



# Relações entre espécies e características ambientais

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**Christian Ricardo Ribeiro**

**Vitor Vieira Vasconcelos**

Universidade Federal do ABC

Março, 2024

São Bernardo do Campo - SP

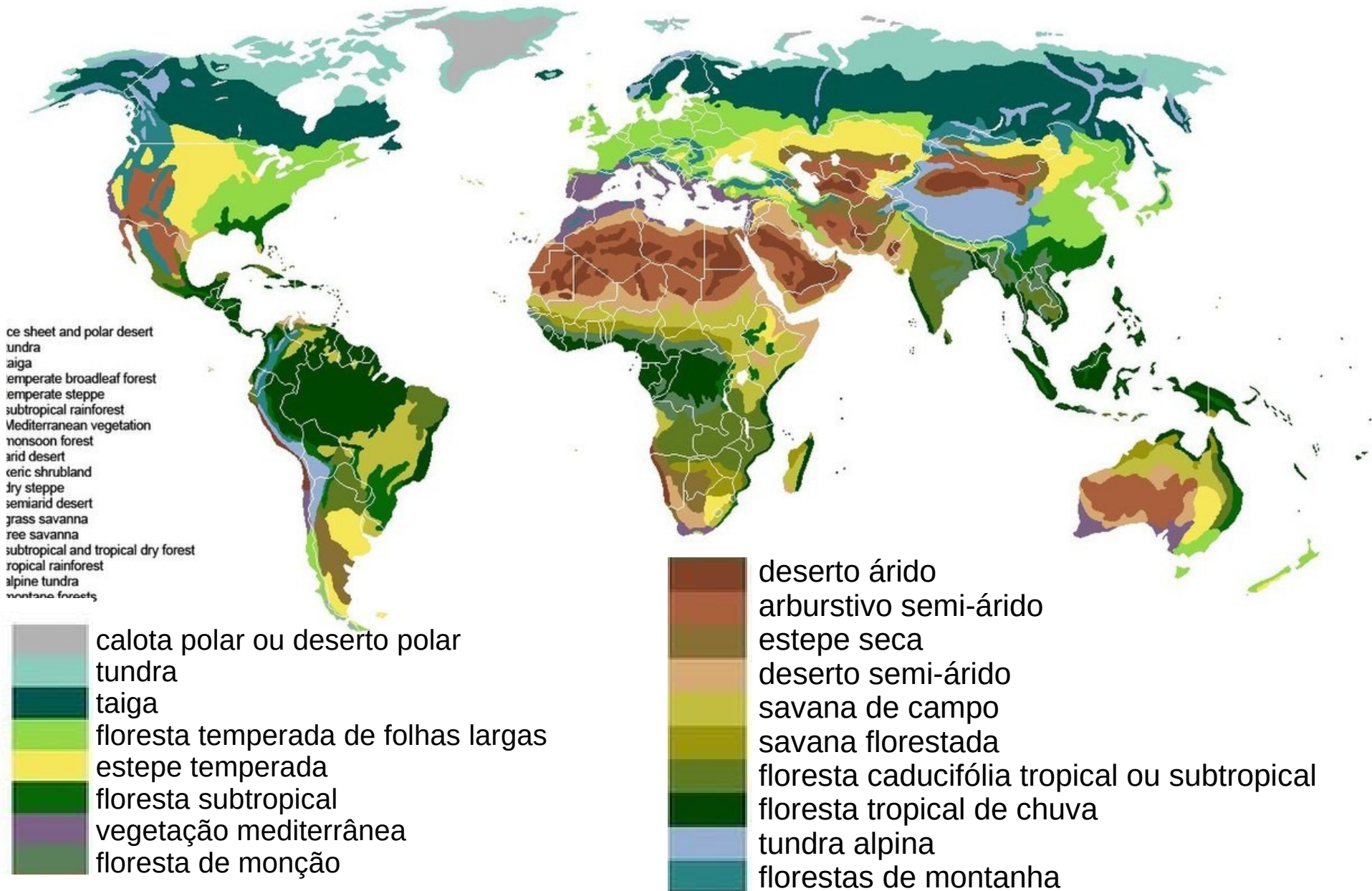
# Objetivos

- Entender as relações entre as distribuições de espécies (biodiversidade) com as características dos ambientes
- Compreender os métodos e técnicas para modelagem de distribuição de espécies

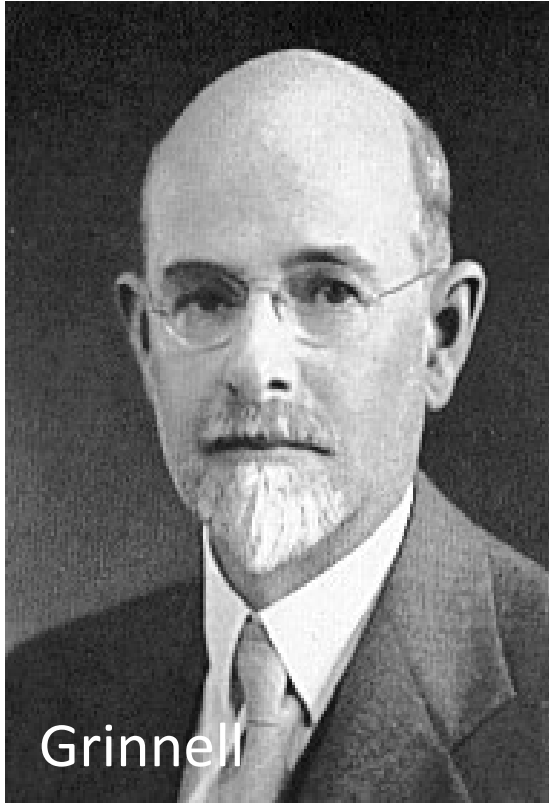
# Por que entender as relações entre as espécies e as características do meio onde vivem?

- Entendimento do funcionamento dos ecossistemas
  - Incluindo agro e socio-ecossistemas.
- Aptidão agrícola e pecuária.
- Espalhamento pragas, de vetores (animais portadores) de doenças e animais exóticos invasores.
- Impactos das mudanças climáticas na distribuição de espécies nativas, cultivadas e invasoras.
- Planejamento de medidas de restauração de ambientes naturais ou manejados.

# Biomas



# Conceito de nicho



Grinnell (1917, 1924): distribuição geográfica dos indivíduos de uma espécie como resposta as variáveis ambientais (temperatura, precipitação, elevação etc).

Elton (1927): nicho como resposta da interação entre espécies

Hutchinson (1957): *nicho* é a soma de todos os fatores ambientais que agem em um determinado organismo, definido como uma região de  $n$  dimensões no hiper-espço.

# Nichos como limites e envelopes

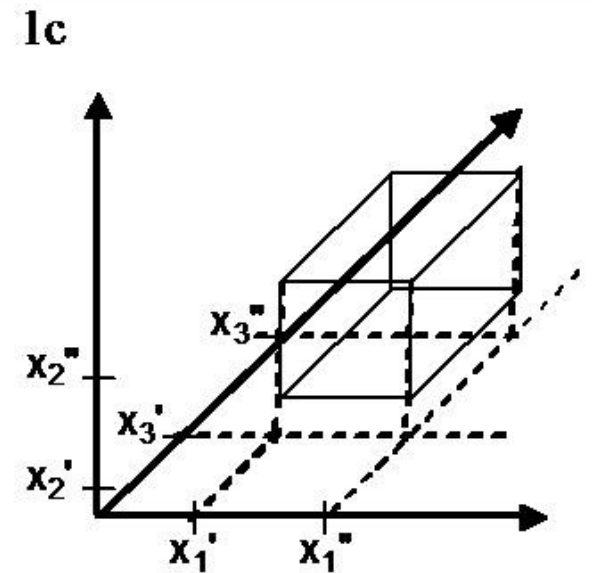
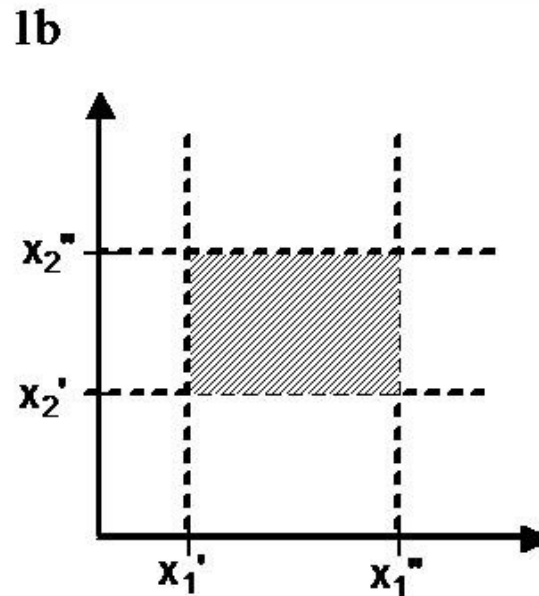
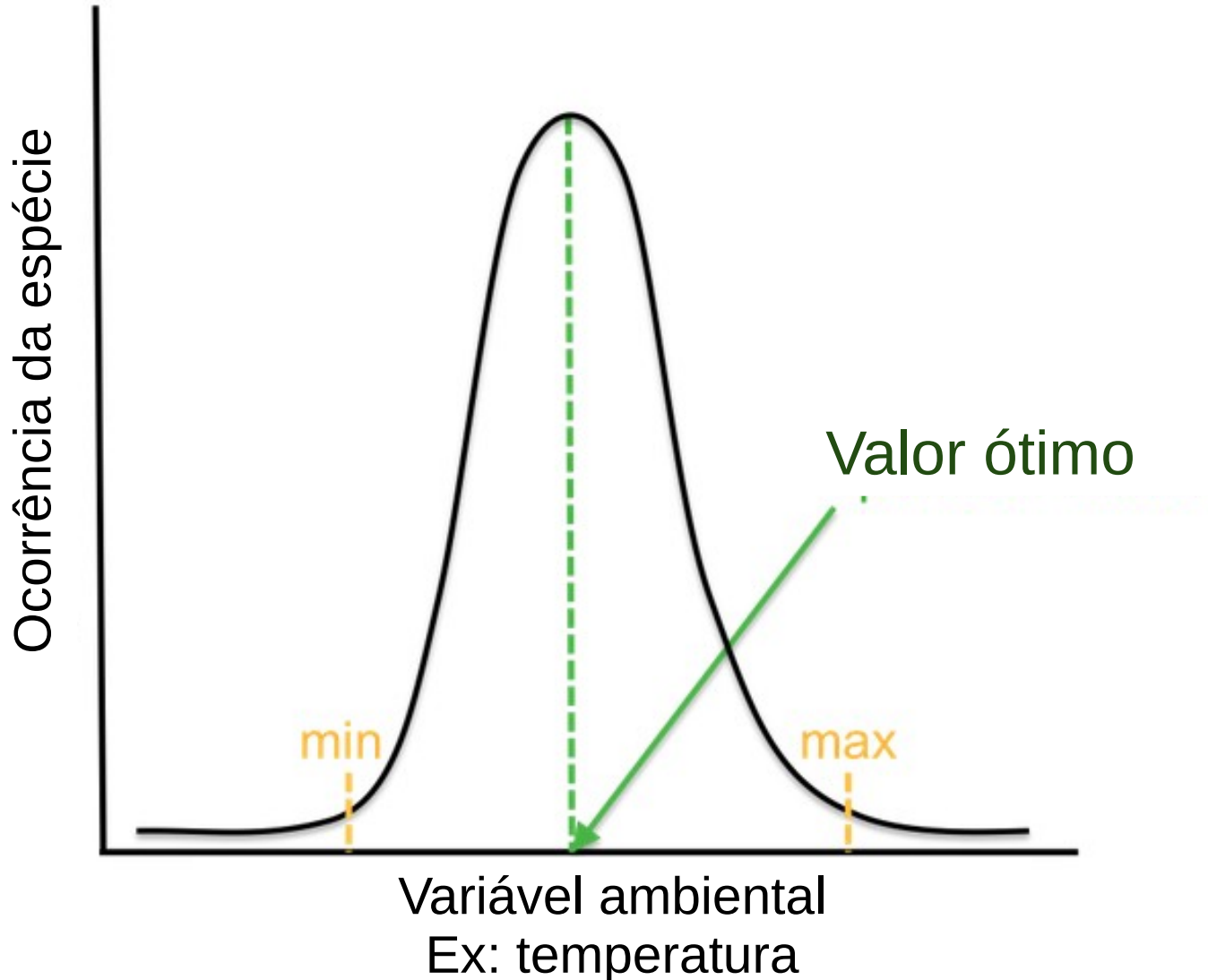
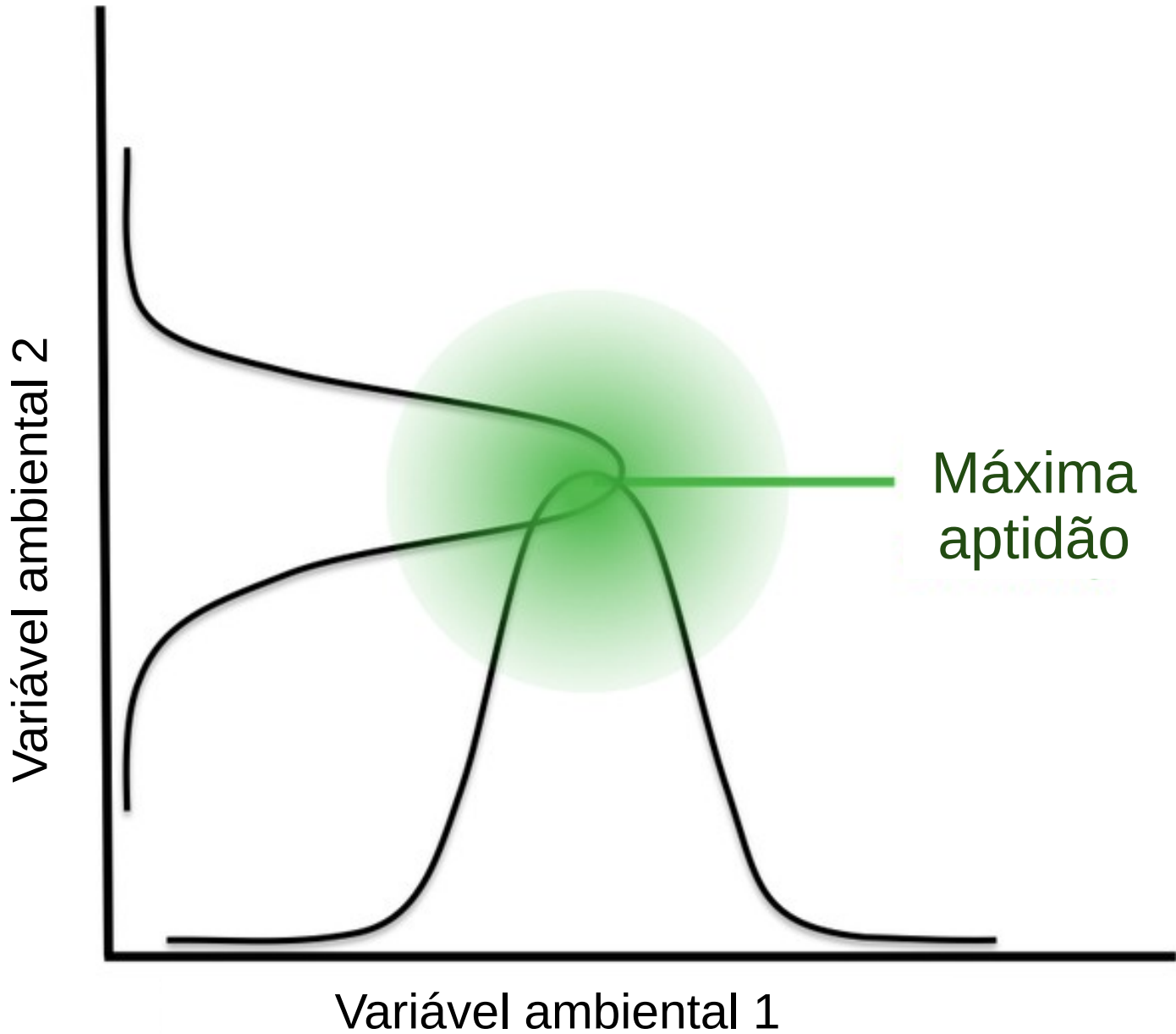


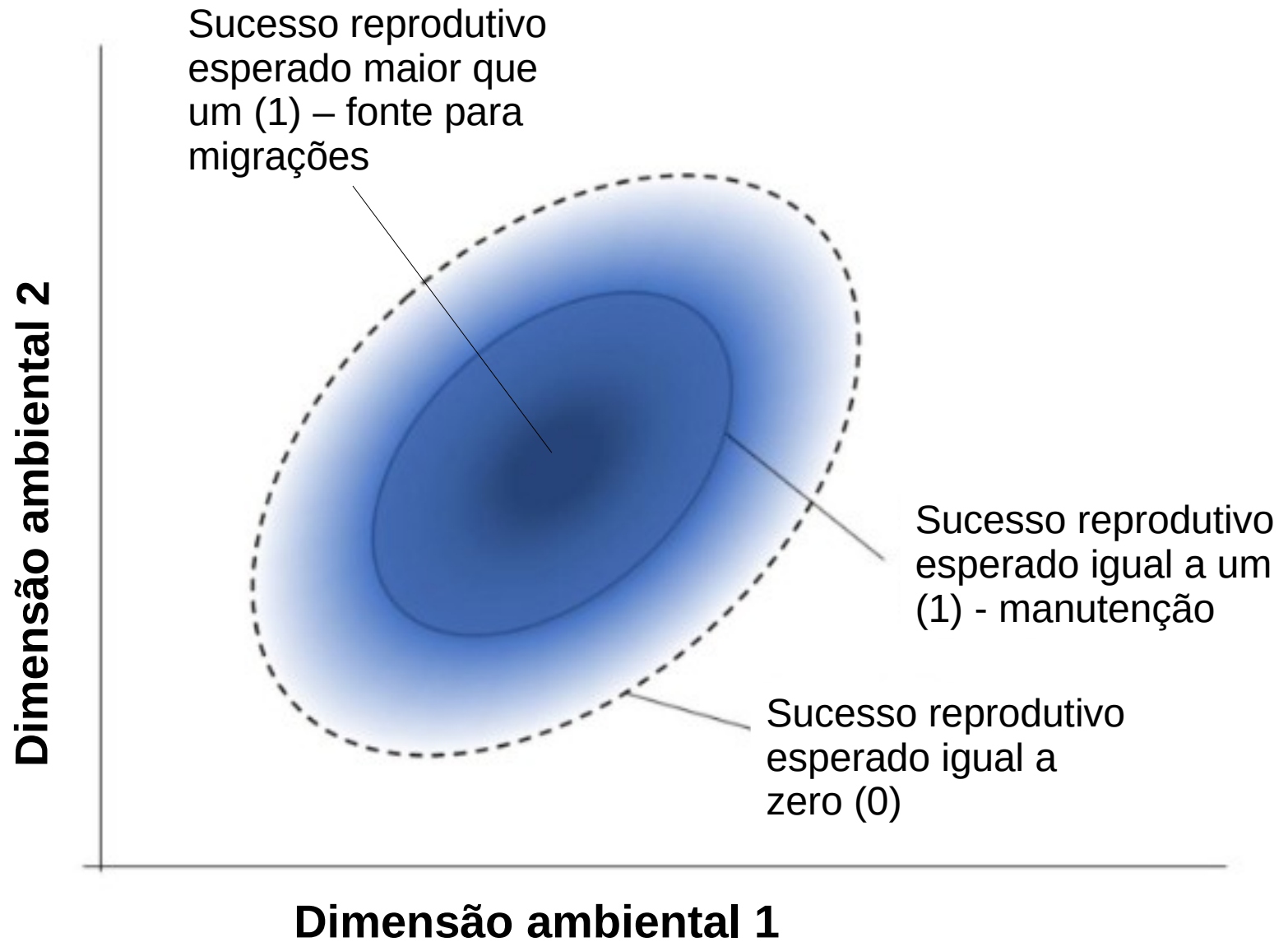
Figura 1: Esquema da definição de nicho ecológico proposta por Hutchinson (1957), para uma (1a), duas (1b) e três (1c) dimensões (variáveis).

