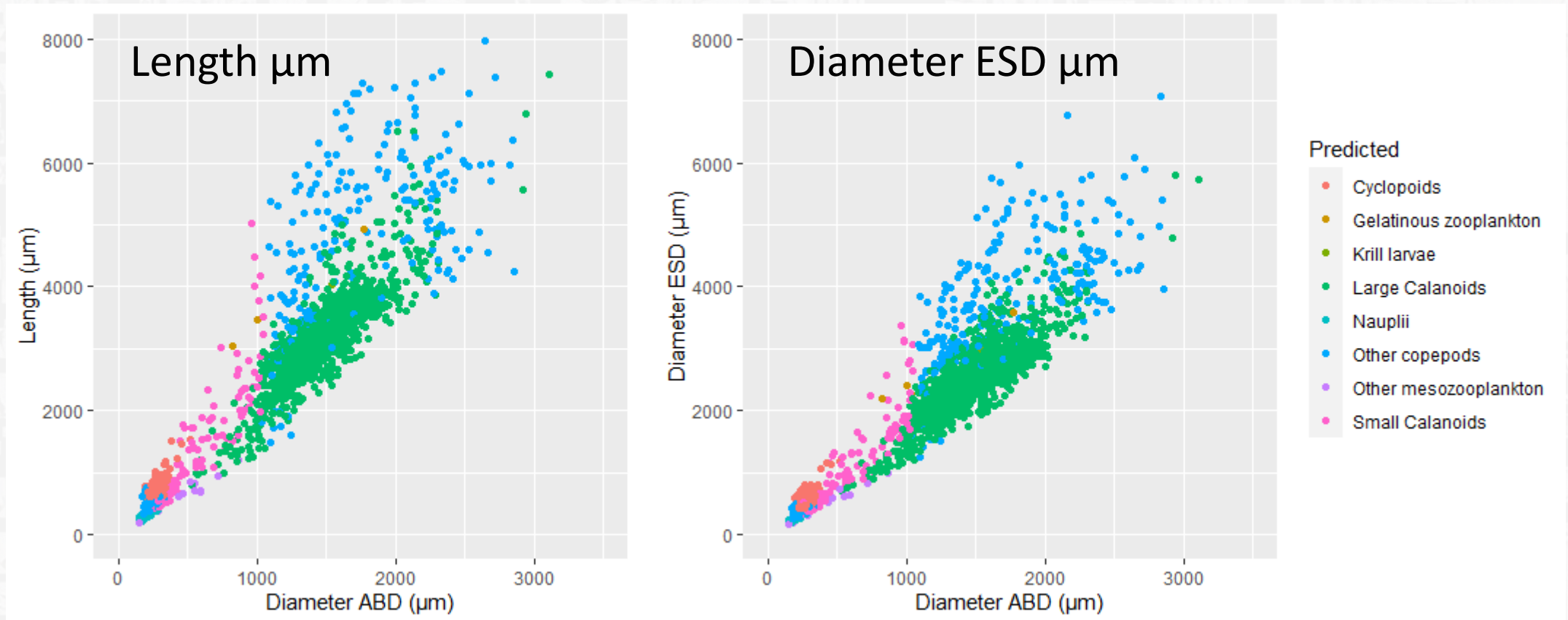


- Predicted
- Cyclopoids
 - Gelatinous zooplankton
 - Krill larvae
 - Large Calanoids
 - Nauplii
 - Other copepods
 - Other mesozooplankton
 - Small Calanoids

Example of data output from single FlowCam runs

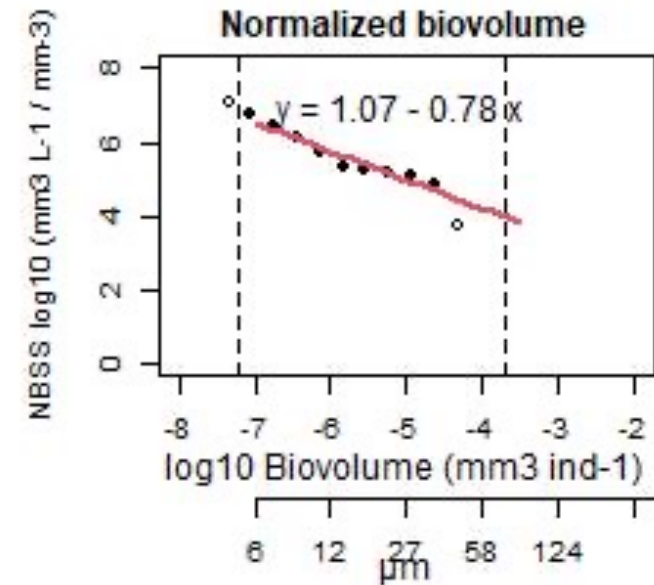
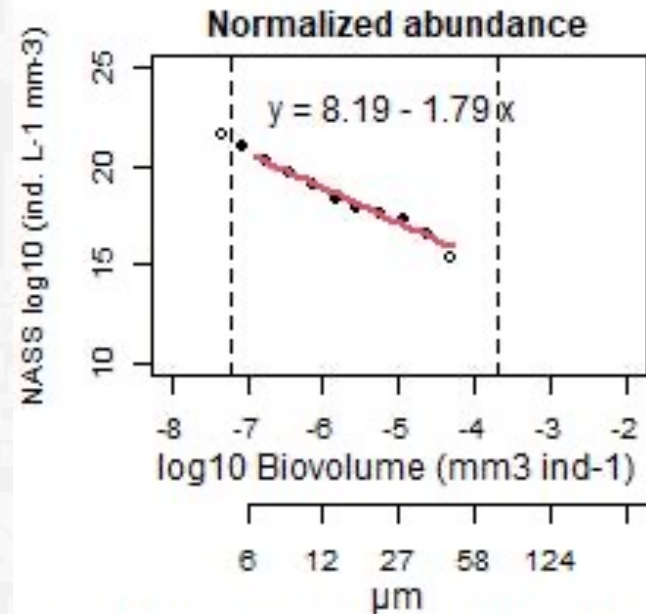
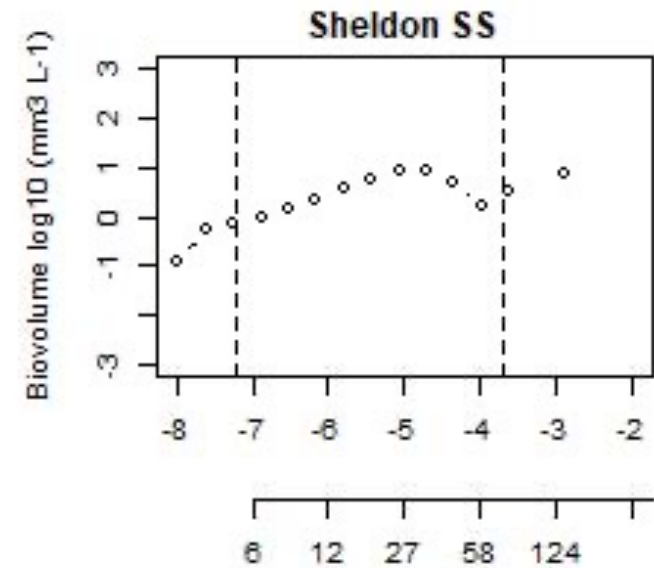
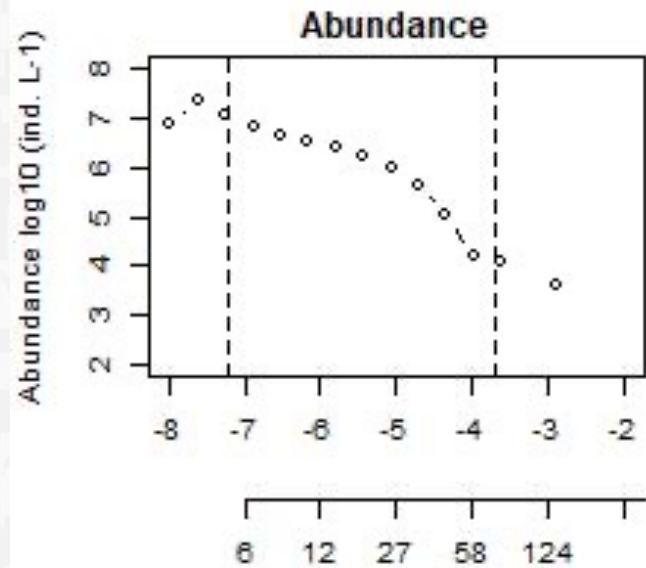
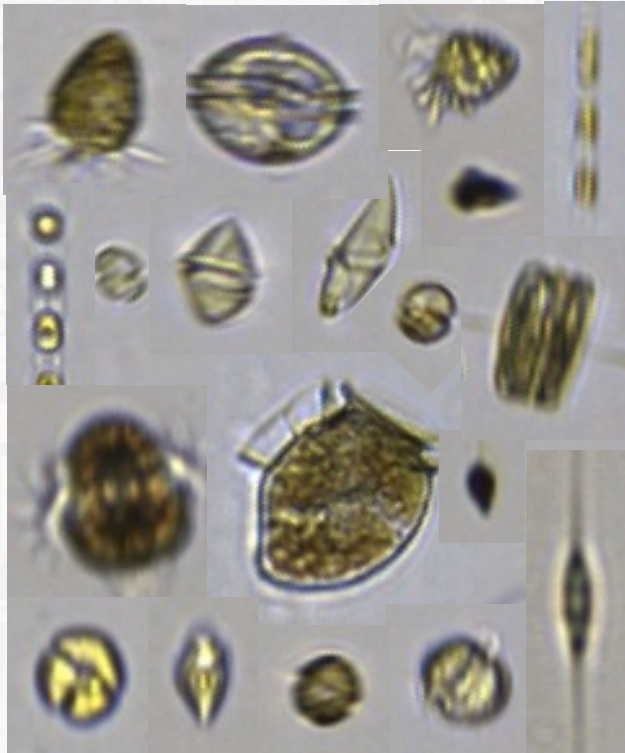
Mesozooplankton sample – WP2 net 180 μ m

