

# Species distribution models (SDM)

AZTI

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# About

This is a short tutorial for constructing species distribution models in R. It describes the whole process from downloading OBIS and GBIF data, generating pseudo-absence data, including environmental data, fitting the model, validating the model and generating the resulting maps for visualization.

The code is available in AZTI's github repository repository and the book is readily available here.



# Chapter 1

## Introduction

Some introduction about SDMs





## Chapter 2

# Libraries

Load libraries that will be used



## Chapter 3

# Presence-absence data

Bla bla bla

### 3.1 Download presence data

Download from GBIF OBIS. Mireia

### 3.2 Create pseudo-absence data

Prevalence 50%

See code from ANICHO (maintaining some space around presences). Leire C.



## Chapter 4

# Environmental data

Bla bla bla

### 4.1 Download from public repositories

Download from Bio-oracle. Guillem le pasa el código a Mireia, que lo sube a github

### 4.2 Operations with rasters (maybe not needed)

We can complete this a bit more later on, though not necessary right now



## Chapter 5

# Prepare final dataset

Bla bla bla

**5.1** Extract environmental data associated to presence-absence data

**5.2** Exploratory plots





## Chapter 6

# Shape Constrained-Generalized Additive Models

One citation is [Citores et al., 2020]

Mention there is an alternative using `mboost` that won't be further developed here.

### 6.1 Model fit

### 6.2 Model selection



## Chapter 7

# Model selection

Bla bla



## Chapter 8

# Model validation

Bla bla

### 8.1 Optimum threshold

### 8.2 k-fold validation



## Chapter 9

# Prediction and maps

predict from fitted models and produce maps





# Bibliography

L. Citores, L. Ibaibarriaga, D. J. Lee, M. J. Brewer, M. Santos, and G. Chust. Modelling species presence–absence in the ecological niche theory framework using shape-constrained generalized additive models. *Ecological Modelling*, 418:108926, 2020.