

# Practical Introduction to Species Distribution Models

Blas M. Benito

EECRG

University of Bergen

Email: [blasbenito@gmail.com](mailto:blasbenito@gmail.com)

*Blas M. Benito*

# **PROGRAM**

# SESSIONS

- **Day 1**
  - Introduction to SDMs
  - The modelling workflow
  - Applications of SDMs
  - Limitations of SDMs
  - Computer programs
  - Preparation of presence records and environmental predictors

# SESSIONS

- **Day 2**
  - Calibrating SDMs with different algorithms: theory and practice
  - GLM, GAM, Random Forest, and Maxent
  - Evaluating SDMs
  - Projection of SDMs on space and time
- **Day 3**
  - Practical workshop
- **Day 4**
  - Presentation of results of the practical workshop
  - Brief introduction to Dynamic SDMs
  - Discussion and conclusions

# INTRODUCTION

# WHAT IS A MODEL?

Simplified description of a physical system that still has properties appearing in the real system (Joly 1988).

Objectives:

- Offer a downsized version of the system
- Allows for a better understanding
- Hypothesis generation and testing
- Exploring “what-if” scenarios
- Forecasting and hindcasting
- Support decision making
- Have fun and learn stuff

# **PRESENCE/ABSENCE AND ECOLOGICAL NICHE**

# LOOKING AT THE SYSTEM

presence / absence



# LOOKING AT THE SYSTEM

presence / absence



abundance



# LOOKING AT THE SYSTEM

presence / absence



abundance



Habitat structure



# LOOKING AT THE SYSTEM

presence / absence



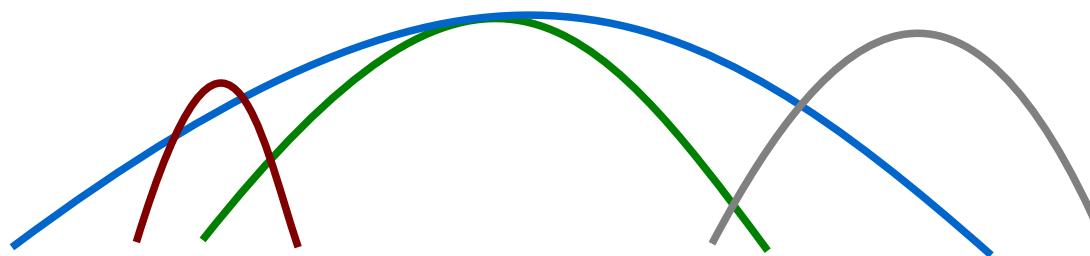
abundance



Habitat structure

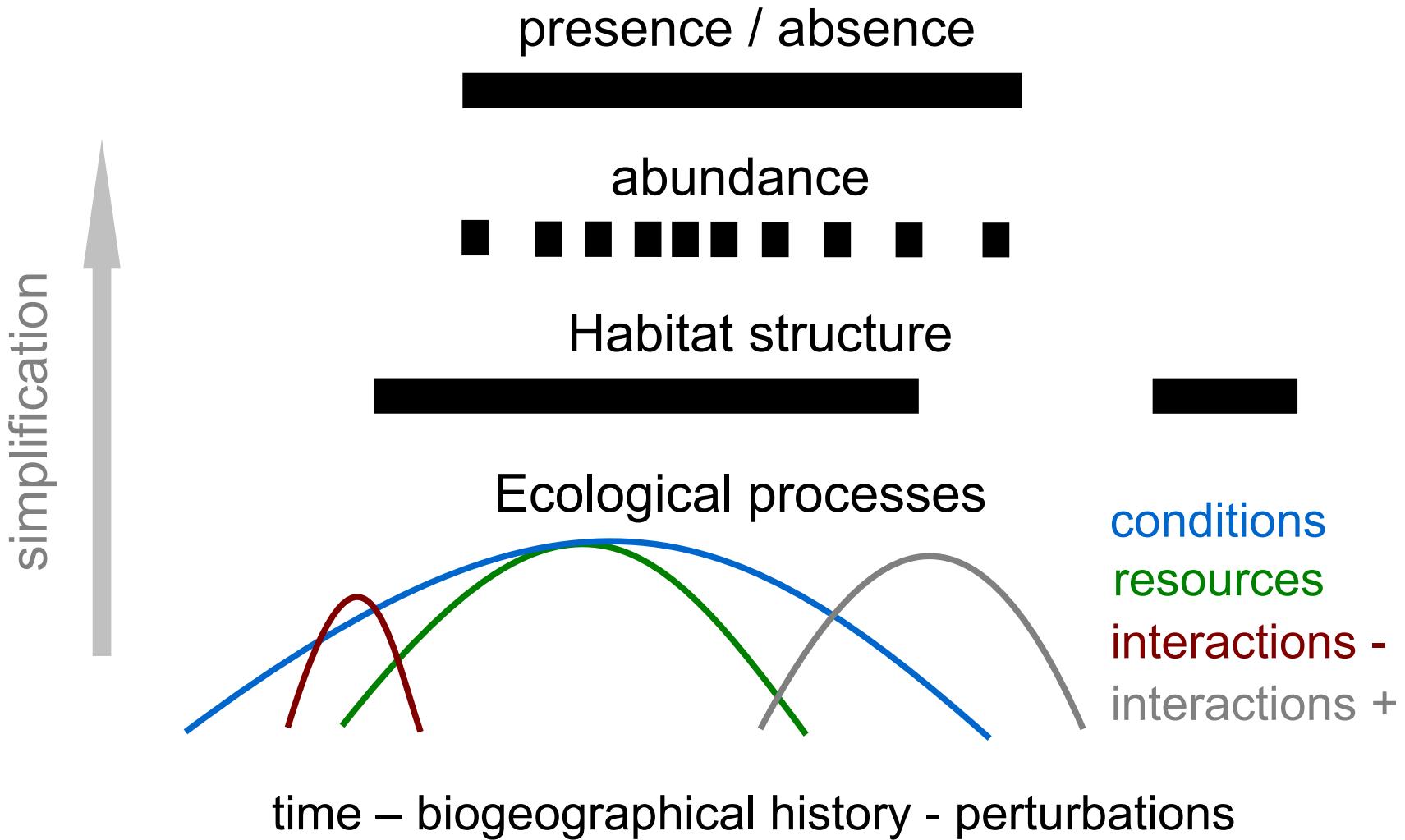


Ecological processes



conditions  
resources  
interactions -  
interactions +

# LOOKING AT THE SYSTEM



# ECOLOGICAL NICHE STRUCTURE



ver McInerny y Etienne 2012 J Biogeogr

**THERE ARE DIFFERENT WAYS TO  
MODEL ECOLOGICAL NICHES**

# CORRELATIVE MODELS



Abiotic factors



presence



= we only know the pattern, not the process

