

Ocean Acidification International Coordination Centre



#### UNIVERSITY OF GOTHENBURG



#### **Basic training course on ocean acidification**

EVT1804704

14-19 March 2022

## Monitoring – How long?



## Chemistry

Upus Natara And Organization	May 2019	
The original version of this Methodology is available as IOC/EC-LI/2	Annex 6	
INDICATOR METHODOLOGY FOR 14.3.1		
Indicator Description 14.3.1 – Average marine acidity (pH) measure representative sampling stations.	ed at agreed suite of	
Target		
14.3: Minimize and address the impacts of ocean acidification, including scientific cooperation at all levels.	g through enhanced	
Tier level		
Tier II - Indicator is conceptually clear, has an internationally establishe standards are available, but data are not regularly produced by countrie	d methodology and es.	
Definition		
This indicator is based on observations that constrain the carbon system capture the variability in ocean acidity at locations providing ocean server this context refers mainly to the four measurable parameters: pH (the co- ions on a logarithmic scale), DIC (total dissolved inorganic carbon), pC pressure), and TA (total alkalinity). Ocean acidification is a reduction in an extended period of typically decades or longer, which is caused prin dioxide from the atmosphere1. Ocean services are the benefits the oce which may be recreational, economic, environmental (by providing coar Average2 as used herein is the equally weighted annual mean.	m, which are required to vices. The carbon system in oncentration of hydrogen O <sub>2</sub> (carbon dioxide partial the pH of the ocean over narily by uptake of carbon van provides to people, stal protection) or cultural.	
A agreed suite of representative sampling stations are sites that: 1) ha frequency adequate to describe variability and trends in carbonate cher information on the exposure of and impacts on marine systems to ocea data of sufficient quality and with comprehensive metadata information data from other sites in the country.	ve a measurement mistry to deliver critical in acidification, 2) provide to enable integration with	
Unit		
pH on total scale		
and/or $pCO_2$ [µatm or ppt], DIC [µmol kg <sup>-1</sup> ], TA [µmol kg <sup>-1</sup> ]		





#### GOA-ON... since 2012



Global Ocean Acidification Observing Network



St. Andrews 2013



Seattle 2012







### A long list of measurements



Global Ocean Acidification Observing Network:

Requirements and Governance Plan

**Second Edition** 



- Level 1 measurements
- Addressing Goal 2 at the broadest scale requires the measurement of biomass or abundance of functional groups, listed below, contemporaneous with the physical and chemical measurements for Goal 1 that achieve at least 'weather' data quality.
- Biomass/abundance of:
- o Phytoplankton
  - o Zooplankton
- o Benthic producers and consumers (shelf seas and nearshore)
  - Etc. etc. etc

Unrealistic Comparison difficult



## Biology Working Group

Global Ocean Acidification Observing Network

## First meeting 2015 (virtual)

**Kirsten Isensee** Sam Dupont Nelson Lagos Luis Valdes Maciej Telszewski Sinead Collins Phil Williamson **Ulf Riebesell** Piero Calosi Yuri Artioli Steve Widdicombe Philip Munday Libby Jewett Nic Bax **Erica Ombres** 

#### 2016 – First meeting in Monaco







Three tasks identified

Task #1 - Inform the chemical monitoring program about the biological needs [Short term]

Task #2 – Evaluate the needs and requirement of a Biological Monitoring Program [Medium/Long term]

Task #3 – **Develop a theoretical framework linking chemical changes to biological response** [Medium/Long term]

#### 2020 – Second meeting in Sweden

## Task #2 – Evaluate the needs and requirement of a Biological Monitoring Program



#### What does OA Biological Impacts monitoring look like?



#### **Expanding to a Global / Regional Comparison**



#### **Identifying Interesting Local Phenomena**



#### **Identifying Interesting Local Phenomena**







- $\checkmark$  We are not here to tell you what to do !
- ✓ There are many reasons to select a site (important, co-location with chemistry, existing infrastructure, etc.)
- $\checkmark$  OK to NOT see any change (if robust data)
- ✓ This sub-module is about how to optimize your chances to see
  OA driven biological changes over a reasonable amount of time

