

Biodiversity Monitoring Stations in the Davis-Baffin Division of the Eastern Arctic Canadian Contributions to the Circumpolar Marine Biodiversity Monitoring Plan



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BACKGROUND

Arctic waters are expected to show the first effects of climate change, including the effects of increasing sea water temperatures and ocean acidification. Baffin Bay and Davis Strait are key areas connecting the Arctic and the North Atlantic Oceans. The long-term time series from the NAFO (Northwest Atlantic Fisheries Organization) Standard Stations off West Greenland show that the observed warming of the North Atlantic Subpolar Gyre has had a tremendous impact on the West Greenland Shelf and off-slope waters (Stein 2010). The distribution of marine fauna in this area could be affected by climate change, including the expansion of ranges northward, and introduction of species (Vongraven 2009). Consequently it is critical to establish baseline data on benthic communities in this area in order to quantify future impacts of climate change. Fisheries and Oceans Canada has established 5 biodiversity monitoring sites in a fishery closure area following the Circumpolar Marine Biodiversity Monitoring Plan.

MEGAFUNA



Benthic megafauna were sampled using a drop camera device (pictured above, right). Photos of the sea floor (pictured above left) were taken approximately every 10 m following transect lines running up-slope. Megafaunal abundance was recorded for all fauna greater than 1 cm.

Large foraminifera (such as *Rhizammina algaeformis*) visible on the sediment surface were qualitatively assessed from each photo based on whether they were absent/rare, sparse, moderate, or abundant.

MACROFAUNA



Macrofauna were sampled from sediment grab samples collected using a mega-box corer and Van Veen grab system deployed at the start or end of each camera station. Sediment samples were sieved into 0.95 cm, 1 mm, and 0.5 mm size fractions. Macrofauna specimens were identified to the lowest taxonomic classification possible.

All pan-Arctic species (i.e., ophiuroids, bivalves, snow crabs) were measured for size-frequency analysis.

MEIOFAUNA & MICROBES

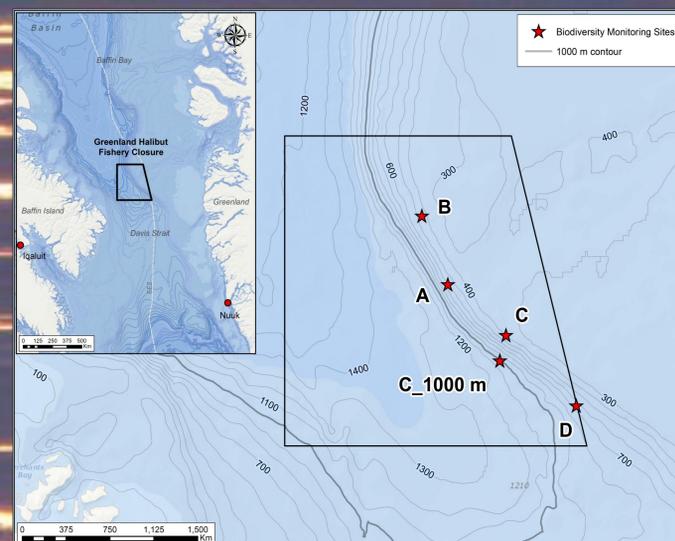
Meiofauna were collected from sediment subsamples collected from three Van Veen grabs per station. Seven sediment subsamples were collected from the top 1 cm of sediment using a modified 60 cc syringe. Samples were preserved in 3 ways: 1) formalin, to assess general meiofauna species composition and abundance, 2) ethanol, to assess foraminifera species composition and abundance, and 3) frozen, for DNA analysis of all meiofauna.

Sediment subsamples for microbes were also collected from the top 1 cm of each of 3 Van Veen grabs per station using a modified 10 cc syringe. Samples were frozen at -80°C.

Priority Focal Ecosystem Components, Parameters, and Indicators Sampled at the Canadian Eastern Arctic Biodiversity Monitoring Sites - Following Gill et al. (2011) and Pomerleau et al. (2014)

| Category | Focal Ecosystem Components | Key Parameters | Indicators |
|--------------------------|----------------------------|--|--|
| Benthic fauna & microbes | Macrofauna & Megafauna | Abundance, Biomass (wet weight), Species composition, Barcoding and other genomics | Community composition, Size-frequency distribution (for selected, mainly pan-Arctic species), Diversity indices (e.g., Shannon, Simpson), Distribution |
| Benthic fauna & microbes | Meiofauna & Microbes | Abundance, Biomass, Species composition, Barcoding and other genomics | Community composition and structure, Diversity indices (e.g., Shannon, Simpson), Distribution |

Location of 5 marine biodiversity monitoring sites within the Greenland Halibut fishery closure (DFO 2007).