Mesozooplankton - Northern Norway – Holmfjord



Area Based Diameter (ABD)

Size structure



Organism Size - Equivalent Circular Diameter

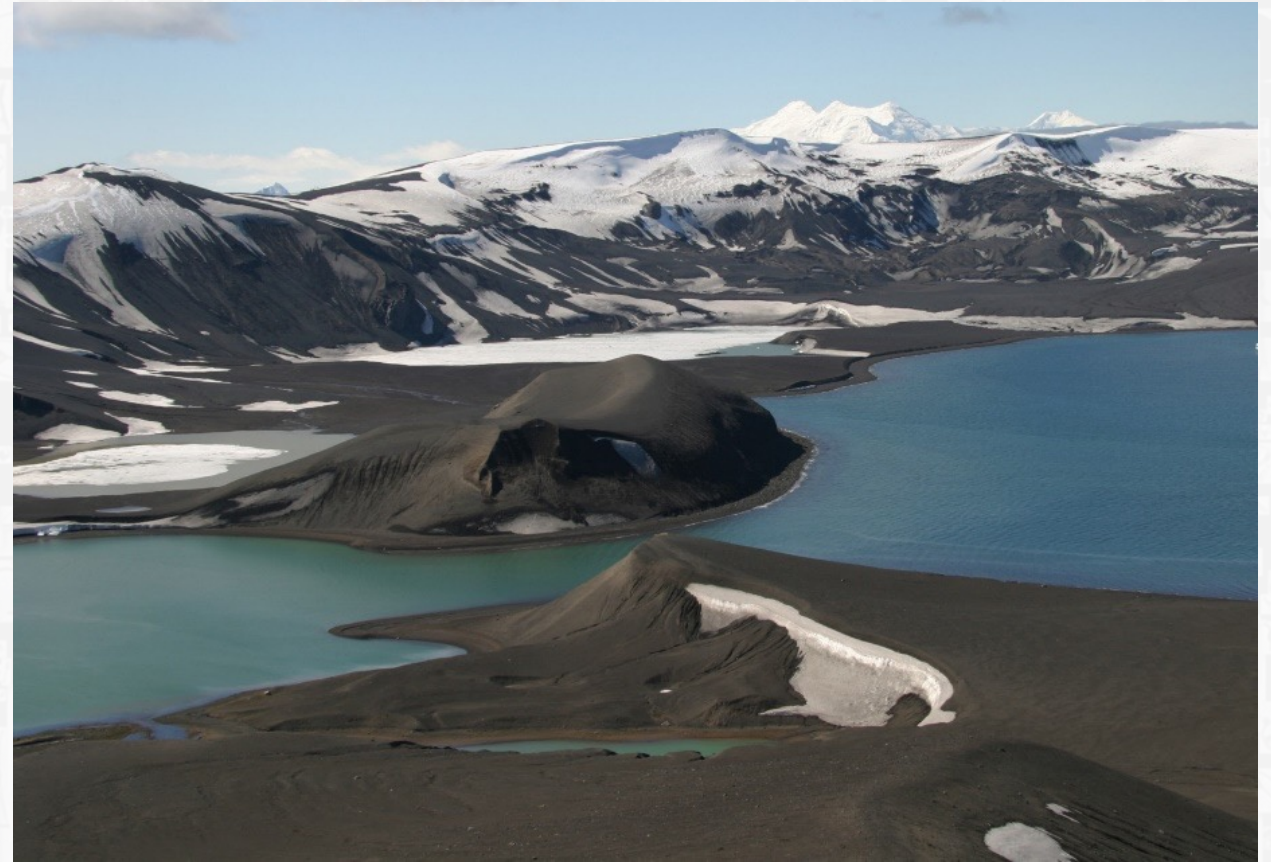
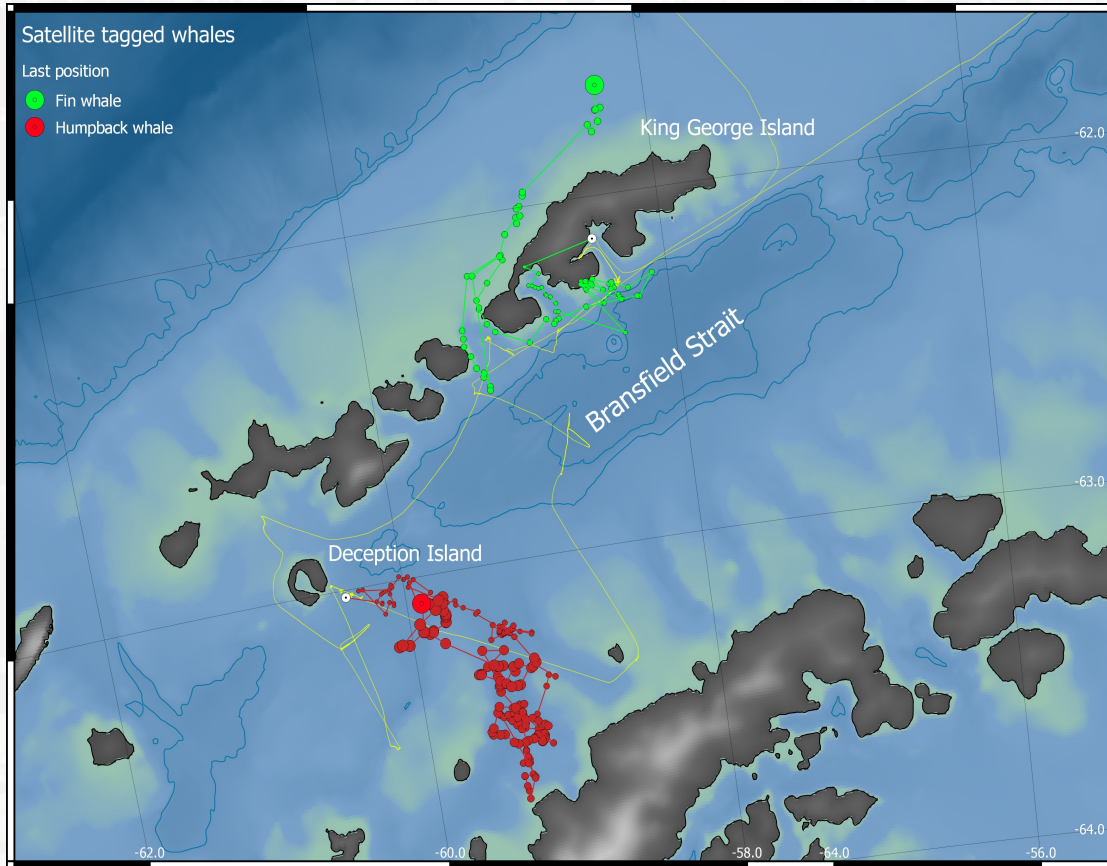
Size Equivalent Circular Diameter

Data valuable even without taxa...