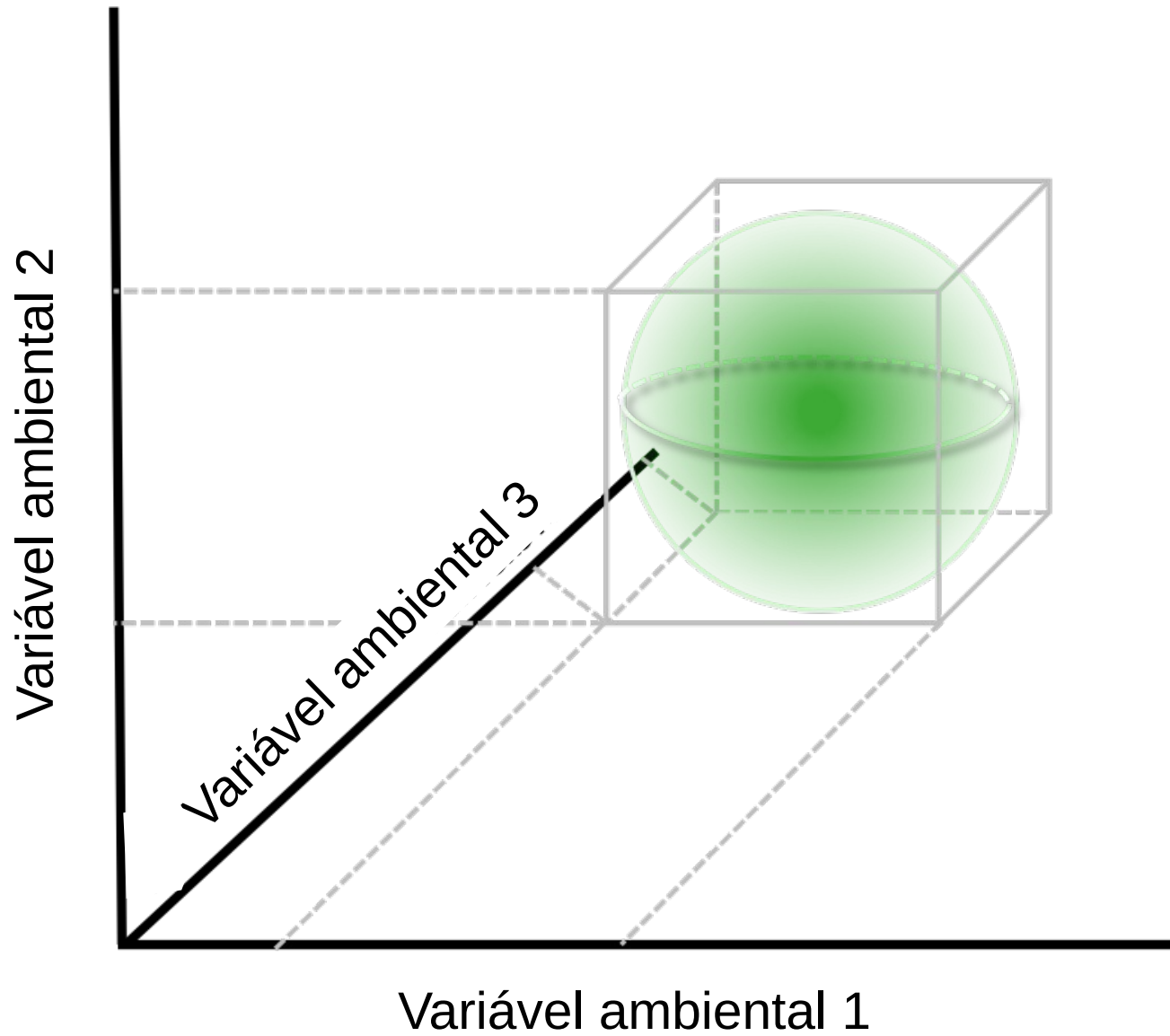
# Nichos como graduação de aptidão





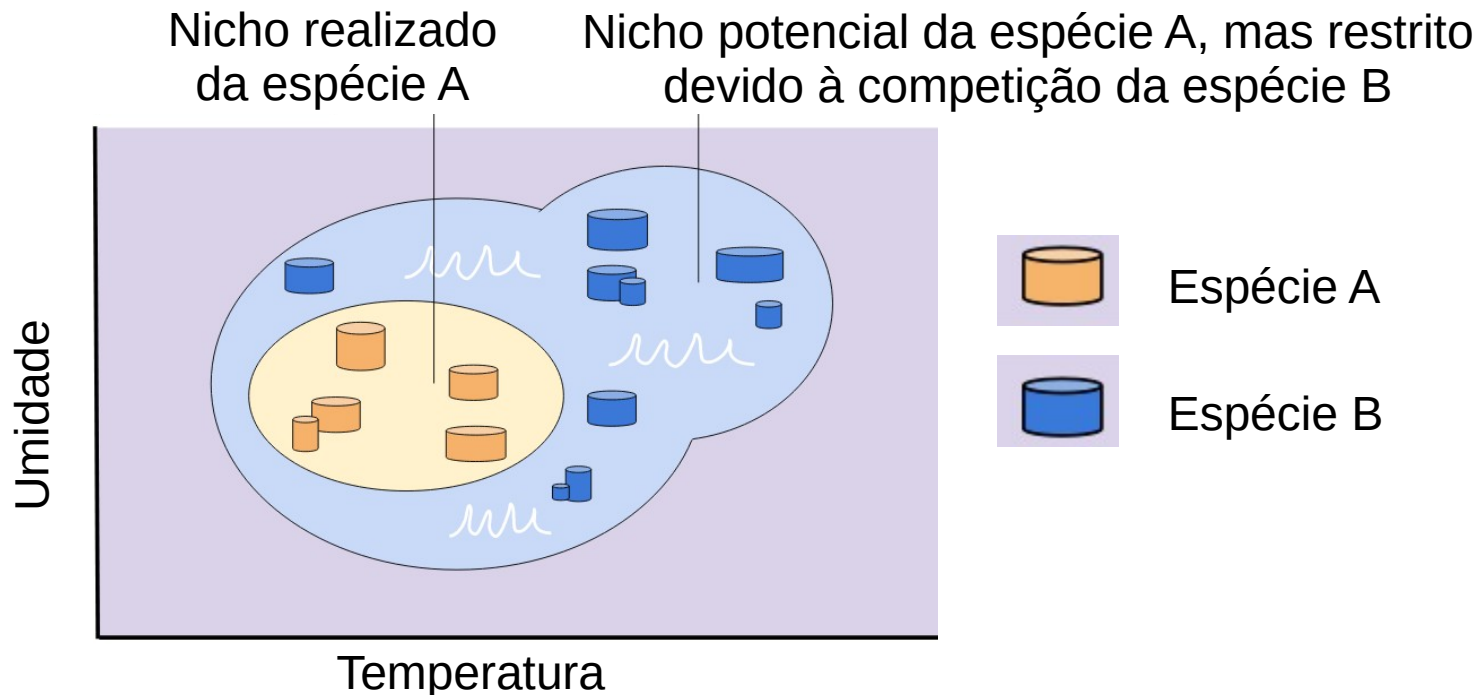


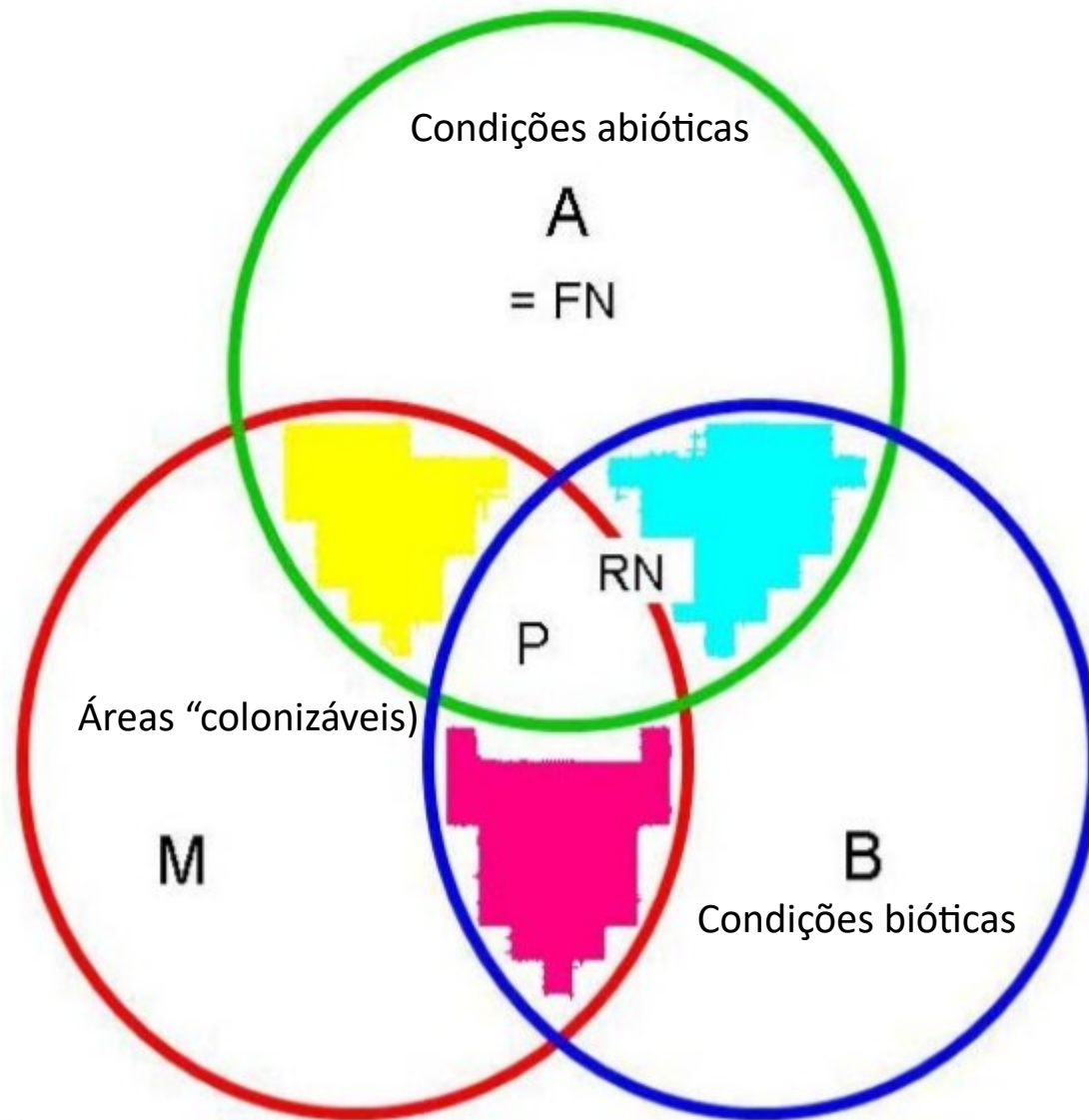




# Nicho fundamental e nicho realizado

O **nicho fundamental** é representado pelas as condições ambientais nas quais as espécies poderiam viver, se não fossem as restrições impostas pelas relações bióticas, que define o **nicho realizado** (Hutchinson 1957).





E a ação humana, como influenciaria nesses processos?

SOBERON, Jorge; PETERSON, A. Townsend. Interpretation of models of fundamental ecological niches and species' distributional areas. Biodiversity Informatics, 2, 2005, pp. 1-10 .

P = distribuição geográfica (potencial) da espécie

# Fitofisionomias por altitude - Humboldt (1806)

Que características ambientais mudam, conforme se sobe a montanha?

Neve

Rochoso

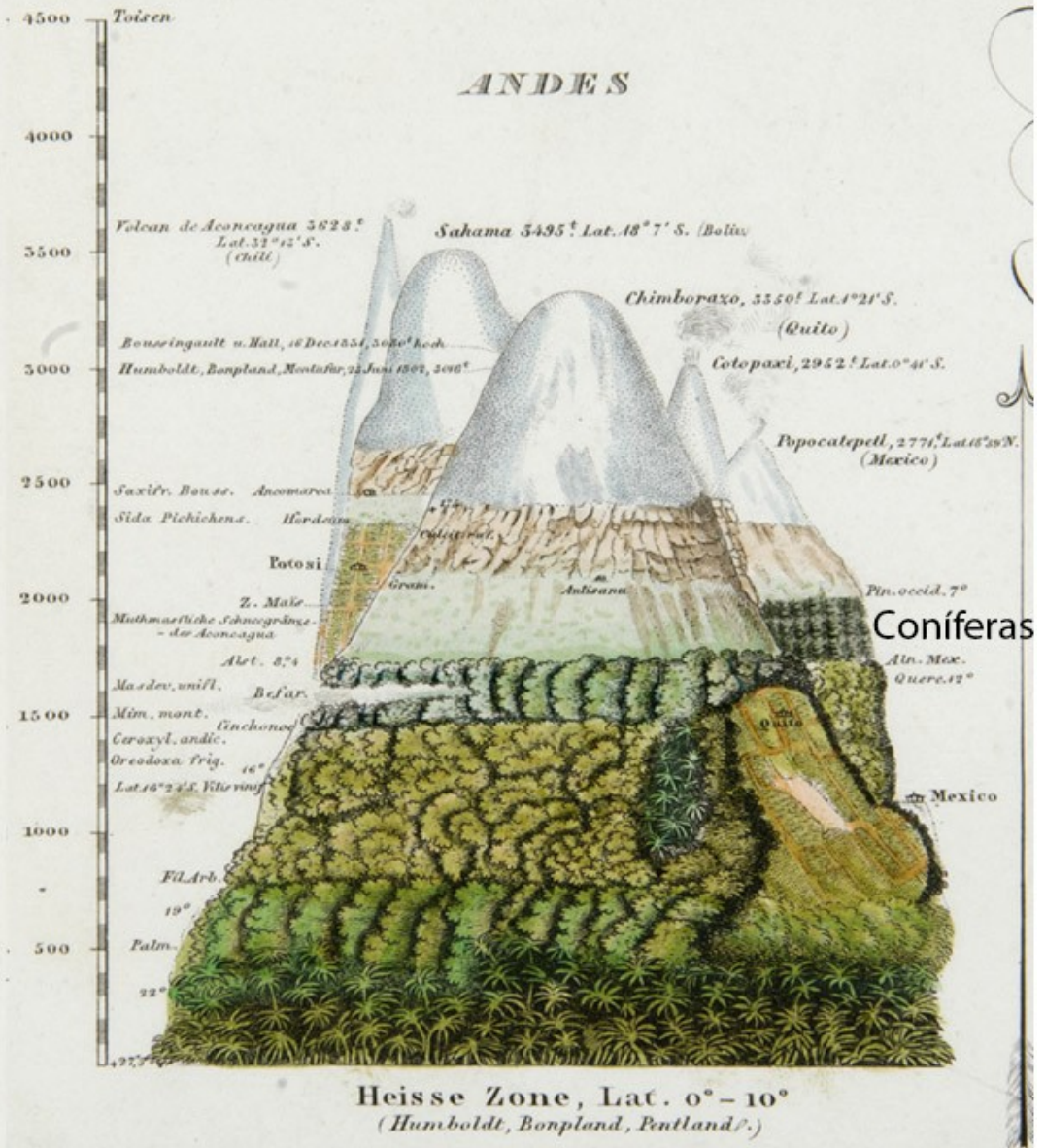
Tundra

Floresta Nebular Altomontana

Floresta Tropical Montana

Floresta Tropical Sub-Montana

Floresta Tropical de Terras Baixas



## Ecosistemas



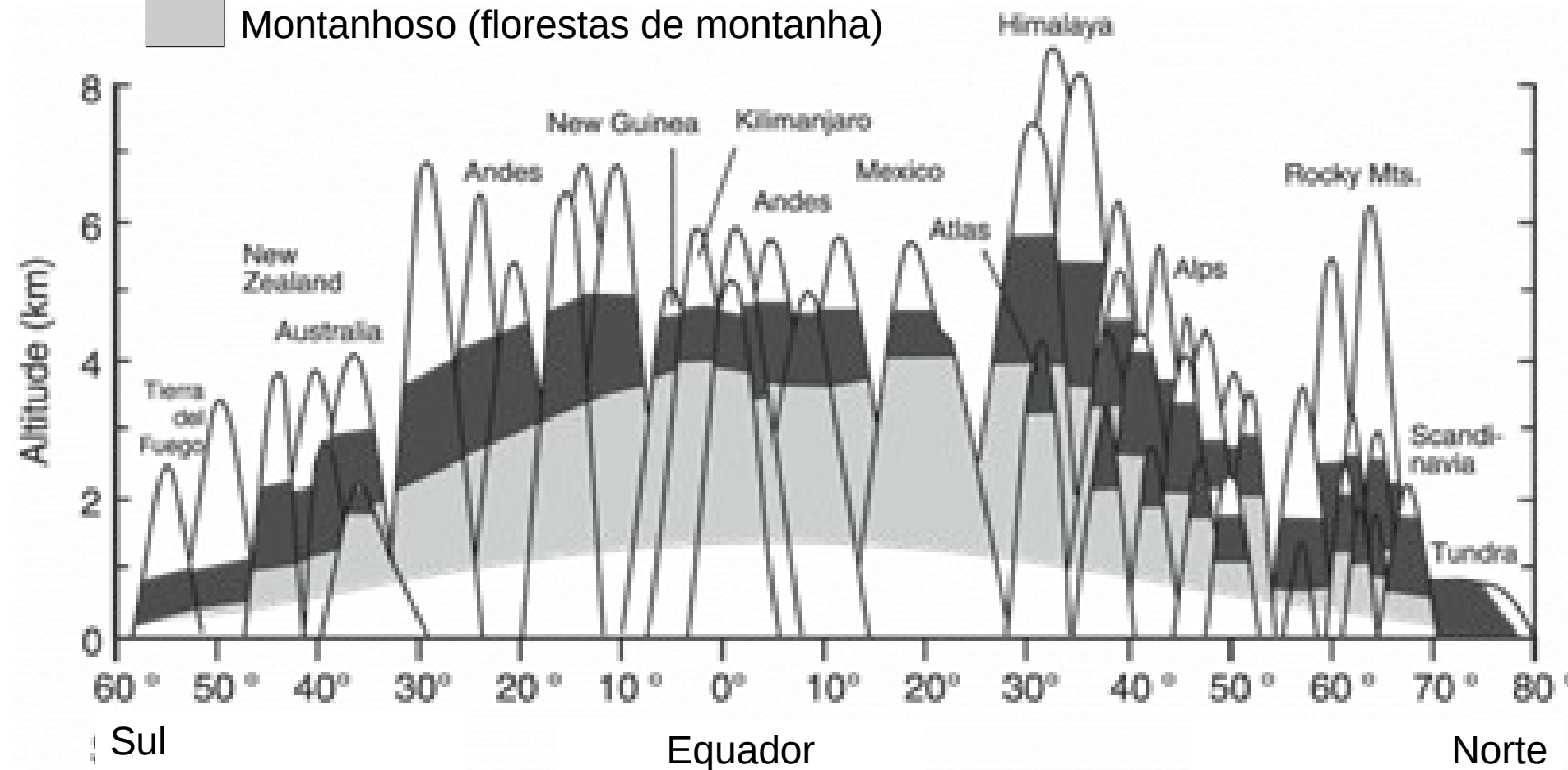
Nival (neve)



Alpino (arbustos)

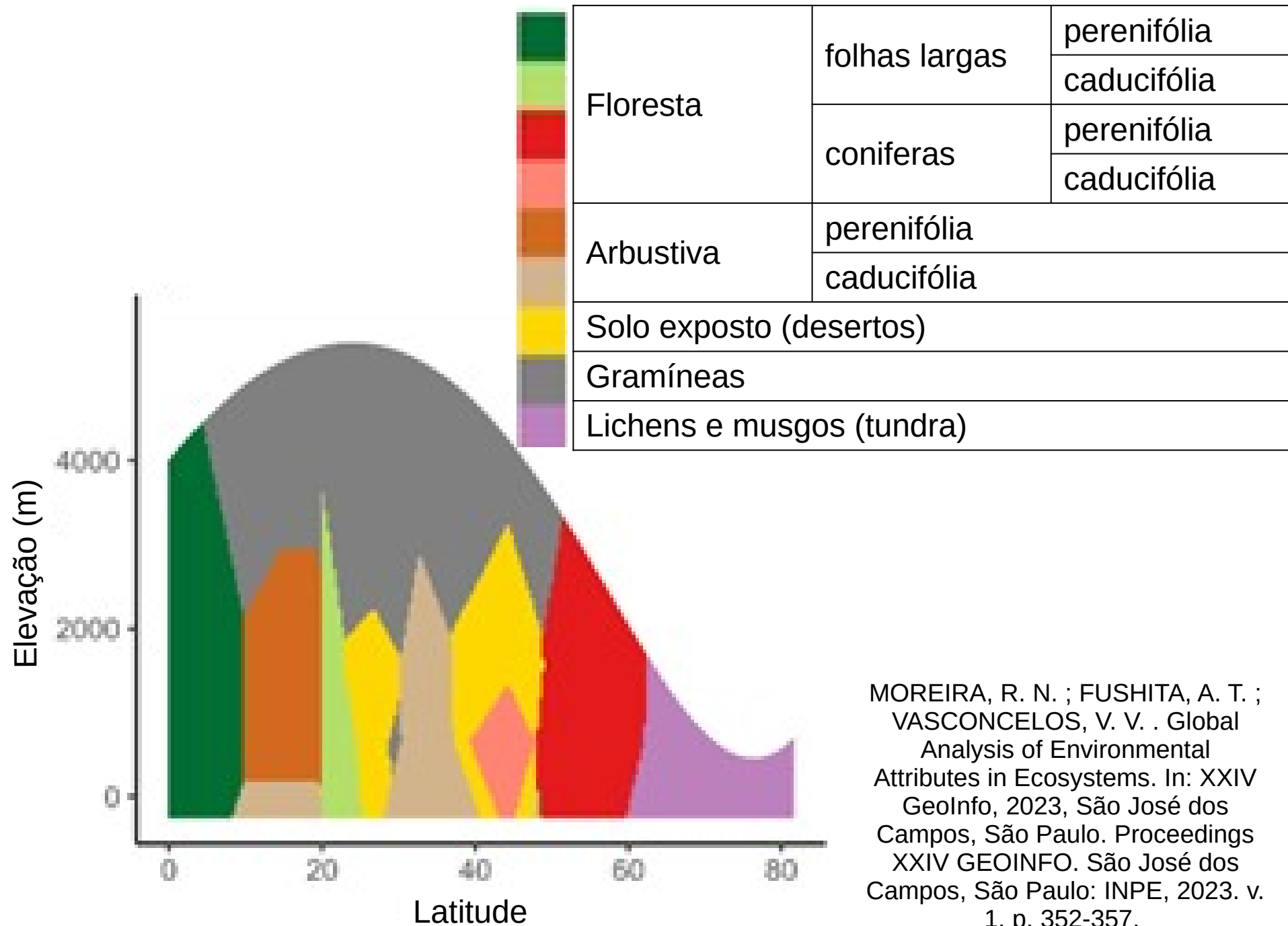


Montanhoso (florestas de montanha)

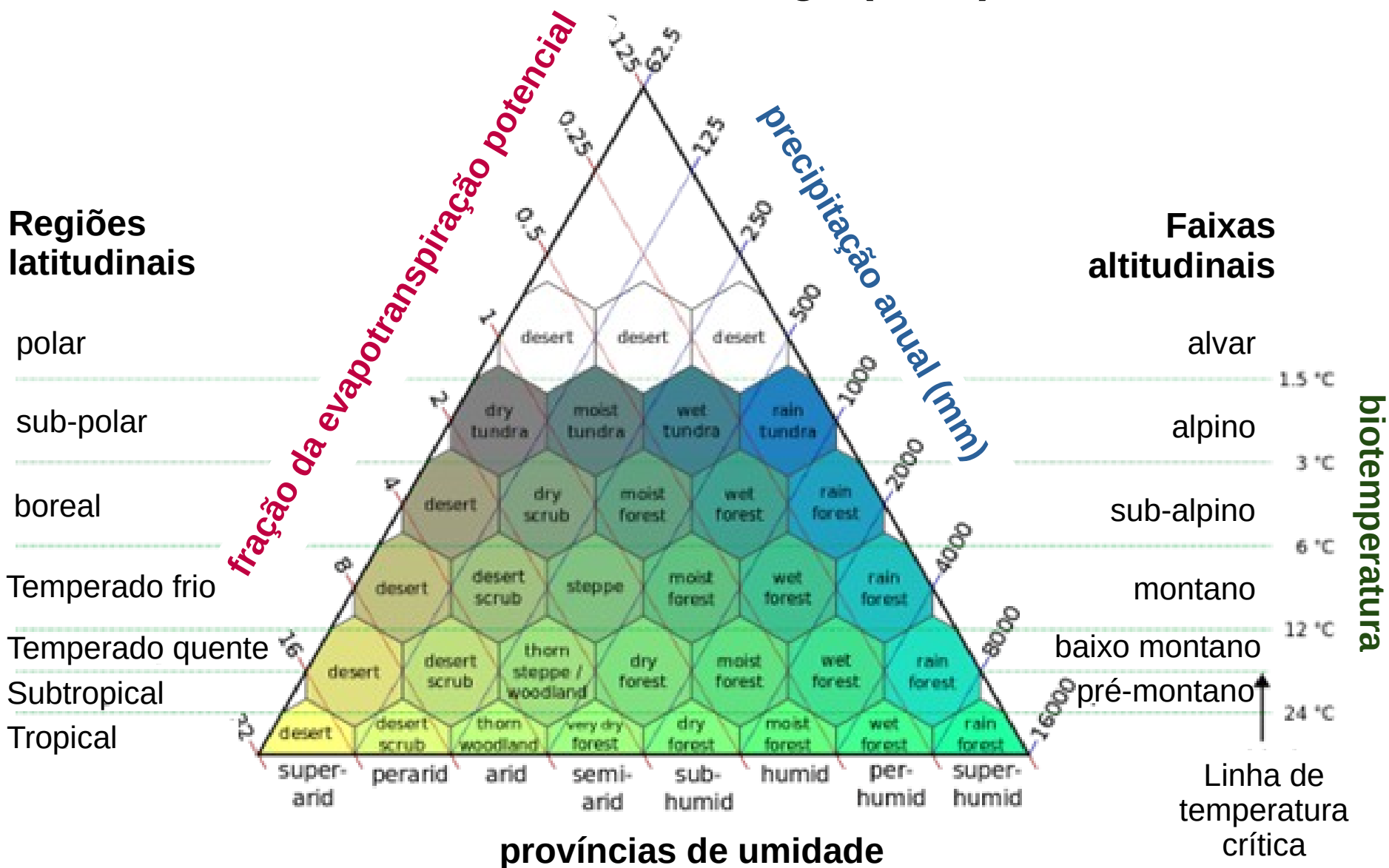


Humboldt, A. von. (1806). Ideen zu einer Physiognomik der Gewächse. Jenaischen Allgemeinen Literatur- Zeitung, 1(62), 489-492. Available in: <<http://www.avhumboldt.de/?p=4100>>

Körner, C. (2003). Alpine Plant Life, 2nd edn. Springer, Berlin.



