# **MarineSDMs**

# Marine Species Distribution Models

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# 1 Introduction

```
# remotes::install_github("debruine/glossary")
library(glossary)
glossary::glossary_path("glossary.yml")
glossary::glossary_popup("click")
glossary::glossary_persistent(TRUE)
# quarto publish gh-pages
```

## 1.1 Background

The best available global distributions are presently AquaMaps (Kaschner et al. 2006; Ready et al. 2010) with supplementation by IUCN RedList range maps<sup>1</sup>. We previously used this to calculate the Biodiversity goal of the Ocean Health Index across all Exclusive Economic Zones (EEZs) (Halpern et al. 2012) and beyond EEZs in the high seas (Visalli et al. 2020) with taxonomic groupings [based on (Tittensor et al. 2010): see gmbi (global marine biodiversity indicators)].

## 1.2 Goals

This book is meant to capture the overview and details of modeling species distributions in the marine environment for the purposes of advancing the status quo of global and U.S. national species distributions along the following dimensions:

#### 1. Space

The current AquaMaps distributions are  $1/2^{\circ}$  (~55 km at equator), whereas the best available global bathymetry is  $1/240^{\circ}$  (< 0.5 km).

#### 2. Time

The current AquaMaps distributions are based on static climatic averages over all seasons, which does not capture temporal dynamics: seasonally within a year, nor long-term climate change trends. This will necessitate sampling the environment contemporaneously

<sup>&</sup>lt;sup>1</sup>IUCN RedList range maps: https://www.iucnredlist.org/resources/spatial-data-download

with species observations before fitting the model and predicting to different environmental snapshots.

#### 3. Environment

Other environmental variables besides the initial physiographic (depth) and oceanographic (temperature, chlorophyll, primary productivity and ice) may elicit an improved statistical fit, related to species' environmental niche. Some candidates include: temperature fronts, eddy kinetic energy, distance from shore, distance from shelf.

#### 4. Biology

Where sufficient observations exist, additional models should be developed highlighting differences between:

- Life stage, e.g. larval vs adult.
- Gender where varies, such as male sperm whales being more cosmopolitan.
- Subpopulations for understanding metapopulation dynamics
- Behavior, such as migrating, feeding or breeding.

By definition MBONMarine Biodiversity Observation Network; see MarineBON.org is a network, so this is inclusive of and meant for all participants.