# MODELOS CORRELATIVOS

## Frontiers in Zoology

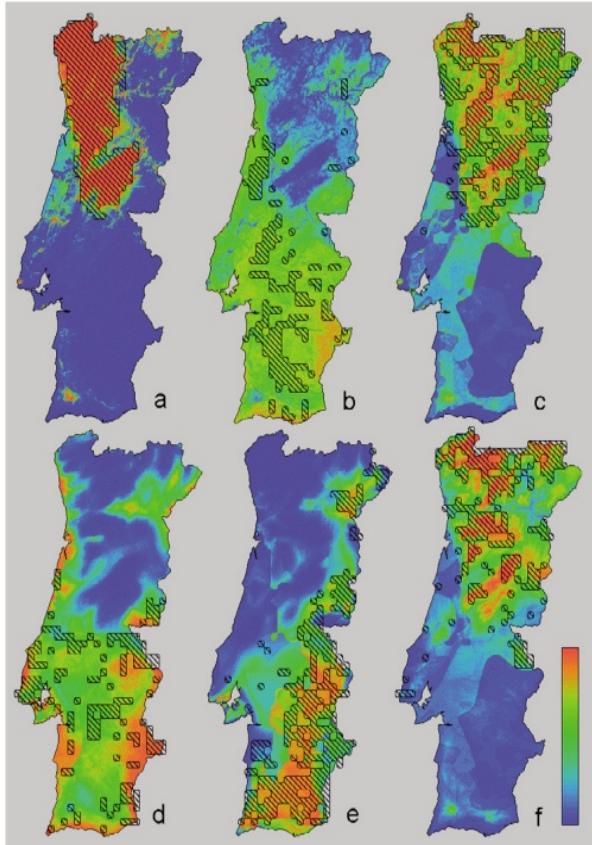


Research

Open Access

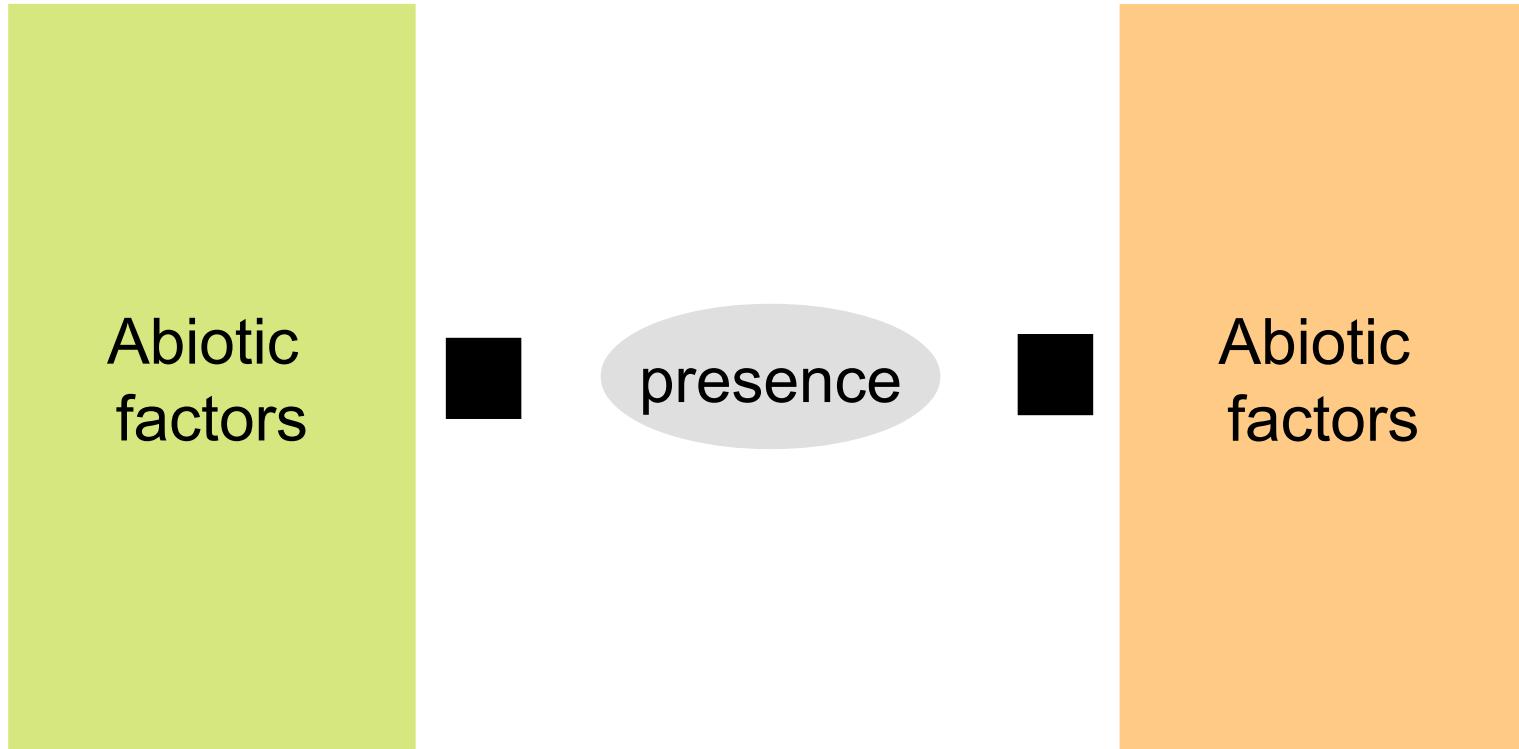
### From descriptive to predictive distribution models: a working example with Iberian amphibians and reptiles

JW Arntzen\*



**Figure 1**  
**Descriptive distribution models for six amphibian species across Portugal.** Descriptive distribution models for six amphibian species across Portugal. Models are derived with stepwise logistic regression analysis of the dependent variable 'presence-absence of the target species' against 13 independent ecological variables (details see text and Table 1). The estimated probability of occurrence (g) ranges from 0 (blue) to 1 (red). Composite colours represent intermediate probabilities as in the colour scale bar. Species are: a) *Chioglossa lusitanica*, b) *Pleurodeles waltl*, c) *Triturus marmoratus*, d) *T. pygmaeus*, e) *Alytes cisternasii* and f) *A. obstetricans*. Recorded presences over the 10 × 10 km UTM-grid are shown by black shadings, after Godinho et al. [34].

# CORRELATIVE MODELS WITH INTERACTIONS



■ = we only know the pattern, not the process

# MODELOS CORRELATIVOS CON INTERACCIONES

Improving species distribution models using biotic interactions:  
a case study of parasites, pollinators and plants

Tereza Cristina Giannini, Daniel S. Chapman, Antonio Mauro Saraiva, Isabel Alves-dos-Santos  
and Jacobus C. Biesmeijer

Ecography 36: 649–656, 2013

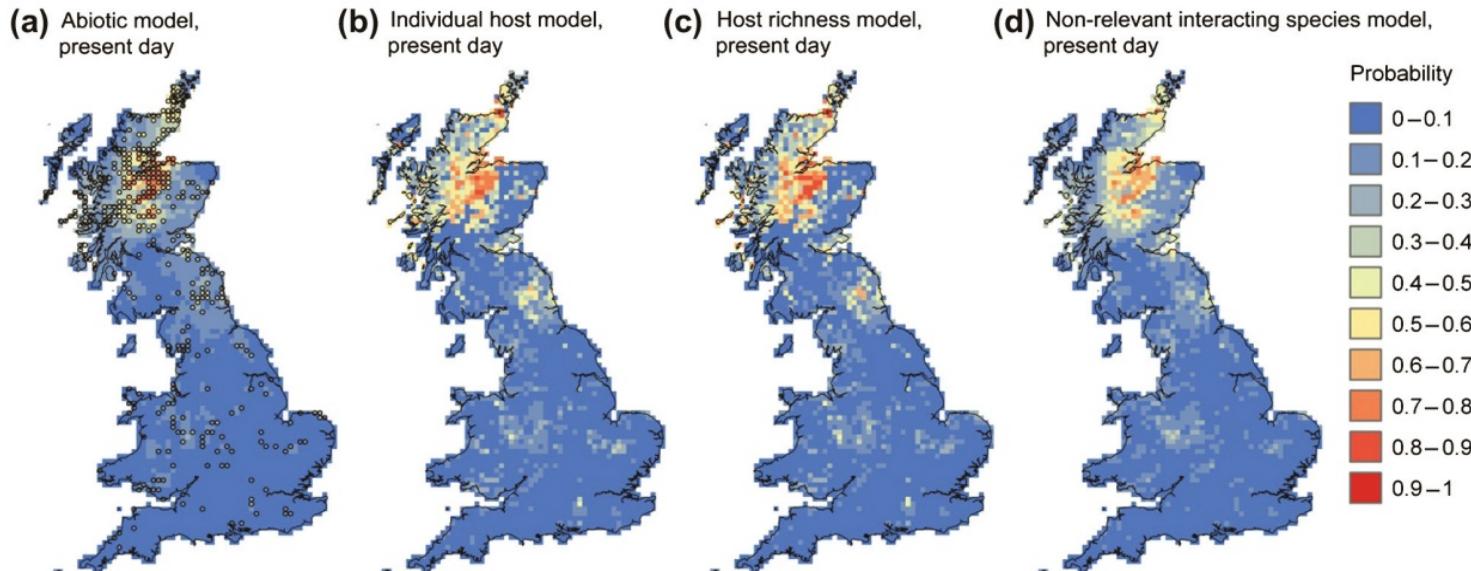
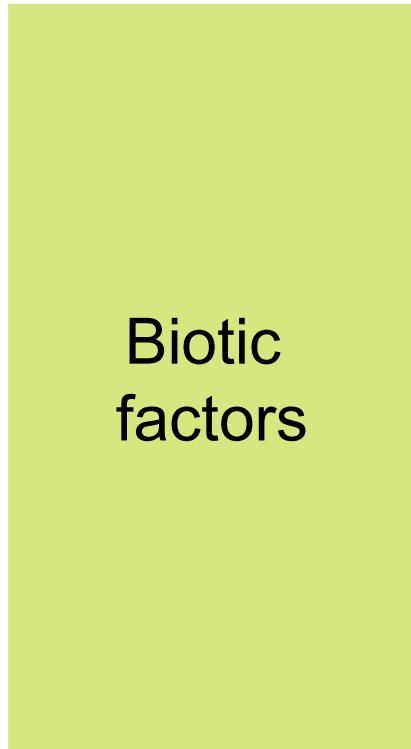


Figure 2. Projections of *Bombus bohemicus* occurrence areas. Top: present climatic conditions using (a) abiotic information only; (b) the most widespread host *B. lucorum*; (c) richness of its three host species (*B. lucorum*, *B. magnus* and *B. cryptarum*); (d) non-relevant interacting species (*B. terrestris*).