Examples of usage

Example of early use of a premature training set and classifier

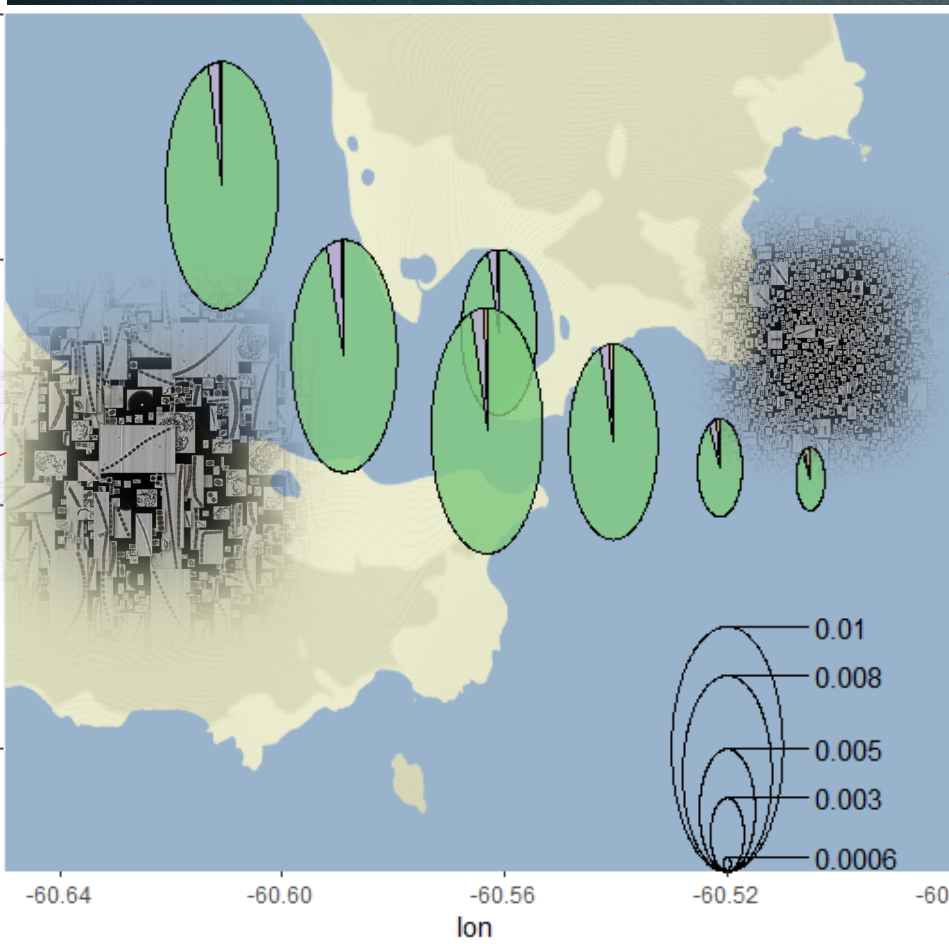
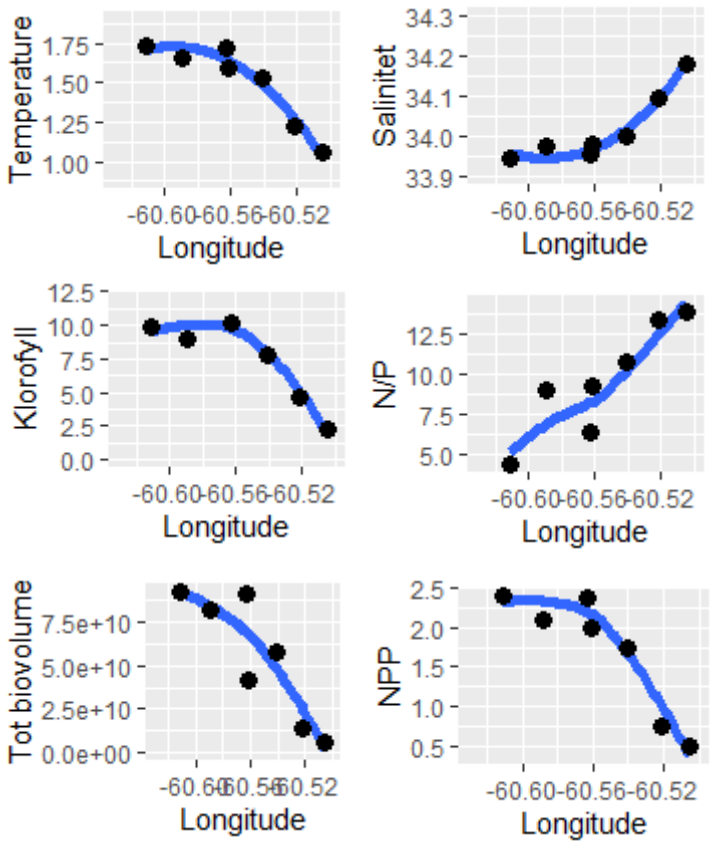
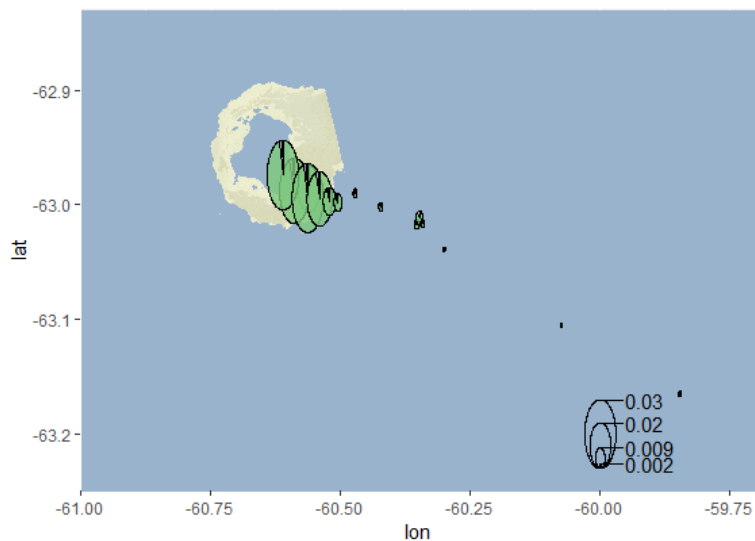


