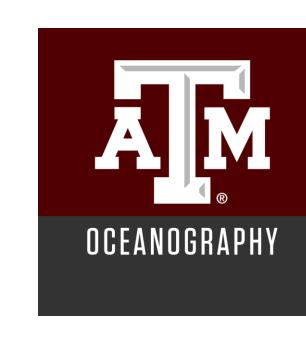
Database Darkly: Why do we care about protistan biodiversity in the deep sea?

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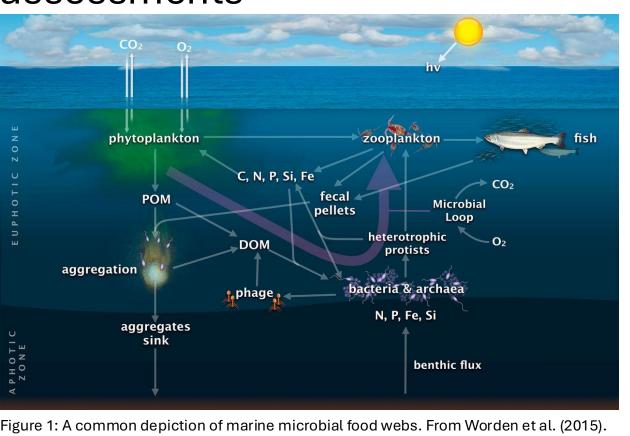
Introduction

What is a protist?

- Protist is a generalized term for single-celled microbial eukaryotes. Protists are essential for ecosystem functions around the planet, with hundreds to thousands of organisms making up a community.
- The domain Eukarya contains all organisms that have a nucleus. This domain comprises a wide range of organisms, including all animals, plants, fungi, and protists.

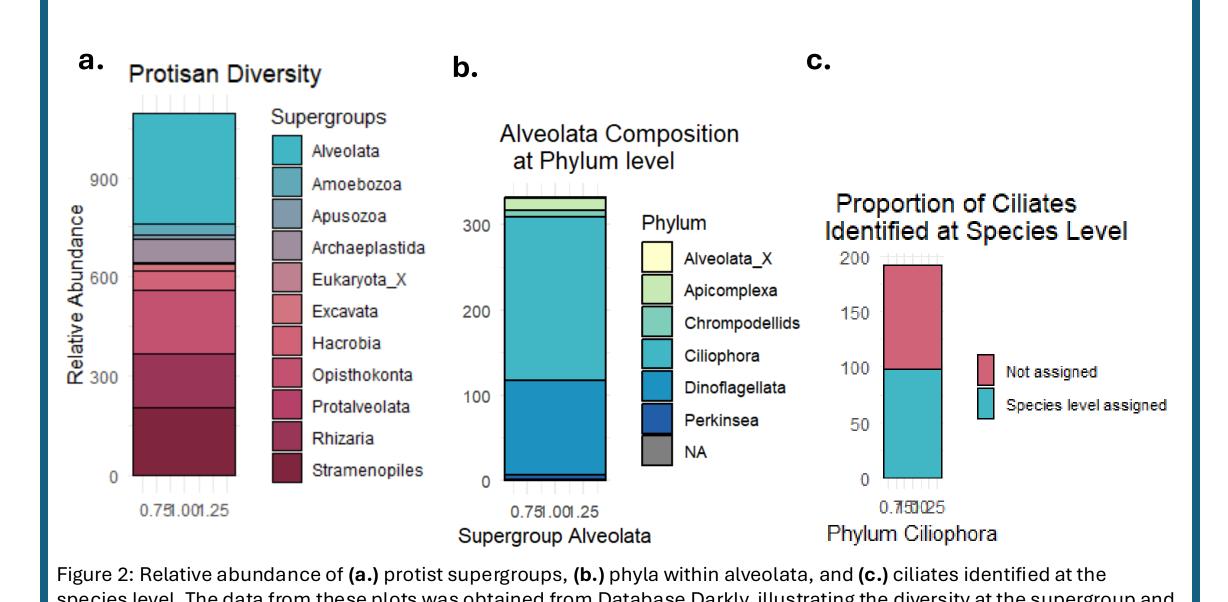
Why do we care about protistan diversity?

- Protists play critical roles in marine ecosystems as primary producers, consumers, decomposers, and trophic links.
- Protistan biomass constitutes a large portion of the living carbon in marine ecosystems. Their rapid metabolic rates underscore their importance in elemental cycling and energy transfer in the ocean.
- Understanding the diversity of marine protists is essential for accurate assessments
- of global biodiversity. Major evolutionary transitions, such as the foundations of multicellularity, occurred within protist lineages.