# ECO-PHYSIOLOGICAL MODELS



presence



= known process (experimentation)

# MODELOS ECO-FISIOLÓGICOS



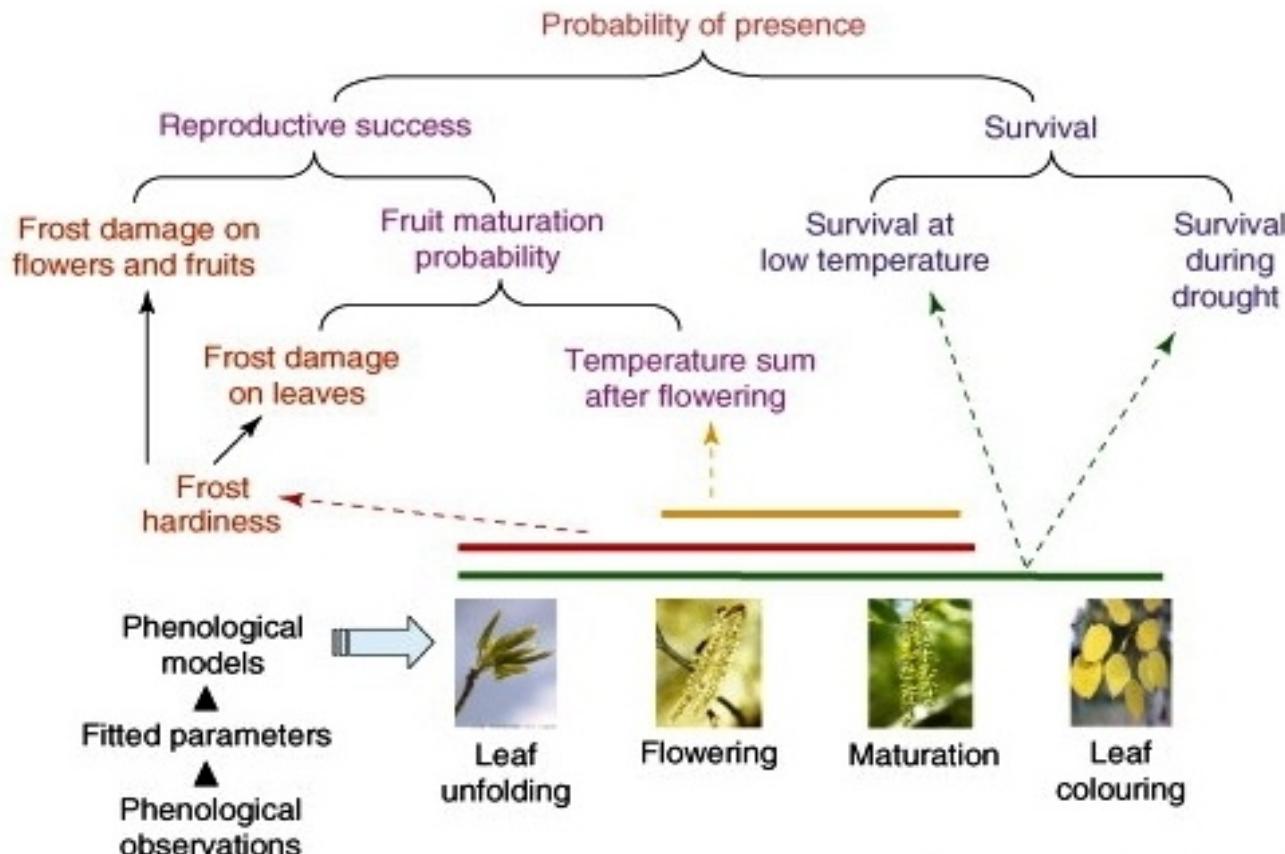
Review

TRENDS in Ecology and Evolution Vol.22 No.7

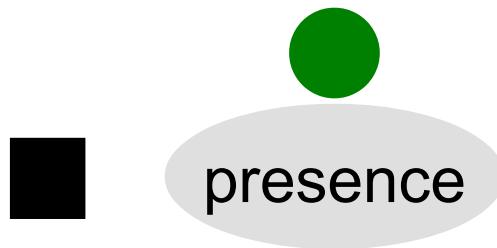
Full text provided by www.sciencedirect.com  
ScienceDirect

## Shifting plant phenology in response to global change

Elsa E. Cleland<sup>1</sup>, Isabelle Chuine<sup>2</sup>, Annette Menzel<sup>3</sup>, Harold A. Mooney<sup>4</sup>  
and Mark D. Schwartz<sup>5</sup>



# HYBRID MODELS



● = known process (population viability)

■ = correlation, only pattern

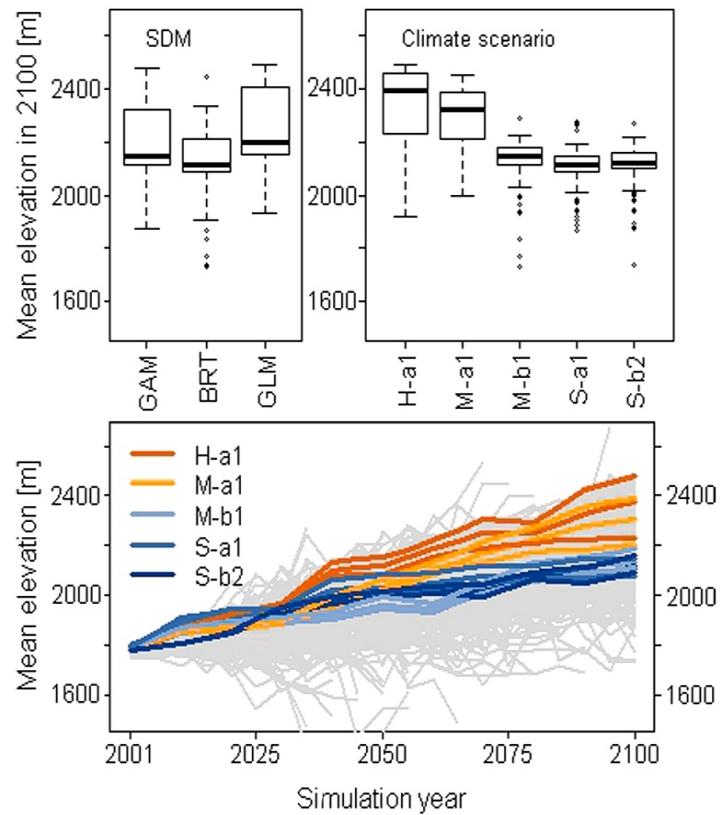
# MODELOS HÍBRIDOS

## Uncertainty in predictions of range dynamics: black grouse climbing the Swiss Alps

Damaris Zurell, Volker Grimm, Eva Rossmannith, Niklaus Zbinden, Niklaus E. Zimmermann and Boris Schröder

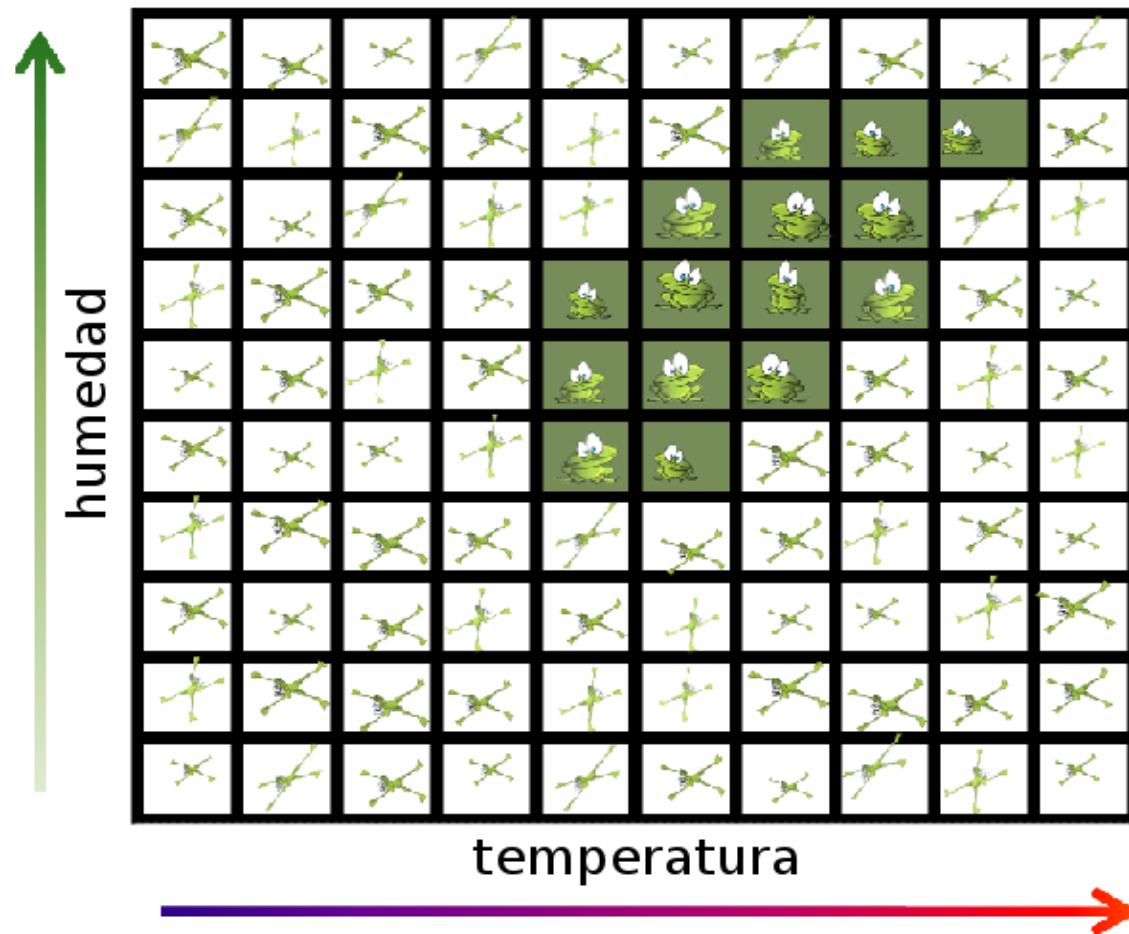
Ecography 34: 001–014, 2011

Figure 2. Mean elevation occupied by black grouse for scenarios of climate change. Bottom: grey lines show mean elevations across all simulations, coloured lines those for default IBM parameterisation (cf. Table 2) across different SDMs and climate scenarios. Top: box-plots depict variation of mean elevations predicted for the end of 21st century (2100) and for different SDMs and climate scenarios.