# Factors modulating biological rate of change

✓ Biological sensitivity

 $\checkmark$  Chemical rate of change

## Biological sensitivity

ecology & evolution

PUBLISHED: 13 MARCH 2017 | VOLUME: 1 | ARTICLE NUMBER: 0084

#### Species-specific responses to ocean acidification should account for local adaptation and adaptive plasticity

Cristian A. Vargas<sup>1,2,3\*</sup>, Nelson A. Lagos<sup>3,4</sup>, Marco A. Lardies<sup>3,5</sup>, Cristian Duarte<sup>3,6</sup>, Patricio H. Manríquez<sup>7</sup>, Victor M. Aguilera<sup>2,8</sup>, Bernardo Broitman<sup>3,7</sup>, Steve Widdicombe<sup>9</sup> and Sam Dupont<sup>10</sup>

## Species sensitivity depends on local conditions (adaptation)

## Biological sensitivity



Species sensitivity depends on other environmental conditions (indirect effects, ecology, evolution, other drivers)

#### Where to monitor to see biological changes?



Chemical rate of change depends on where you are



Factors modulating biological rate of change





## How to estimate how long to monitor to see (robust) changes?

#### Use experimental data

### Example: Gullmarsfjord, Sweden



### Rate of chemical change



Marine Acidification On effects and monitoring of marine acidification in the seas surrounding Sweden

Editor Pin Andersson Co-authors Bertil Häkansson, Johan Häkansson, Elisabeth Sahisten Swedish Meteorological and Hydrological Institute Oceanographic Unit Jonathan Havenhand, Mike Thornyke, Sam Dupont Gothenburt University. Sven Lovice Contre for Manie Science



Kattegat surface water (0-25m)

-0.0044 pH unit / year

## Step 1: turn time into pH



## Step 1: turn time into pH



## Step 1: turn time into pH







## Biological sensitivity (e.g. blue mussels)



Limitations:

- Experimental design
- Adaptation / Acclimation
- Ecological interactions
- $\circ$  Modulating factors





Effect size (in this case, same thing)

Effect size (%) = 100 x  $e^{-e(5.837 \times (pH - 7.765))}$ 

















2040

2020

#### What can be expected



**Biological observation** (projected)

2080

2100

2060

Year



Not linear
 Need a wide range of pH
 to have the full curve



Not possible to calculate real non-linear rate of change with incomplete dataset

In that example, need 200 years of data

#### 10 years of data



Linear regression (R<sup>2</sup>=0.97)

Effect size = 0.0332 x Time - 67.043

Biological rate: 0.033 % / year

#### 40 years of data



Linear regression (R<sup>2</sup>=0.99)

Effect size = 0.4266 x Time - 871.03

Biological rate: 0.427 % / year

"Maximum" linear growth



Linear regression (R<sup>2</sup>=0.99)

Effect size = 0.9196 x Time – 1681

Max biological rate: 0.9196 % / year (linear)



Estimate the observed maximum rate of change after different duration of biological monitoring

## Rate of biological change vs duration of monitoring



- Reach saturation

## Need >80 years of data for a robust evaluation

Caution: Factors modulating biological rate of change

✓ Biological sensitivity

 $\checkmark$  Chemical rate of change



For this exercise we assumed that both were **constants** BUT both can vary over time

#### **Biological sensitivity**



# Higher the sensitivity = shorter the monitoring



The higher the species sensitivity, the faster you observe a robust rate of change

## Higher the sensitivity = shorter the monitoring



Robust data after:

# Faster the chemical rate = shorter the monitoring







IF the goal is to observe robust estimate of biological rate of change, prioritize locations with high biological sensitivity and high chemical rate of change





- $\checkmark$  We are not here to tell you what to do !
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- ✓ This sub-module is about how to optimize your chances to see
  OA driven biological changes over a reasonable amount of time

#### What specific Biological Indicators to measure?

- Insisting that everyone measures the same small sub-set of biological parameters is counterproductive.
- It restricts the scope of research, inhibits new discoveries and is not inclusive to all, in terms of either geography, local scientific interests or research capability.
- In the natural world it is unlikely that biological changes will be solely driven by OA so no biological measure is a perfect indicator of OA impact.
- Our approach would be to allow researchers to chose from a wide variety of parameters and methods, provided the chosen methodologies are robust and the data collected contribute to 1 of 5 overarching OA impact themes.
- Our 5 proposed impact themes focus on key biological processes for which we have some mechanistic understanding to suggest they are potentially sensitive to changes in carbonate chemistry.

#### **Calcifying Organisms and Calcification**

#### Rationale:

- $\circ$  Calcium carbonate (CaCO<sub>3</sub>) structures for structural support and protection is used by many marine organisms.
- Changes in seawater chemistry have been shown to strongly affect calcium carbonate structures and biomineralization rates.
- $\circ~$  Studies on calcifiers is perhaps the most mature area of OA impact research.