Example of use of a premature training set and classifier

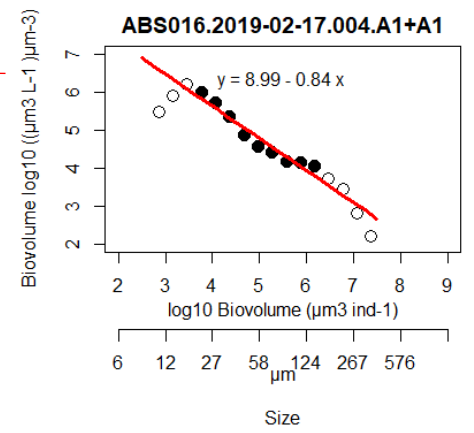


Deception Island – Antarctica

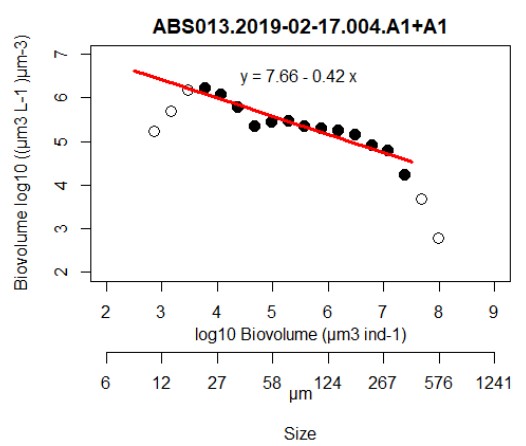
Deception Island



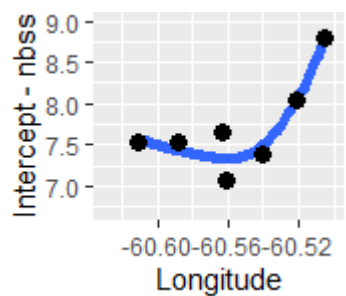
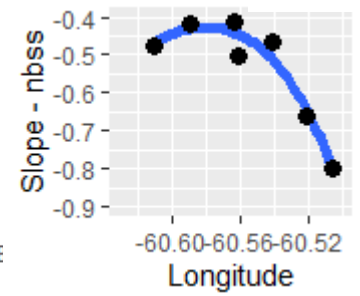
Biomassestørrelsesfordeling ute



Biomassestørrelsesfordeling inni



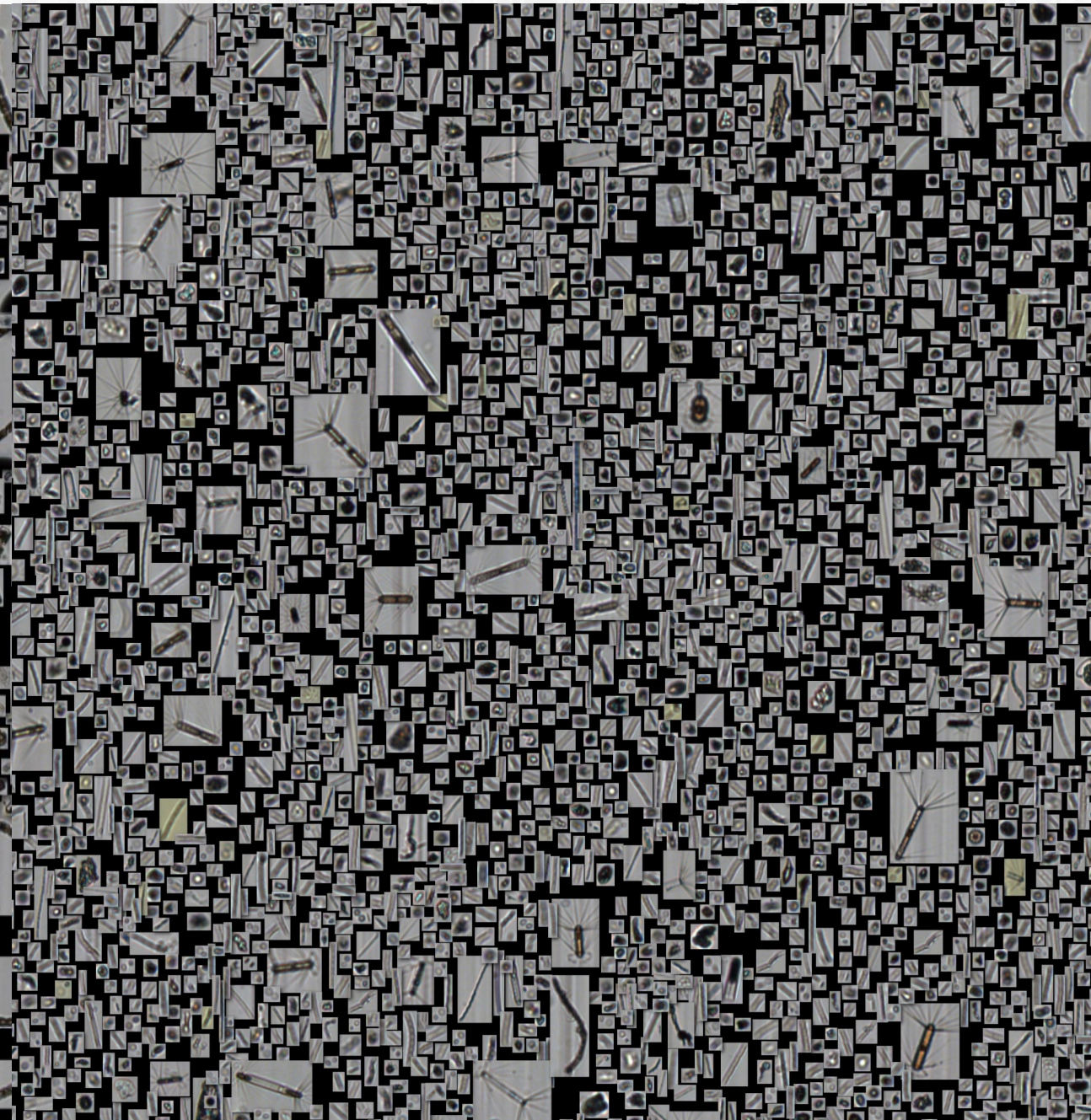
- Autotrophs
- Heterotrophs
- Mixotrophs