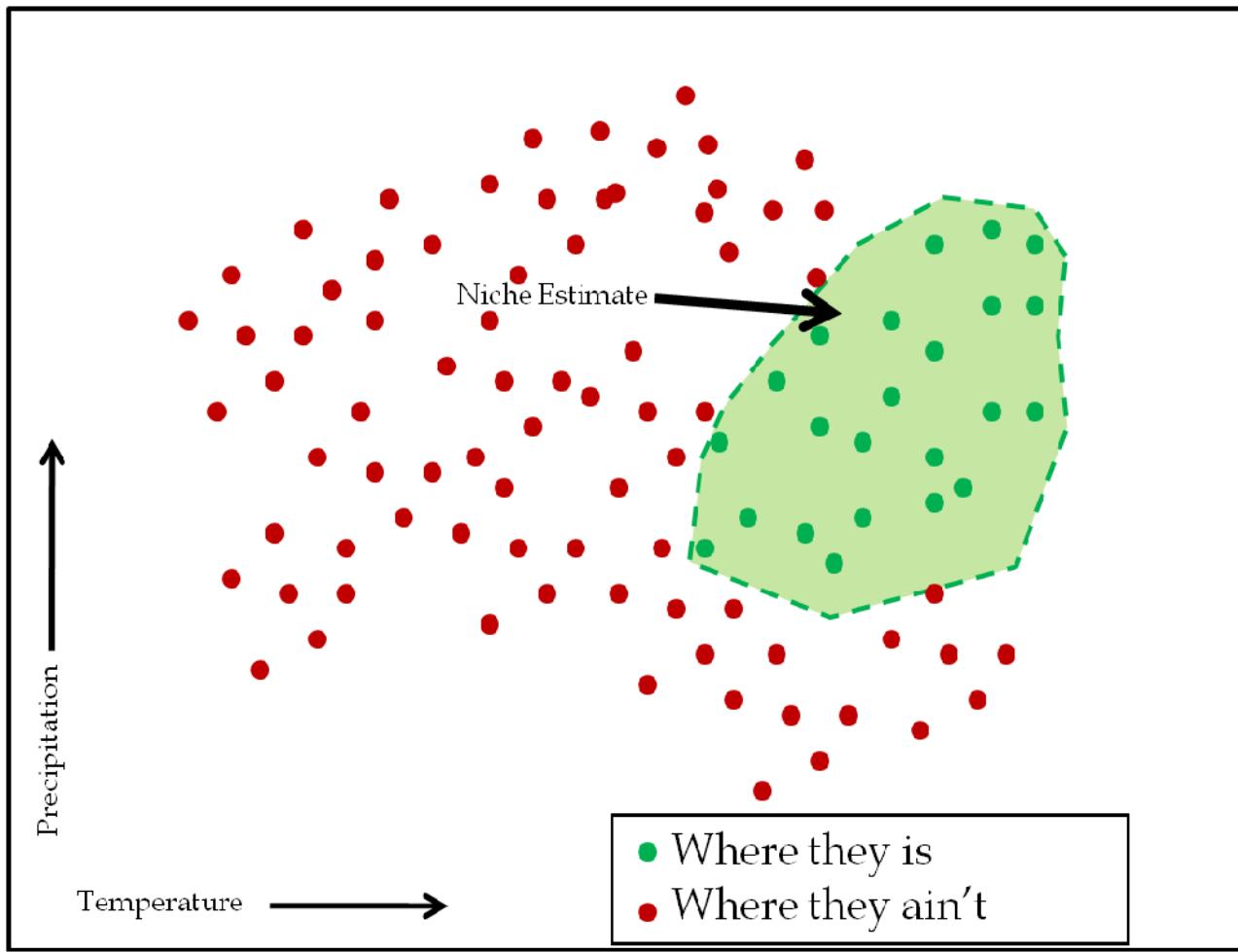


# **SPECIES DISTRIBUTION MODELLING**

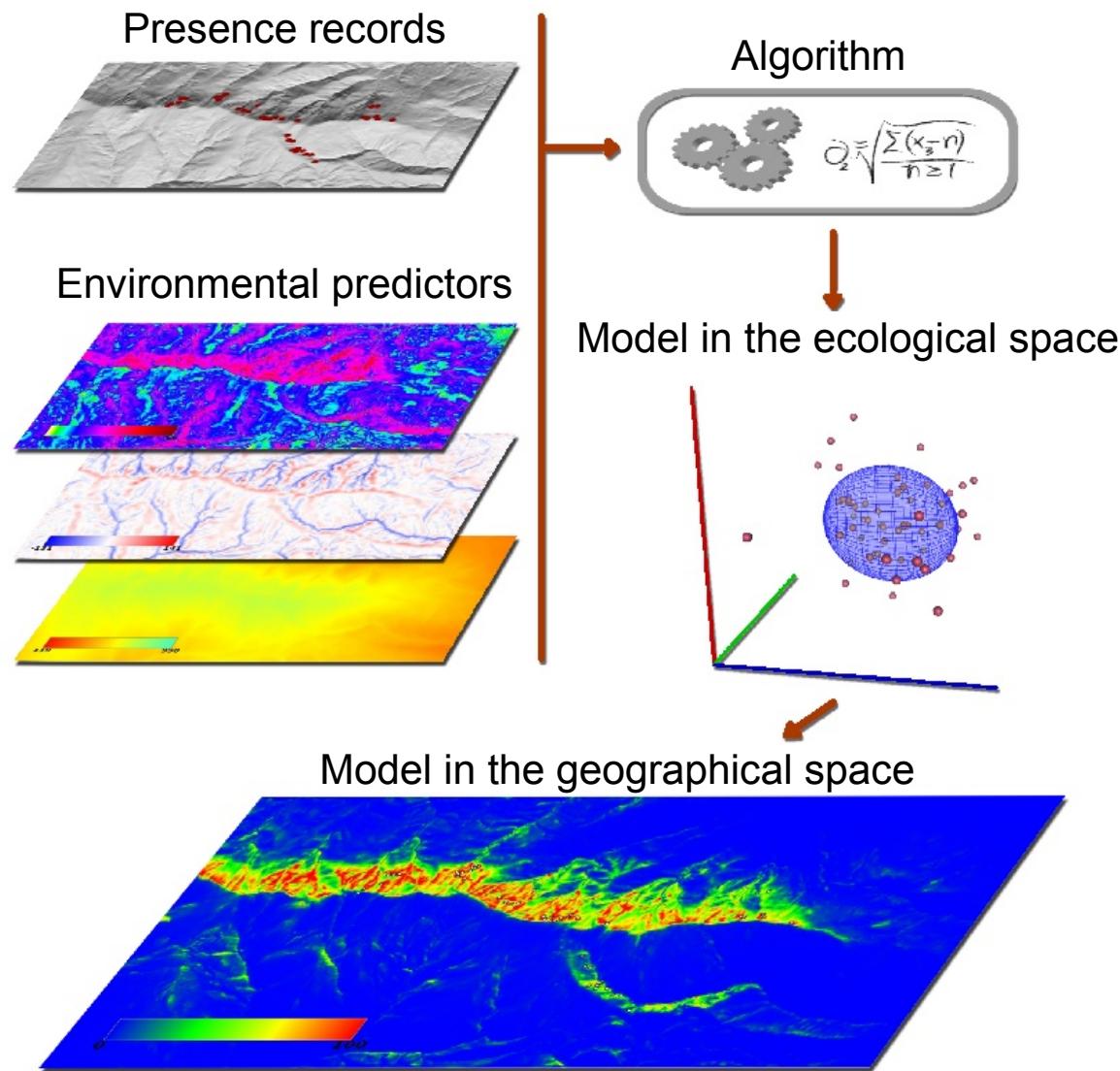
# EXPERIMENTAL APPROACH



# CORRELATIVE APPROACH



# CORRELATIVE APPROACH



# KEY ASSUMPTIONS

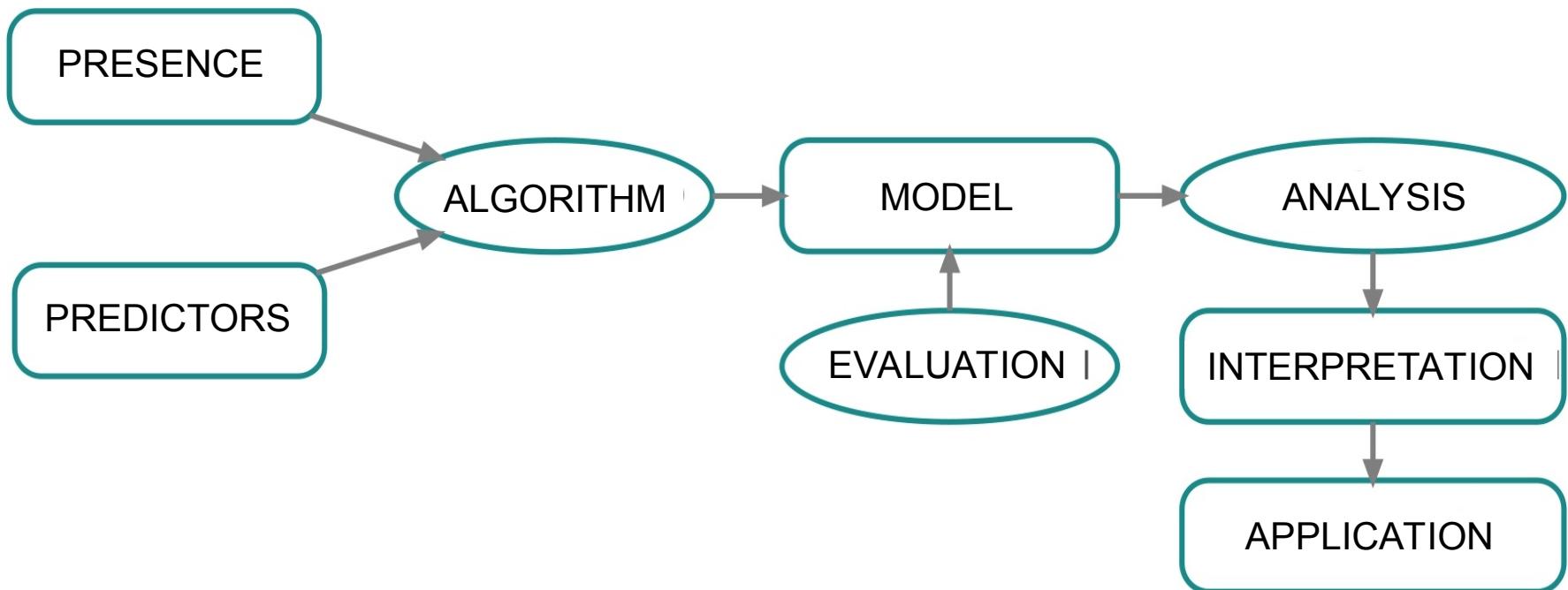
- SPECIES ARE IN EQUILIBRIUM WITH CLIMATE:* The species is present in every climatically suitable area, and absent from every unsuitable one.
- ECOLOGICAL NICHE IS STATIC IN TIME:* Niche dimensions and hypervolume's shape remains constant over time.

