

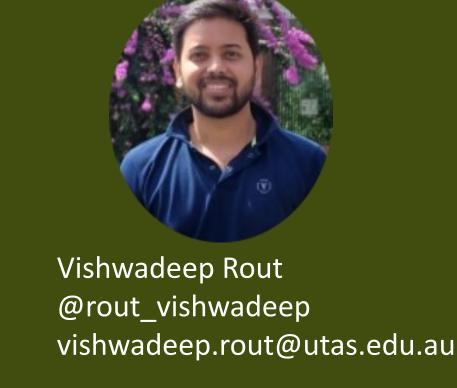
Probing ancient Antarctic krill populations

Vishwadeep Rout^{1,3}, Leonie Suter^{1,2}, Kerrie Swadling^{1,2}, So Kawaguchi^{1,2} and Linda Armbrecht^{1,3}

¹ Institute for Marine and Antarctic Studies (IMAS), University of Tasmania, Hobart, TAS, Australia

² Australian Antarctic Division (AAD), Kingston, TAS, Australia

³ Australian Centre for Excellence in Antarctic Science (ACEAS)



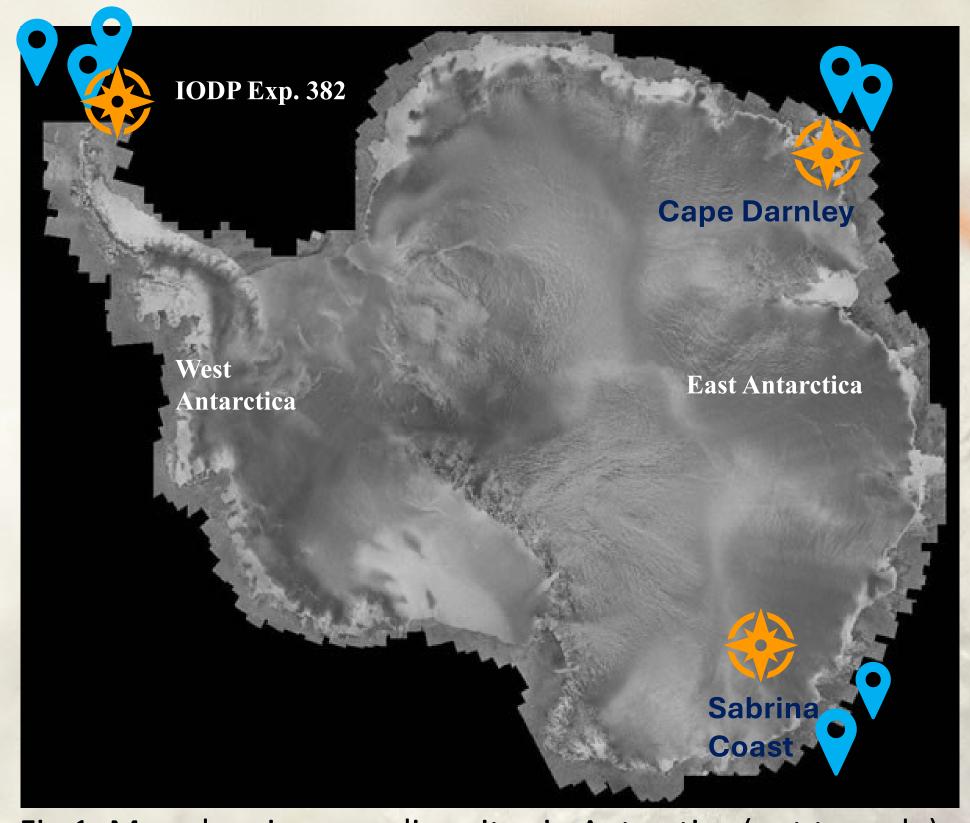


Fig.1. Map showing sampling sites in Antarctica (not to scale).

What is SedaDNA? SedaDNA as a paleoproxy?

SedaDNA is DNA from dead organisms that have sunk from the ocean to the seafloor, forming a record of "who" has inhabited the ocean in the past.

It allows us to study past marine biodiversity comprehensively across different levels of the food web uncovering wide-scale community shifts as a response to past climatic change. Note: SedaDNA is ancient DNA (aDNA) found in sediments (Fig. 4).