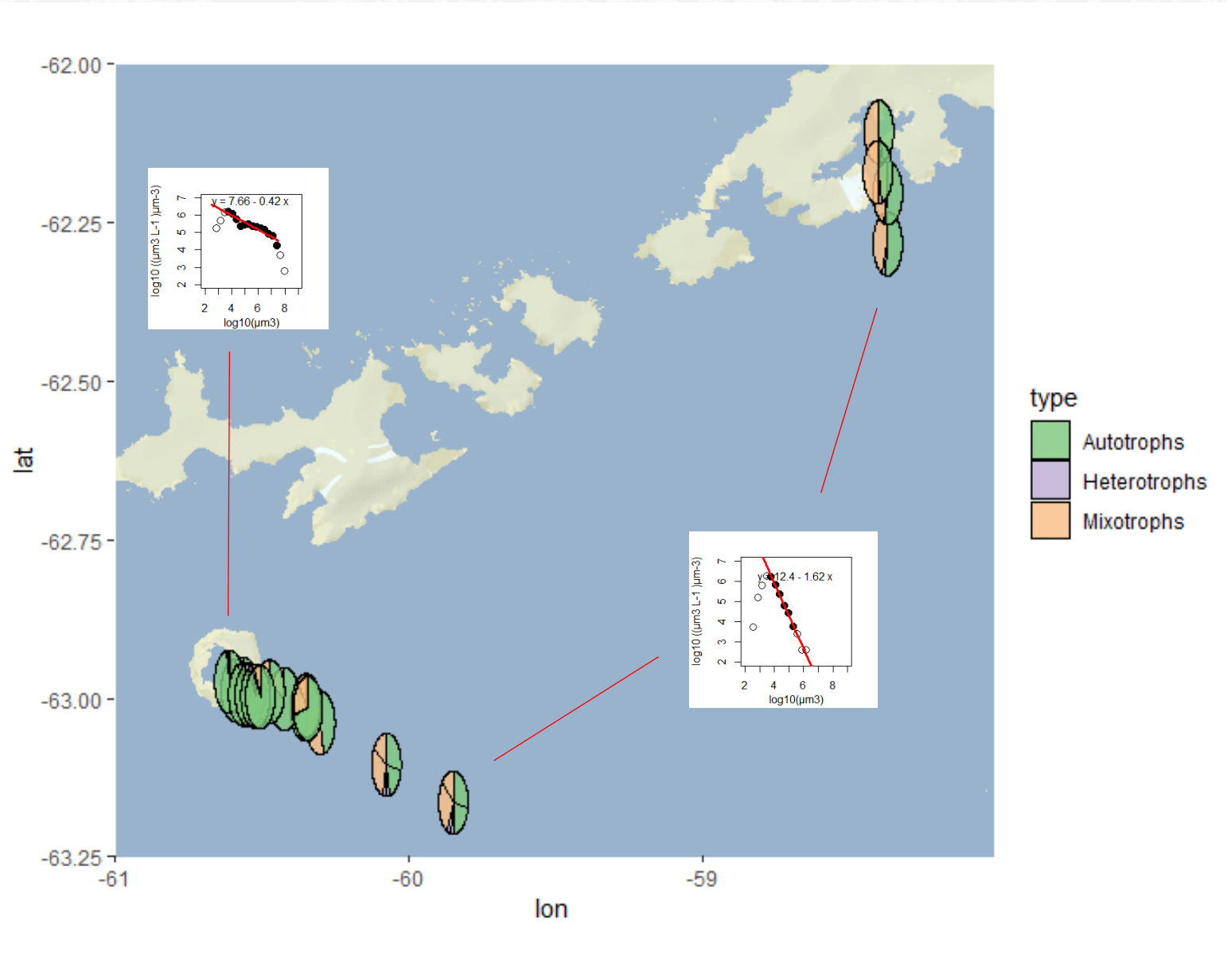
Within the Deception Island



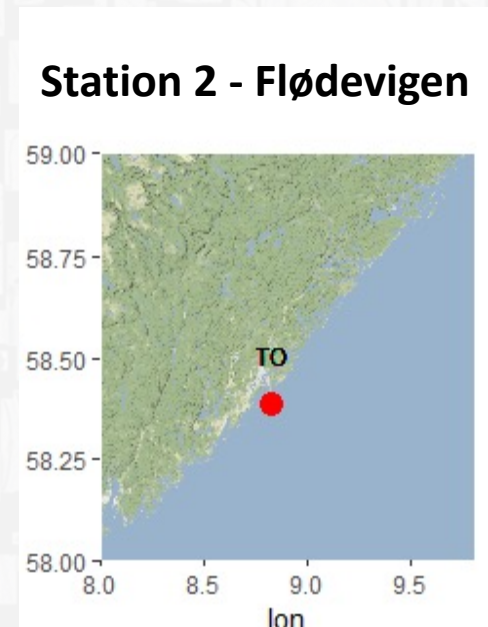
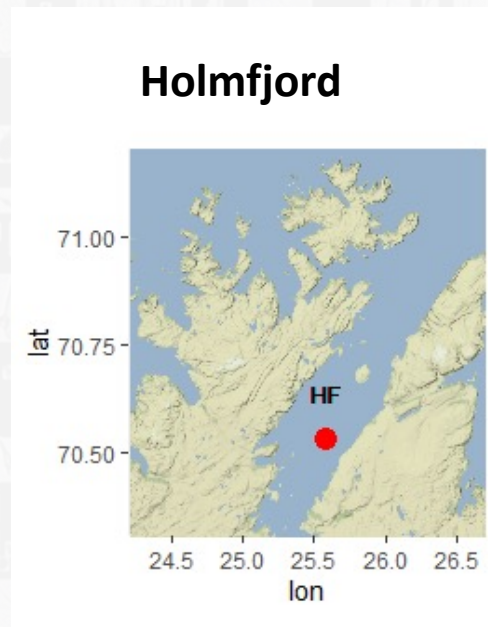
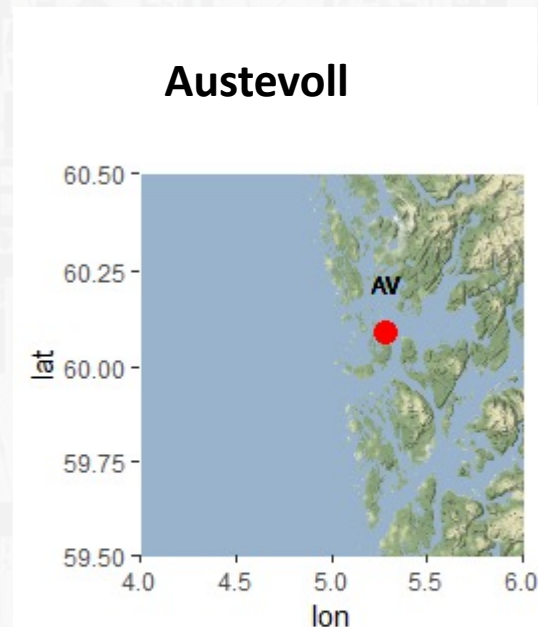
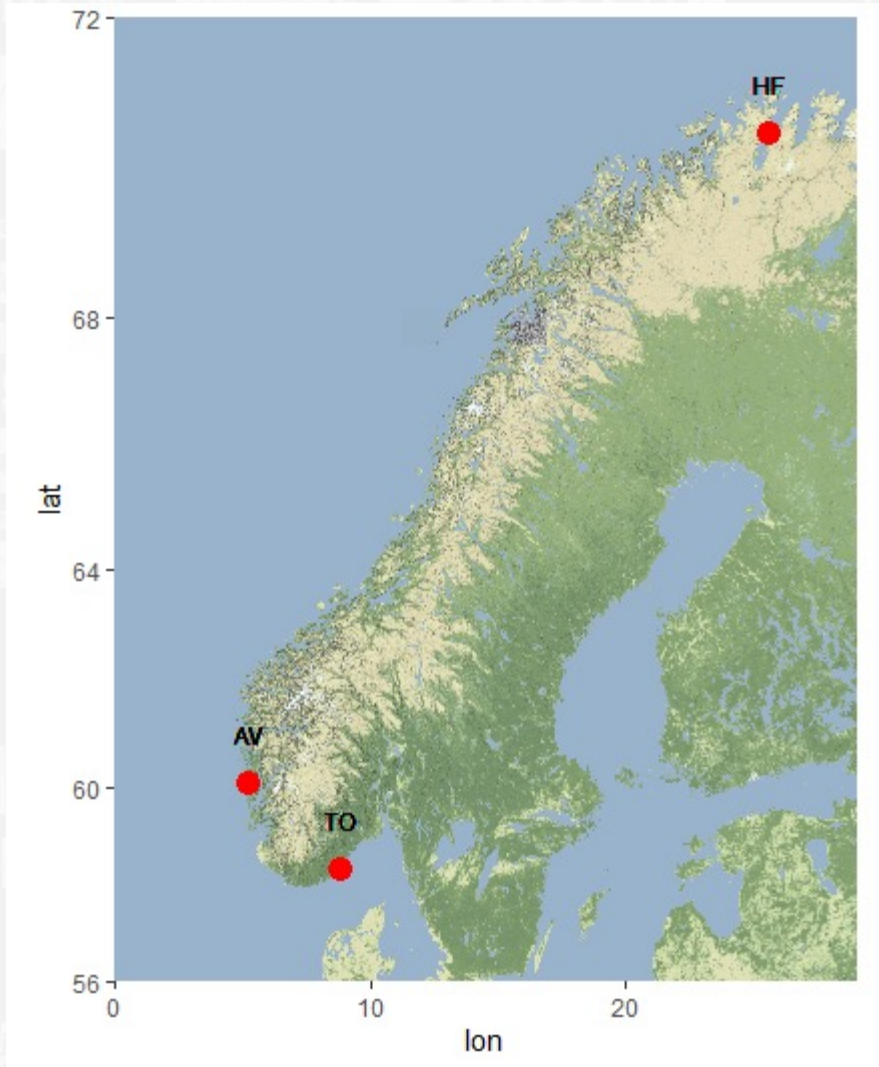
Just outside the opening of the entrance into the caldera