# **THE MODELLING WORKFLOW**

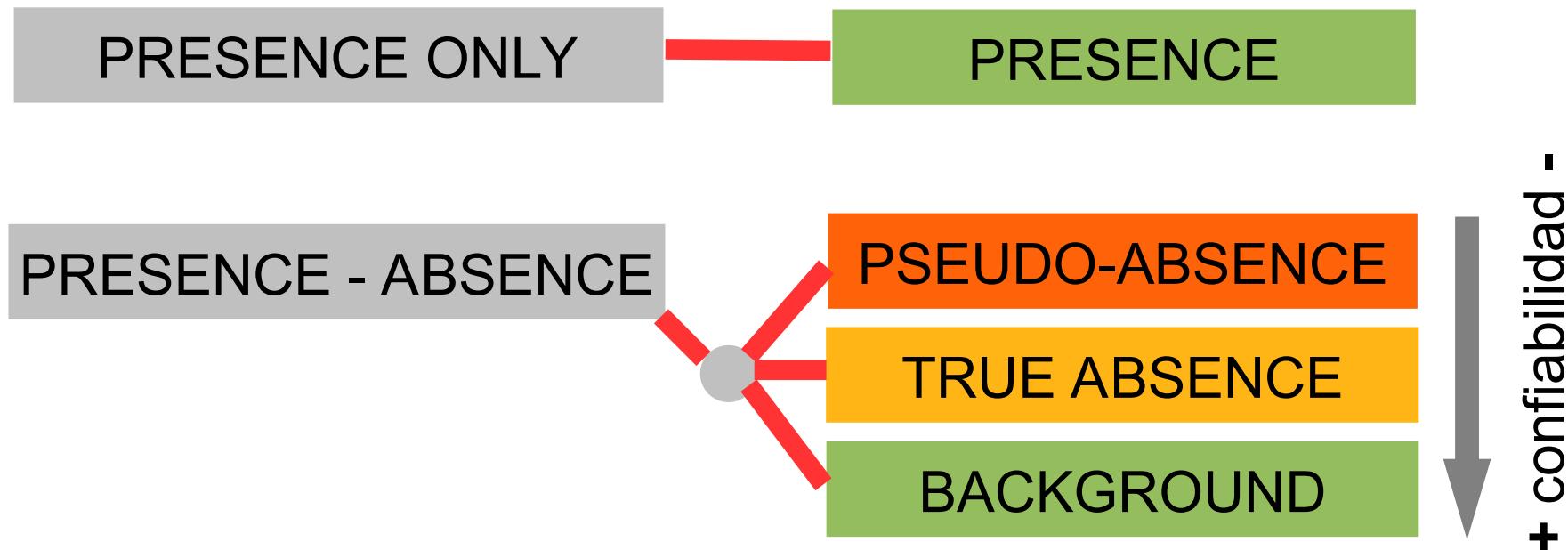
# THEORETICAL FORMULATION

- Define objectives
- Gather knowledge on the target species
- Select meaningful predictors
- Select the right tools
- Draw a workflow, have a plan!

# FLUJO DE TRABAJO



# TYPES OF PRESENCES



# PRESENCE ONLY

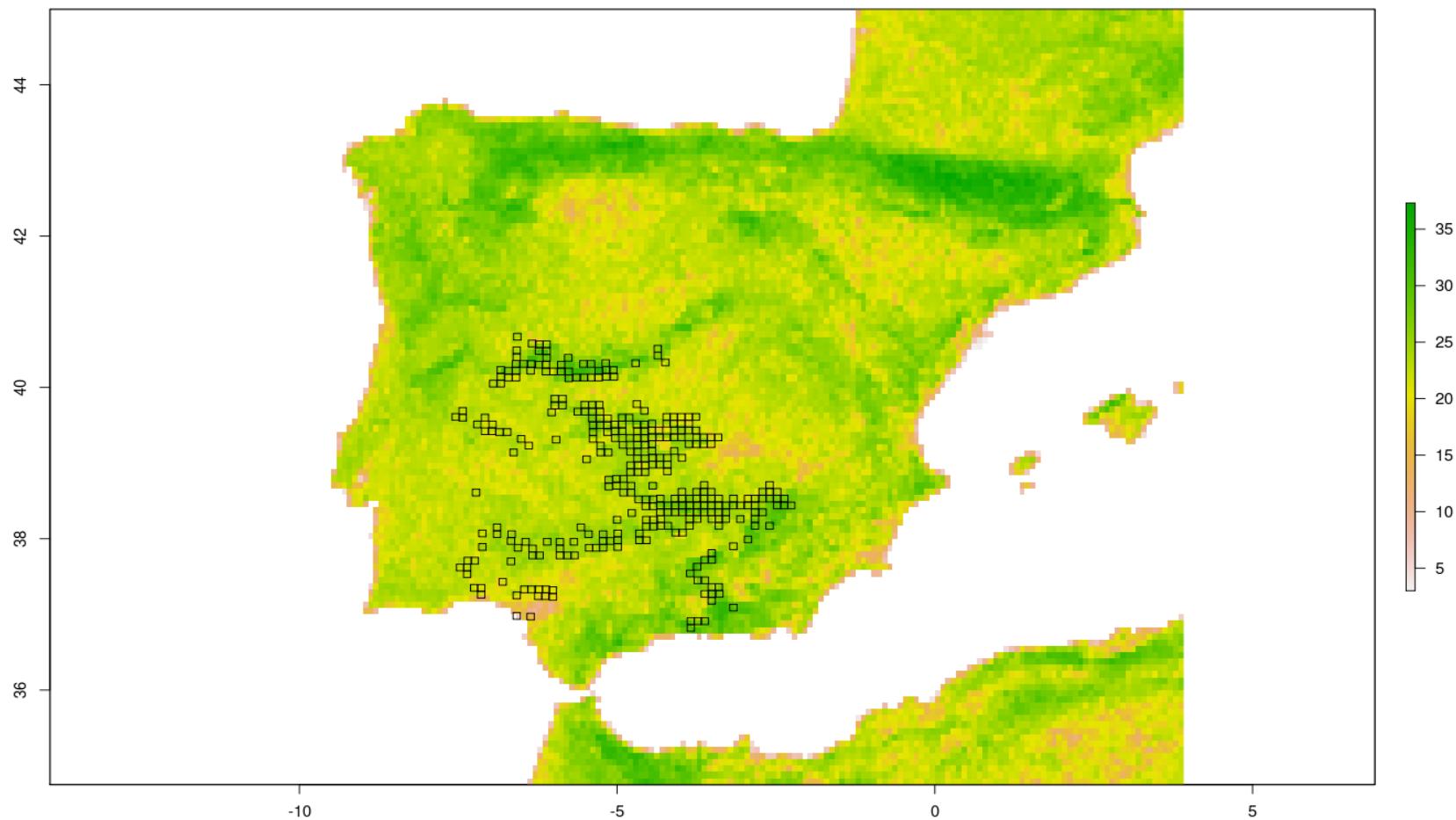
## PROS

- High availability (GBIF and the likes)

## CONS

- Difficult to compute probability of presence
- Sampling bias usually unknown
- Be careful with coordinate precision!