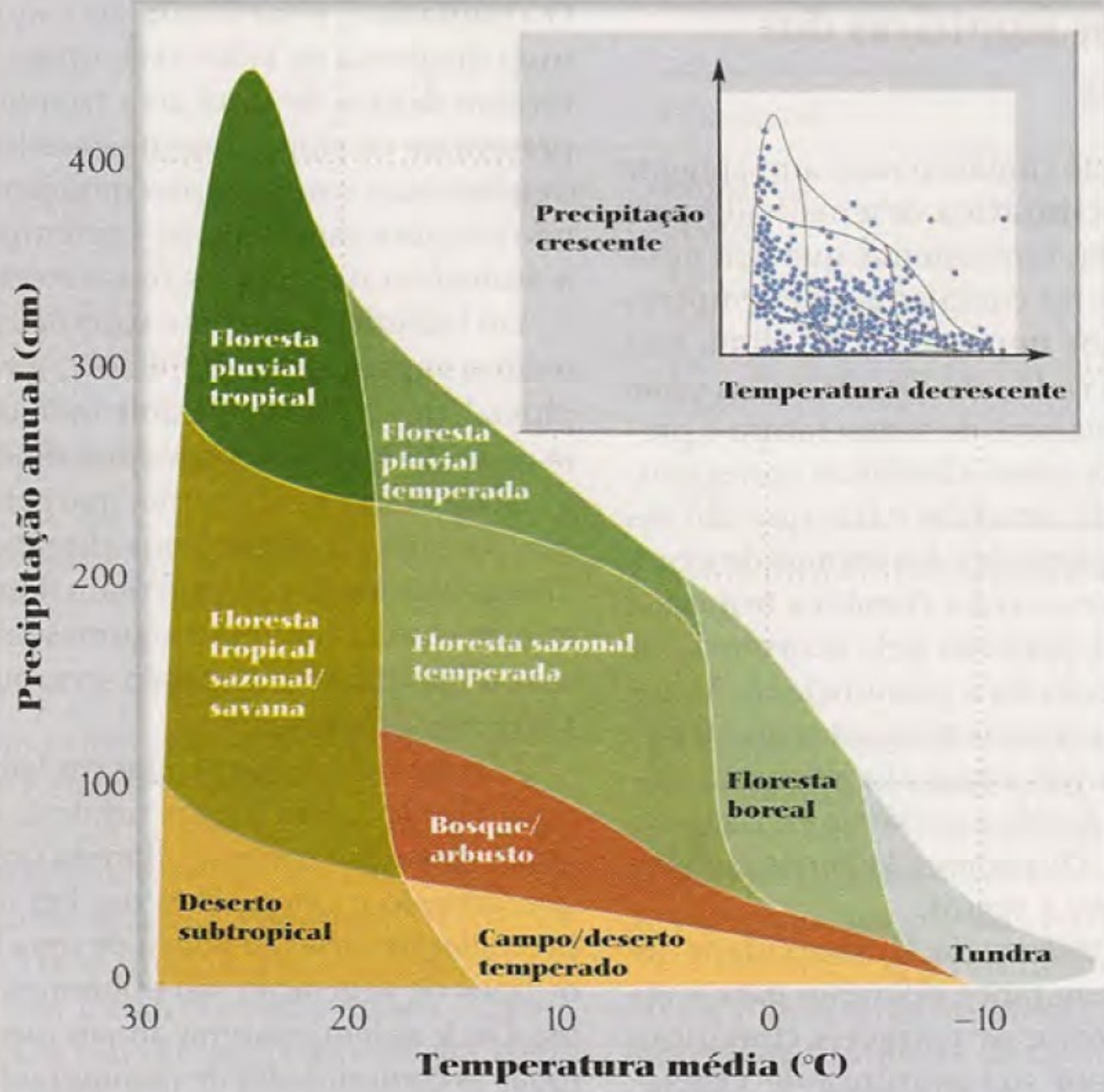
# Zonas de Vida - Holdridge (1947)



Holdridge, L.R. (1947). Determination of world plant formations from simple climate data. *Science*, 105, 367–368.  
 Halasz, P. (2007). Holdridge Life Zone Classification scheme. Available at:  
[https://commons.wikimedia.org/wiki/File:Lifozones\\_Pengo.svg](https://commons.wikimedia.org/wiki/File:Lifozones_Pengo.svg), accessed on 11/4/2018.



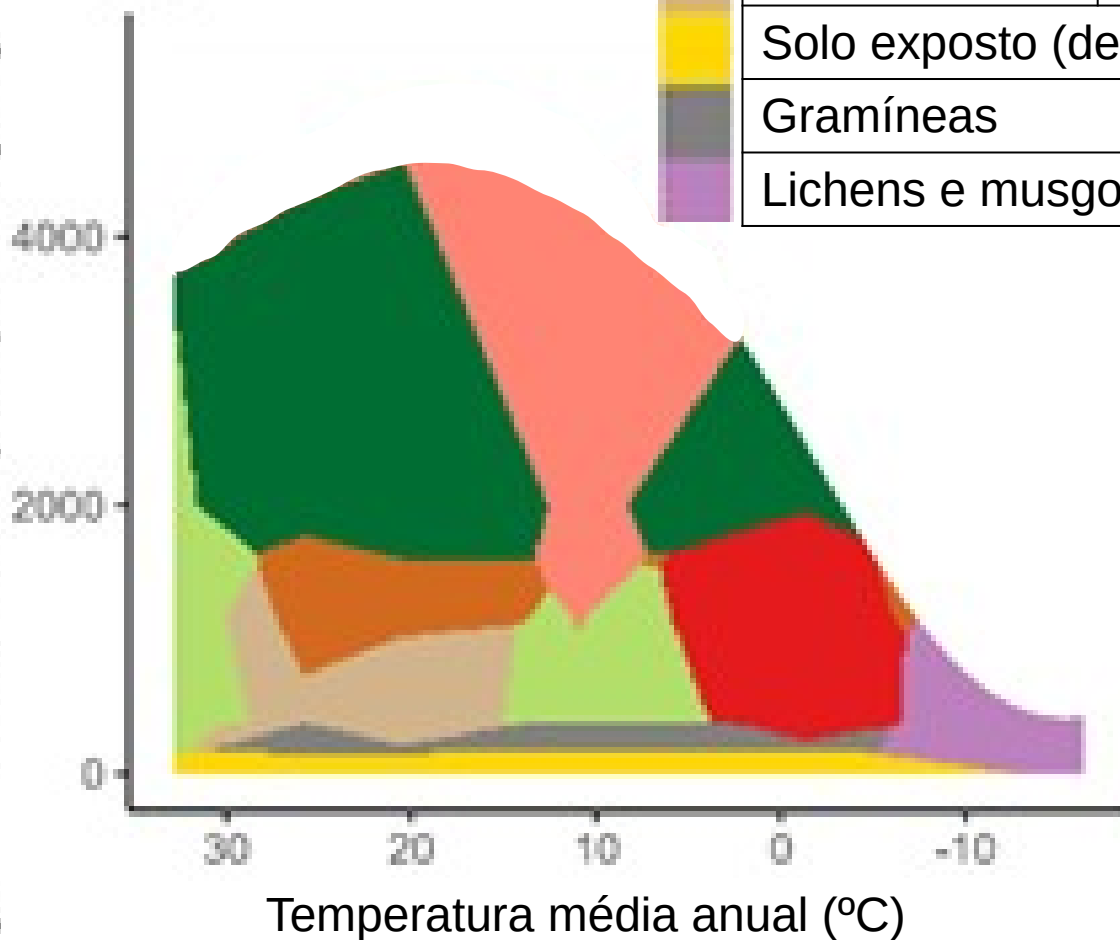
# Divisão de biomas com base na precipitação e temperatura Whittaker (1975)



Whittaker, R. H. Communities and Ecosystems, 2nd ed., Macmillan, New York (1975).

RICKLEFS, Robert E. The economy of nature. Macmillan, 2008.

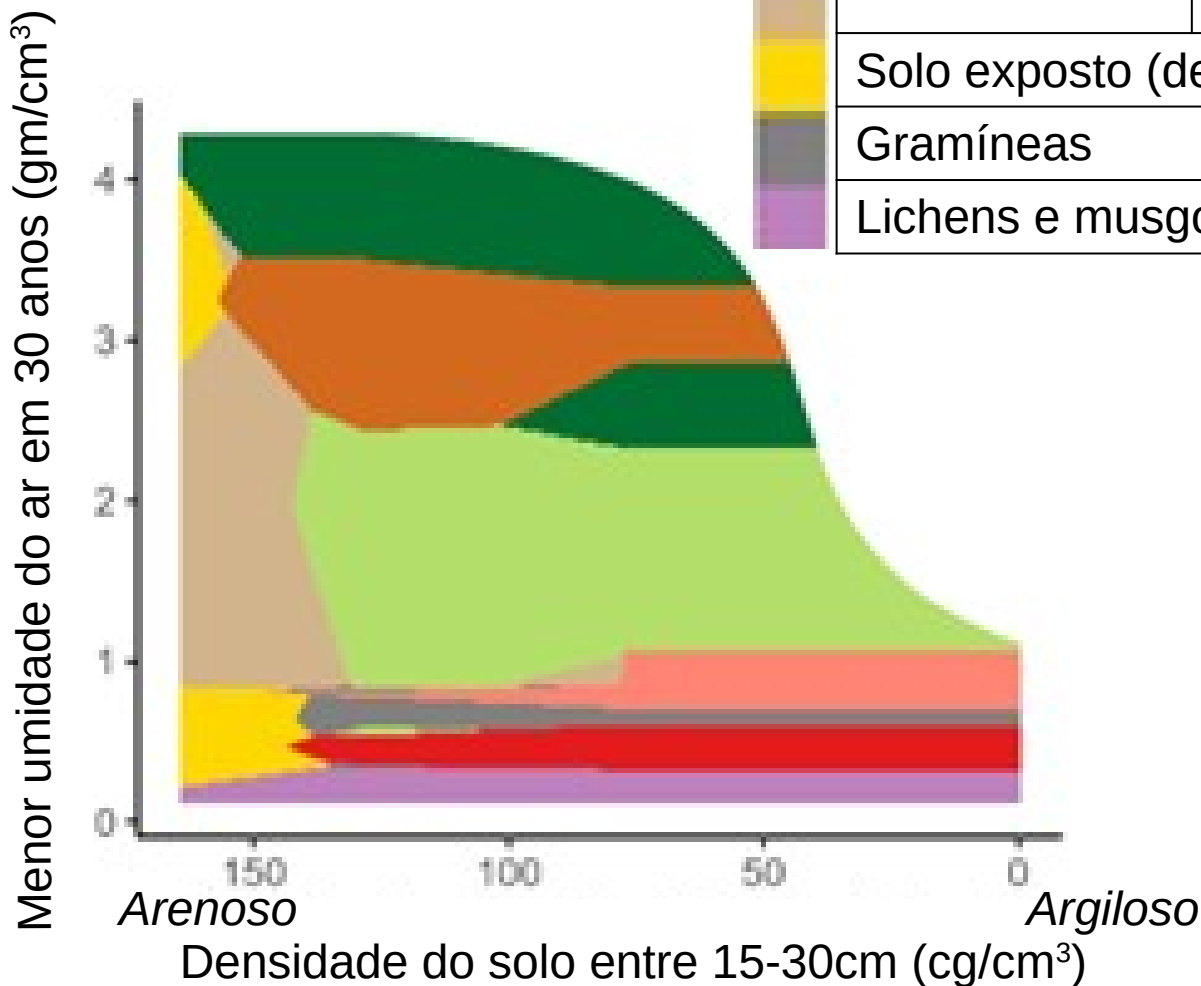
Precipitação anual (mm)



Floresta	folhas largas	perenifólia
		caducifólia
	coníferas	perenifólia
		caducifólia
Arbustiva	perenifólia	
	caducifólia	
Solo exposto (desertos)		
Gramíneas		
Lichens e musgos (tundra)		

MOREIRA, R. N. ;  
FUSHITA, A. T. ;  
VASCONCELOS, V. V. .  
Global Analysis of  
Environmental Attributes in  
Ecosystems. In: XXIV  
GeoInfo, 2023, São José  
dos Campos, São Paulo.  
Proceedings XXIV  
GEOINFO. São José dos  
Campos, São Paulo: INPE,  
2023. v. 1. p. 352-357.

# Melhor combinação para explicação dos ecossistemas



Floresta	folhas largas	perenifólia
		caducifólia
	coníferas	perenifólia
		caducifólia
Arbustiva	perenifólia	
	caducifólia	
Solo exposto (desertos)		
Gramíneas		
Lichens e musgos (tundra)		

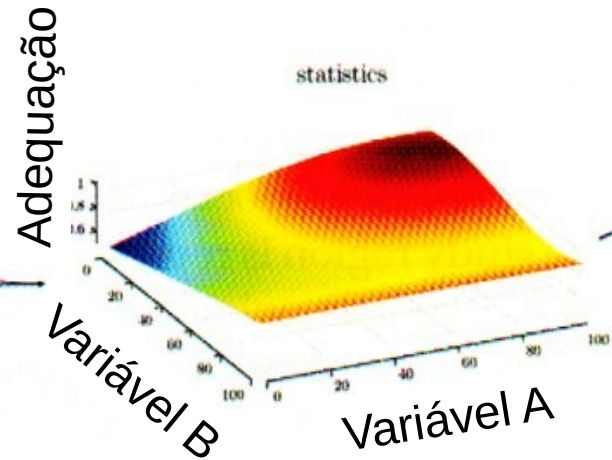
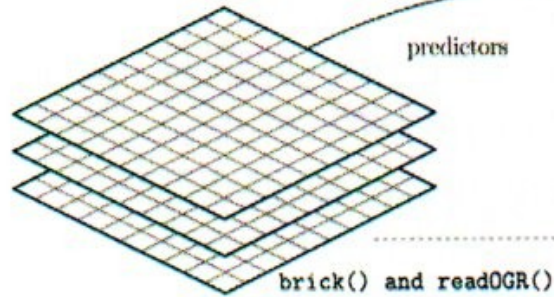
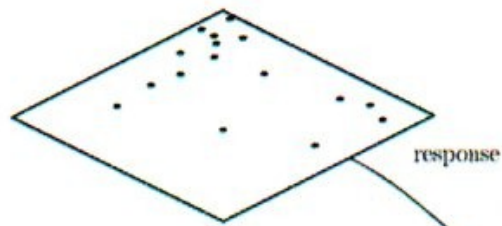
MOREIRA, R. N. ; FUSHITA, A. T. ; VASCONCELOS, V. V. . Global Analysis of Environmental Attributes in Ecosystems. In: XXIV GeoInfo, 2023, São José dos Campos, São Paulo. Proceedings XXIV GEOINFO. São José dos Campos, São Paulo: INPE, 2023. v. 1. p. 352-357.

# Objetivos

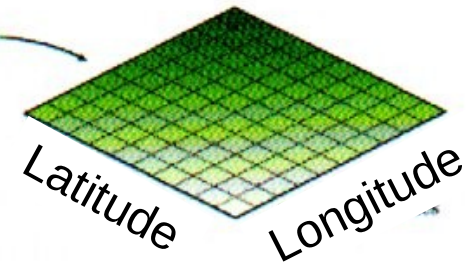
- Entender as relações entre as distribuições de espécies (biodiversidade) com as características dos ambientes
- **Compreender os métodos e técnicas para modelagem de distribuição de espécies**

# Modelagem de processos pontuais

Ocorrência dos pontos no espaço



Cor expressa probabilidade de ocorrência



Distribuição espacial das características explicativas

Modelo relacionando as variáveis com a ocorrência dos pontos

Mapa com a probabilidade de ocorrência dos pontos

# Modelagem de processos pontuais

<b>Eventos pontuais</b>	<b>Variáveis explicativas</b>	<b>Resultado</b>
Observações de espécies	Características do ambiente (temperatura, umidade, etc.)	Probabilidade de ocorrência de espécies
Ocorrência de deslizamentos de terra	Características dos geossistemas (declividade, rocha, solo, etc.) e de ocupação	Probabilidade de ocorrência de deslizamentos
Ocorrência de queimadas	Características climáticas (temperatura, chuva), sociais (proximidade de rodovias) e de vegetação (biomassa)	Probabilidade de ocorrência de queimadas
Notificações de crimes	Características socioeconômicas (renda, concentração de pessoas, distância a delegacias)	Probabilidade de ocorrência de crimes

# Modelagem de processos pontuais

<b>Eventos pontuais</b>	<b>Variáveis explicativas</b>	<b>Resultado</b>
Notificações de casos de dengue	Clima propício para o vetor (temperatura, umidade, etc.), locais de reprodução (lixo acumulado, ferro velho, etc.), porcentagem de casas sem tela nas janelas	Probabilidade de ocorrência da doença

# <https://posit.co/download/rstudio-desktop/>



PRODUCTS ▾

SOLUTIONS ▾

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PRICING

## 1: Install R

RStudio requires R 3.3.0+. Choose a version of R that matches your computer's operating system.

[DOWNLOAD AND INSTALL R](#)

## 2: Install RStudio

[DOWNLOAD RSTUDIO DESKTOP FOR WINDOWS](#)

Size: 215.66 MB | [SHA-256: D3C03C42](#) | Version: 2023.12.1+402 | Released: 2024-01-29

# <https://cran.r-project.org/bin/windows/base/>

## R-4.3.2 for Windows

[Download R-4.3.2 for Windows](#) (79 megabytes, 64 bit)

[README on the Windows binary distribution](#)

[New features in this version](#)



# Wallace

Interface gráfica para modelagem de  
distribuição e espécies

<https://wallaceecomod.github.io/>

Wallace Ecological Modeling App

[About](#) [Resources](#) [Dev Team](#) [Acknowledgments](#)

*Gulf of  
Panama*



## Wallace

a flexible platform for reproducible modeling of species niches and distributions

[</> CRAN](#)

[GITHUB](#)

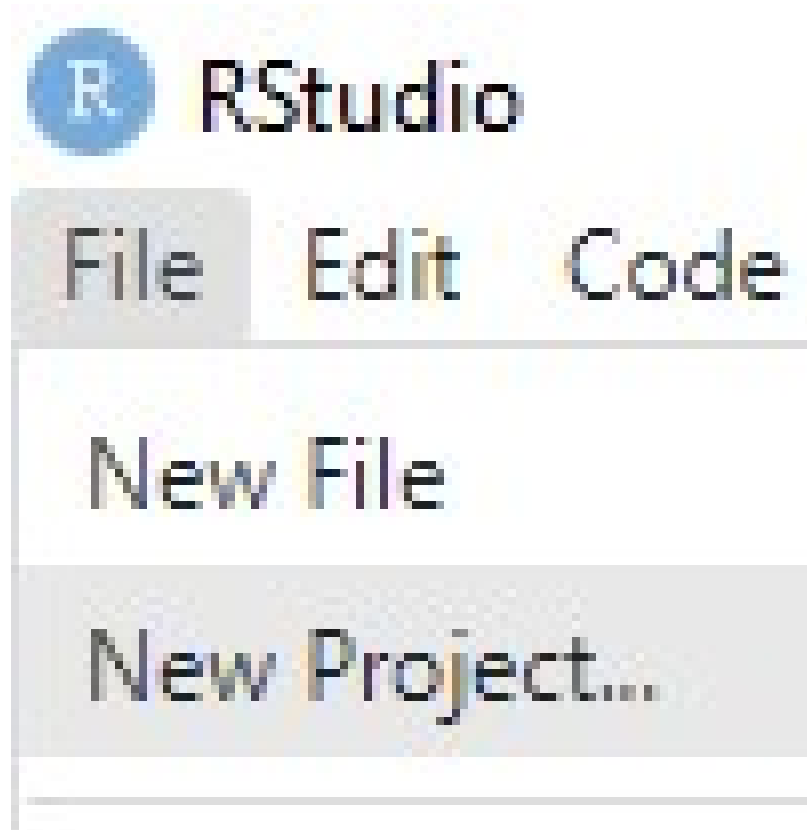
[EMAIL](#)

[FORUM](#)

[VIGNETTE](#)

- Copiar o arquivo de aula para uma pasta
- Script para chamar abrir a interface gráfica Wallace, dentro do RStudio
  - wallace.R




- Abrir o Rstudio
- Menu File → New Project



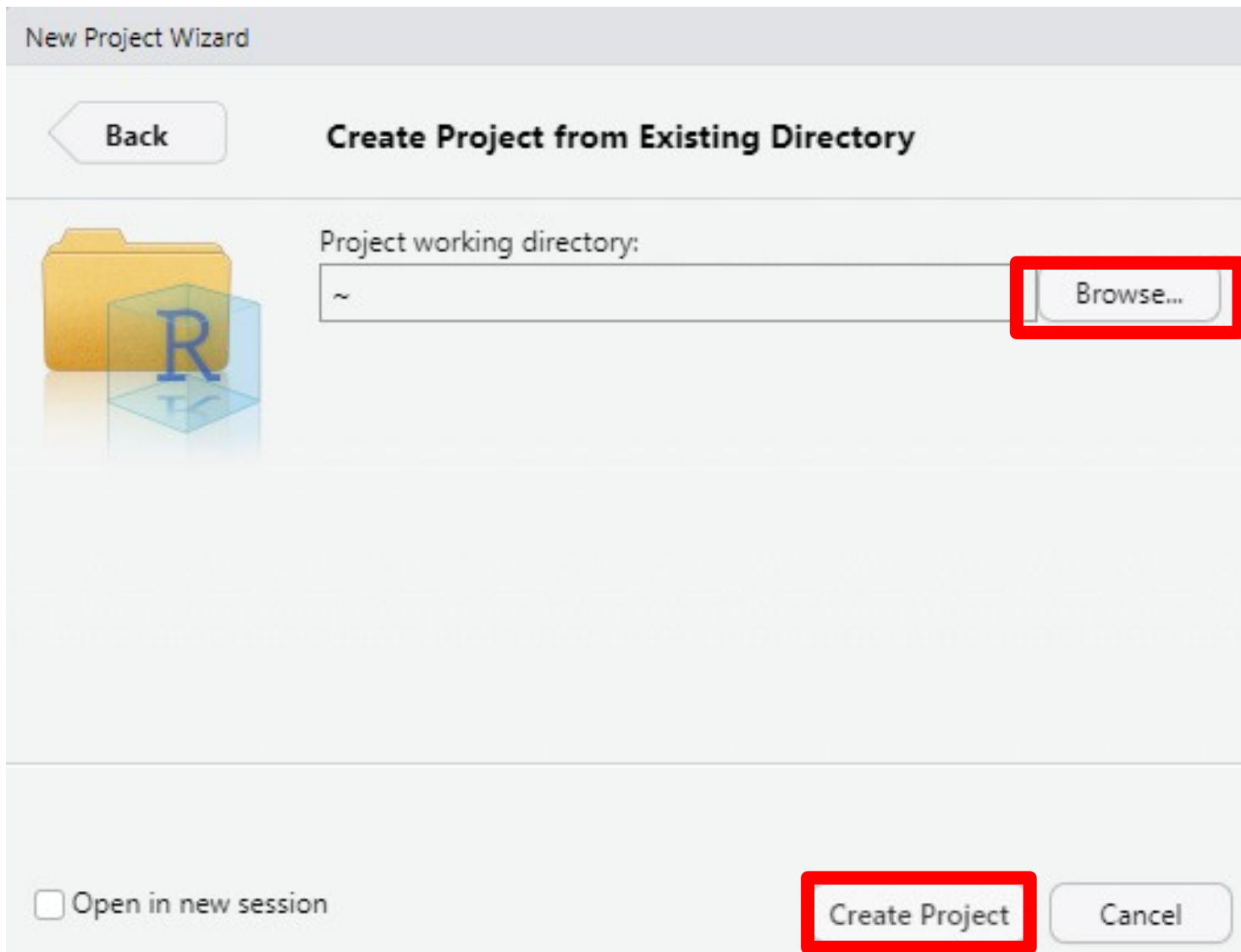
# • Create project -> Existing

New Project Wizard

## Create Project

- **New Directory**  
Start a project in a brand new working directory >
- **Existing Directory**  
Associate a project with an existing working directory >
- **Version Control**  
Checkout a project from a version control repository >

- Escolha a pasta onde colocou o arquivo de aula
- Create project



# Interface do RStudio

The image shows the RStudio interface with three main panels highlighted by red text:

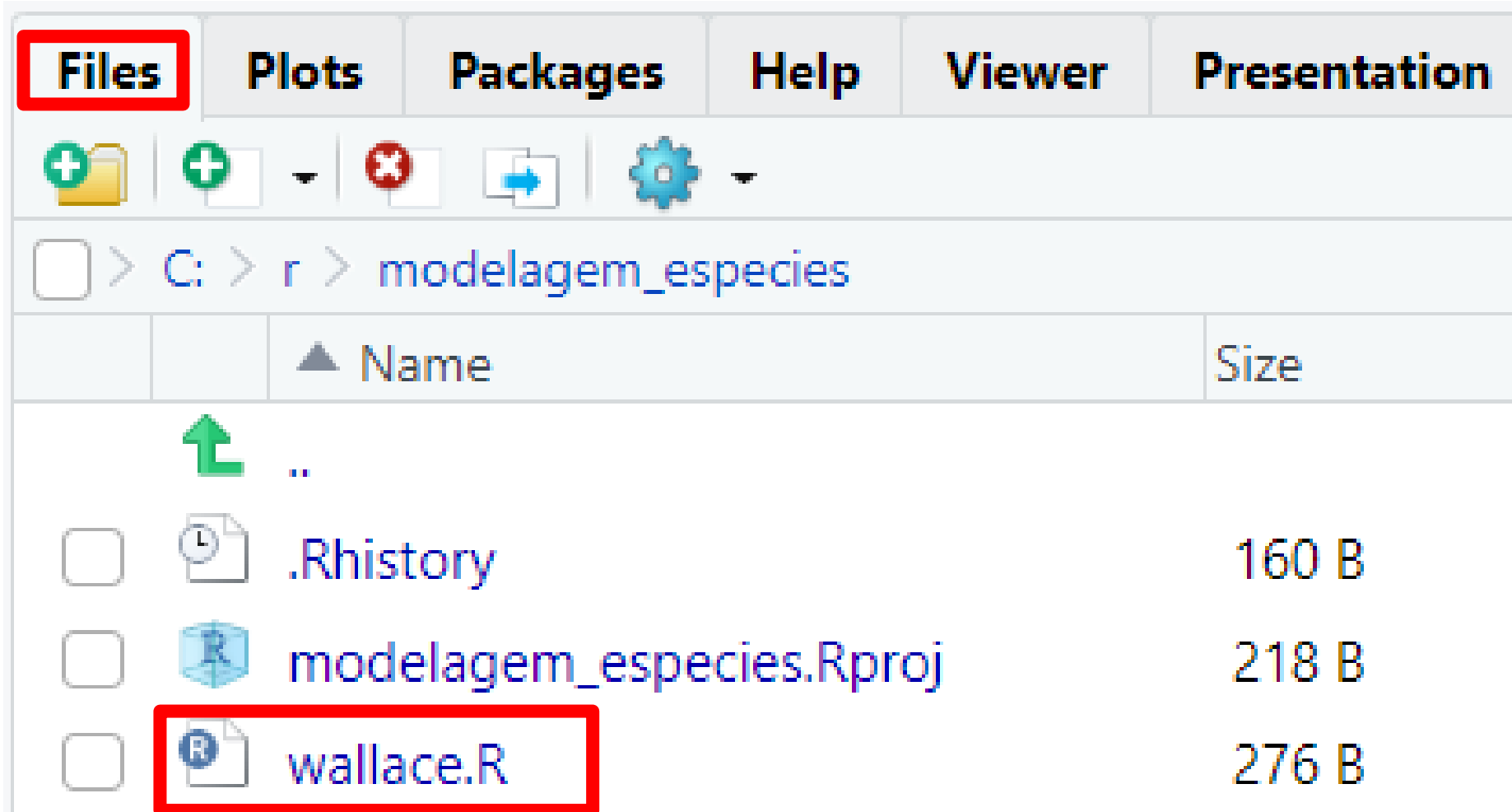
- Scripts:** The top-left panel shows an R script named 'wallace.R' with the following code:

```
1 # Install wallace from CRAN
2 install.packages("wallace")
3 install.packages("markdown")
4
5 # OR install wallace from GitHub
6 install.packages("remotes")
7 remotes::install_github("wallaceEcoMod/wallace")
8
9 # THEN load wallace
10 library(wallace)
11 # AND Run wallace
12 run_wallace()
```
- Objetos:** The top-right panel, titled 'Environment', shows 'Environment is empty'.
- Arquivos:** The bottom-right panel, titled 'Files', shows a file explorer view of the 'modelagem\_especies' directory with the following files:

Name	Size	Modified
..		
.Rhistory	160 B	Mar 6, 2024, 1:09 A
modelagem_especies.Rproj	218 B	Mar 10, 2024, 3:58
wallace.R	276 B	Mar 10, 2024, 3:58
- Resultados:** The bottom-left panel, titled 'Console', shows the R prompt and the following text:

```
R is a collaborative project with many contributors.
Type 'contributors()' for more information and
'citation()' on how to cite R or R packages in publications.
Type 'demo()' for some demos, 'help()' for on-line help, or
'help.start()' for an HTML browser interface to help.
Type 'q()' to quit R.
> |
```

- Na janela inferior direita, escolha a aba “Files”
- Clique no script “wallace.R”



- Clique no botão “Run”



R modelagem\_especies - RStudio

File Edit Code View Plots Session Build Debug Profile Tools Help

+ - + - + - Go to file/function | Addins -

wallace.R x

← → Source on Save 🔍 ✨ 📄



↻ Source - ☰

```
1 # Install Wallace from CRAN
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12:14 (Top Level) ▾

R Script ▾

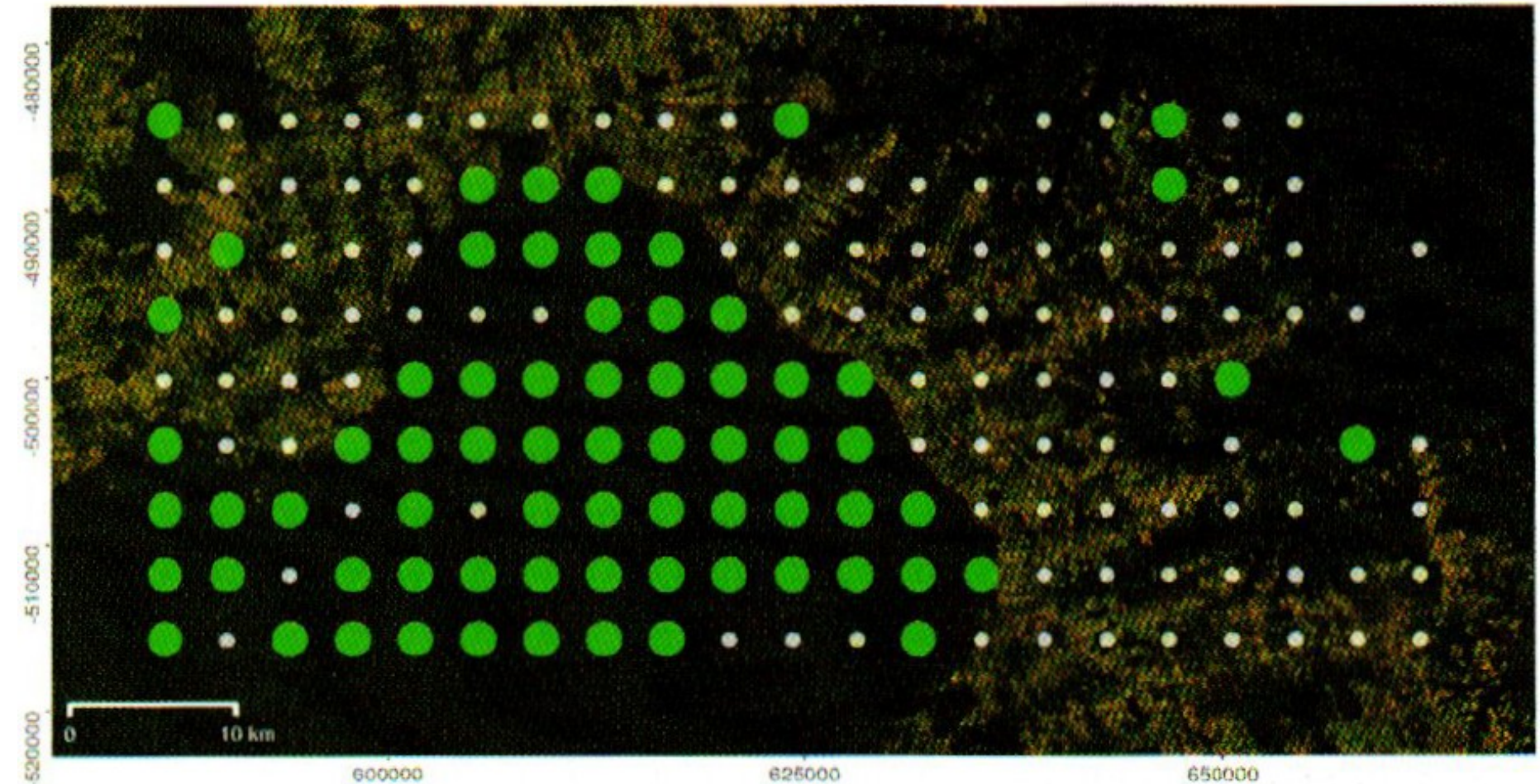


# Pressupostos dos modelos de distribuição potencial de espécies

- 1) Uma espécie está em equilíbrio com as características físicas do ambiente em que ocorre.
- 2) A amostragem do espaço geográfico (registros de ocorrência da espécie) é uma amostra que representa o espaço ambiental ocupado pela espécie.
- 3) Conservação do nicho: a propensão de espécies em manter características herdadas referente ao nicho ao longo do tempo evolutivo

# Modelos de levantamento de campo

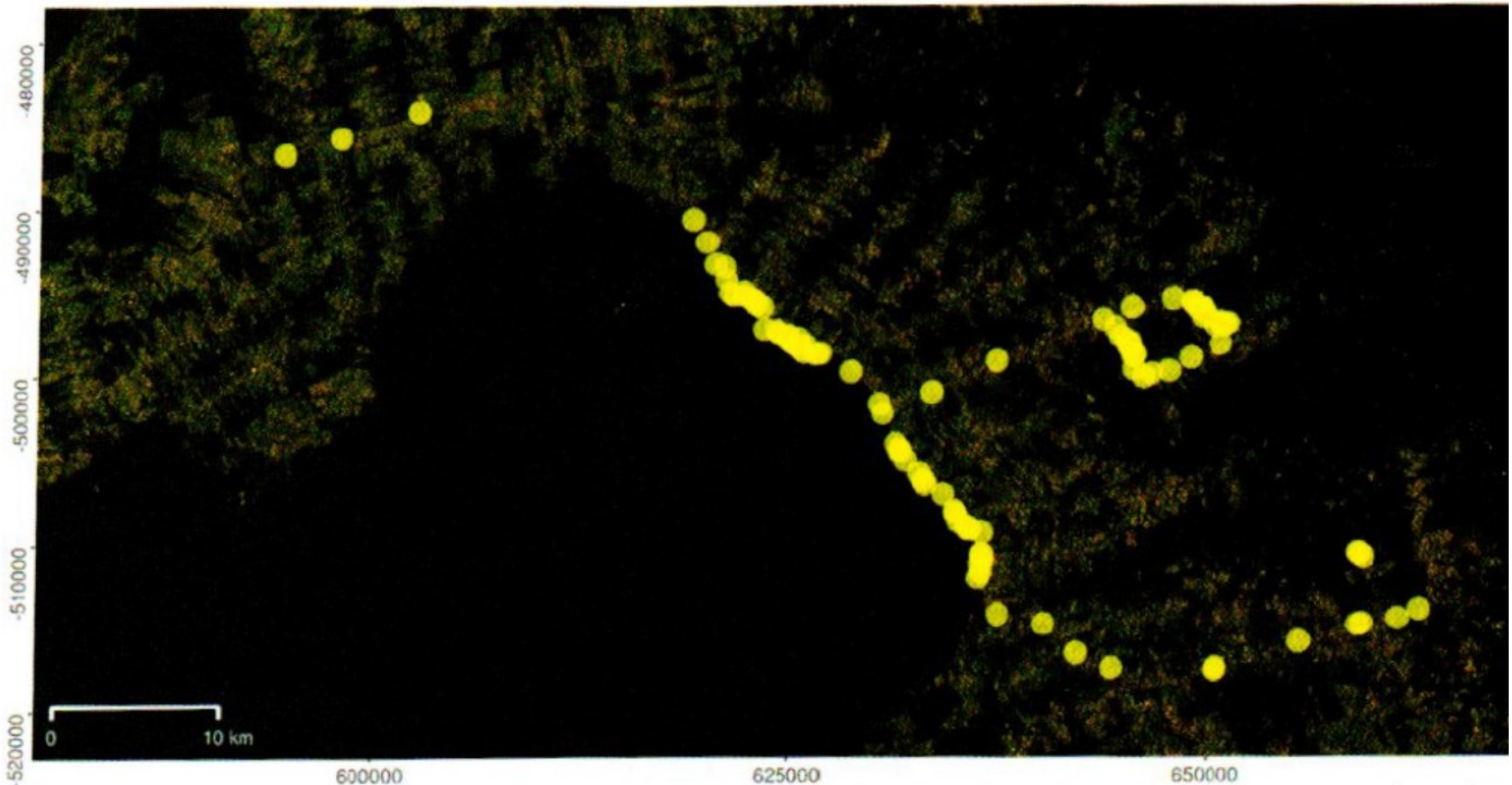
Ideal: ocorrência vs. não ocorrência, sistemático 



Reineking, B. Et al., Modelling species distributions. In: Wegmann M. et al. (org). Remote sensing and GIS for Ecologists. Pelag. 2016

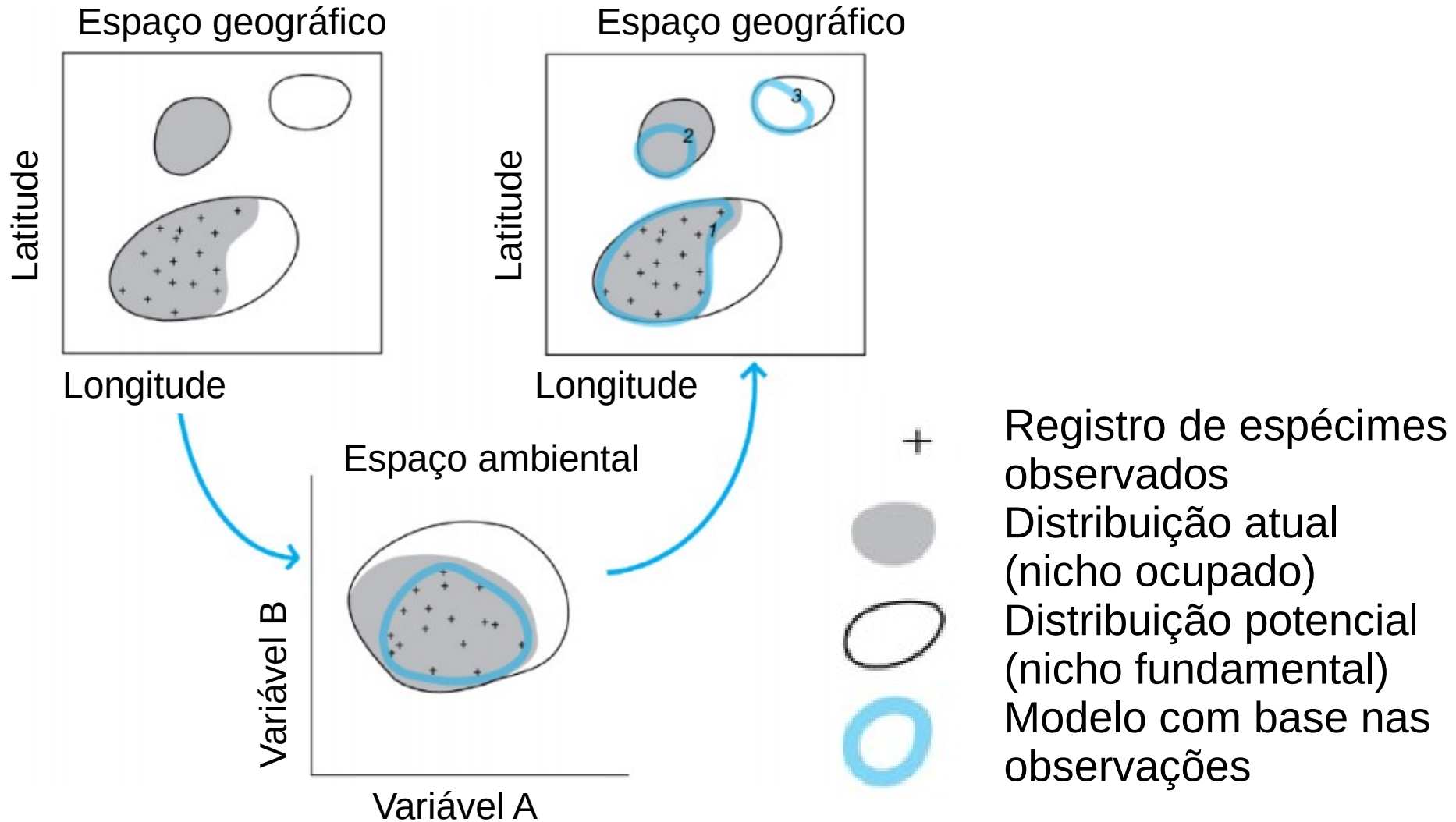
# Modelos de levantamento de campo

Usual: somente ocorrência, em locais de fácil acesso



Reineking, B. Et al., Modelling species distributions. In: Wegmann M. et al. (org). Remote sensing and GIS for Ecologists. Pelag. 2016

# Distribuição no espaço geográfico x espaço ambiental



Pearson, R. G. (2007). Species' distribution modeling for conservation educators and practitioners. *Synthesis*. American Museum of Natural History, 50, 54-89.

# GBIF – Global Biodiversity Information Facility

<https://www.gbif.org/>



Get data

How-to

Tools

Community

About

GBIF | Global Biodiversity Information Facility

## Free and open access to biodiversity data

OCCURRENCES

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DATASETS

PUBLISHERS

RESOURCES

| Search



What is GBIF?

About GBIF Brazil

# Modelagem de Espécie Invasora

- Javali (*Sus scrofa*)
- Altamente adaptável a diversos ambientes
- Destroi ecossistemas nativos, plantações e criações de porcos



# Tela Inicial

[Intro](#)[Occ Data](#)[Env Data](#)[Process Occs](#)[Process Envs](#)[Env Space](#)[Partition Occs](#)[Model](#)[Visualize](#)[Transfer](#)[Reproduce](#)[Support](#)

## WORKFLOW

*Wallace* (v2.1.2) currently includes ten components, or steps of a possible workflow. Each component includes two or more modules, which are possible analyses for that step.

### Components:

#### 1. Obtain Occurrence Data

- Query Present Database
- User-specified Occurrences

#### 2. Obtain Environmental Data

- WorldClim
- EcoClimate
- User-specified Environmental Data

#### 3. Process Occurrence Data

- Select Occurrences on Map
- Remove Occurrences by ID
- Spatial Thin

#### 4. Process Environmental Data

- Select Study Region by Extent
- Draw Study Region
- User-specified Study Region

#### 5. Characterize Environmental Space

- Environmental Ordination

[About](#)[Team](#)[How To Use](#)[Load Prior Session](#)

## What is *Wallace*?

Welcome to *Wallace*, a flexible application for reproducible ecological modeling, built for community expansion. The current version of *Wallace* (v2.1.2) steps the user through a full niche/distribution modeling analysis, from data acquisition to visualizing results.

The application is written in [R](#) with the web app development package [shiny](#). Please find the stable version of *Wallace* on [CRAN](#), and the development version on [Github](#). We also maintain a [Wallace website](#) that has some basic info, links, and will be updated with tutorial materials in the near future.

*Wallace* is designed to facilitate spatial biodiversity research, and currently concentrates on modeling species niches and distributions using occurrence datasets and environmental predictor variables. These models provide an estimate of the species' response to environmental conditions, and can be used to generate maps that indicate suitable areas for the species (i.e. its potential geographic distribution; Guisan & Thuiller 2005; Elith & Leathwick 2009; Franklin 2010a; Peterson et al. 2011). This research area has grown tremendously over the past two decades, with applications to pressing environmental issues such as conservation biology (Franklin 2010b), invasive species (Ficetola et al. 2007), zoonotic diseases (González et al. 2010), and climate-change impacts (Kearney et al. 2010).

Also, for more detail, please see our initial publication in *Methods in Ecology and Evolution* and our follow-up in *Ecography*.

Kass J. M., Vilela B., Aiello-Lammens M. E., Muscarella R., Merow C., Anderson R. P. (2018). *Wallace*: A flexible platform for reproducible modeling of species niches and distributions built for community expansion. *Methods in Ecology and Evolution*, 9(4): 1151-1156. DOI: [10.1111/2041-210X.12945](https://doi.org/10.1111/2041-210X.12945)

Kass, J.M., Pinilla-Buitrago, G.E, Paz, A., Johnson, B.A., Grisales-Betancur, V., Meenan, S.I., Attali, D., Broennimann, O., Galante, P.J., Maitner, B.S., Owens, H.L., Varela, S., Aiello-Lammens, M.E., Merow, C., Blair, M.E., Anderson R.P. (2022). *wallace 2*: a shiny app for modeling species niches and distributions redesigned to facilitate expansion via module contributions. *Ecography*, 2023(3): e06547. DOI: [10.1111/ecog.06547](https://doi.org/10.1111/ecog.06547).



Aba: Occ data (Ocorrências)

Módulo:  
"Query database"

Base de dados: GBIF

Nome científico da espécie:  
Sus scrofa

Máximo número de  
ocorrências:  
300

**Component: Obtain Occurrence Data** ?

**Modules Available:**

Query Database (Present)

User-specified

---

**Module: Query Database (Present)** ?

**R packages:** BIEN, spocc

---

**Choose Database**

GBIF  VertNet  BIEN

Keep only occurrences with uncertainty values

Include Data Source Citations

**Enter species scientific name**

Sus scrofa

**Set maximum number of occurrences**

300

Query Database



Species menu

Sus\_scrofa

Log window

Please find messages for the user in this log window.
> Sus scrofa | Total gbif records returned [300] out of [373827] total (limit 300). Records without coordinates removed [1]. Duplicated records removed [11]. Remaining records [288].

Map

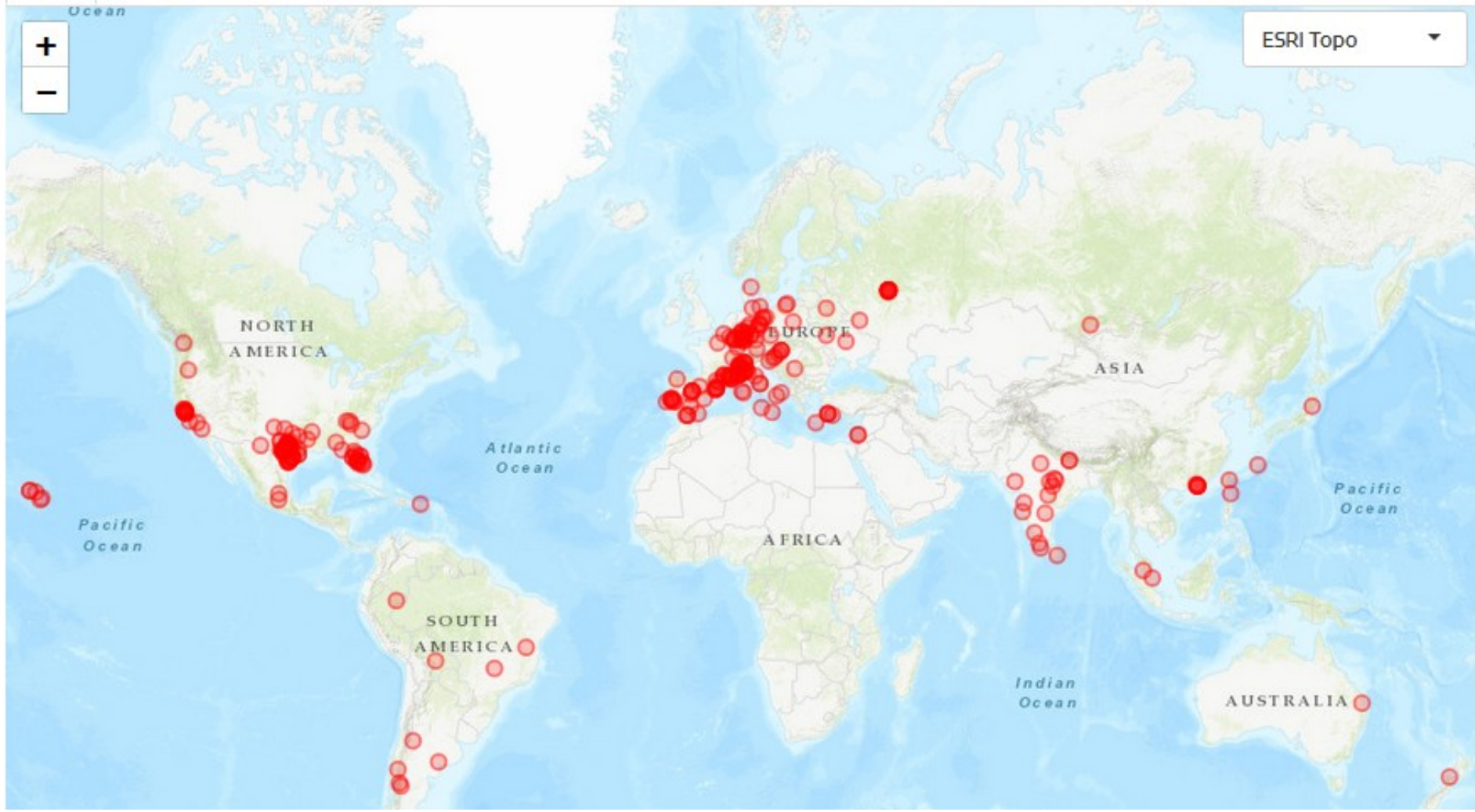
Occurrences

Results

Component Guidance

Module Guidance

Save



ESRI Topo

- Aba “Occurrence”
- Observe as coordenadas, países, ano, tipo e instituição

Map **Occurrences** Results Component Guidance Module Guidance Save


Show  entries Search:

occID	scientific_name	longitude	latitude	country	state_province	locality	year	record_type	catalog_number	institution_code
1	<i>Sus scrofa</i> Linnaeus, 1758	10.23	56.85	Denmark			2024	HUMAN_OBSERVATION	Arter_099e732a-1622-40b6-a0d3-b10201190bc5	MST-and-NHMD
2	<i>Sus scrofa</i> Linnaeus, 1758	13.17	52.57	Germany			2024	HUMAN_OBSERVATION	1463b494-d855-5376-aa12-20375cf69298	https://artenfinder.net
3	<i>Sus scrofa</i> Linnaeus, 1758	2.17	42.75	France	Languedoc-Roussillon		2024	HUMAN_OBSERVATION	195454565	iNaturalist
4	<i>Sus scrofa</i> Linnaeus, 1758	27.72	53.97	Belarus	Minsk		2024	HUMAN_OBSERVATION	195466145	iNaturalist
5	<i>Sus scrofa</i> Linnaeus, 1758	-95.26	29.07	United States of America	Texas		2024	HUMAN_OBSERVATION	195524959	iNaturalist
6	<i>Sus scrofa domesticus</i> Linnaeus, 1758	-121.63	37.67	United States of America	California		2024	HUMAN_OBSERVATION	195530288	iNaturalist
7	<i>Sus scrofa</i> Linnaeus, 1758	8.23	40.94	Italy	Sardegna		2024	HUMAN_OBSERVATION	195551470	iNaturalist
8	<i>Sus scrofa</i> Linnaeus, 1758	-99.61	31.68	United States of America	Texas		2024	HUMAN_OBSERVATION	195570096	iNaturalist
9	<i>Sus scrofa domesticus</i> Linnaeus, 1758	-42.22	-14.5	Brazil	Bahia		2024	HUMAN_OBSERVATION	195582670	iNaturalist

*Note: To save your session code or metadata, use the Reproduce component*

## Save Session

By saving your session into an RDS file, you can resume working on it at a later time or you can share the file with a collaborator.

 Save Session


## Download Data

Download data/results from analyses from currently selected module

Download original  
occurrence data

 CSV file

Download current table

 CSV file

Download all data

 CSV file

# Variáveis comumente utilizadas para modelagem de distribuição de espécies

- Clima (temperatura, precipitação)
  - Médias anuais
  - Sazonalidade
- Topografia
  - Elevação
  - Declividade
  - Curvatura
- Solos
  - Classes de solo
  - Textura (arenoso, silteoso, argiloso)
  - Nutrientes

# Variáveis comumente utilizadas para modelagem de distribuição de espécies

- Ocorrências de outras espécies
  - Tipo de vegetação, para prever animais
  - Predador e presa
  - Competição
- Pressões antrópicas
  - Ocupação do solo
  - Densidade humana ou de vias de locomoção
  - Desmatamento
  - Poluição do ar, água ou solo

- Aba “EnvData”
  - Dados ambientais
- Módulo:
  - WorldClim Bioclimes
- Resolução
  - 10 arcmin
- Marque:
  - “Save to memory for faster processing and save/load option”

The screenshot shows the QGIS interface with the 'Env Data' component selected in the top navigation bar. The component is titled 'Component: Obtain Environmental Data'. Under 'Modules Available', 'WorldClim Bioclimes' is selected. The 'Module: WorldClim Bioclimes' section shows 'R packages: raster'. The 'Select WorldClim bioclimatic variable resolution' dropdown is set to '10 arcmin'. The checkbox 'Save to memory for faster processing and save/load option' is checked. The 'Select bioclim variables' dropdown shows 'bio01, bio02, bio03, bio04, bio05, bio06, bio07, bio08, bio09, bi'. The 'Batch' checkbox is unchecked. The 'Load Env Data' button is visible at the bottom.

**Component: Obtain Environmental Data** ⓘ

**Modules Available:**

- WorldClim Bioclimes
- ecoClimate
- User-specified

---

**Module: WorldClim Bioclimes** ⓘ

**R packages:** raster

---

**Select WorldClim bioclimatic variable resolution**

10 arcmin

Save to memory for faster processing and save/load option

**Select bioclim variables**

bio01, bio02, bio03, bio04, bio05, bio06, bio07, bio08, bio09, bi

Batch

Load Env Data

# WorldClim

<https://worldclim.org/>

Variáveis climáticas:

Histórico

Presente

Projeções climáticas

Variáveis bioclimáticas

## WorldClim

Maps, graphs, tables, and data of the global climate

[Download](#)



# Variáveis bioclimáticas

BIO1 = Temperatura média anual

BIO2 = Amplitude térmica diurna (Média mensal (max temp - min temp))

BIO3 = Isotermalidade (BIO2/BIO7) ( $\times 100$ )

BIO4 = Sazonalidade da temperatura (desvio padrão  $\times 100$ )

BIO5 = Temperatura máxima do mês mais quente

BIO6 = Temperatura mínima do mês mais frio

BIO7 = Amplitude térmica anual (BIO5-BIO6)

BIO8 = Temperatura média do quadrimestre mais úmido

BIO9 = Temperatura média do quadrimestre mais seco

BIO10 = Temperatura média do quadrimestre mais quente

BIO11 = Temperatura média do quadrimestre mais frio

BIO12 = Precipitação anual

BIO13 = Precipitação do mês mais úmido

BIO14 = Precipitação do mês mais seco

BIO15 = Sazonalidade da precipitação (coeficiente de variação)

BIO16 = Precipitação do quadrimestre mais úmido

BIO17 = Precipitação do quadrimestre mais seco

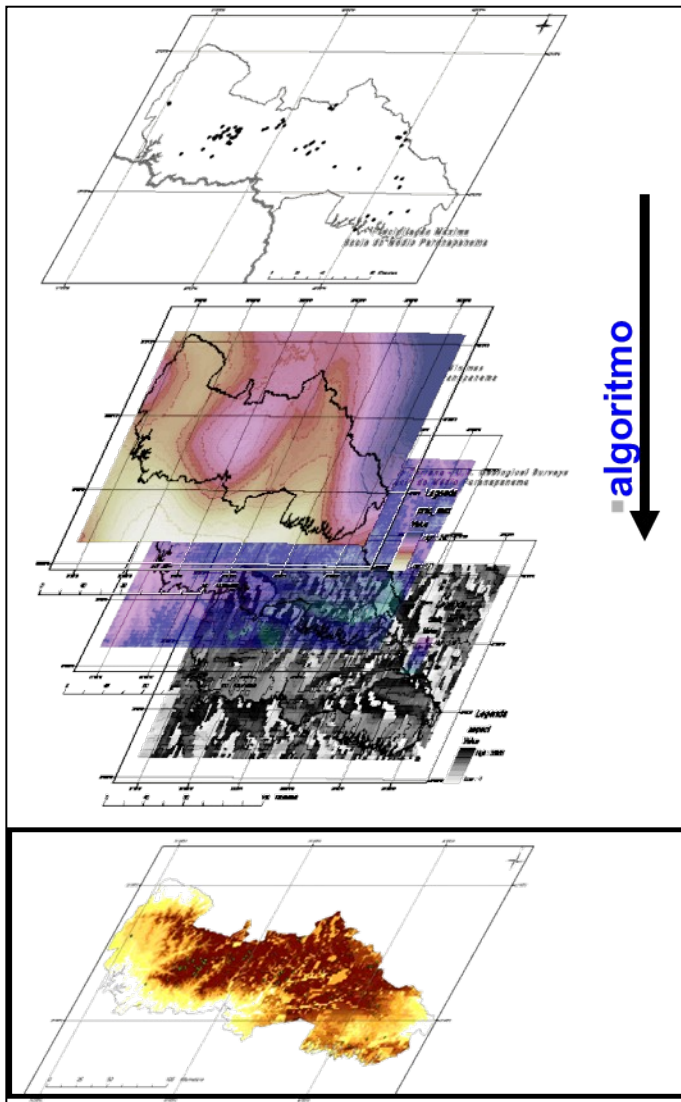
BIO18 = Precipitação do quadrimestre mais quente

BIO19 = Precipitação do quadrimestre mais frio

Booth, T. H., Nix, H. A., Busby, J. R., & Hutchinson, M. F. (2014). BIOCLIM: the first species distribution modelling package, its early applications and relevance to most current MAXENT studies. *Diversity and Distributions*, 20(1), 1-9.



# Exemplo de tabela de dados, com os valores ambientais adicionados, utilizada no procedimento de modelagem



algoritmo



	pa	bio1	bio12	bio16	bio17	bio5	bio6	bio7	bio8
1	1	183	988	374	110	323	66	257	240
2	1	226	1395	455	177	327	126	202	265
3	1	192	2160	833	267	262	107	155	219
4	1	191	1200	516	140	276	91	186	220
5	1	209	1330	501	190	300	103	197	237
6	1	205	1354	541	165	296	99	197	233
7	1	208	1403	539	180	296	104	193	236
8	1	222	1444	697	30	291	139	152	226
9	1	229	1217	580	111	309	146	163	243
10	1	186	1934	550	444	289	87	201	187
11	1	199	1436	779	46	284	97	186	218
12	1	257	1267	568	52	333	156	177	269
13	1	194	1919	533	443	295	99	197	208
14	1	165	2041	575	460	271	50	221	168
15	1	195	1449	750	61	281	79	201	219
16	1	197	1500	720	97	281	96	185	223
17	1	188	1530	724	101	272	98	174	213
18	1	223	1475	583	153	302	144	158	250
19	1	196	1484	741	90	273	86	186	218
20	1	258	2691	1077	276	317	215	102	259
21	1	262	3406	1242	238	309	219	90	252
22	1	202	2026	611	426	275	140	135	202
23	1	236	1072	476	46	306	162	145	235
24	1	236	1072	476	46	306	162	145	235
25	1	225	2622	1183	84	295	164	131	222
26	1	192	2160	833	267	262	107	155	219
27	1	136	1746	578	312	200	78	122	131
28	1	267	1912	851	54	322	223	99	262
29	1	167	3159	1168	312	224	111	113	166
30	1	167	3159	1168	312	224	111	113	166
31	1	261	3561	1376	315	321	216	105	258
32	1	236	1072	476	46	306	162	145	235
33	1	216	3185	1063	443	277	160	117	212
34	1	259	2876	1119	163	315	214	101	253
35	1	267	1912	851	54	322	223	99	262
36	1	262	2483	933	229	329	201	127	256
37	1	258	2691	1077	276	317	215	102	259
38	1	167	3159	1168	312	224	111	113	166
39	1	258	2691	1077	276	317	215	102	259
40	1	202	1751	530	337	315	79	236	232
41	1	265	2109	865	195	325	221	105	265
42	1	261	3561	1376	315	321	216	105	258
43	1	167	3159	1168	312	224	111	113	166
44	1	167	3159	1168	312	224	111	113	166
45	1	228	3218	1418	153	303	161	141	225

(Siqueira, 2014)



**Please, check Log window for  
more information**

OK

- Ocorrências removidas porque estavam na mesma célula (pixel) da base do WorldClim




### Log window