Suitable conditions for preservation of sedaDNA?

Polar deep-ocean environments suitable particularly sedaDNA for preservation. Low temperatures, low oxygen concentrations, and the absence of UV radiation are beneficial.

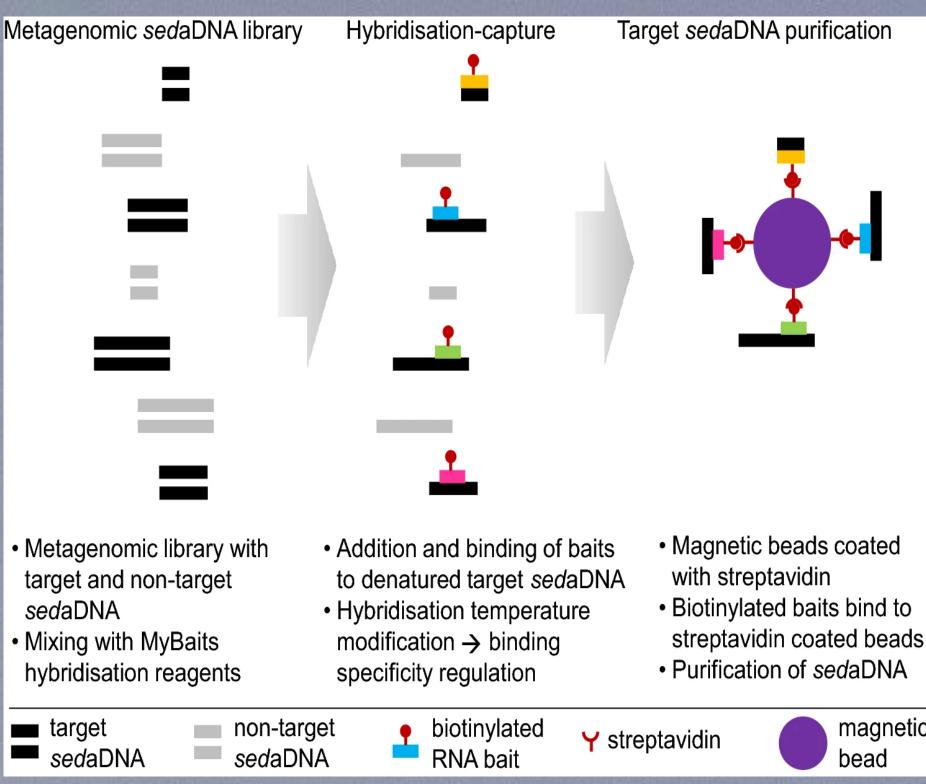


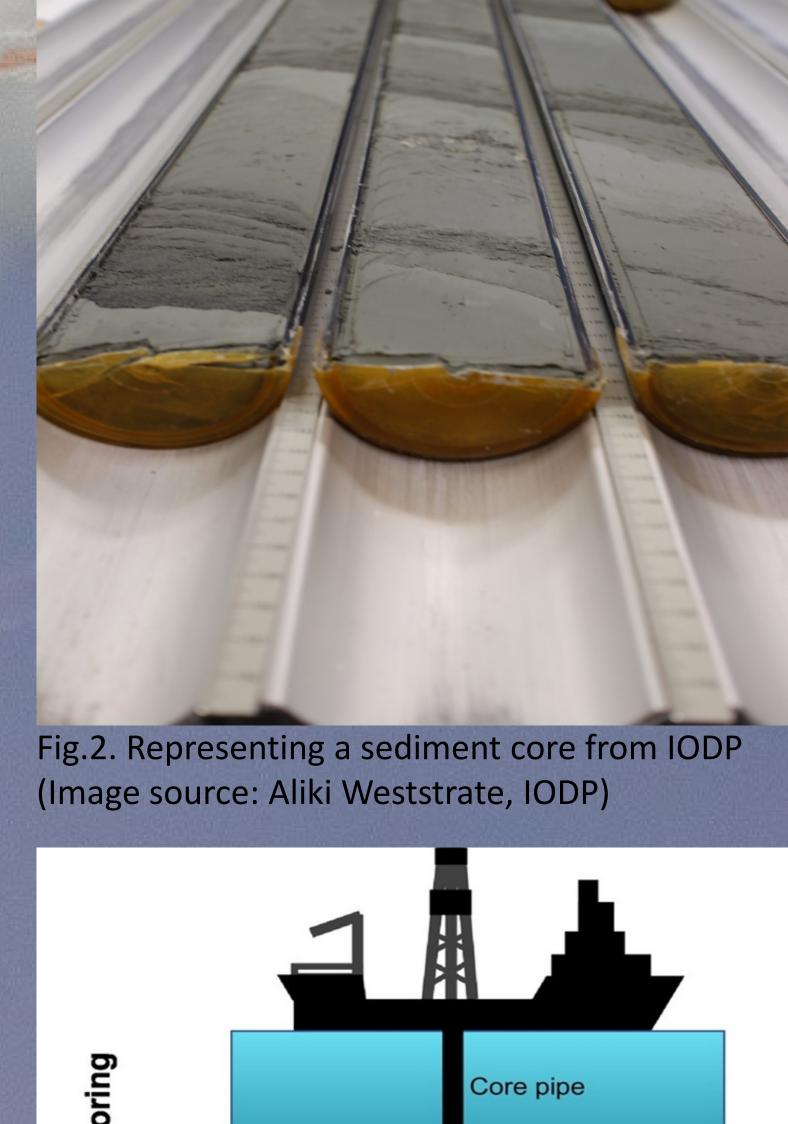
Fig.3. Schematic of hybridisation capture applied to marine sedimentary ancient DNA (sedaDNA) in three main steps (Armbrecht, et al., 2021).

Background

Antarctic krill, vital in Southern Ocean ecosystems, face threats from climate change such as ocean warming and acidification, reduced sea-ice habitat and overfishing. Understanding their response to past environmental changes is crucial for predicting their adaptability to ongoing climate change and aiding conservation efforts. To date, such paleo studies targeting krill have been impossible, as krill leave no fossil records.

Our Goal

To investigate Antarctic krill population dynamics over hundreds of thousands of years by looking for their sedimentary ancient DNA (sedaDNA) from Southern Ocean sediment cores using an RNA-based hybridization-



Core liner with core Decontaminated Decontaminated core-liner core-cutter aDNA sample Sediment core (core centre) tracer samples (core centre and periphery) 00 Negative control Samples

ACGTACGT ACGTACGT Fig.4. Key steps involved in acquiring deep sea sediment cores, subsampling, DNA extraction, aDNA preparation for sequencing and data generation (Armbrecht, et al., 2019).

