Feeding rates on and nutritional content of algal species collected in Antarctica, Fiji, and California.

Website: https://www.bco-dmo.org/dataset/739345
Data Type: Other Field Results, experimental

Version: 1

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Project

» <u>Detecting genetic adaptation during marine invasions</u> (Genetic Adaptation Marine Inv)

Contributors	Affiliation	Role
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Abstract

Feeding rates on and nutritional content of algal species collected in Antarctica, Fiji, and California.

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Coverage

Spatial Extent: N:38.32 E:-64.05 S:-64.77 W:-123.04

Temporal Extent: 1996 - 2014

Dataset Description

Feeding rates on and nutritional content of algae.

Methods & Sampling

These data are described in detail in Demko, A. M., C. D. Amsler, M. E. Hay, J. D. Long, J. B. McClintock, V. J. Paul, and E. E. Sotka. 2017. Declines in plant palatability from polar to tropical latitudes depend on herbivore and plant identity. Ecology 98:2312–2321. We quantified the relative palatability of 50 seaweeds by offering each generalist herbivore a pairwise choice between each test seaweed and a control in artificial feeding assays with finely ground lyophilized tissue. For each assay, 8 g of freeze-dried and ground powder (ground via Wiley Mill) of one experimental seaweed and the Ulva control were rehydrated with 28 mL distilled water and mixed with 72 mL molten agar (2% by mass). Seaweed mixes were then poured into side-by-side lanes in a mold on window screen (1 9 2 mm squares) in a thickness of approximately 2 mm. After cooling, the screen was then cut into strips with approximately 80 squares of each food type separated by 2 cm. Individual strips were then isolated with approximately 30 separate crabs and 50 separate urchins, and removed before the entirety of either food was consumed or until 24–30 h had elapsed. Replicates in which <10% or >95% of all food offered

was consumed were removed before statistical analysis because of their low power to infer feeding choice. Our sample size for each seaweed-herbivore combination ranged from n = 10 to 46. Palatability of each seaweed to each herbivore was quantified as the proportion of the experimental seaweed consumed divided by the total consumption of experimental and control seaweed within a replicate (%T).

Data Processing Description

BCO-DMO Data Processing Notes:

- -Replaced NA and blank cells with nd
- -Reformatted column names to comply with BCO-DMO standards

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Data Files

File

palatability_algae.csv(Comma Separated Values (.csv), 10.54 KB)

MD5:83900be5f93bdf589b7776c478814f40

Primary data file for dataset ID 739345

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Related Publications

Demko, A. M., Amsler, C. D., Hay, M. E., Long, J. D., McClintock, J. B., Paul, V. J., & Sotka, E. E. (2017). Declines in plant palatability from polar to tropical latitudes depend on herbivore and plant identity. Ecology, 98(9), 2312–2321. doi:10.1002/ecy.1918

Methods

Results

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Parameters

Parameter	Description	Units
Division	Division of alga	unitless
Order	Order of alga	unitless
Family	Family of alga	unitless
GenSpec	Genus and species of alga	unitless
Location	Antarctica (64.77 deg S, 64.05 deg W), California (32.72 deg N, 117.16 deg W, and 38.32 deg N, 123.04 deg W) or Fiji (18.00 deg S, 179.00 deg E)	unitless
Collection_Site	Local site name	unitless
Zone	Polar, temperate, or tropical	unitless
Latitude_of_Collection	Absolute degrees latitude	decimal degrees
Mean_Distribution_Latitude	A mean of the latitude where that seaweed genus has been collected; as recorded in http://gbif.org	decimal degrees
Ash_Free_Dry_Mass	Ash-free dry mass	percent per drymass
Percent_Protein	Percent protein	percent per drymass
Percent_Carbon	Percent Carbon	percent per drymass
Percent_Nitrogen	Percent Nitrogen	percent per drymass
Carbon_Nitrogen	Carbon to Nitrogen ratio	ratio of absolute milligrams
Percent_Phenolics	Percent Phenolics	percent per drymass
Crabs_PercentT	The proportion of the treatment seaweed consumed divided by the total consumed in an assay (control + treatment); results for the crab Mithraculus sculptus	percent
Urchins_PercentT	The proportion of the treatment seaweed consumed divided by the total consumed in an assay (control + treatment); results for the urchin Echinometra lucunter	percent
Wet_Mass	The wet mass of collected material	grams
Dry_Mass	The dry mass of collected material	grams
WM_DM	Ratio of wet mass to dry mass	grams

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Project Information

Detecting genetic adaptation during marine invasions (Genetic Adaptation Marine Inv)

Coverage: Estuaries of NW and NE Pacific; estuaries of NW and NE Atlantic

Description from NSF award abstract:

Biological introductions, defined as the establishment of species in geographic regions outside the reach of their natural dispersal mechanisms, have dramatically increased in frequency during the 20th century and are now altering community structure and ecosystem function of virtually all marine habitats. To date, studies on marine invasions focus principally on demographic and ecological processes, and the importance of evolutionary processes has been rarely tested. This knowledge gap has implications for management policies, which attempt to prevent biological introductions and mitigate their impacts. The Asian seaweed *Gracilaria vermiculophylla* has been introduced to every continental margin in the Northern Hemisphere, and preliminary data indicate that non-native populations are both more resistant to heat stress and resistant to snail herbivory. The project will integrate population genetics, field survey and common-garden laboratory experiments to comprehensively address the role of rapid evolutionary adaptation in the invasion success of this seaweed. Specifically, the PIs will answer the following. What is the consequence of introductions on seaweed demography and mating systems? How many successful introductions have occurred in North America and Europe? Where did introduced propagules originate? Do native, native-source and non-native locations differ in environmental conditions? Do native, native-source and non-native populations differ in phenotype?

The intellectual merit of this project is based on three gaps in the literature. First, while biological invasions are widely recognized as a major component of global change, there are surprisingly few studies that compare native and non-native populations in their biology or ecology. Native and non-native populations will be surveyed in a similar manner, allowing assessment of differences in population dynamics, mating system, epifaunal and epiphytic communities, and the surrounding abiotic and biotic environment. Second, *G. vermiculophylla* exhibits a life cycle typical of other invasive species (including some benthic invertebrates), yet we still lack data on the effects of decoupling the haploid and diploid stages on genetic structure, and in turn, on the evolvability of their populations. Finally, this project will provide unequivocal evidence of an adaptive shift in a marine invasive. To our knowledge, such evolutionary change has been described previously for only a complex of marine copepod species. *G. vermiculophylla* will serve as a model for understanding evolution in other nuisance invasions, and perhaps lead to novel methods to counter future invasions or their spread.

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Funding

Funding Source		Award	
	NSF Division of Ocean Sciences (NSF OCE)	OCE-1357386	

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