```
bio05, bio06, bio07, bio08, bio09, bio10, bio11, bio12, bio13,  
bio14, bio15, bio16, bio17, bio18, bio19 at 10 arcmin resolution.  
> ! WARNING : Sus scrofa | Removed 63 localities that shared the  
same grid cell. occIDs: 34, 42, 52, 58, 70, 71, 75, 93, 94, 95, 99,  
100, 103, 105, 106, 109, 138, 139, 140, 142, 143, 148, 149, 151,
```

Aba:

“Occurrences”

Restaram 225  
ocorrências

Map **Occurrences** Results  Component Guidance  Module Guidance  Save

Show  entries

occID	scientific_name	longitude	latitude	country	state_province	locality
1	<i>Sus scrofa</i> Linnaeus, 1758	10.23	56.85	Denmark		
2	<i>Sus scrofa</i> Linnaeus, 1758	13.17	52.57	Germany		
3	<i>Sus scrofa</i> Linnaeus, 1758	2.17	42.75	France	Languedoc-Roussillon	
4	<i>Sus scrofa</i> Linnaeus, 1758	27.72	53.97	Belarus	Minsk	
5	<i>Sus scrofa</i> Linnaeus, 1758	-95.26	29.07	United States of America	Texas	
6	<i>Sus scrofa domesticus</i> Linnaeus, 1758	-121.63	37.67	United States of America	California	
7	<i>Sus scrofa</i> Linnaeus, 1758	8.23	40.94	Italy	Sardegna	
8	<i>Sus scrofa</i> Linnaeus, 1758	-99.61	31.68	United States of America	Texas	
9	<i>Sus scrofa domesticus</i> Linnaeus, 1758	-42.22	-14.5	Brazil	Bahia	
10	<i>Sus scrofa</i> Linnaeus, 1758	8.47	45.91	Italy	Piemonte	

Showing 1 to 10 of 225 entries


- Aba “Save”

Map Occurrences Results  Component Guidance  Module Guidance  Save

*Note: To save your session code or metadata, use the Reproduce component*

### Save Session

By saving your session into an RDS file, you can resume working on it at a later time or you can share the file with a collaborator.


 Save Session

### Download Data

Download data/results from analyses from currently selected module

Download environmental variables (Select download file type)

GeoTIFF ▼

 ZIP file

- Aba: Process Occs
- Selecione “Spatial thin”
- Distância: 5km

[Intro](#)[Occ Data](#)[Env Data](#)[Process Occs](#)

## Component: Process Occurrence Data



### Modules Available:

Select Occurrences On Map

Remove Occurrences By ID

Spatial Thin

### Module: Spatial Thin ?

R packages: *spThin*

The minimum distance between occurrence locations (nearest neighbor distance) in km for resulting thinned dataset. Ideally based on species biology (e.g., home-range size).

#### Thinning distance (km)

Batch

Thin Occurrences

## Log window

157, 169, 177, 179, 181, 186, 191, 192, 193, 196, 199, 202, 206,  
208, 212, 224, 225, 226, 228, 229, 234, 236, 242, 243, 244, 245,  
246, 257, 262, 263, 270, 271, 274, 275, 283, 294, 299, 300.

> ***Sus scrofa*** | Worldclim variables ready to use.

> ***Sus scrofa*** | Total records thinned (5 km) to 220 localities

# Fontes de incerteza para modelagem de processos pontuais

- Previsão para fora da área de estudo ou para locais com características diferentes das amostradas
  - Ex: expansão potencial de espécies invasoras, vetores de doença, plantios agrícolas
- Previsões para outros períodos de tempo (passado, futuro)
  - Ex: efeito de mudanças climáticas

- As relações seriam as mesmas?
- Outras variáveis e processos ambientais e sociais seriam relevantes?
- As projeções são confiáveis?



- Aba “Process Envs”
- Marque “Select Study Region”
- Extensão:
  - “Minimum convex polygon”
- Buffer:
  - 1 km



Intro

Occ Data

Env Data

Process Occs

Process Envs

### Component: Process Environmental Data ?

#### Modules Available:

Select Study Region

Draw Study Region

User-specified Study Region

#### Module: Select Study Region by Extent ?

R packages: *sp*, *sf*

#### Step 1: Choose Background Extent

##### Background Extents:

bounding box

minimum convex polygon

point buffers

##### Study region buffer distance (degree)

1

Batch


Select

Map

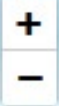
Occurrences

Results

 Component Guidance

 Module Guidance

 Save



ESRI Topo



- Amostragem aleatória de pontos para comparação:
  - 220, igual ao número de ocorrências

## Step 2: *Sample Background Points*

Mask predictor rasters by background extent and sample background points

No. of background points

220

Batch


Sample

*Note: To save your session code or metadata, use the Reproduce component*

## Save Session

By saving your session into an RDS file, you can resume working on it at a later time or you can share the file


**⚠ The current session data is large, which means the downloaded file may be large and the download might**

 Save Session

## Download Data


Download data/results from analyses from currently selected module

Download shapefile of  
background extent


 ZIP file

Download predictor rasters  
masked to background  
extent (Select download file  
type)

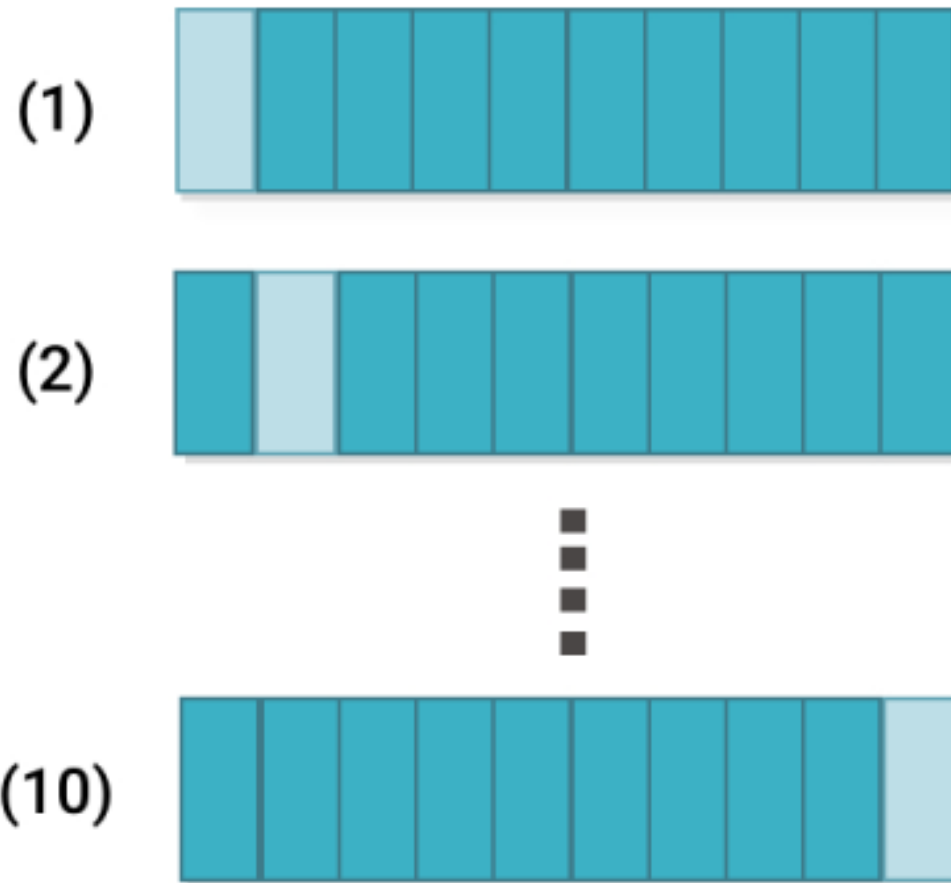
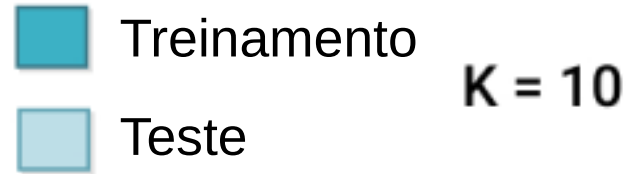
GeoTIFF ▼

 ZIP file

Download sample  
background points

 CSV file

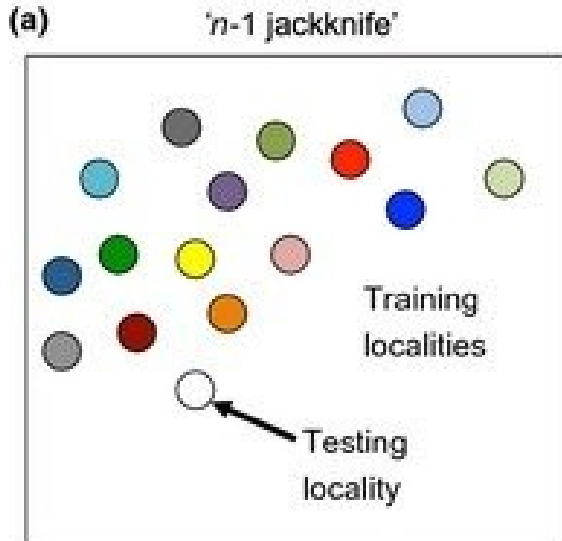
# Validação cruzada



# Métodos de particionamento

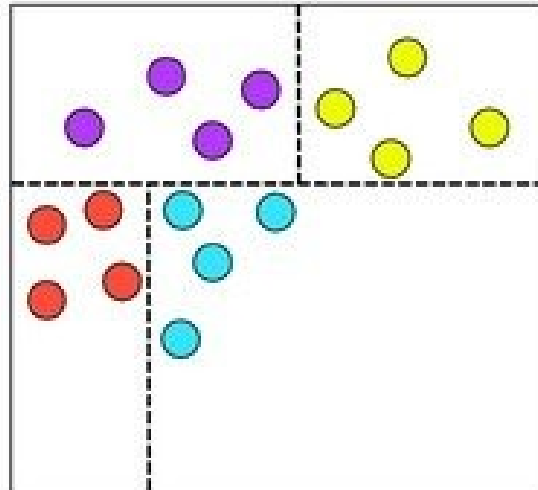
## Aleatório

'n-1 jackknife'



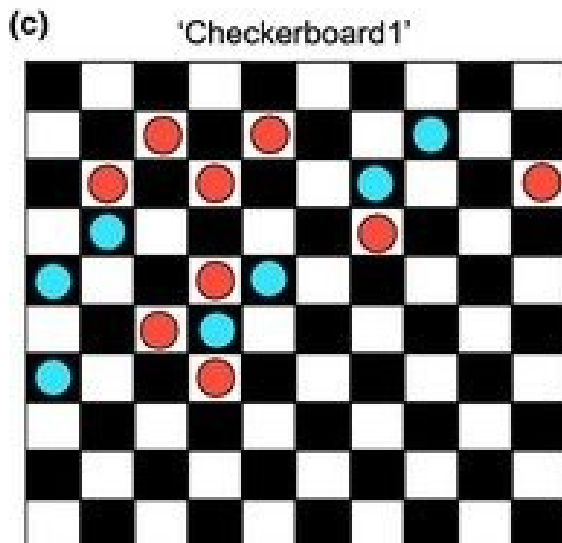
## Agrupamento espacial

(b) 'Block'



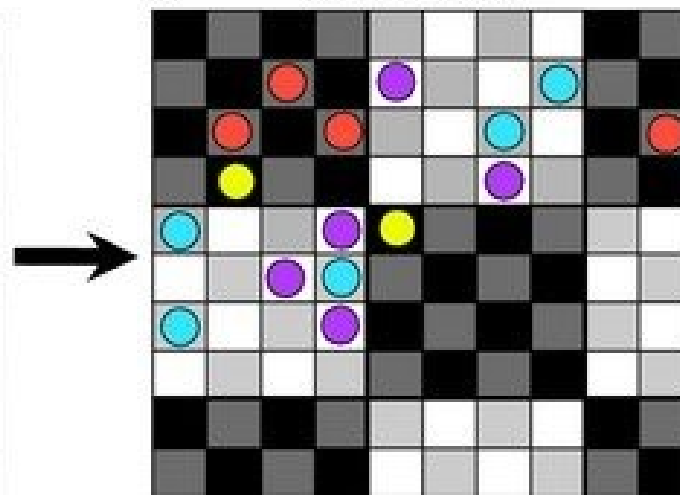
## Tabuleiro 1

'Checkerboard1'



## Tabuleiro 2

(d) 'Checkerboard2'



Muscarella, R. et al. (2014). ENM eval: An R package for conducting spatially independent evaluations and estimating optimal model complexity for Maxent ecological niche models. *Methods in ecology and evolution*, 5(11), 1198-1205.

[Intro](#)[Occ Data](#)[Env Data](#)[Process Occs](#)[Process Envs](#)[Env Space](#)[Partition Occs](#)

## Component: Partition Occurrence Data ?

### Modules Available:

Non-spatial Partition

Spatial Partition

### Module: Spatial Partition ?

R packages: *ENMeval*

### Options Available:

Checkerboard 2 (k = 4)

### Aggregation Factor

2

Batch

Partition

### Species menu

Sus\_scrofa

Map

Occurrences

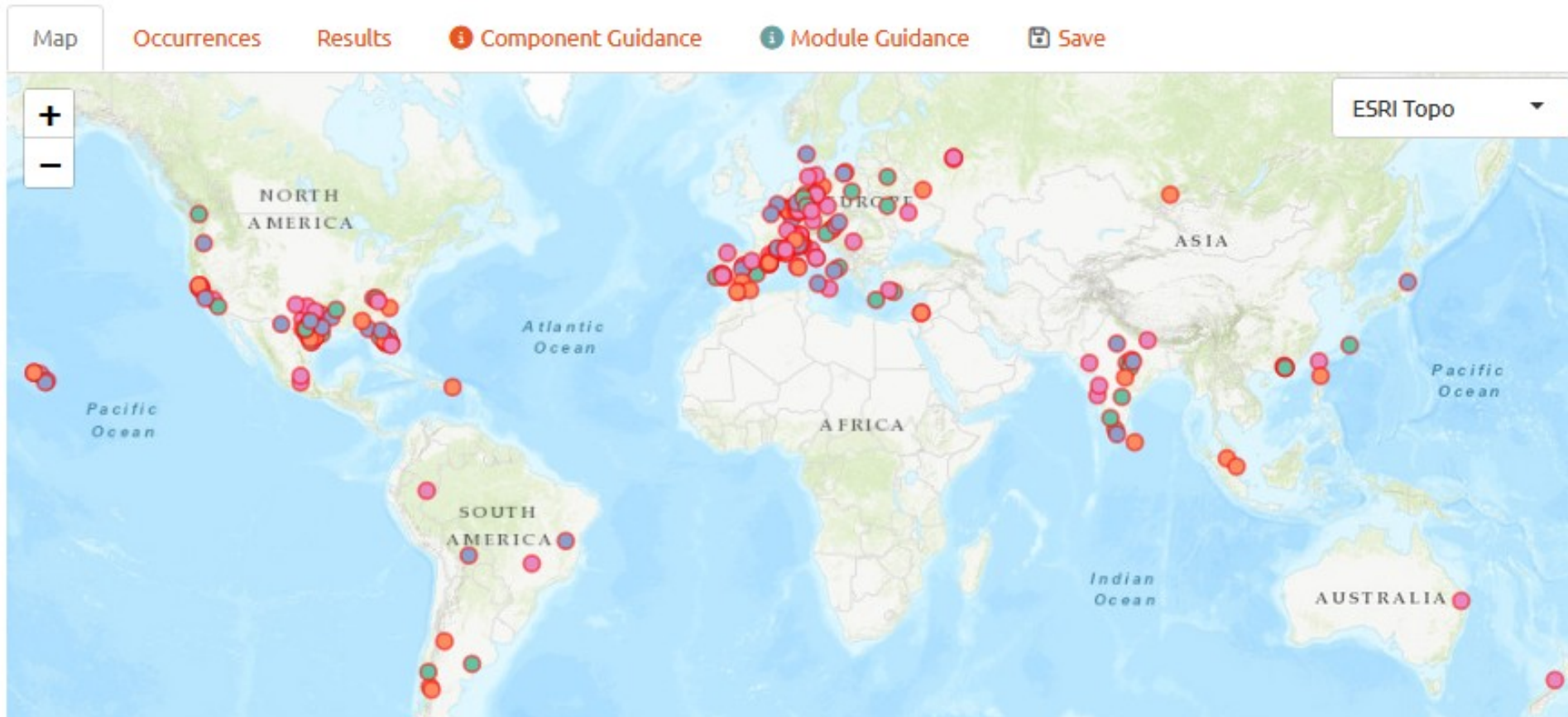
Results

+

-



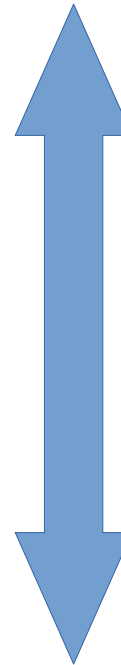
Cada um dos 4 grupos (cores) será usado para calibrar o modelo, e os 3 outros serão usados para validação





# Modelos de processos pontuais

- Envelopes
- Regressão linear
- Regressão não-linear
- Aprendizado de máquinas



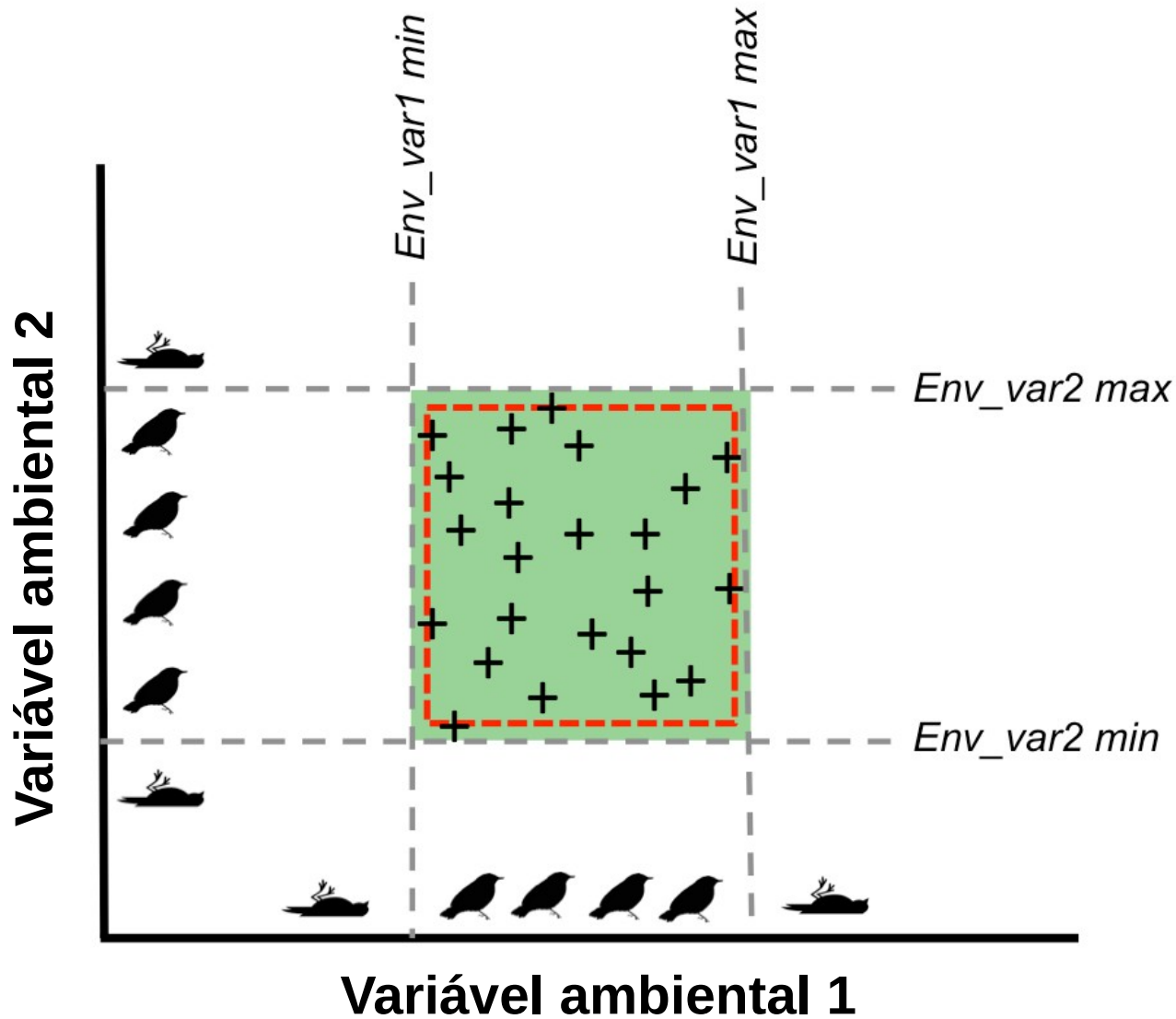
**Mais simples**

Pior ajuste  
Extrapolação mais confiável

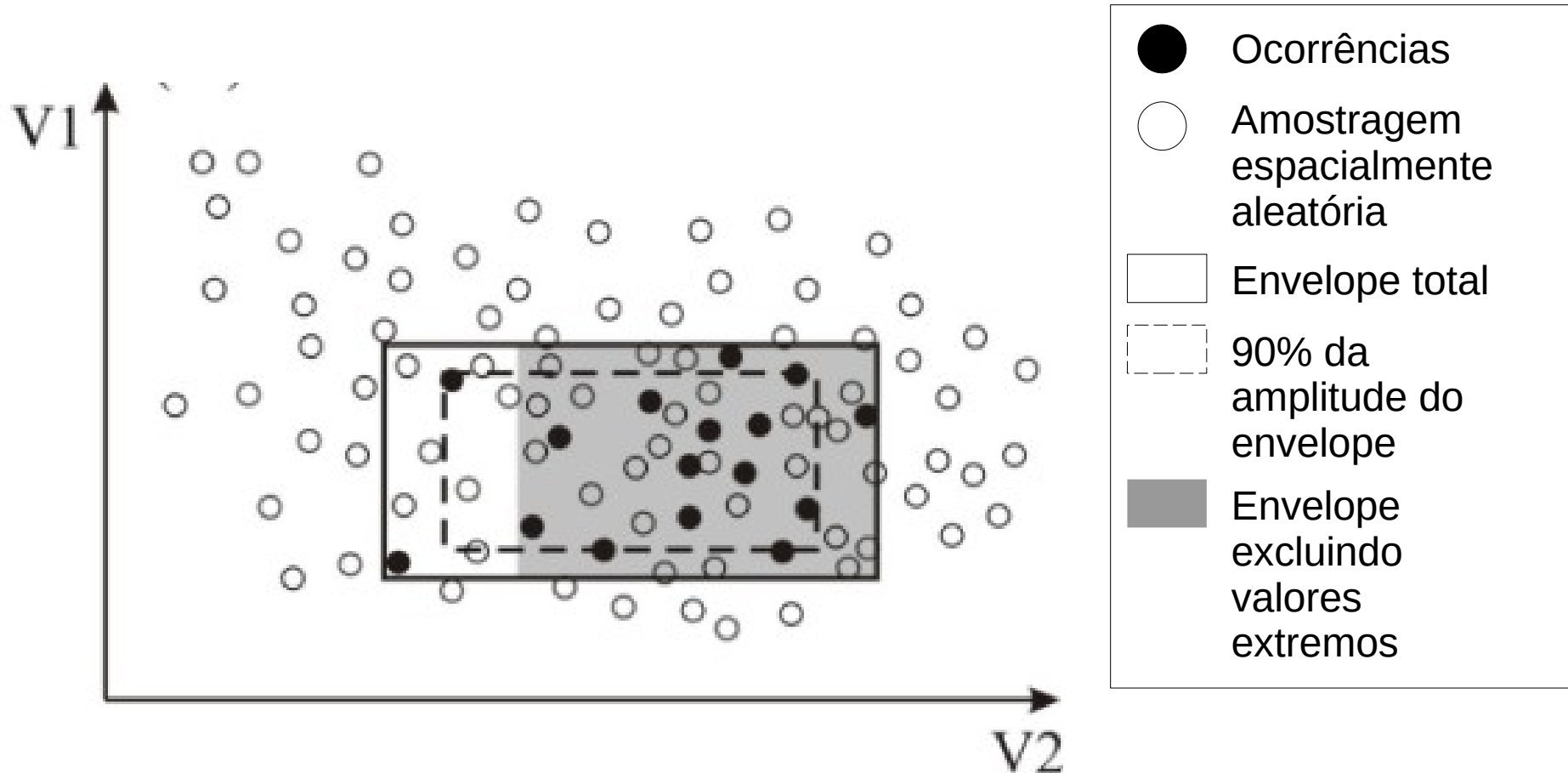
**Mais complexos**

Melhor ajuste  
Extrapolação menos confiável

# Envelope



# BIOCLIM (Método de Envelope)



Lima-Ribeiro, M. D. S., & Diniz-Filho, J. A. F. (2012). Modelando a distribuição geográfica das espécies no passado: uma abordagem promissora em paleoecologia. *Revista Brasileira de Paleontologia*, 15(3), 371-385.

- Aba “Model”
- Modelo: BIOCLIM (Envelope)

## Component: Build and Evaluate Niche Model



### Modules Available:

- Maxent
- BIOCLIM**

### Module: BIOCLIM

R packages: *ENMeval*, *dismo*

Batch

**Run**

### Species menu

Sus\_scrofa

Map

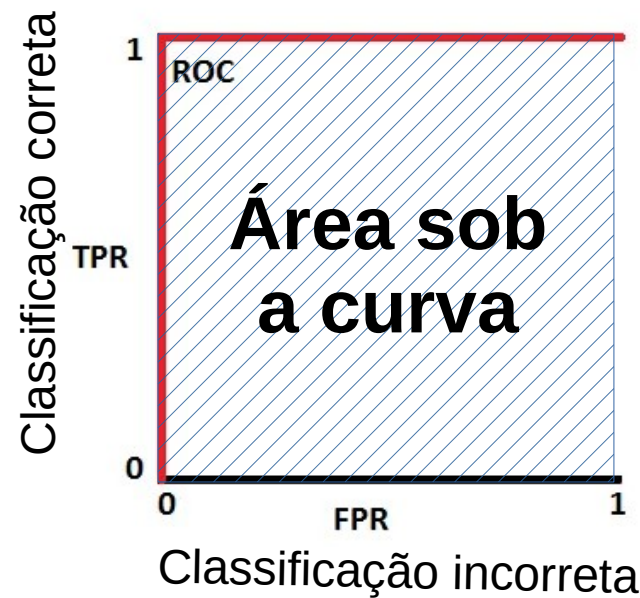
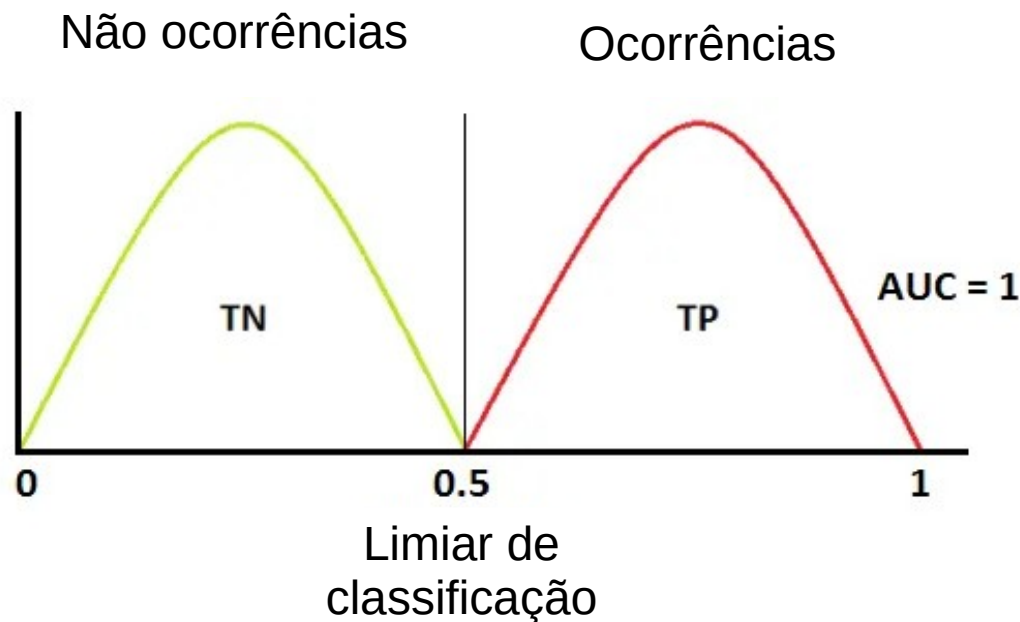
Occurrences

Results



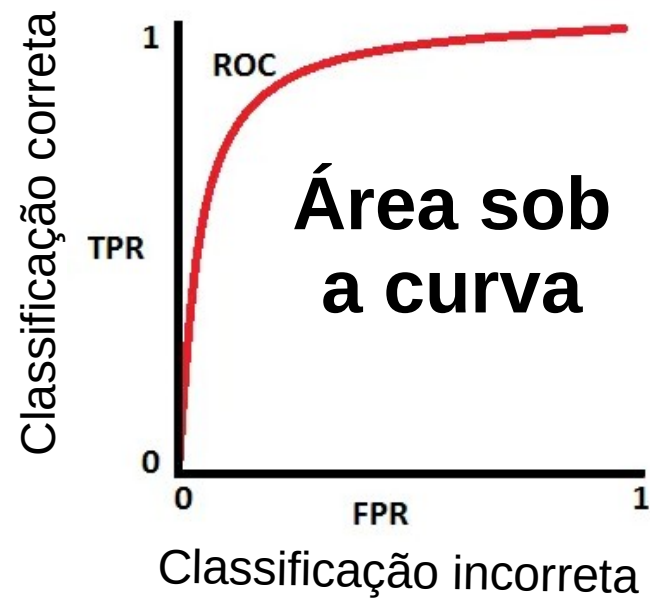
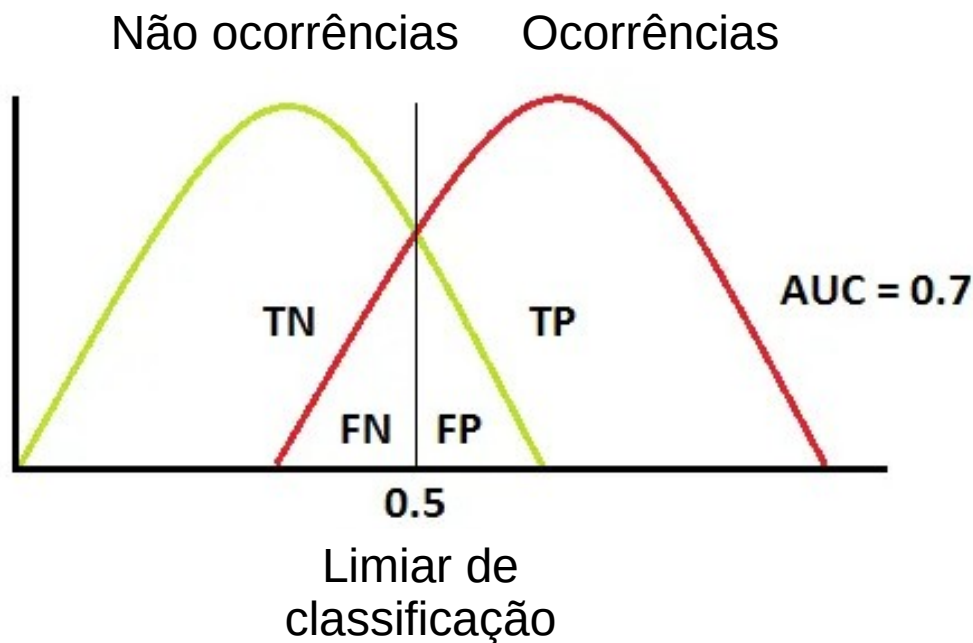
# Area sob a curva (AUC – Area Under Curve)

## Identificação perfeita de



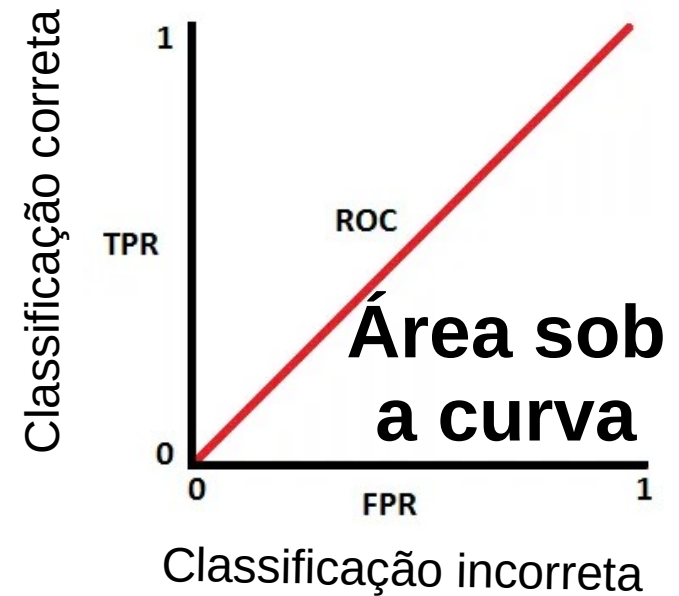
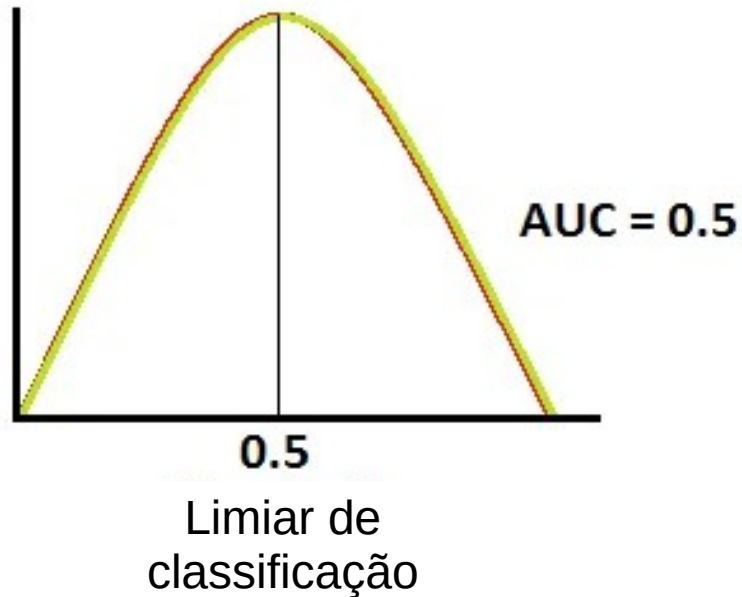
# Area sob a curva (AUC – Area Under Curve)

## Identificação parcial de



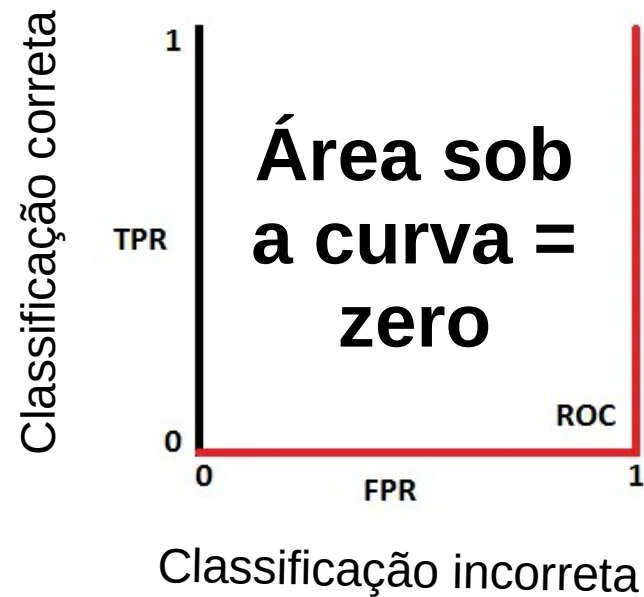
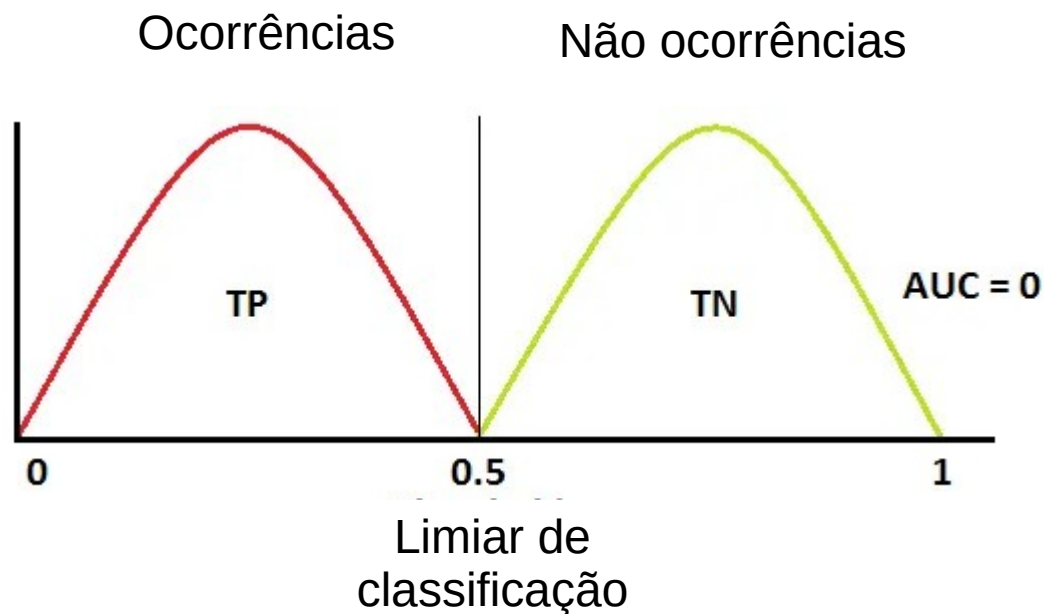
# Area sob a curva (AUC – Area Under Curve)

## Identificação aleatória



# Area sob a curva (AUC – Area Under Curve)

Identificação inversa de





- Aba “Results”
- Qualidade da calibração: auc.train
- Qualidade da validação: auc.val.avg

### Species menu

Sus\_scrofa

### Select model:

bioclim

### Log window