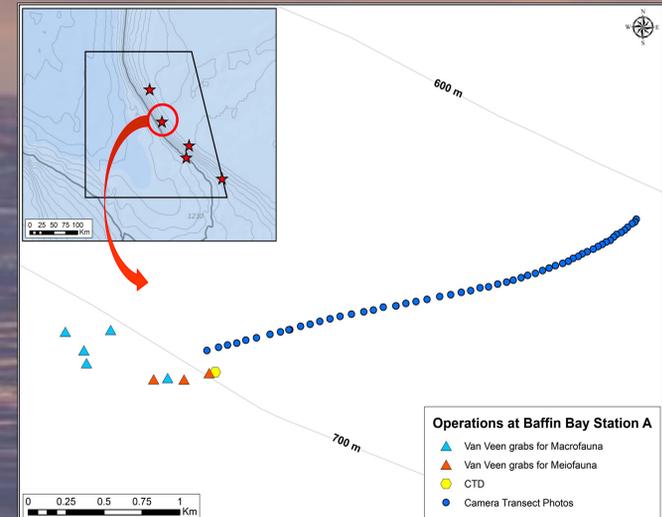
Summary of the camera operations conducted at 5 marine biodiversity monitoring sites in the Greenland Halibut fishery closure area (black box) for the collection of information on megafauna. CTDs and Van Veen Grab samples were conducted at each station for the collection of data on megafauna, meiofauna/microbes, and water column characteristics.

| Station | Camera position (dd) | | Depth (m) | | Number of photos |
|----------|----------------------|----------------------|-----------|-----|------------------|
| | Start | End | Start | End | |
| A | 67.7769/ -59.0593 | 67.7694/ -59.0850 | 593 | 705 | 58 |
| B | 67.9986/ -59.2658 | 67.9944/ -59.3086 | 404 | 457 | 74 |
| C | 67.6126/ -58.5469 | 67.6002/ -58.5708 | 229 | 482 | 102 |
| C_1000 m | 67.5297/ -58.6014 | 67.5436/ -58.5904 | 974 | 879 | 72 |
| D | 67.3829/ -57.9260 | 67.3707/ -57.9145 | 634 | 648 | 68 |

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Operations at the Baffin Bay Station A biodiversity monitoring site. Photographic transects were conducted to sample megafauna; Van Veen grabs for macrofauna and meiofauna/microbes, and CTDs for water column characteristics.



DNA BARCODING

DNA-based identifications using a portion of the cytochrome oxidase subunit I (COI) mitochondrial protein-coding gene have been employed across broad taxonomic ranges (Herbert et al. 2003) and provide an especially useful tool in cases where external morphology may not provide sufficient traits for full taxonomic resolution, especially in areas where the species are not well known. This approach has been popularly coined "DNA-barcoding". Unique or unknown specimens from each sampling station were targeted for DNA barcoding purposes. Each specimen was either subsampled for tissue, or the entire specimen was preserved in 95% ethanol. Genomic DNA was extracted using DNeasy®Tissue Kit (Qiagen), then PCR-amplified with one of several primer pairs to obtain a fragment of the mitochondrial cytochrome oxidase subunit I gene.

SPECIMEN PHOTOGRAPHS



An attempt was made to photograph all macrofaunal taxa identified from the grab samples. When possible these were taken at sea, prior to preservation, to capture natural colour.

CURRENT STATUS AND FUTURE PLANS

It is expected that all data from the 5 marine biodiversity monitoring sites will be extracted by April 2015. Additional camera transects and grab samples were taken in other parts of Davis Strait, and west of Resolution Island. Full sampling according to the CBMP protocols were undertaken at some of these locations. These may be analyzed in future depending upon funding. All data from the Greenland Halibut closure area sites will be made available through the Arctic Council Data Portal (Pomerleau et al. 2014).

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