Classification to trophic group

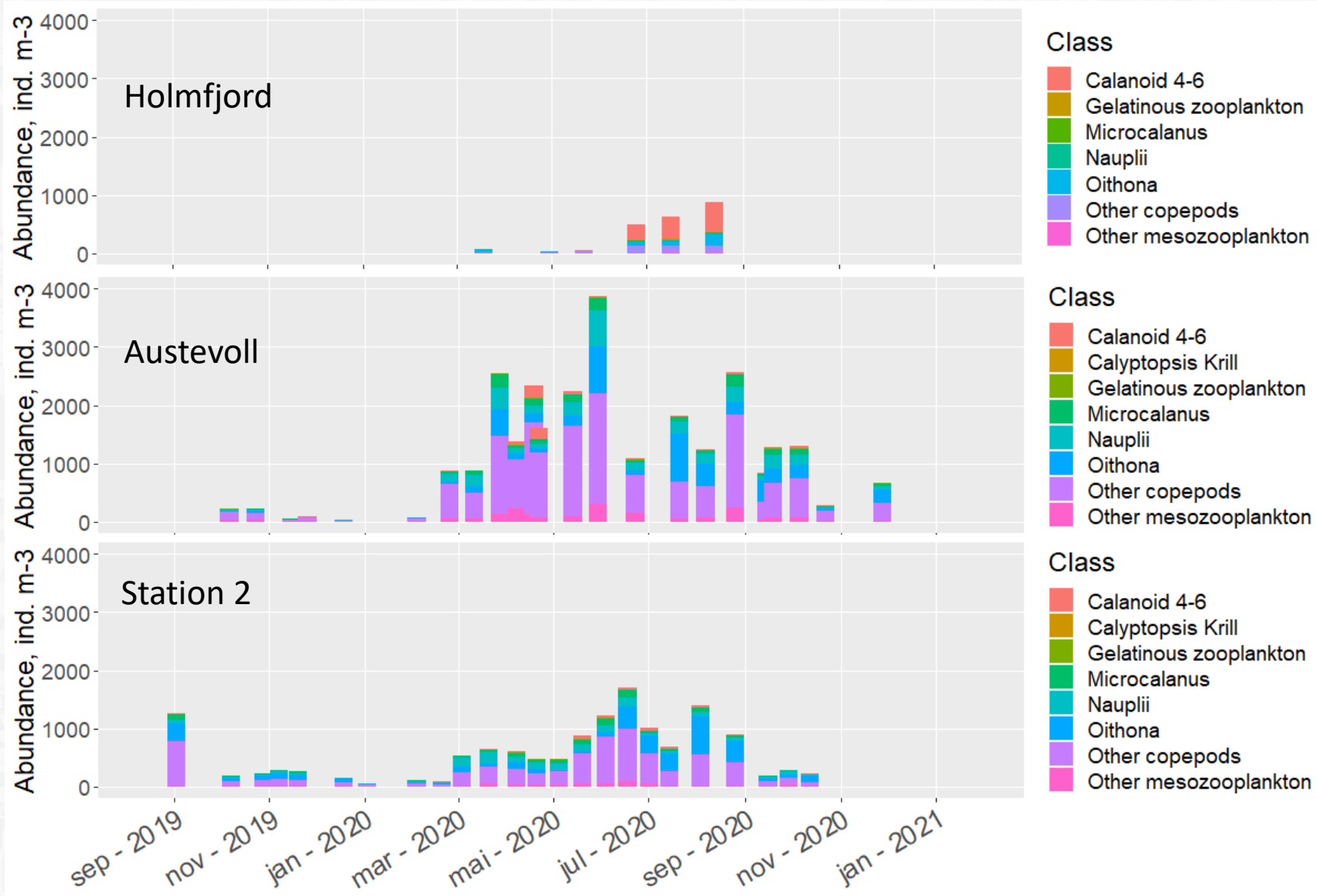
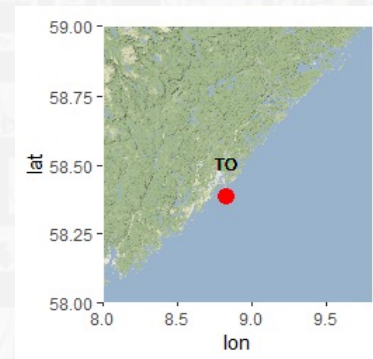
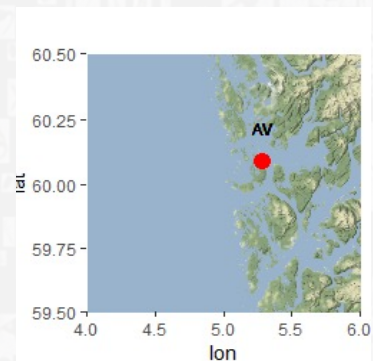
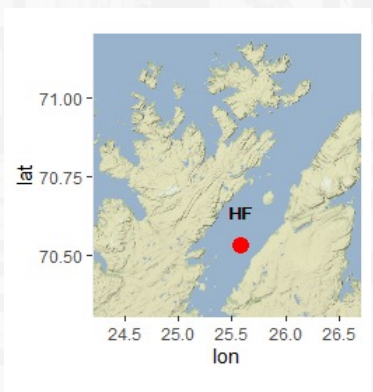


IMR coastal monitoring stations for chemistry, physics and plankton abundance, biovolume and size structure (by FC)

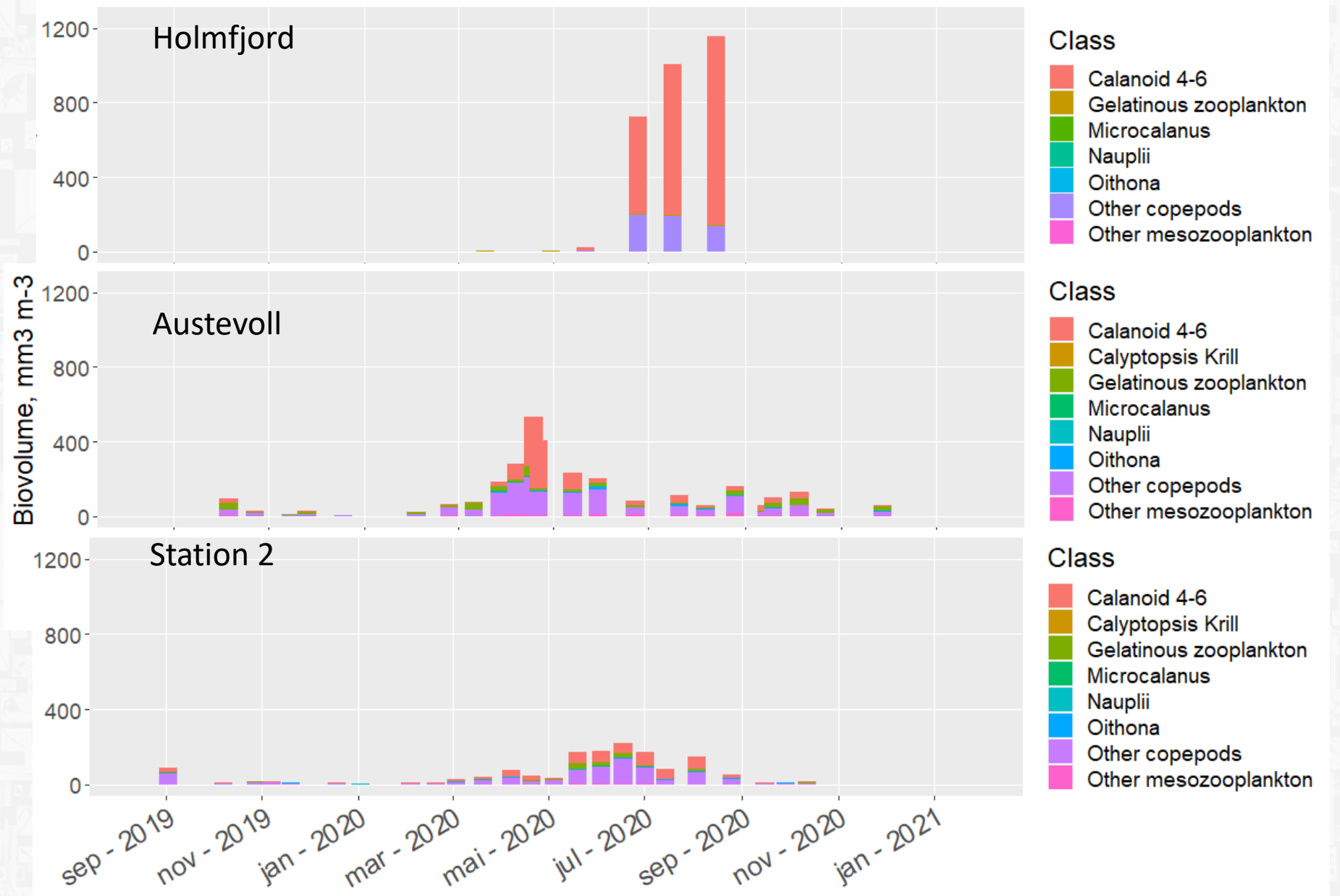
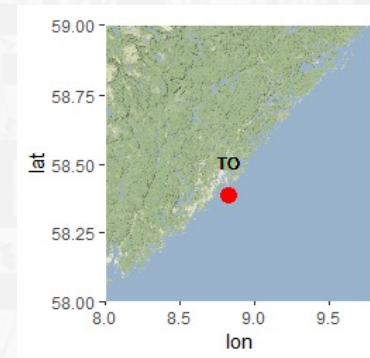
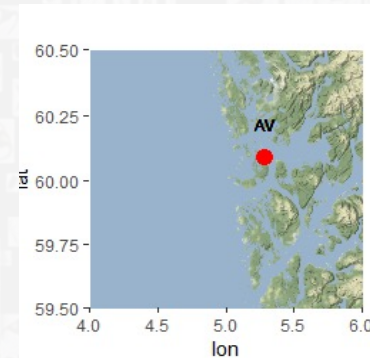
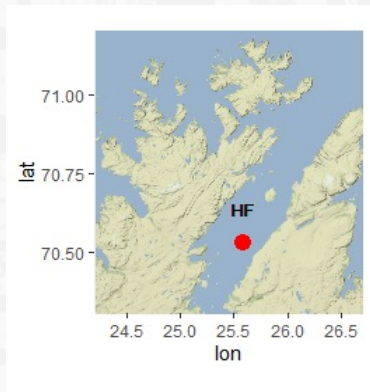