```
370536 total p  
> Sus scrofa | C  
with aggregati  
> Sus scrofa | E  
results.
```

Map

Occurrences

Results

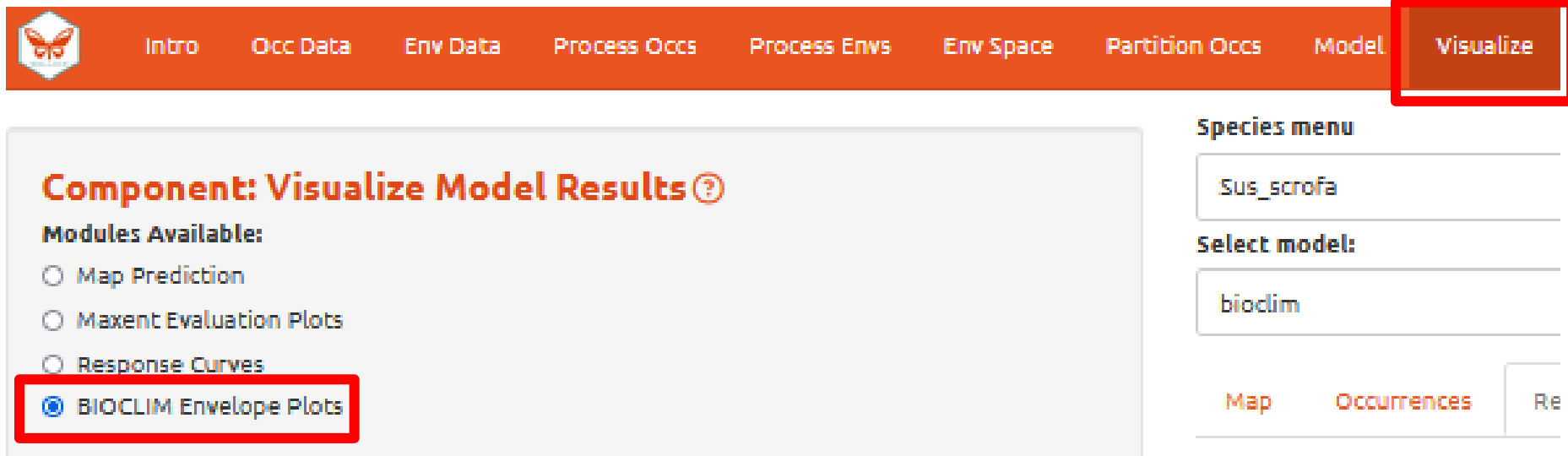
 Component Guidance

 Modu

*Evaluation statistics: full model and partition averages*

	auc.train	cbi.train	auc.diff.avg	auc.diff.sd	auc.val.avg
1	0.878	0.746	0.036	0.034	0.836

- Aba “Visualize”
- Selezione: “Bioclim Envelope Plots”



The screenshot displays the software's navigation bar and the 'Visualize Model Results' component. The 'Visualize' button in the top bar is highlighted with a red box. In the 'Modules Available' section, the 'BIOCLIM Envelope Plots' option is selected and also highlighted with a red box. The 'Species menu' on the right shows 'Sus\_scrofa' and the 'Select model:' dropdown is set to 'bioclim'. At the bottom right, there are tabs for 'Map', 'Occurrences', and 'Re'.

**Component: Visualize Model Results** ⓘ

**Modules Available:**

- Map Prediction
- Maxent Evaluation Plots
- Response Curves
- BIOCLIM Envelope Plots**

**Species menu**

Sus\_scrofa

**Select model:**

bioclim

Map Occurrences Re

- Alterando os eixos, observam-se as combinações entre cada variável
- Verde = Observações
- Vermelho = amostragem aleatória

● BIOCLIM Envelope Plots

Module: BIOCLIM Envelope Plots ?

R packages: *dismo*

Pick a bioclimatic variable number for each axis

Axis 1

1

Axis 2

2

Set threshold

0.9

Module Developers: *Jamie M. Kass, Robert Muscarella, Bruno Vilela, Robert P. Anderson*

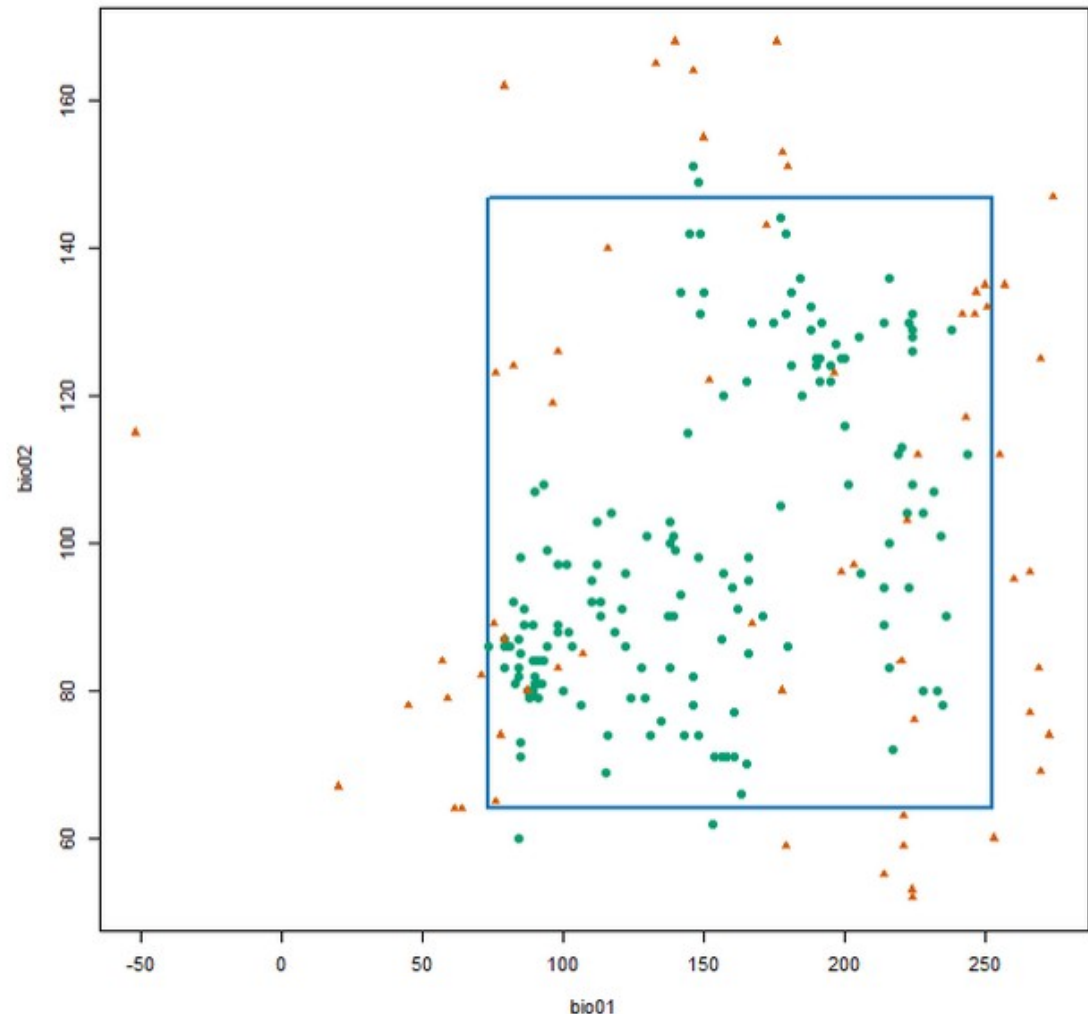
*dismo* references

**Species Distribution Modeling**

Package Developers: *Robert J. Hijmans, Steven Phillips, John Leathwick and Jane Elith*

[CRAN](#) | [documentation](#)

Map Occurrences Results **Component Guidance** Module Guidance Save



- Seleziona: “Map Prediction”
- Plot

## Component: Visualize Model Results



### Modules Available:

Map Prediction

Maxent Evaluation Plots

Response Curves

BIOCLIM Envelope Plots

---

### Module: Map Prediction

R packages: *dismo*

---

### Set threshold

No threshold 

Plot

- Vermelho: maior probabilidade
- Azul: menor probabilidade

#### Species menu

Sus\_scrofa

#### Select model:

bioclim

#### Log window

