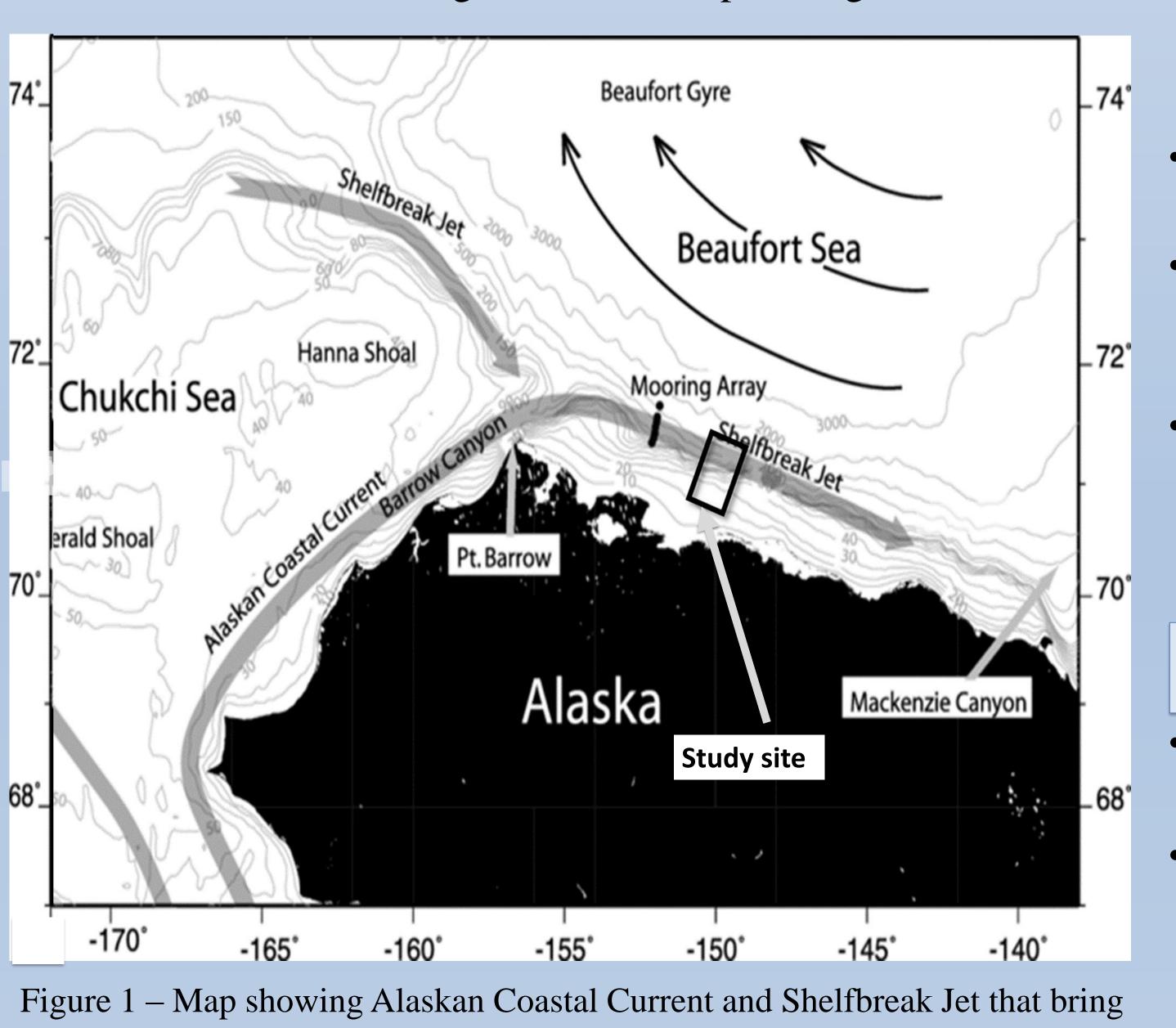


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#### Introduction

- Alexandrium tamerense is a well studied dinoflagellate • known for its ability to produce the neurotoxin that causes paralytic shellfish poisoning (John et al. 2014).
- Until 1970 A. tamerense was only found in Europe, North • America, and Japan but has been increasingly found all over the globe, and as a result toxic blooms have been increasing (Lilly et al. 2007).
- Alexandrium is characteristically found in temperate and subtropical regions (Lilly et al. 2007) and as the Arctic warms, there is considerable concern that it may be expanding into the Arctic.
- Warm Pacific water flows into the Alaskan Beaufort Sea from the Alaskan Coastal Current and is brought onto the Beaufort Sea shelf by the Shelfbreak Jet, shown in Figure 1 (Spall et al. 2008).
- Under certain wind conditions, the Shelfbreak Jet reverses ulletresulting in upwelling delivering nutrient rich, cold water to the region. (Pickart et al. 2011).
- study combined multiple methods to This lacksquareAlexandrium abundance in the Beaufort shelf region collected before, during, and after an upwelling event.



warm Pacific water to the Beaufort Sea Shelf (Pickart et al. 2011). The study site is displayed with a black rectangle.

Acknowledgements: Funding for this work was provided by awards from the Jeffress Trust Awards Program in Interdisciplinary Research to PDC, the Jacques S. Zaneveld endowed scholarship to SVE, and the Weston Howland Jr. Postdoctoral Scholarship and WHOI Access to Sea Fund to KL. Our fieldwork was supported by NSF grant PLR-160391 to Carin Ashjian and Joel Llopiz (Woods Hole Oceanographic Institution). FlowCAM (Fluid Imaging Technologies; loaned to the project by the NSF United States Antarctic Program).

# Alexandrium in the Arctic: Are harmful algae spreading as the Arctic warms? Sveinn V. Einarsson<sup>1</sup>, Kate Lowry<sup>2,3</sup>, Carin Ashjian<sup>3</sup>, Robert Pickart<sup>3</sup>, P. Dreux Chappell<sup>1</sup>