Seawater samples (500 ml) from 5 meter depth were obtained from Niskin bottles every 2-4 weeks at three coastal stations and fixed in acidic Lugol (2% final concentration). The samples were imaged using a FlowCam (20x magnification, 800 μ m flowcell) and the imaged volume ranged from \sim 100-130 ml), resulting in a detection limit of \sim (1/0.1), or 10 individuals L⁻¹.

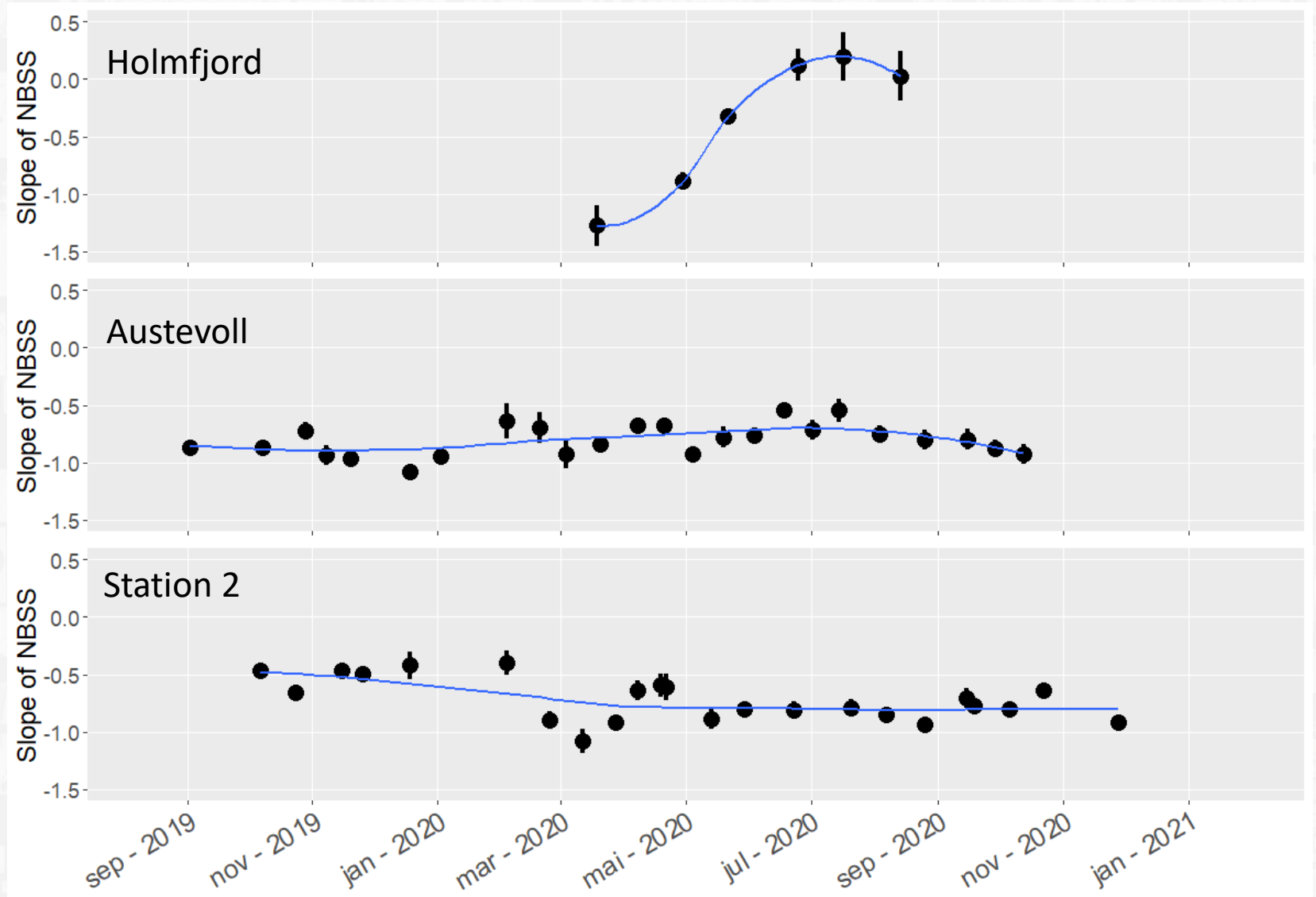
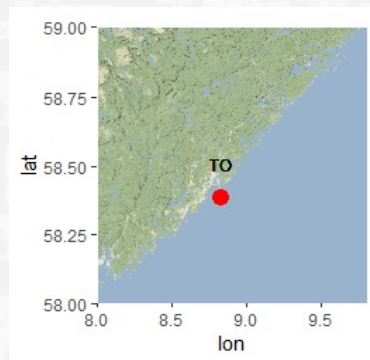
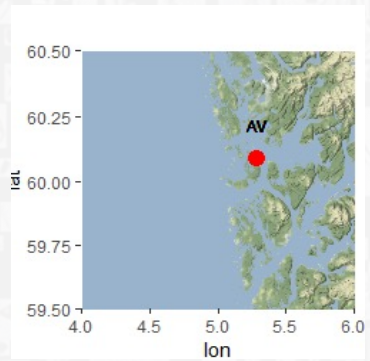
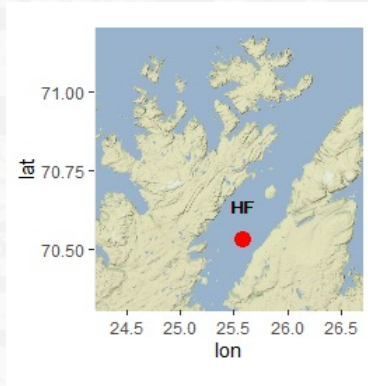
Mesozooplankton abundance 180 – 2000 μm



