

Abstract

The biodiversity of certain areas can be negatively affected by human influence in the environment - this includes the introduction of invasive species (Day, 2016). Our group has chosen to study the effect of boat traffic on biodiversity in Cold Spring Harbor. We hypothesized that the introduction of invasive species via boat traffic has negatively impacted Cold Spring Harbor wildlife by harming organisms native to the area. To test our hypothesis we collected information concerning the regularity of recreational and commercial boating throughout the past century (starting with the publication of the Davenport paper in 1904). We arranged to obtain this information from various sources including: an interview from the staff of the the Cold Spring Harbor Whaling Museum, public records, the Davenport paper, and scientists from the Cold Spring Harbor Laboratories. However we were not able to collect this information from these sources due to the recent outbreak of Covid -19. Despite the fact that we were unable to collect data from outside sources, we were able to use our own data we collected earlier this year to create a partial conclusion of our research. However, we were not able to determine any correlation between boat traffic and the biodiversity of Cold Spring Harbor. Our data measures the current biodiversity of the area by collecting and documenting organisms found in Cold Spring Harbor and barcoding their DNA. Barcoding the DNA of the collected organisms would have allowed us to accurately identify them and compare the most prevalent organisms of the area currently to the most prevalent species during the time of the Davenport study. However we were unable to visit the DNA Learning Center to do this due to recent social distancing regulations. By doing this we would have been able to determine if there have been any significant increases or decreases in biodiversity in Cold Spring Harbor.

Research

In order to prove that a direct correlation exists between boat traffic and biodiversity, our team had to research and compare the trend in Cold Spring Harbor boat traffic to the trend in biodiversity. This information was not easily accessible online- we had to conduct our own outside research to try to gain a better understanding of our topic.

CHANGES IN C.S.H. BIODIVERSITY:

Davenport Paper published 1904

time.

This paper was written in 1904 and gives a brief summary of biodiversity in Cold Spring Harbor at that

Cold Spring Harbor Laboratories' Biodiversity Records The Davenport paper was helpful, although it was somewhat vague. To gain a clearer understanding of past biodiversity, our group was looking

contacting the CSH laboratory for more detailed biodiversity records.

B.L.I. Collection Our group, along with the rest of our

class, collected organisms at the

Spit in Cold Spring Harbor in

partnership with Barcode Ll. We created a detailed form illustrating

our collective results and used this preadsheet to compare the

xistence of certain species from the

CHANGES IN C.S.H. BOATING:

Schoolwide Survey

Our group searched extensively for official online boating but unfortunately there was very limited information. Therefore, we it necessary to find this A through our school's student body. We created a survey to get a sense of boat ownership numbers among

the families attending Cold Spring Harbor High School.

CSH Whaling Museum Interview

Cold Spring Harbor is well known for once being a whaling community in the 1900s. We planned to email a staff member of the Whaling Museum to ask for an interview and hopefully educate ourselves about whaling in the harbor and the development of boat traffic levels throughout the last century, but the

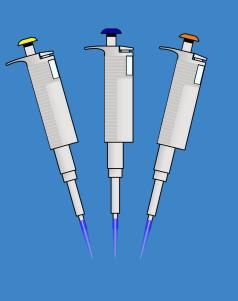
pandemic had their offices closed.

Transportation of Invasive Species via Boat Traffic in Cold Spring Harbor and Its Effect on Biodiversity

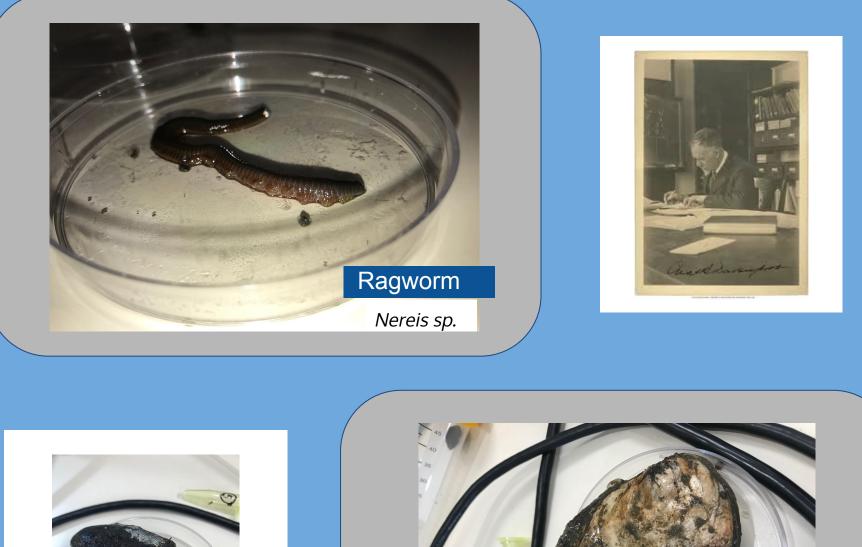
Materials

The Davenport Paper,

published in 1904, was needed to compare our experiment results. We also used various tools to collect our species such as filters for sand and dirt, and shovels.