evaluate

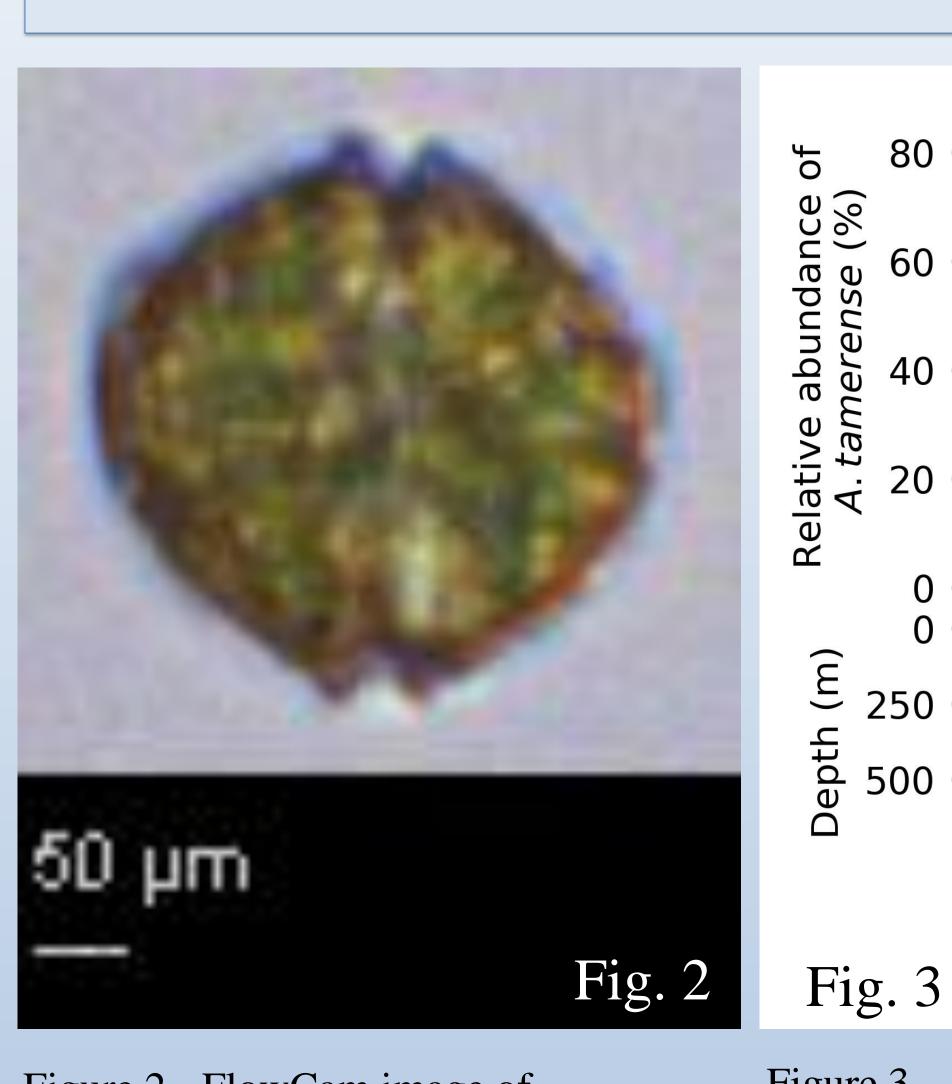


Figure 2 - FlowCam image of Alexandrium found before upwelling.

# Fig. 3 (a)

### Discussion

- Here we have shown 3 methods of monitoring Alexandrium: FlowCam analysis, quantitative PCR, and relative abundance of 18S sequences. Each method showed an increased abundance before upwelling on the Beaufort Sea shelf and shelfbreak.
- It does seem that warm Pacific water is potentially bringing Alexandrium into the Arctic where it hasn't been found before.
- Curiously Alexandrium decreased dramatically during and after upwelling, which would be a source of nutrients that might otherwise be expected to increase abundance.
- A possible explanation is that the Shelfbreak Jet is known to reverse during upwelling winds (Pickart et al. 2011). Shutting down the import of warm Pacific water, and taking away the warmer water *Alexandrium* is found in.

  - Sequence Variants identified using BLAST (Altschul et al. 1990.
  - qPCR assay for A. tamerense followed Hosoi-Tanabe and Sako, 2005.

Re	eferences:
•	John et al. 2014. 10.1016/j.protis.2014.10.001
•	Lilly et al. 2007. https://doi.org/10.1111/j.1529-8817.2007.00420.x
•	Spall et al. 2008. https://doi.org/10.1175/2007JPO3829.1
•	Pickart et al. 2011. 10.1016/j.pocean.2010.11.005
•	Liao et al. 2015. https://doi.org/10.1038/srep16630