capture technique (Fig.3).

What are 'baits'? 'Baits' (analogous to the baits used in a fishing context) are short RNA probes designed to be complementary to any chosen DNA sequences (e.g., specific genes of a target organism). By binding to the target sequence, the baits can capture DNA fragments of interest and make

Application of the 'krill baits':

The krill baits will be applied to sedaDNA extracts from the East Antarctic Sabrina coast (IN2017 V01), Cape Darnley (IN2023_V01), and West Antarctic Iceberg Alley (IODP Exp. 382). Refer to Fig. 1 (map on the top left).

sequencing more efficient and cost-effective (Fig.3).

References:

- Armbrecht, L., 2022. Sedimentary ancient DNA (sedaDNA) as a new paleo proxy to investigate organismal responses to past environmental changes in Antarctica
- CORE SCIENCE #1 WHAT IS CORING AND WHAT DOES IT TELL US? by Aliki Weststrate JOIDES Resolution, On board Expedition 375
- Armbrecht, et al., 2021. Hybridisation capture allows DNA damage analysis of ancient marine eukaryotes. Scientific Reports, 11(1), p.3220
- Armbrecht, et al., 2019. Ancient DNA from marine sediments: Precautions and considerations for seafloor coring, sample handling and data generation. Earth-Science Reviews, 196, p.102887 Acknowledgement: The background image is taken by Pete Harmsen/Australian Antarctic Division.



Image: ©Pete Harmsen/Australian Antarctic Division