Methodology For our biodiversity study we went to the southeastern side of the spit at low tide. Using shovels, buckets, strainers and nets in groups of 3, accompanied by a scientist from the Cold Spring Harbor Laboratories, our group searched for organisms in our designated area. When we found an organism, we recorded its location and habitat, the time we found it, and the name of its species(this was determined with the help of the knowledgeable scientists that accompanied us in our collection). We also took notes of certain species that appeared to be particularly abundant, as well as the environment in which the organism was found. To find the organisms, we searched in the mud with the shovel to look for oysters and clams, turned over large pieces of driftwood or rocks to uncover crabs and looked around on the ground for other organisms. During our collection, we also gathered data on the quality of the water. We measured the pH (using pH strips), turbidity (using a turbidity tube), temperature (using digital thermometers in degrees Celsius) and filter samples of the water we collected back at the DNALC. After our collection, we went to the DNA lab and documented our samples and photographed them. They then were placed in ethanol to preserve them so we could barcode them later on. As a class, we shared all of our results via a Google spreadsheet to consolidate the data we collected. This made one complete data set and made barcoding our collected samples easier and more organized.



Ribbed Mussel

Geukensia demissa

Oyster

Tools Used to Prepare

Organisms for Barcoding

1 microscope camera

Micropipettes

• 6 beakers

• Ethanol

• 12 test tubes

Crassatrea virginice

Results

4 G4-001	Mollusca	Atlantic oyster drill	Urosalpinx sp	40.86721 N	73.46692 W
4 G4-002	Crustacea	Blue crab	Callinectes sapidus	40.86721 N	73.46692 W
4 G4-003	Mollusca	Ribbed mussel	Geukensia demissa	40.86703 N	73.46402 W
4 G4-004	Mollusca	Mud snail	Lyanassa obsoleta	40.86712 N	73.46395 W
G4-005	Platyhelminthes	Ragworm	Nereis sp	40.86811 N	73.46162 W
4 G4-006	Mollusca	Oyster	Crassatrea virginice	40.86813 N	73.46260 W
G4-007	Crustacea	Barnacles	Balanidae sp	40.86912 N	73.46301 W
4 G4-008	Mollusca	Jingle shell	Anomia simplex	40.86912 N	73.46301 W
G4-009	Crustacea	Fiddler Crab	Uca sp	40.86912 N	73.46301 W
G4-010	Platyhelminthes	Small worm	Cerebratulus lactus	40.86912 N	73.46301 W
4 G4-011	Crustacea	Mud crab	Dyspanpeus sayi	40.86912 N	73.46301 W
4 G4-012	Mollusca	Soft shell clam	Mya sp	40.86912 N	73.46301 W

The scientific and common names of the organisms found by our group during our biodiversity collection are shown in the table above (Figure 1).

Sea Stars: The Importance of DNA Barcoding

After combining and analyzing the results of our class' data collections, we found that a different group, Group #2, had discovered a starfish, scientifically identified as *asterias forbesi*. However, upon researching the organism, we found that several sites seem to think that asterias forbesi is actually extinct. Having access to DNA barcoding is so important because it can solve these issues and prove that *asterias forbesi* is still in existence and can be found throughout the northwest Atlantic coast.

Mollusca: The Phylum That Continued to Thrive

Our research found that one particular phylum of organisms studied in our collection seemed to be particularly successful: mollusca. Our group believes this is because the increasing boat traffic did not have a detrimental effect on these organisms- in fact, certain organisms like mussels or barnacles even find shelter on these boats.

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Conclusion

commendations

cate Boaters on Spread of AIS

- rict Boating in Designated Nature Areas
- ts must be checked for wildlife/foreign material pre re-entering the Long Island Sound.
- -distance travel must be reported to the county ore departure.

knowledgements

code L.I. & Cold Spring Harbor DNA Learning Center Our project was done in coordination with the Barcode L.I. program. d Spring Harbor DNA Lab

ne instructors at the DNA Lab taught our groups to isolate DNA and w to barcode organisms. This experiment would not have been ossible without their help.