## 1.3 Inspirations

• AquaMaps.org

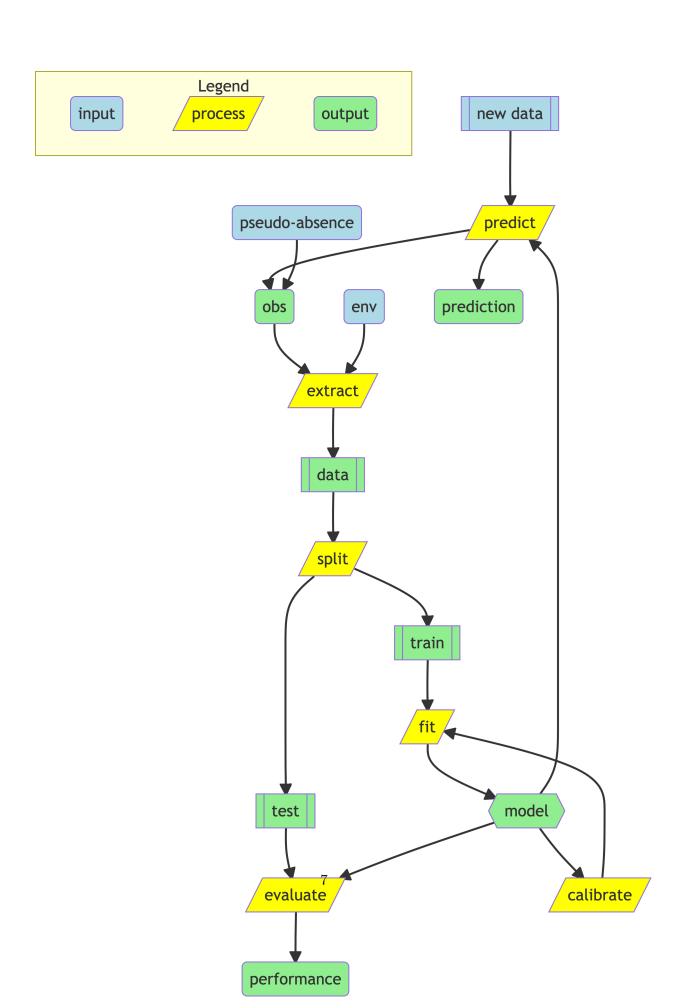
Lots of work. To be continued

• R package dismo: sdm.pdf -> raster sdm -> terra sdm

•

# Part I

# **Process**



# 2 Prep Data

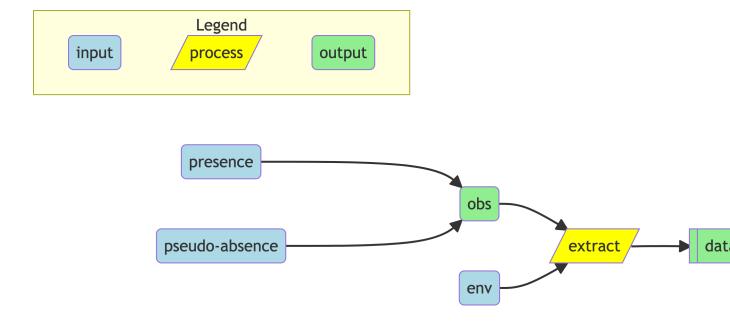


Figure 2.1: Diagram of SDM data preparation for model fitting.

#### • obs

observations: occurrences from OBIS; masked by FAO regions defined by AquaMaps (Skyttner 2020)

- presence
  - OBIS: species occurrence
- absence
  - OBIS not-species, but same family
- env
  - environment
- tbl

table of observations (presence and absence) with environmental values

## 2.0.1 Environmental Predictors

## 2.0.2 Physiographic

- depth Bathymetric Depth
- d2coast Distance to Coast
- d2shelf Distance to Shelf

## 2.0.3 Time Varying

• vgpm Vertically integrated primary Productivity model

## 2.0.3.1 Depth & Time Varying

- Temperature
- Salinity

# 3 Model

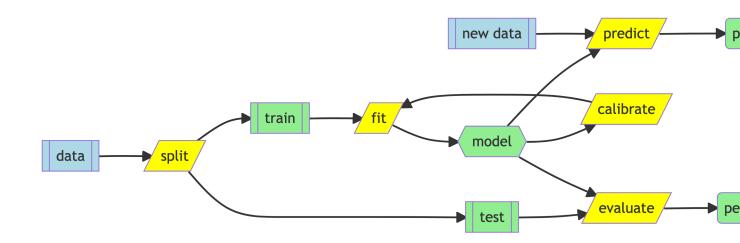


Figure 3.1: Diagram of SDM Modeling processes.

# 4 Mosaic

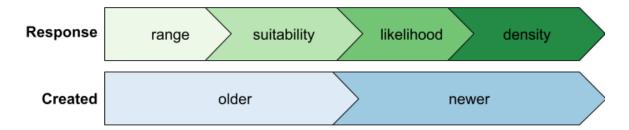


Figure 4.1: Hierarchy of preferred model outputs based on response type and age.

This is illustrated well by Figure 4.1.

# **5** Software

## 5.1 R

Most packages have not yet migrated from using the deprecated raster R package to the new terra package, except for biomod2 (ref?).

#### • eks

Tidy and Geospatial Kernel Smoothing for spatially filtering outlier observations

• test

## 5.2 Python

#### • Xarray

Xarray makes working with labelled multi-dimensional arrays in Python simple, efficient, and fun!