# PRESENCE ONLY



# PRESENCE - ABSENCE

## PROS

- Allow to compute probability of presence
- Provide info on sampling bias

## CONS

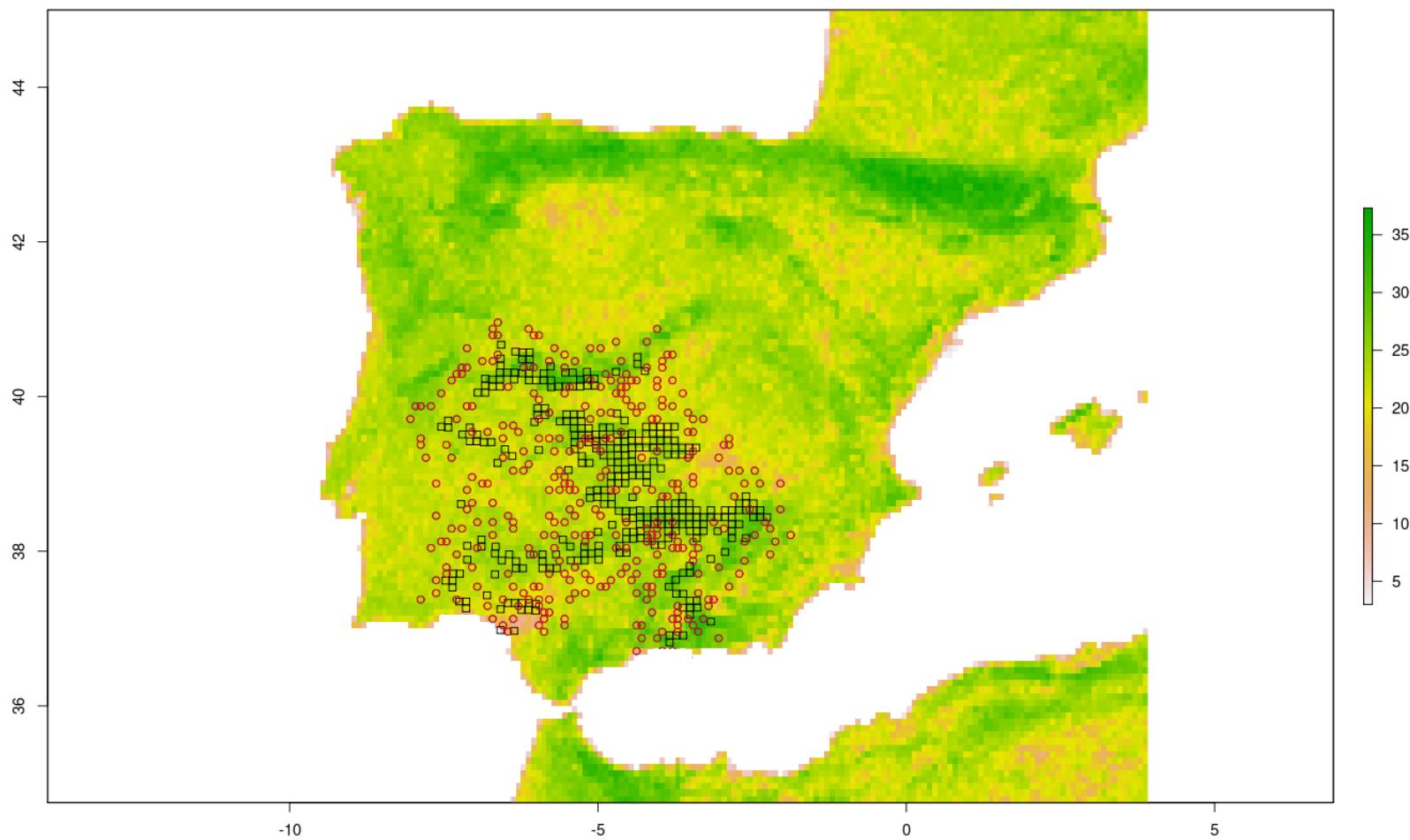
- How do you differentiate a real absence from false one (cryptic species, short lived individuals, etc)?
- Low availability

# NATURE OF ABSENCES

- **Contingency:** area is suitable, but there are dispersal restrictions, historical reasons, or temporal perturbations
- **Environmental:** absence of suitable habitat
- **Methodological:** due to sampling bias

Lobo et al. 2010

# ABSENCE



# PSEUDO-ABSENCE

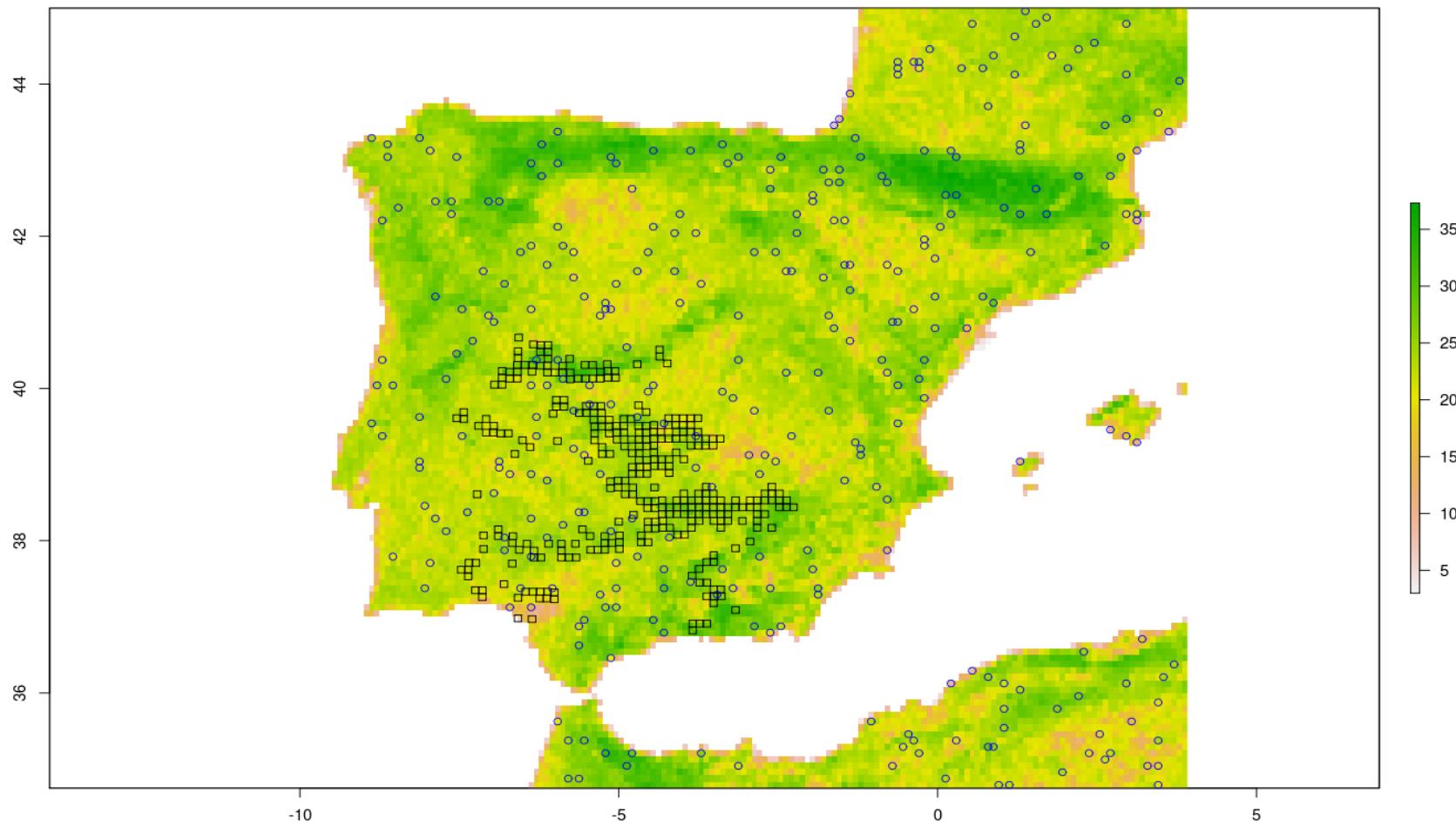
## PROS

- Easy to generate

## CONS

- They are not real absences
- What ecological criteria should we use?
- Don't take sampling bias into account