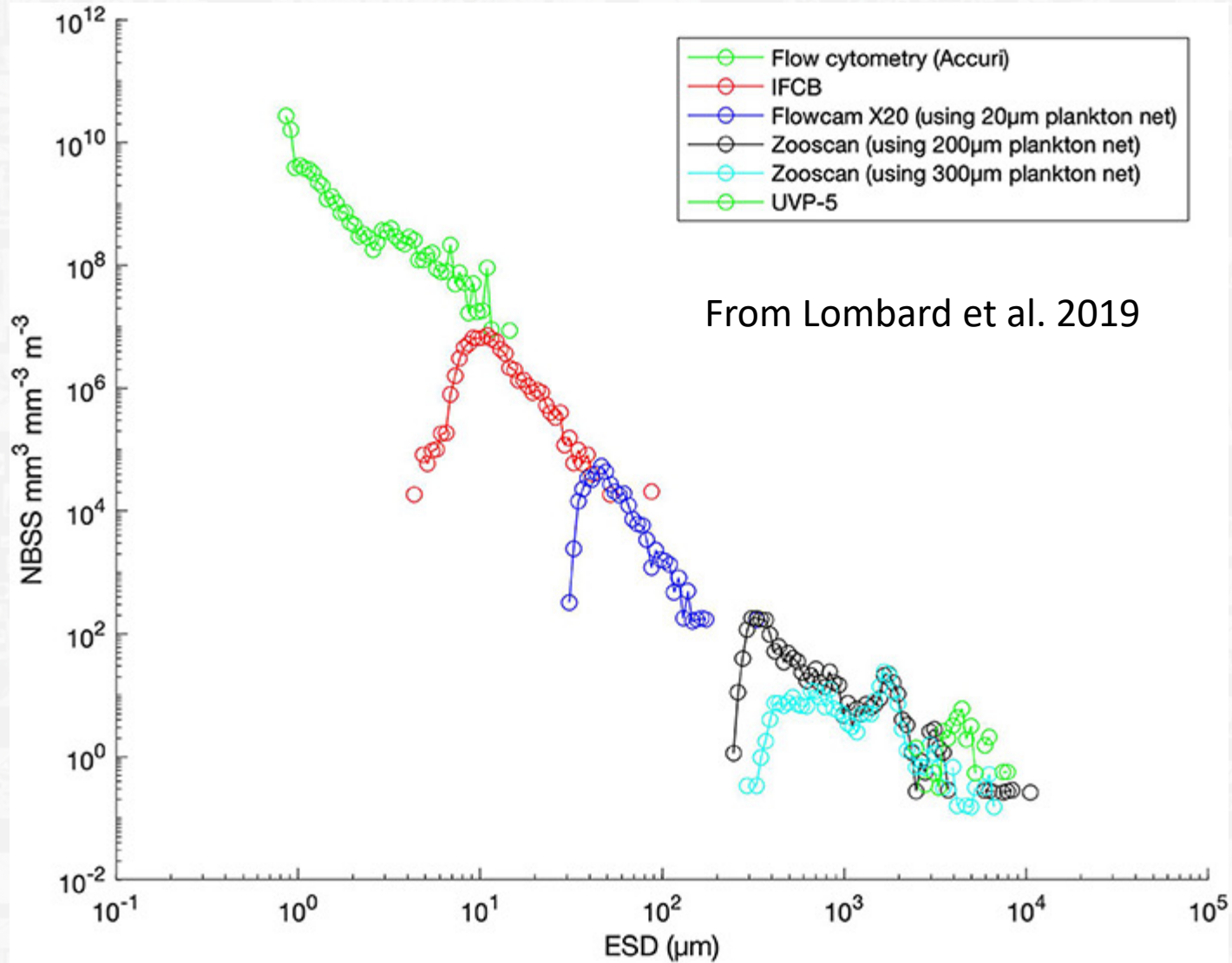
Mesozooplankton biovolume from 180 – 2000 μm



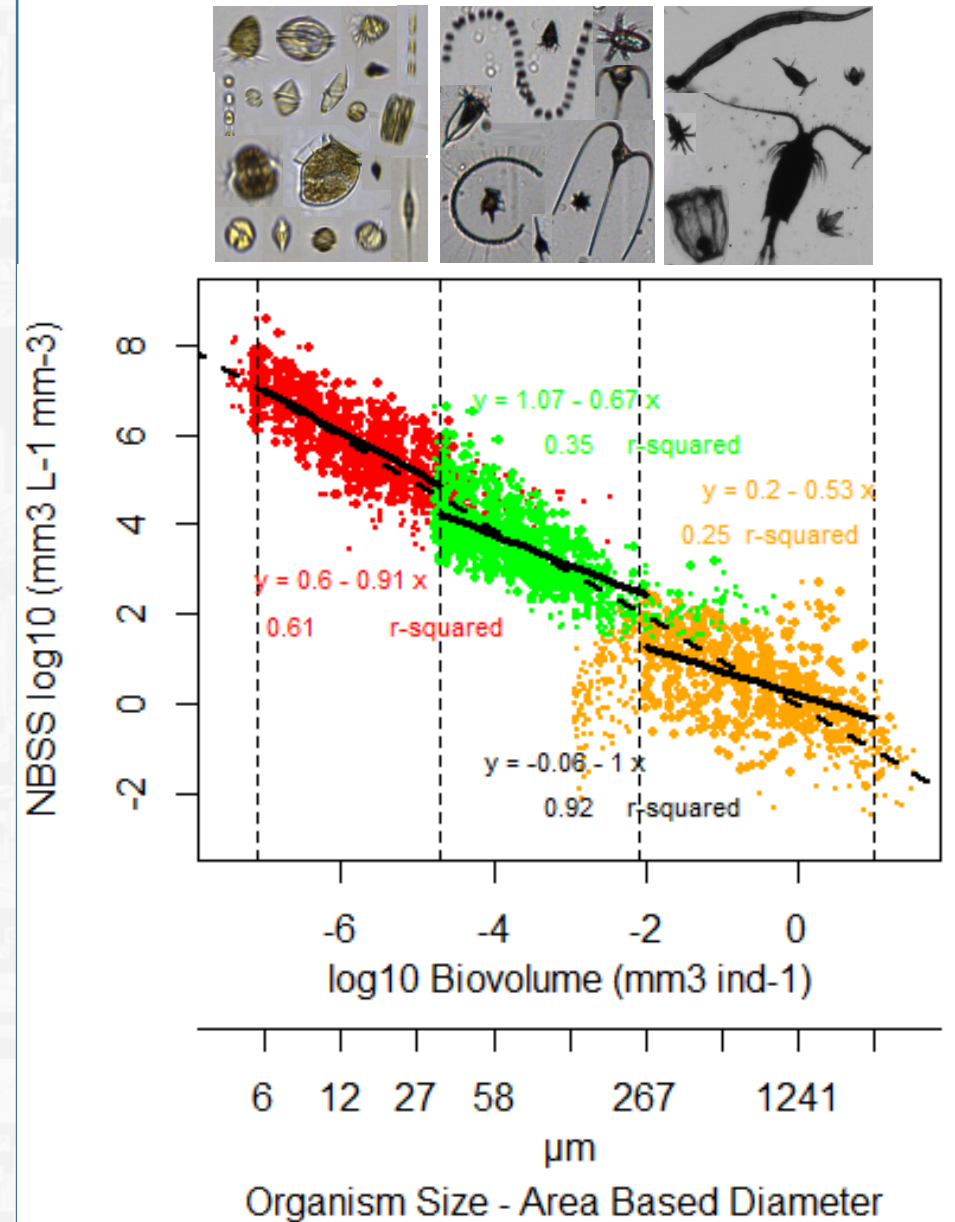
Mesozooplankton size structure 180 – 2000 μm - Slope



Inspiration to how we may combine results obtained from multiple flowcams



Size structure may guide us in merging the results from different magnification settings and instruments in terms of abundances and biovolumes of taxa across a large range of organism sizes





Thanks to:

Eva Álvarez, National Institute of Oceanography and applied geophysics, Trieste, Italy

Gayantonia Franzé, IMR

Magnus Reeve, IMR

Mona Ring Kleiven, IMR

Hege Mathisen, IMR

for help with annotating images,

Kjell Gundersen, Tone Falkenhaug and Lars Johan

Naustvoll (all IMR) for great patience and Ketil Malde

for recent initiative to take the data to the next level