# 5.3 Google Earth Engine

#### • XEE

XEE is a new Python package for Earth Engine that provides a set of functions to facilitate the use of Earth Engine API. It is designed to be used in Jupyter notebooks and Google Colab. Documentation



Figure 5.1: Source: Kernel density estimates for tidy and geospatial data in the eks package

# 6 Organize

### 6.1 Partners

- Confirmed
  - AquaMaps

 $distribution \ maps \ for \ over \ 33,500 \ species \ of \ fishes, \ marine \ mammals \ and \ invertebrates$ 

contacts:

\* Gabriel Reygondeau (g.reygondeau@oceans.ubc.ca)

#### - DisMAP

Distribution Mapping and Analysis Portal contacts:

- \* Melissa Karp (melissa.karp@noaa.gov)
- \* Roger Griffis (roger.b.griffis@noaa.gov)

#### - MBON

 $\begin{tabular}{ll} \it Marine Biodiversity Observation Network \\ \it contacts: \end{tabular}$ 

- \* Ben Best
- \* Tylar Murray
- \* Dan Otis
- \* Frank Muller-Karger

#### - OBIS

Ocean Biodiversity Information System contacts:

- \* Candido Principe De Souza, Silas (s.principe@unesco.org)
- \* Ward Appeltans (w.appeltans@unesco.org)
- Proposed

#### - FishGLOB

fish biodiversity under global change (global bottom trawl) contacts:

\* Malin Pinsky

#### - MGEL

Marine Geospatial Ecology Lab, Duke contacts:

- \* Jason Roberts
- \* Patrick Halpin

#### - PMEL

 $Pacific\ Marine\ Environmental\ Lab,\ NOAA$  contacts:

- \* Heather Welch
- \* Elliot Hazen

## 6.2 Portal

• Candidate website:

MarineSpeciesMaps.org (BDB registered domain)

- similar to:
  - \* MarineRegions.org spatial authority
  - \* MarineSpecies.org taxonomic authority

## 6.3 Timeline

### 6.3.1 Meetings

2024-01-23

Marine Biodiversity Summit in DC (organized by Gabrielle, Emmett)

• 2024-03-21

SDM meeting in New Orleans (organized by BDB; so far AquaMaps)

# 6.3.2 Sequence

1.

# References

- Halpern, Benjamin S., Catherine Longo, Darren Hardy, Karen L. McLeod, Jameal F. Samhouri, Steven K. Katona, Kristin Kleisner, et al. 2012. "An Index to Assess the Health and Benefits of the Global Ocean." *Nature*. https://doi.org/10.1038/nature11397.
- Kaschner, K., R. Watson, A. W. Trites, and D. Pauly. 2006. "Mapping World-Wide Distributions of Marine Mammal Species Using a Relative Environmental Suitability (RES) Model." *Marine Ecology Progress Series* 316 (July): 285–310. https://doi.org/10.3354/meps316285.
- Ready, Jonathan, Kristin Kaschner, Andy B. South, Paul D. Eastwood, Tony Rees, Josephine Rius, Eli Agbayani, Sven Kullander, and Rainer Froese. 2010. "Predicting the Distributions of Marine Organisms at the Global Scale." *Ecological Modelling* 221 (3): 467–78. https://doi.org/10.1016/j.ecolmodel.2009.10.025.
- Skyttner, Markus. 2020. "Aquamapsdata: Curated Data from AquaMaps.org." https://github.com/raquamaps/aquamapsdata.
- Tittensor, Derek P., Camilo Mora, Walter Jetz, Heike K. Lotze, Daniel Ricard, Edward Vanden Berghe, and Boris Worm. 2010. "Global Patterns and Predictors of Marine Biodiversity Across Taxa." *Nature* 466 (7310): 1098–1101. https://doi.org/10.1038/nature09329.
- Visalli, Morgan E., Benjamin D. Best, Reniel B. Cabral, William W. L. Cheung, Nichola A. Clark, Cristina Garilao, Kristin Kaschner, et al. 2020. "Data-Driven Approach for Highlighting Priority Areas for Protection in Marine Areas Beyond National Jurisdiction." Marine Policy, March, 103927. https://doi.org/10.1016/j.marpol.2020.103927.

# **A** Glossary

```
glossary::glossary_table(as_kable=F) |>
knitr::kable("pipe", escape = F, row.names = F)
```

# Part II Explorations

# **AquaMaps Downscaled**