# PSEUDO-ABSENCE



# BACKGROUND

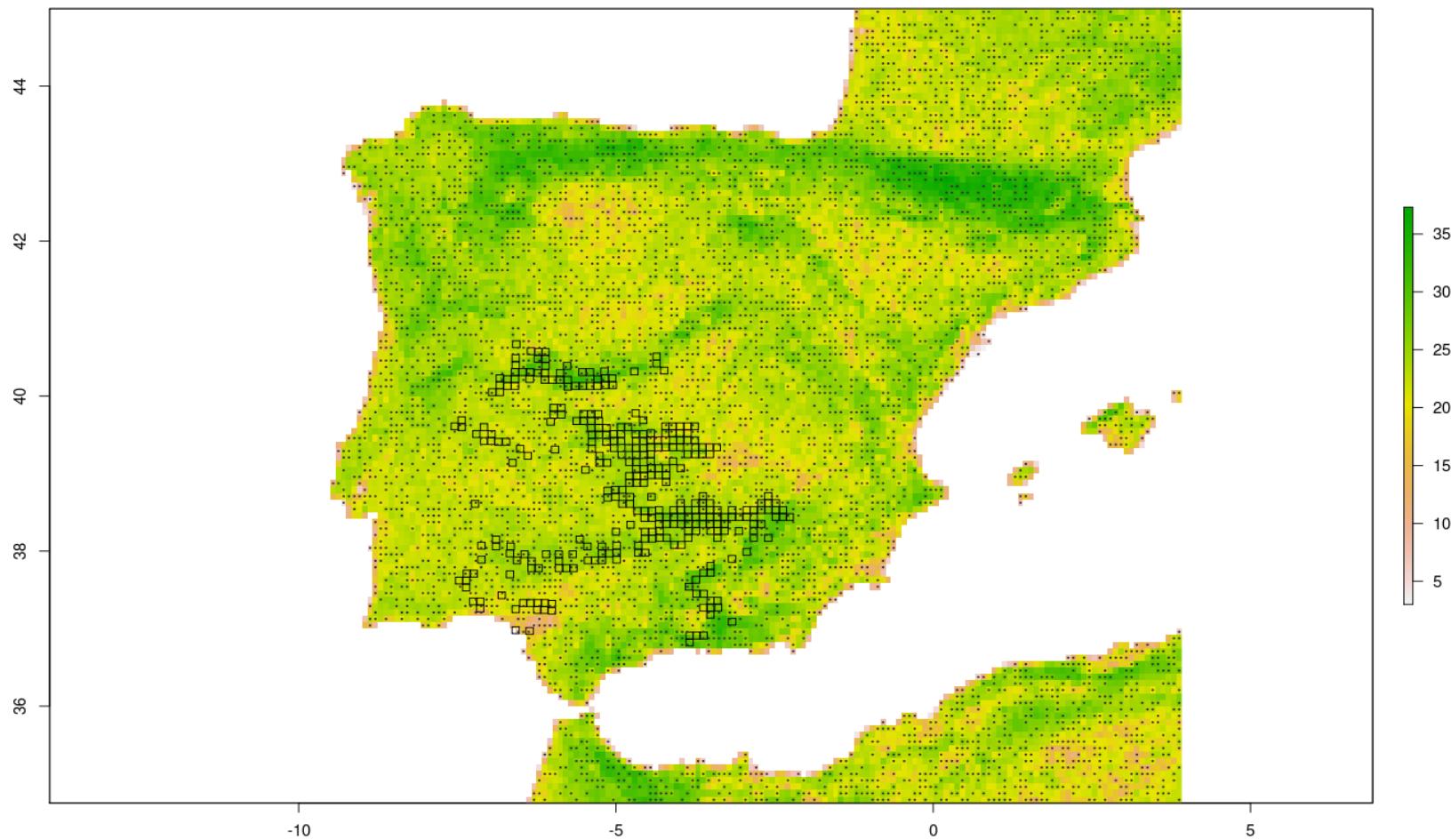
## PROS

- Easy to generate
- No issues if they overlap with presences (in fact, they should overlap!)

## CONS

- Some methods don't work well with them (Random Forest, Support Vector Machines)
- Required weighting to be used in regression models

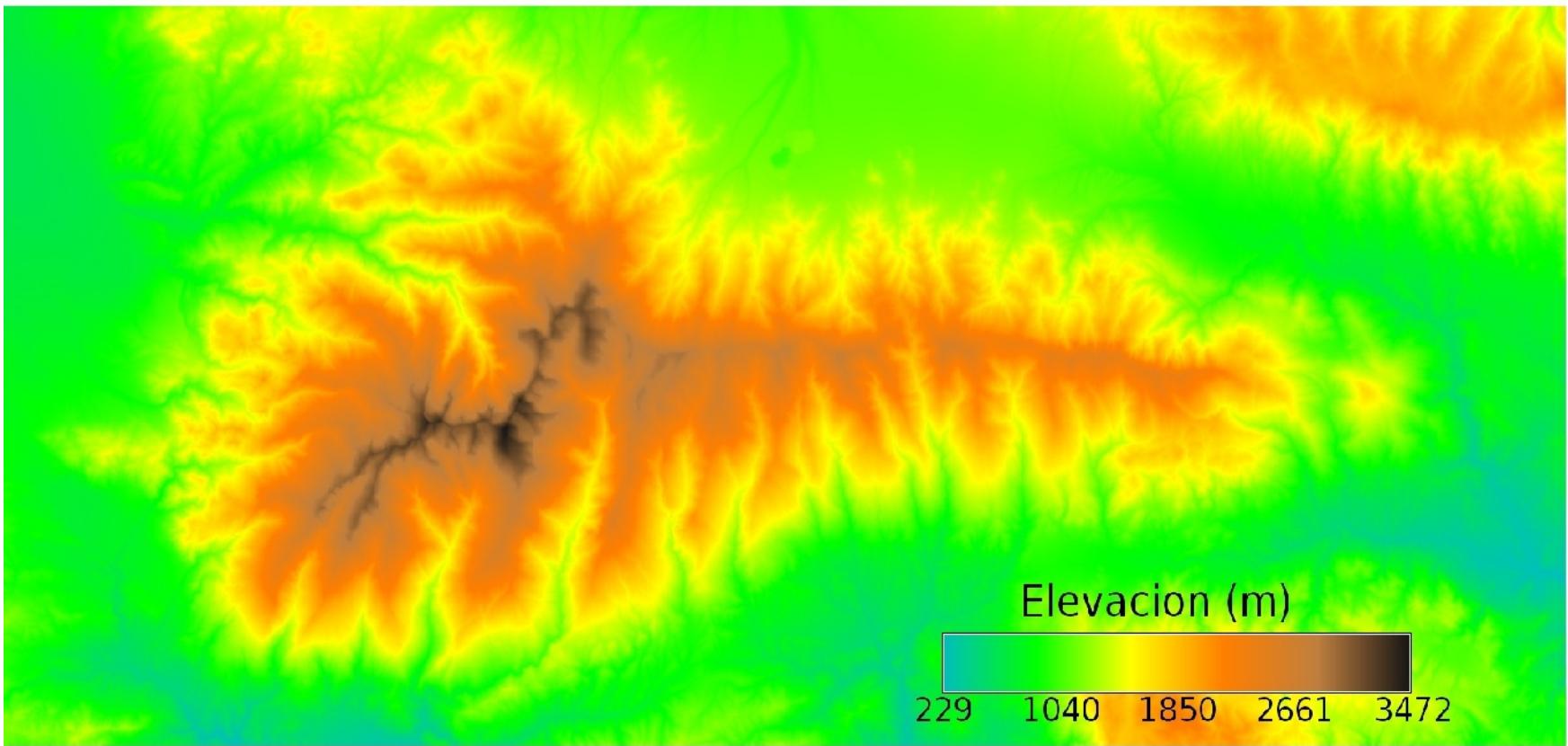
# BACKGROUND:



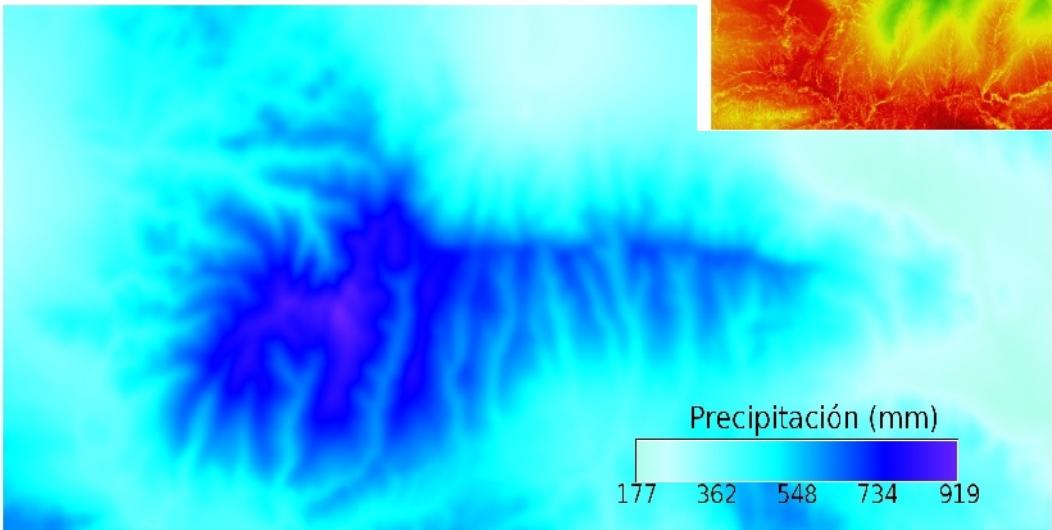
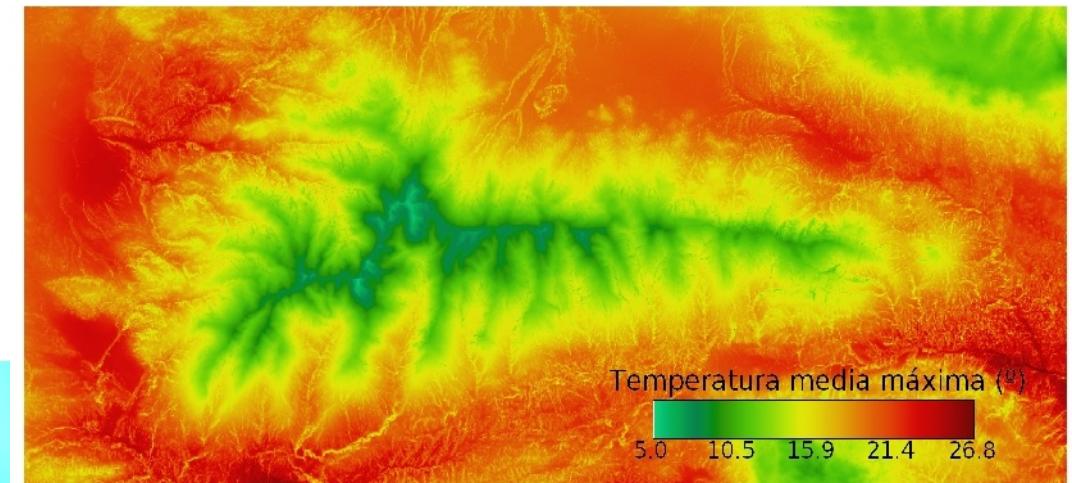
# ENVIRONMENTAL PREDICTORS