- Relative prevalence and success of calcifying organisms within an ecosystem. [Changes in biomass/abundance/percent cover of biocalcifying species, compared to non-calcifying species. Inorganic to organic biomass ratios (PIC:POC) of individual organisms, populations or whole assemblages.]
- **Calcified Biostructure Morphology.** [State, structure and mineral composition of calcified biostructures. Calcified structure morphology or function such as weight, density, damage or abnormality, dissolution severity, or strength.]
- **Rates of Calcification Calcifying.** [Measured rates of calcification or dissolution on organismal to ecosystem scales.]

#### Autotrophs and Primary Production (Photosynthesis)

#### Rationale:

- $\,\circ\,\,$  Primary production serves as the foundation for all marine ecosystems and CO\_2 is the fundamental fuel for photosynthesis .
- Yes, temperature and nutrients are very important, but increasing CO<sub>2</sub> can still modulate primary production via interactions with metabolically-costly carbon concentrating mechanisms as well as a broader suite of physiological and biochemical factors, including proton pumps, cellular membrane potential, enzyme activity, and energy partitioning.
- $\circ$  CO<sub>2</sub> and pH can have indirect effects, such as regulating flagellar motion (and thus cell motility and access to nutrients) and the chemical availability of essential micronutrients.

- **Biomass/Standing Stock**. [Total chlorophyll a concentration, phytoplankton cell abundance, microphytobenthos, macroalgae or seagrass biomass]
- **Productivity Rates.** [Carbon fixation rates, individual or community growth or production rates.]
- **Phenology.** [Onset or duration of blooms or other rapid growth periods.]

#### Heterotrophs and Secondary Production (Growth and Energetics)

#### Rationale:

- Long-term exposure to elevated pCO<sub>2</sub> levels increases metabolic energy demand in many marine organisms, to support increased acid-base balance, increased calcification costs, and increased physical activities.
- Additional energy expenditures leave less energy available to invest in other key processes, including protein synthesis and growth.
- For humans, reductions in marine secondary production may have profound implications for fisheries and aquaculture.

- **Biomass / Standing Stock.** [Changes in total heterotrophic community or individual abundance and biomass, body average size, percent cover.]
- **Productivity Rates.** [Rates of individual or community growth or secondary production, from *in-situ* techniques or algorithms.]
- **Phenology.** [Onset, appearance or duration of specific life stages (e.g. time spent in plankton, inter-moult times.]

#### **Biodiversity and Community Structure (Mortality)**

#### Rationale:

- $\circ~$  Ocean Acidification has been seen to cause mortality in many species.
- Sensitivity of marine organisms varies enormously between species allowing OA to act directly as a strong selective pressure, decreasing biodiversity directly through species loss and altering community structure.
- Or indirectly through altered competition, trophic interactions, decreasing energy generation and flows, degrading critical biogenic habitats, or increasing susceptibility to pathogens and diseases.

- **Taxonomic Diversity / Community Composition.** [Identity and quantity (number or biomass) of the species/taxa/groups present within a community or assemblage at any given time. Used to calculate a wide variety of indices, each focussing on specific aspects of community structure and biodiversity.]
- Functional or Trait Diversity. [Requires the collection of species abundance/biomass data as described above, but now the individual organisms are grouped, not by their taxonomic relationships, but by their shared functional, ecological or behavioural traits. The same biodiversity metrics as used for describing taxonomic diversity can then also be applied to these aggregated data to generate estimates of functional or trait diversity.]

#### **Genetic Changes**

#### Rationale:

- Ocean Acidification drives phenotypic and genetic changes (natural selection)
- This can be specific (OA signature)
- Genetic signal dependent on diversity, strength and direction of the pressure

- Neutral genetic variation. [= non-functional genetic variation. Population genetic parameters (e.g. number of alleles, heterozygosity, effective population size, inbreeding and population divergence) using classic molecular markers (e.g., allozymes, microsatellites or mtDNA) or through high-throughput sequencing approaches.]
- **Functional genetic variation.** [= functional genetic variation. Using quantitative trait locus (QTL), genome-wide association studies (GWAS), genome scan via restriction-site-associated DNA taqs (RAD-seq), and RNA sequencing (RNA-seq).]
- *Mutation rates.* [for few loci (including genes of particular interest) or from whole-genomes using high-throughput sequencing technologies.]