Sample key for Symbiodinium species and abundance in Dendrogyra cylindrus from 3 regions of the Florida Reef Tract 2015 - 2016 (EMUCoReS project)

Website: https://www.bco-dmo.org/dataset/700101 Data Type: Other Field Results, experimental Version: Version Date: 2017-05-17

Project

» RAPID: A hyper-thermal anomaly in the Florida Reef Tract: An opportunity to explore the mechanisms underpinning patterns of coral bleaching and disease (EMUCoReS)

Contributors	Affiliation	Role
<u>Rodriguez-Lanetty,</u> <u>Mauricio</u>	Florida International University (FIU)	Principal Investigator
Lirman, Diego	University of Miami Rosenstiel School of Marine and Atmospheric Science (UM-RSMAS)	Co-Principal Investigator
<u>Richardson, Laurie</u>	Florida International University (FIU)	Co-Principal Investigator
Copley, Nancy	Woods Hole Oceanographic Institution (WHOI BCO-DMO)	BCO-DMO Data Manager

Table of Contents

- <u>Coverage</u>
- Dataset Description
 - Data Processing Description
- Data Files
- <u>Parameters</u>
- Deployments
- Project Information
- Funding

Coverage

Spatial Extent: N:25 E:-80.4 S:24.47 W:-81.75 **Temporal Extent**: 2015-04-01 - 2016-04-30

Dataset Description

This dataset includes a key to samples of *Dendrogyra cylindrus* which were analyzed for 23S on an Illumina platform from three sites in the Florida Reef Tract, one each in the upper, middle and lower Keys in 2015-2016. Tube numbers are matched to sample name, status (pre-bleach, pale, bleached, recovery), region, locations, sampling time point, total number of colonies and number of colonies sampled.

Data Processing Description

BCO-DMO Processing Notes:

- added conventional header with dataset name, PI name, version date
- modified parameter names to conform with BCO-DMO naming conventions

Data Files

 File

 sample_key_Dcyl_2015_16.csv(Comma Separated Values (.csv), 4.40 KB)

 MD5:4970e86ec7c61bedff8641e450cebf0d

 Primary data file for dataset ID 700101

[table of contents | back to top]

Parameters

Parameter	Description	Units
date_sampled	date sampled	unitless
site	collection site	unitless
tube	sample tube	unitless
sample_ID	sample identifier ((yyyymmdd.site.colony-num)	unitless
depth	sampling depth	meters
total_num_colonies	total number of Dendrogyra cylindrus colonies	colonies
num_colonies_sampled	number of Dendrogyra cylindrus colonies sampled	colonies
lat	latitude; north is positive	decimal degrees
lon	longitude; east is postivie	decimal degrees

[table of contents | back to top]

Deployments

Coral_Bleaching_FRRP

Website	https://www.bco-dmo.org/deployment/640250
Platform	shoreside Florida_Coral_Reefs
Start Date	2014-01-01
End Date	2015-08-20
Description	Coral reef surveys as part of the project "RAPID: A hyper-thermal anomaly in the Florida Reef Tract: An opportunity to explore the mechanisms underpinning patterns of coral bleaching and disease". Single location entered: Florida Reef Tract, 24.8684, -80.6435 in order to 'ground' the datasets.

[table of contents | back to top]

Project Information

RAPID: A hyper-thermal anomaly in the Florida Reef Tract: An opportunity to explore the mechanisms underpinning patterns of coral bleaching and disease (EMUCoReS)

Coverage: Florida Reef Tract (24.868358, -80.643495)

Coral reefs are among the most biologically diverse and economically important ecosystems on the planet. However, coral reefs are in a state of global decline due to effects of climate change, disease outbreaks, and other stressors. Mass coral bleaching events, a breakdown of the association between corals and their symbiotic algae, are predicted to become more frequent and severe in response to climate change, and it is expected that subsequent disease outbreaks will become more common. Beginning in August 2014, nearly all coral species in the Florida Reef Tract have undergone severe bleaching, in some cases followed by coral mortality and/or disease outbreaks. This widespread, thermal-induced event presents a unique time-sensitive opportunity to explore the mechanisms underpinning the patterns of coral bleaching, disease, and recovery. The mechanisms linking patterns of bleaching, disease, mortality, and recovery remain relatively unexplored. This research will explore the influences that genotype combinations of host polyps, their algal symbionts, and associated bacterial have on bleaching/disease likelihood and recovery/mortality predisposition of coral specimens. By providing a mechanistic understanding of the processes that underlie coral bleaching and subsequent recovery this research will contribute to measures in support of preserving this invaluable natural resource. The study will further involve students from diverse backgrounds as well as provide project internship opportunities for high school students. A web based radio blog will disseminate project results and other relevant developments to the broad audiences

Mass coral bleaching events are predicted to become more frequent and severe in response to climate change, and it is expected that subsequent disease outbreaks will become more common. The lack of a baseline genetic datasets for coral holobionts prior to previous natural bleaching events has hindered our understanding of recovery patterns and physiological tolerance to thermal stress, also known as coral bleaching. An extensive pre-thermal stress baseline of genotypic identity of coral hosts, Symbiodinium, and associated bacterial community offers a unique opportunity to analyze changes associated with current bleaching event along the Florida coastline and to document holobiont compositions most and least resistant/resilient to bleaching and disease. Repeated sampling of the same coral colonies will allow the investigators to compare holobiont composition before, during and after bleaching of both healthy and diseased individuals. This bleaching event is a time-sensitive natural experiment to examine the dynamics of microbes (Symbiodinium and bacteria) associated with affected colonies, including their potential influence on disease susceptibility and resistance of reef corals. This effort would constitute the first time that high throughput sequencing of coral, Symbiodinium endosymbiont, and the coral-associated bacterial community genotypes are together used to explain patterns of disease, recovery, and mortality following natural bleaching. This study will likely change the way investigators study emerging wasting diseases of keystone species that define marine benthic communities.

[table of contents | back to top]

Funding

Funding Source	Award
NSF Division of Ocean Sciences (NSF OCE)	<u>OCE-1503483</u>
NSF Division of Ocean Sciences (NSF OCE)	<u>OCE-1503430</u>

[table of contents | back to top]