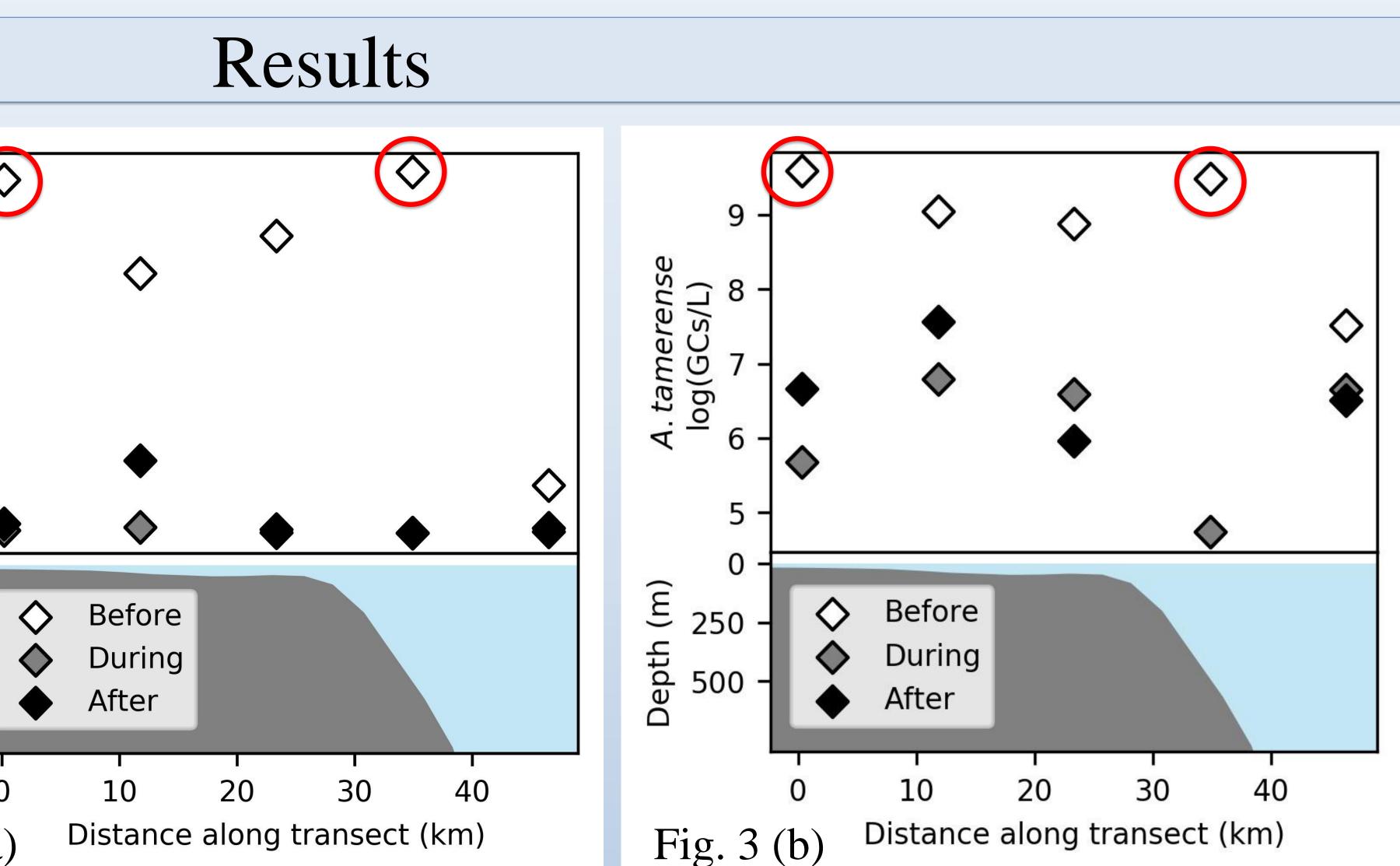


Figure 3 - (a) Relative abundance of A. tamerense 18S sequences as a proportion of total 18S phytoplankton sequences (dinoflagellates, diatoms, and haptophytes) (b) qPCR abundance of A. tamerense.

#### Thoughts for the future Arctic is getting warmer (Liao et al. 2015) and sea ice is

- retreating more each year.
- Beaufort sea upwelling is strongest when there is less sea ice (Schulze and Pickart 2012) and the winds that drive the upwelling in the Beaufort Sea are becoming more frequent (Pickart et al. 2013).
- As the Arctic warms, Alexandrium needs to be monitored in this area due to its ability to produce neurotoxins that can accumulate in shellfish and fish, and as a result can effect humans, other mammals, and seabirds after consuming contaminated prey (Anderson et al. 2012).

## Methods

For FlowCAM (Fluid Imaging Technologies) imaging, samples were pre-filtered using 100 µm Nitex mesh and 5 mL of filtered sample were run at two different magnifications (100x and 40x) to assess the microplankton community.

The V9 variable region of 18S rRNA was amplified from DNA extracted using the Dneasy kit (Qiagen) from 4L of filtered seawater using 1391F/EukBR primer pair (Stoeck et al. 2010) modified for Illumina sequencing and sequenced at the ODU MiSeq Sequencing facility. DNA sequences were analyzed using DADA2 (Callahan et al. 2016). Amplicon

• Schulze and Pickart 2012. 10.1029/2012JC007985 • Pickart et al. 2013. http://dx.doi.org/10.1016/j.dsr.2013.01.007 • Anderson et al. 2012. 10.1016/j.hal.2011.10.012 • Stoeck et al. 2010. 10.1111/j.1365-294X.2009.04480.x • Callahan et al. 2016. 10.1038/nmeth.3869 • Altschul et al. 1990. http://dx.doi.org/10.1016/S0022-2836(05)80360-2

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