```
> Sus scrofa | Occurrences partitioned by checkerboard 2 method with aggregation factor 2.  
> Sus scrofa | BIOCLIM ran successfully and output evaluation results.  
> Sus scrofa | BIOCLIM model prediction plotted.
```

Map

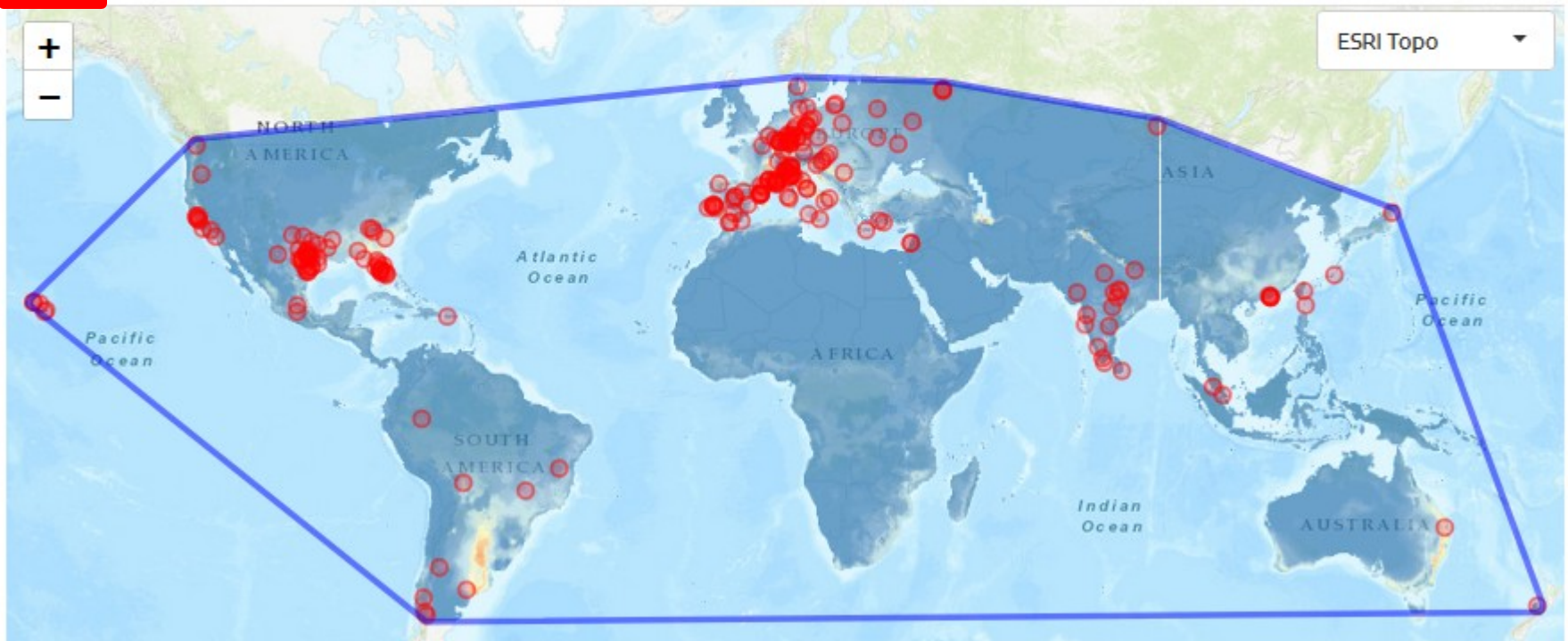
Occurrences

Results

Component Guidance

Module Guidance

Save



- Aba: Transfer
- “Transfer to new Time

- Região de estudo:  
“Same extent”
- Create

Intro Occ Data Env Data Process Occs Process Envs Env Space Partition Occs Model Visualize **Transfer**

**Component: Model Transfer** ⓘ

Modules Available:

- Transfer to New Extent
- Transfer to New Time**
- Transfer to User Environments
- Calculate Environmental Similarity

---

**Module: Transfer to New Time** ⓘ

R packages: *dismo*

---

**Step 1: Choose Study Region**

Select method

Same extent

You will use the same extent

**Create**

Species menu

Sus\_scrofa

Select model:

bioclim

Map Occurrences Results ⓘ Component Guide

- Base das variáveis de mudança climática:

WorldClim

- Período: 2070

- Modelo climático: ACCESS1-0

- Cenário RCP: 4.5 (tendencial)

## Step 2: Transfer

Transfer model to extent (red)

Select source of variables

WorldClim  ecoClimate

Select time period

2070

Select global circulation model

ACCESS1-0

Select RCP

4.5

Set threshold

No threshold

Transfer





## Component: Model Transfer

### Modules Available:

- Transfer to New Extent
- Transfer to New Time
- Transfer to User Environments
- Calculate Environmental Similarity

---

## Module: Calculate Environmental Similarity

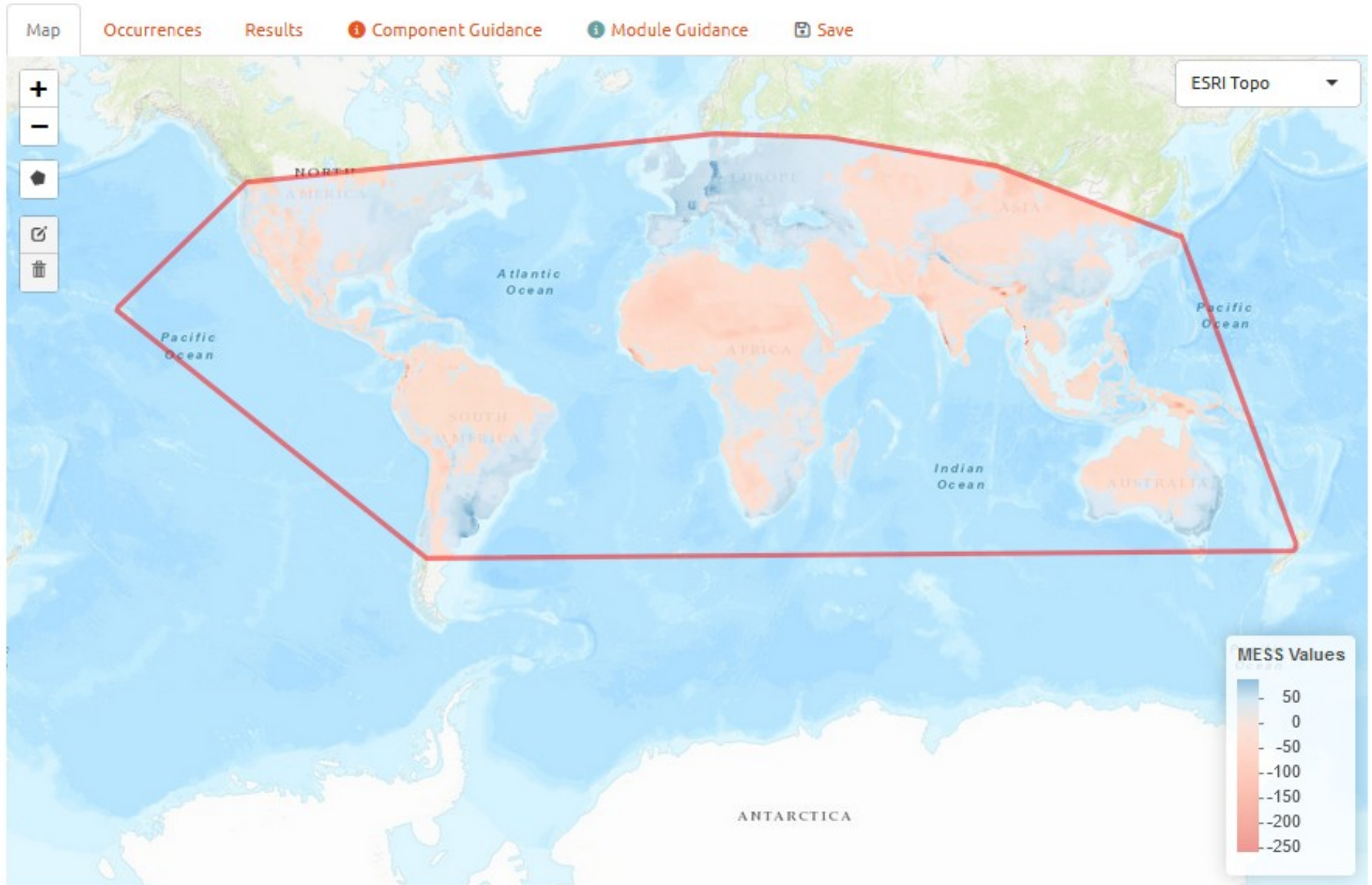
R packages: *dismo*

---

Calculate MESS for current extent

Calculate MESS


- Azul: maior similaridade
- Vermelho: maior mudança




*Note: To save your session code or metadata, use the Reproduce component*

## Save Session

By saving your session into an RDS file, you can resume working on it at a later time or you can share the file with a collaborator.


 The current session data is large, which means the downloaded file may be large and the download might take a long time.

 Save Session

## Download Data


Download data/results from analyses from currently selected module

Download shapefile of extent of transfer

 ZIP file


Download environmental variables of transfer (Select download file type)

GeoTIFF ▼

 ZIP file

Download transfer (Select download file type\*\*)

GeoTIFF ▼

 Transfer file

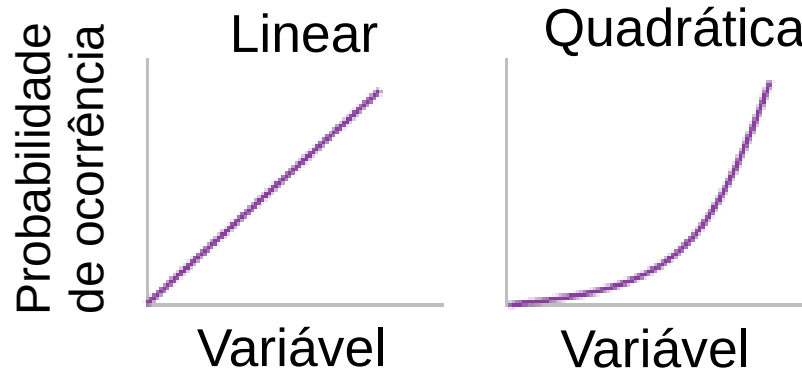
Download MESS (Select download file type\*\*)

GeoTIFF ▼

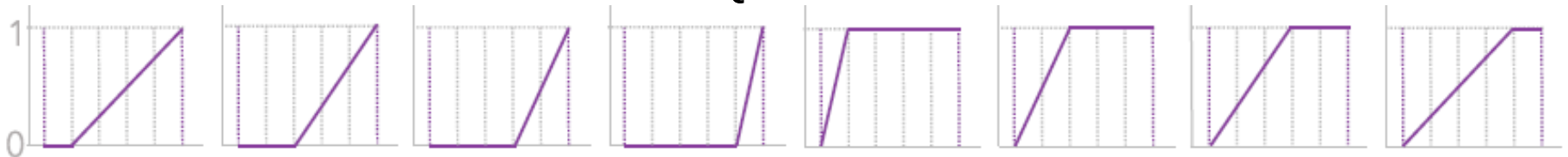
 MESS file

# Maxent

- Testa várias combinações de influências de variáveis.
- Compara: Previsão X Penalidade de complexidade
- Relações: lineares, quadráticas, quebras, produtos



## Quebras



**Produto:** Probabilidade de ocorrência = Variável A x Variável B



## Component: Build and Evaluate Niche Model ?

### Modules Available:

 Maxent BIOCLIM

### Module: Maxent ?

R packages: *ENMeval*, *dismo*, *maxnet*

( **NOTE** : see module guidance for troubleshooting tips if you are experiencing problems.)

### Select algorithm

 maxnet  maxent.jar

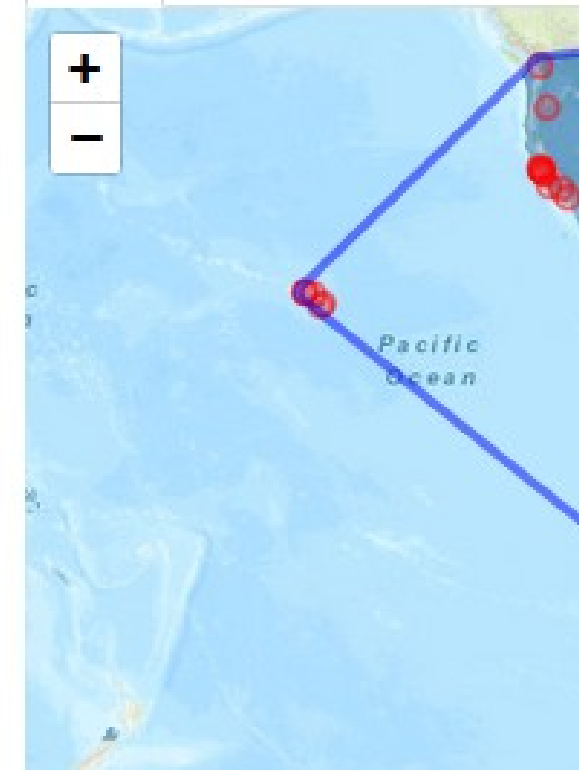
### Select feature classes *(flexibility of modeled response)*

key: L inear, Q uadratic, H inge, P roduct

 L  LQ  H  LQH  LQHP

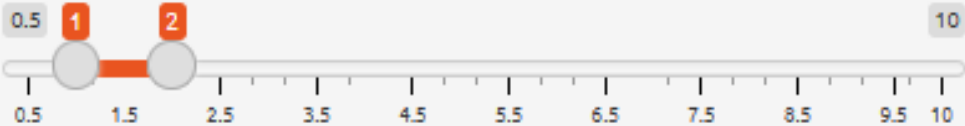
### Species menu

### Select model:

[Map](#)[Occurrences](#)[Results](#)

- Está usando variáveis categóricas?
  - Não
- Clamping (não extrapolar para valores de variáveis além dos registrados para as espécies)
  - Sim
- Processamento paralelo:
  - Sim (deixe o número de processadores [cores] como automaticamente – varia para cada computador)

Select regularization multipliers (*penalty against complexity*)



Multiplier step value

1

Are you using a categorical variable?

NO

Clamping?

TRUE

Parallel?

TRUE

Specify the number of cores (max. 4)

3

Batch

Run

# Atividade

- Compare os modelos Maxent e Bioclim quanto a:
  - Validação da qualidade do modelo.
  - Mapa de predição
  - Projeção para mudanças climáticas
  - Mapa de similaridade ambiental da projeção

- Aba “Results”
- Qualidade da calibração: auc.train
- Qualidade da validação: auc.val.avg

### Species menu

Sus\_scrofa

### Select model:

fc.LQ\_rm.1

### Log window