The Eukaryotic Tree of Life

- Represents the eukaryotic biodiversity on earth, underscoring how all life relates back to a common ancestor.
- Historically, relationships were determined based on morphological traits. However, microscopy cannot adequately observe the smallest protists (picoeukaryotes). Furthermore, organisms may appear morphologically similar yet unrelated.
- Modern molecular tools such as metabarcoding and meta'omics allow access to genetic information.
- These tools are especially helpful since many marine protists are uncultivable, making it nearly impossible to observe morphology and behavior.
- Nonetheless, there are still challenges in constructing the tree as genomic sequencing alone cannot fully reveal the underlying complexity of protistan physiology.



phylum levels. The third plot highlights the challenges in identifying protists at the species level.

Eukarya

Parasitic Protists and their

Photosynthetic past

Apicomplexans are parasitic protists that contain

an organelle that was once believed to be able to

undergo photosynthesis, the apicoplast. While it

no longer has the signature green hue and ability

to photosynthesize, it is necessary for the

However, the discovery of chromera velia, a

photosynthetic protist, connected the relict

organelle of the apicomplexans to their past. This

organism had a photosynthetic plastid while also

found in apicomplexans. Additionally, the shared

having the UGA-Trp codon, which has only been

psbA gene indicates the missing link between

apicomplexans and dinoflagellates

parasite's survival.

In the history of life, we trace a profound transition from a world dominated by prokaryotes to one enriched with complex eukaryotic life forms. This evolutionary event underscores the deep interconnections between various forms of life.

By documenting species from hydrothermal vents, Database Darkly helps us understand the ecological roles of ancient protists, providing a crucial link between early eukaryotic life and modern biodiversity.

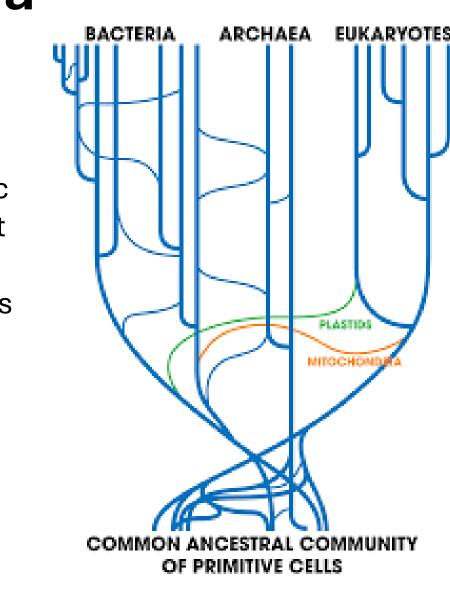


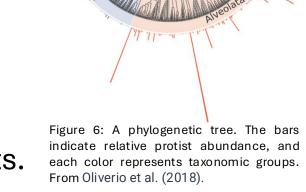
Figure 3: 2005 tree of life showing horizontal

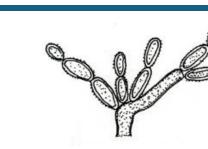
Regarding Geothermal and **Hydrothermal Protists**

Protistan communities consist of a wide variety of organisms. In extreme environments, one way to predict this variability is by pH and/or temperature. As pH decreases, microbial species diversity often follows. In geothermal springs, the Alveolata supergroup is found most consistently across springs and is often the most abundant.

Eukaryotes have adapted to survive in these environments because they altered their internal processes during the genetic transfer of proteins, allowing them to keep their internal pH stable.

Additionally, the anaerobic *ciliate* Trimyema minutum thermophilium has been found at high-temperature springs and is also associated with shallow sea hydrothermal vents.





Deep Sea Fungi



Research on the diversity and ecological role of deep-sea fungi was relatively rare until the early 2000s. What has been discovered in the last few decades suggests potential for biotechnical and biomedical applications.

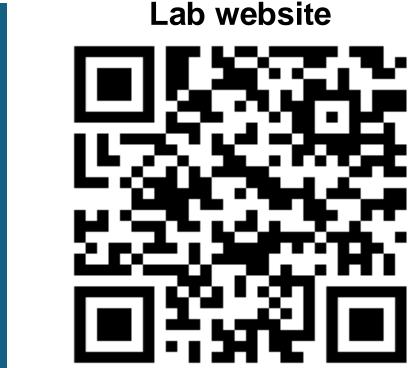
The extreme environment of the deep sea requires organisms to adapt physiologically. Deep-sea fungi regulate and modify gene expression to accommodate the low-temperature, high-pressure environment. These alterations yield specialized proteins and secondary metabolites with industrial and medical applications.

Expanding our knowledge on deep-sea fungi would not be possible without scientific collaboration, including the use of databases such as Database Darkly.

Methods

- The goal of this project was to create a database that compiled information from a deep-sea hydrothermal vent molecular survey (Hu et al. 2023) with existing databases.
- Several databases were used to cross-reference any unknown entries that may be listed under different taxonomic names.
- Protistan taxonomic naming structure is notoriously challenging to document and often changes with new genetic surveys. Therefore, we worked to include other names species may have been known by in the past
- The **PR2 (Protist Ribosomal Reference) database** is a collection of three 18S rRNA databases that aim to provide annotated sequences from domain to species. It currently contains over 220,000 sequences focused on protists but also contains metazoa, fungi, and plant sequences.
- marine organisms and their names. This database was used as a reference for any other name that an organism could go by, which then would widen our research parameters.