```
GCM ACCESS1-0 under RCP 4.5.
> Sus scrofa | Generated MESS map.
> Sus scrofa | Maxent ran successfully and output
results for 2 models (Regularization multiplier 1
Feature classes: [LQ]), using maxnet with clamp
```

Map Occurrences **Results** Component Guidance Module Guidance Save

Evaluation **Lambdas**

*Evaluation statistics: full model and partition averages*

	fc	rm	tune.args	auc.train	cbi.train	auc.diff.avg	auc.diff.sd	auc.val.avg
1	LQ	1	fc.LQ_rm.1	0.9	0.925	0.033	0.012	0.876
2	LQ	2	fc.LQ_rm.2	0.893	0.941	0.033	0.019	0.871



- Aba “Lambdas”
- Coeficientes (influência) das variáveis

Map

Occurrences

Results

 Component Guidance

 Module Guidance

Evaluation

Lambdas

### *Maxent Lambdas File*

```

      bio05          bio08          bio09          bio13          bio17
7.058144e-04  3.952946e-03 -1.304549e-04  3.037027e-03  5.250937e-04
      bio18          bio19      I(bio02^2)      I(bio03^2)      I(bio04^2)
-5.391945e-04  2.400944e-03 -8.206660e-06 -2.882603e-04 -1.814565e-08
      I(bio07^2)      I(bio08^2)      I(bio11^2)      I(bio12^2)      I(bio15^2)
-5.036809e-08 -3.286015e-06 -2.623125e-05 -1.217354e-07 -3.695331e-05
      I(bio17^2)      I(bio19^2)
2.840267e-06 -2.215379e-06

```

# Variáveis bioclimáticas

BIO1 = Temperatura média anual

BIO2 = Amplitude térmica diurna (Média mensal (max temp - min temp))

BIO3 = Isotermalidade (BIO2/BIO7) ( $\times 100$ )

BIO4 = Sazonalidade da temperatura (desvio padrão  $\times 100$ )

BIO5 = Temperatura máxima do mês mais quente

BIO6 = Temperatura mínima do mês mais frio

BIO7 = Amplitude térmica anual (BIO5-BIO6)

BIO8 = Temperatura média do quadrimestre mais úmido

BIO9 = Temperatura média do quadrimestre mais seco

BIO10 = Temperatura média do quadrimestre mais quente

BIO11 = Temperatura média do quadrimestre mais frio

BIO12 = Precipitação anual

BIO13 = Precipitação do mês mais úmido

BIO14 = Precipitação do mês mais seco

BIO15 = Sazonalidade da precipitação (coeficiente de variação)

BIO16 = Precipitação do quadrimestre mais úmido

BIO17 = Precipitação do quadrimestre mais seco

BIO18 = Precipitação do quadrimestre mais quente

BIO19 = Precipitação do quadrimestre mais frio

Booth, T. H., Nix, H. A., Busby, J. R., & Hutchinson, M. F. (2014). BIOCLIM: the first species distribution modelling package, its early applications and relevance to most current MAXENT studies. *Diversity and Distributions*, 20(1), 1-9.

- ABA “Visualize”
- Response curves
- Variável: bio05 (Temperatura máxima do mês mais quente)



Intro

Occ Data

Env Data

Process Occs

Process Envs

Env Space

Partition Occs

Model

Visualize

## Component: Visualize Model Results ?

### Modules Available:

- Map Prediction
- Maxent Evaluation Plots
- Response Curves
- BIOCLIM Envelope Plots

### Module: Plot Response Curves ?

R packages: *dismo*, *maxnet*

### Select variable

bio05

### Species menu

Sus\_scrofa

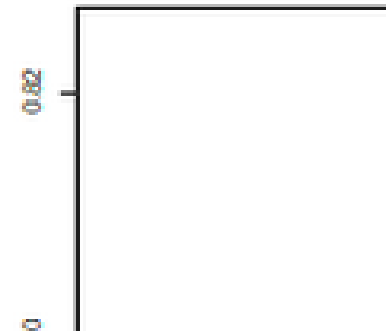
### Select model:

Fc.LQ\_rm.1

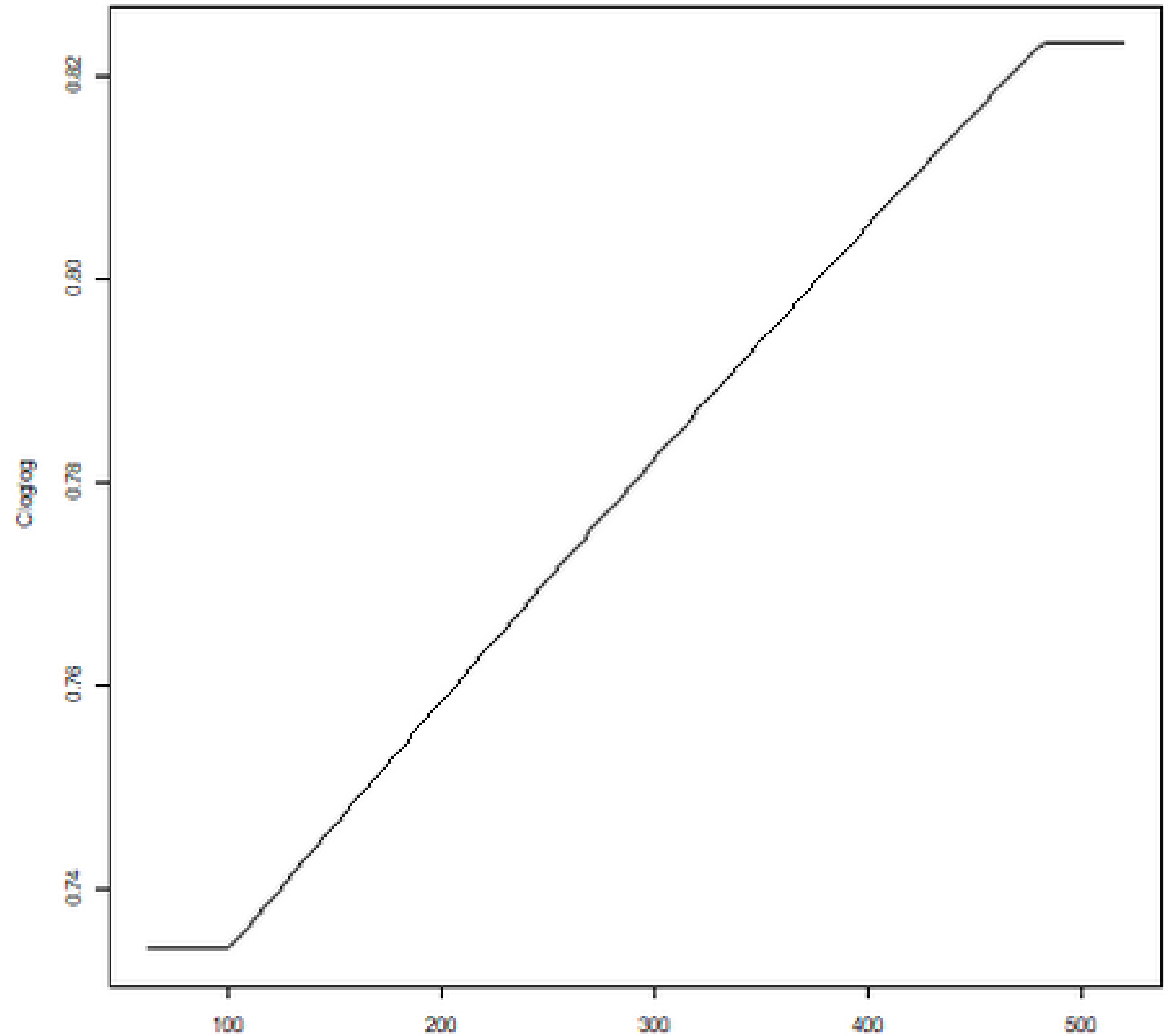
Map

Occurrences

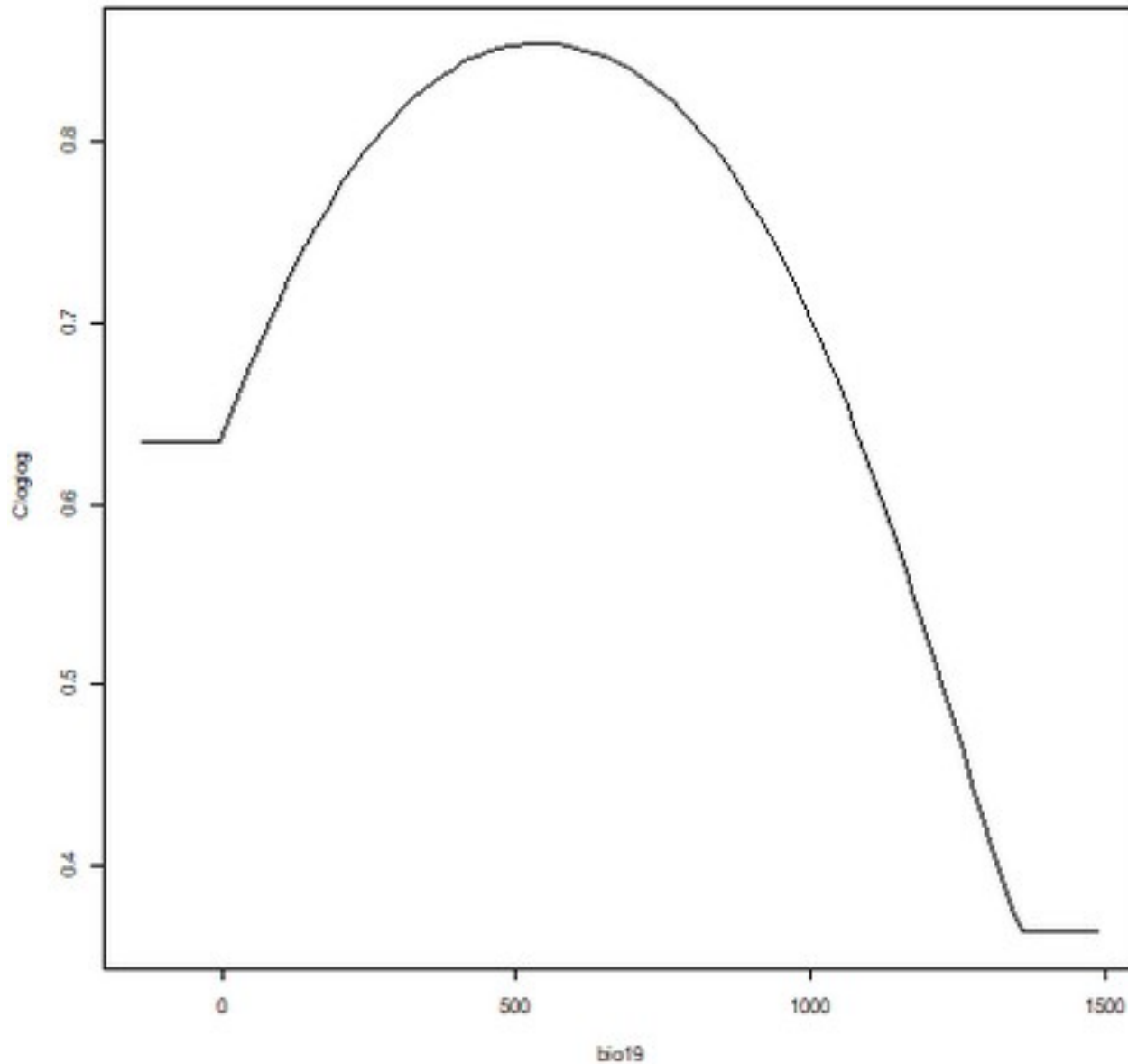
Re



- Aba  
“Results”

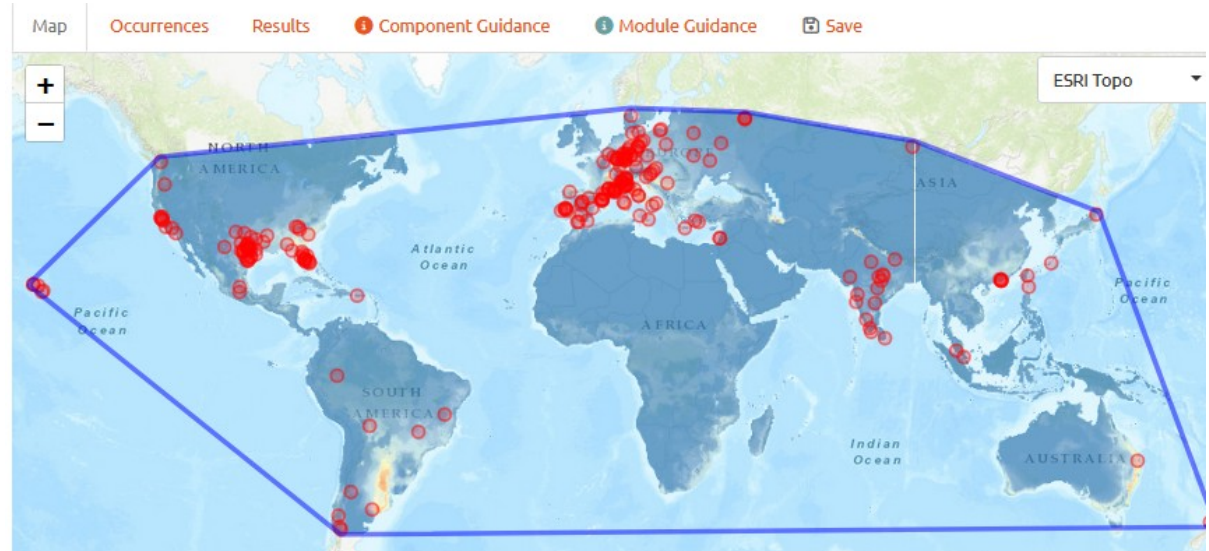


- Altere para “bio19” (precipitação no quadrimestre mais frio)

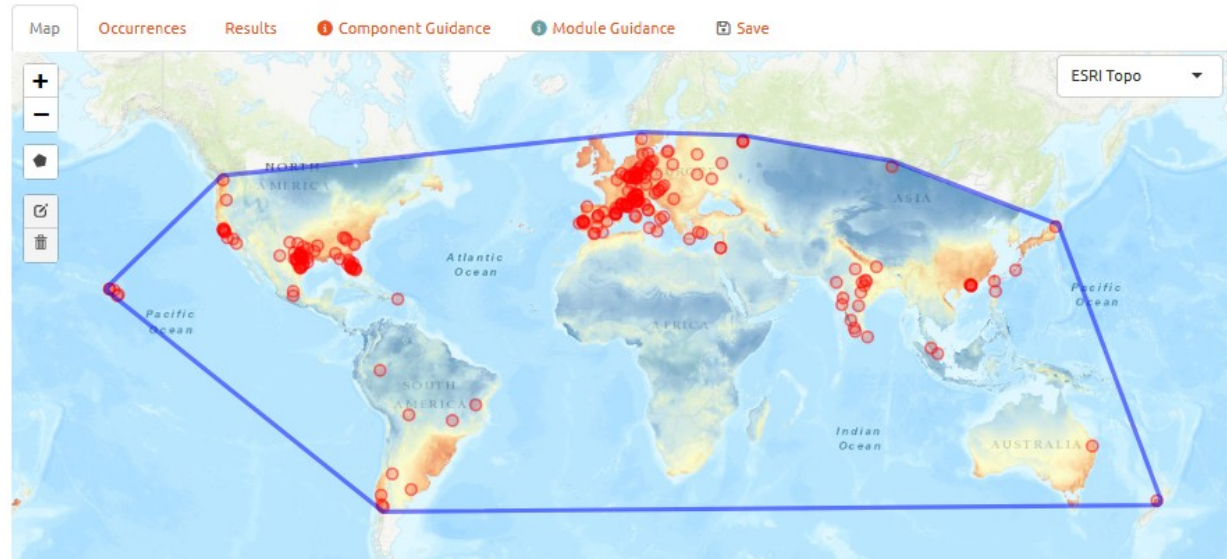


- Visualize o mapa de predição do Maxent
  - O padrão espacial está diferente do mapa do BioClim?

BioClim



Maxent



- Faça a predição do Maxent para o futuro
- Visualize o mapa de similaridade ambiental da predição

Predição



Similaridade ambiental



- Aba “Reproduce”
- Escolha “Session code”
- Download



## Component: Reproduce

### Modules Available:

Session Code

Metadata


Reference Packages

### Module: Download Session Code ?

R packages: *rmarkdown*, *knitr*

#### Select download file type

Rmd

 Download Session Code

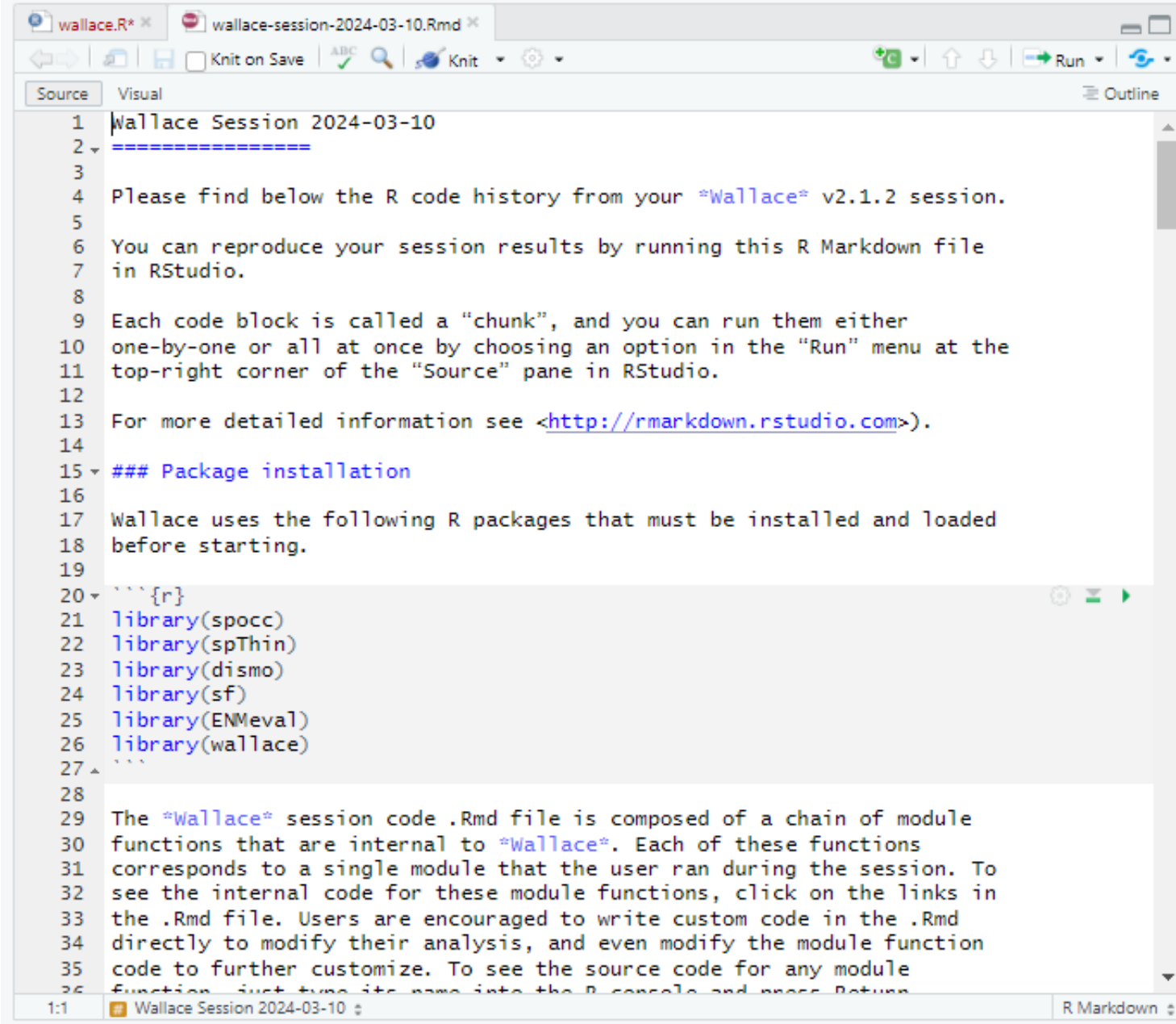
## Module: *Session Code*

### BACKGROUND

Over the decade of the 2010s, scientific practice increasingly the area of modeling species niches/distributions has advanced community-driven standards (see Fitzpatrick et al. 2021 for an et al. 2019), standardized metadata frameworks (RMMS, Metadata reporting (ODMAP, Zurell et al. 2020). These tools facilitate the model, indicating whether it meets minimal standards for application leveraging *EIMeal 2.0* and *rangeModelMetadata*, *Wallace* now also form the basis of ODMAP reporting) and allows the user *Wallace* promotes documentation and downstream assessment information that includes sources of input data, methodology (Module: *Download Session Code*) is a file that can be re-run in and dependent packages). Many intermediate and advanced *Additionally, Wallace* now provides citations of the particular *Reference Packages*).

Via the *Session Code* module, the user can download files that executable code that can reproduce them). This functionality





Source Visual Outline

```
1 Wallace Session 2024-03-10
2 =====
3
4 Please find below the R code history from your *Wallace* v2.1.2 session.
5
6 You can reproduce your session results by running this R Markdown file
7 in RStudio.
8
9 Each code block is called a "chunk", and you can run them either
10 one-by-one or all at once by choosing an option in the "Run" menu at the
11 top-right corner of the "Source" pane in RStudio.
12
13 For more detailed information see <http://rmarkdown.rstudio.com>).
14
15 ### Package installation
16
17 Wallace uses the following R packages that must be installed and loaded
18 before starting.
19
20 ```{r}
21 library(spocc)
22 library(spThin)
23 library(dismo)
24 library(sf)
25 library(ENMeval)
26 library(wallace)
27 ```
28
29 The *Wallace* session code .Rmd file is composed of a chain of module
30 functions that are internal to *Wallace*. Each of these functions
31 corresponds to a single module that the user ran during the session. To
32 see the internal code for these module functions, click on the links in
33 the .Rmd file. Users are encouraged to write custom code in the .Rmd
34 directly to modify their analysis, and even modify the module function
35 code to further customize. To see the source code for any module
36 function, just type its name into the R console and press Return.
```

1:1 Wallace Session 2024-03-10 R Markdown

# Muito obrigado!



Dúvidas, comentários, sugestões?

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