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Database Darkly website



Acknowledgements: Kayla Nedd, Alexis Adams, Julie Huber, Harriet Alexander, Arianna Krinos, Rika Anderson, Maria Pachiadaki, Ginny Edgcomb, Erica Herrera, Jeff Seewald, Susan Lang, Chip Brier, Chris German, Sean Sylva, Amy Smith, & Gretta Serres

Funding: National Science Foundation, Center for Dark Energy Biosphere Investigations, & the TAMU Undergraduate Research Program



The WoRMS (World Register of Marine Species) database provides a comprehensive list of

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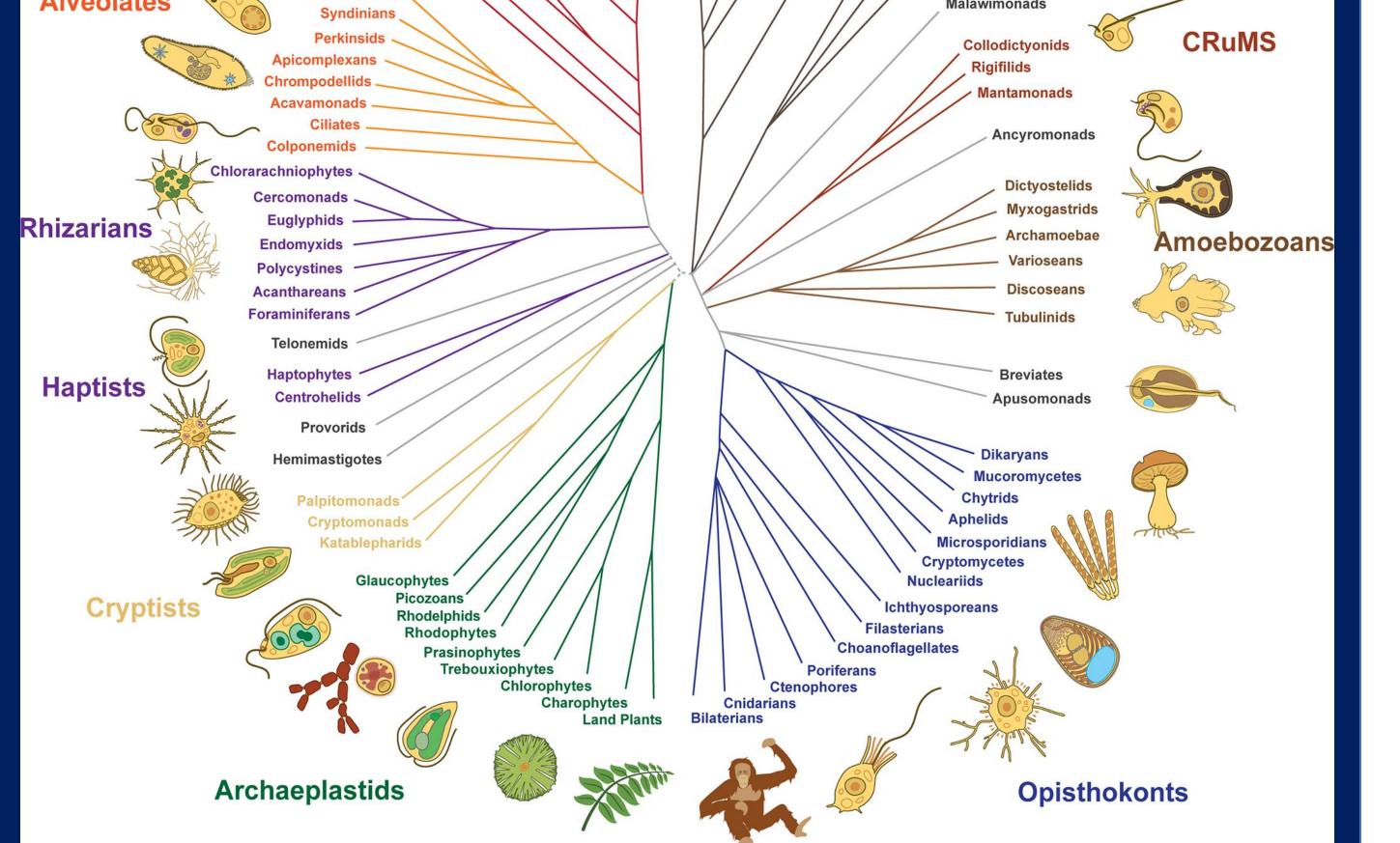


Figure 7: Eukaryotic Tree of Life. From Keeling, PJ & Eglit, Y (2023).