- Main features
  - Effect on species distribution
  - Available as a raster GIS map
  - Same extension and resolution
  - Low collinearity
- Matching resolution with the accuracy of presence data
- Watch for interactions among predictors

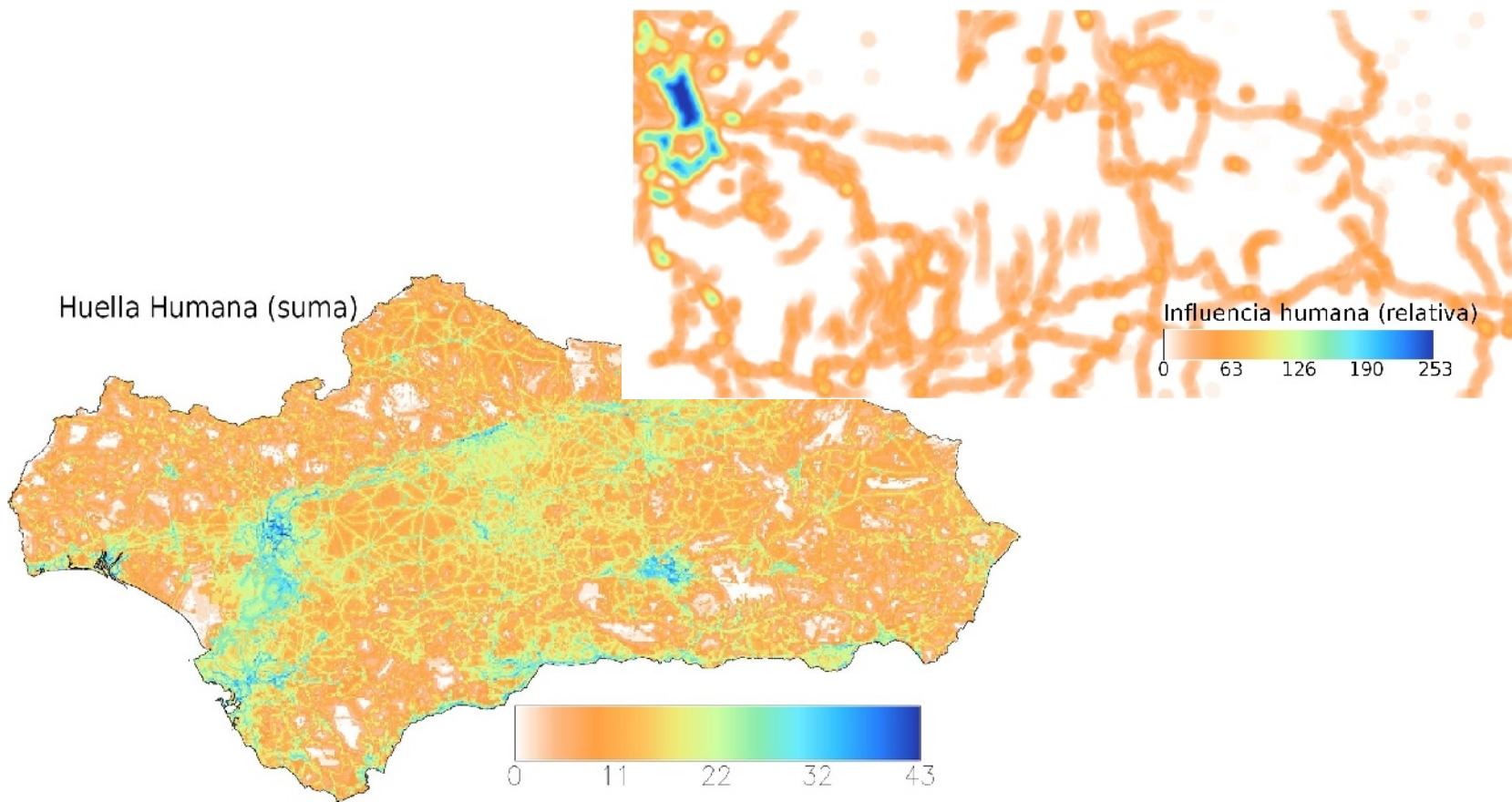
# TOPOGRAPHY



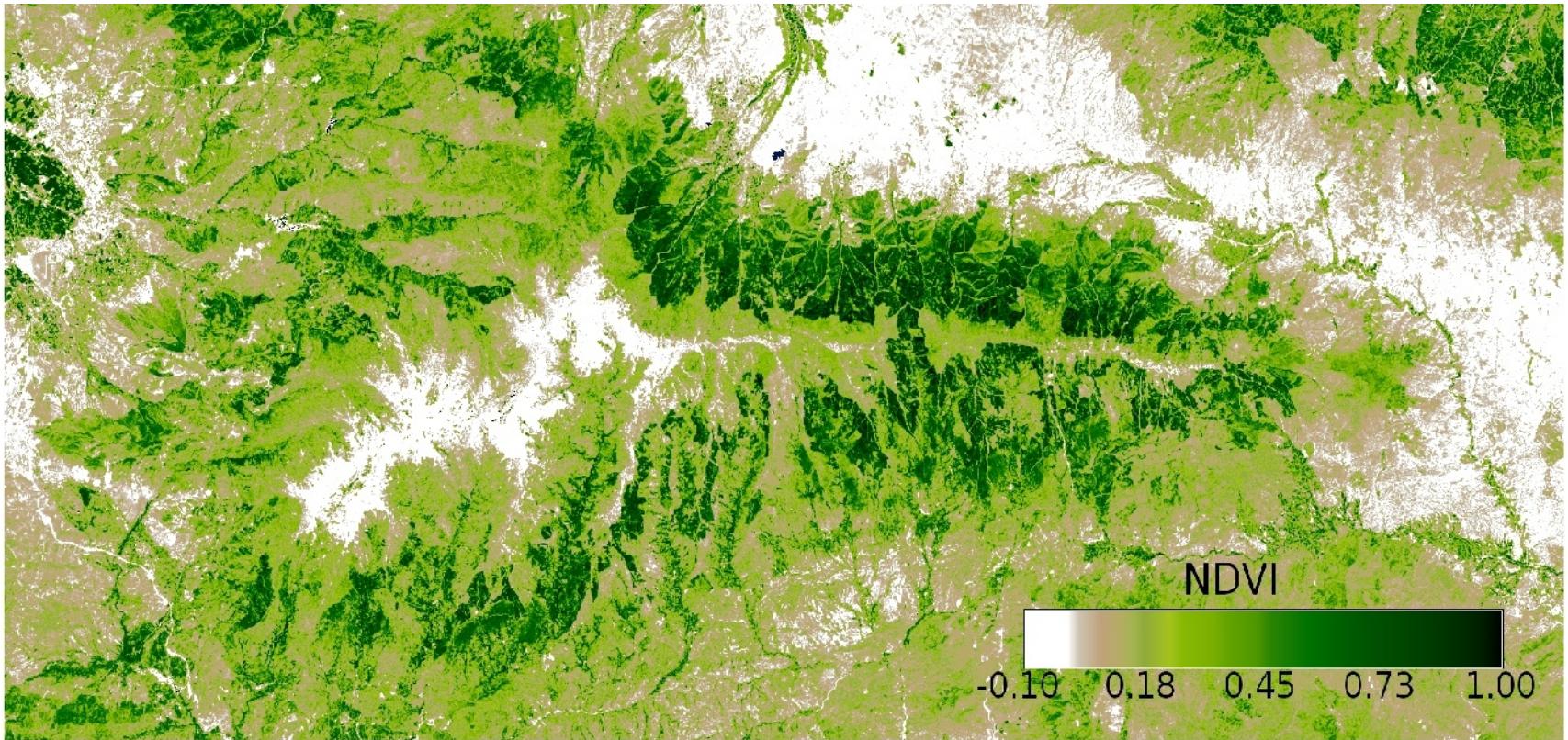
# CLIMATE



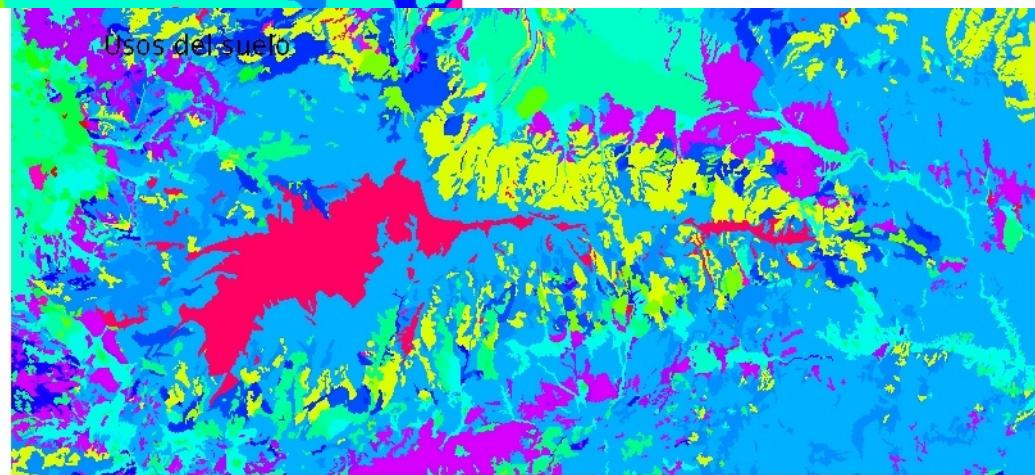
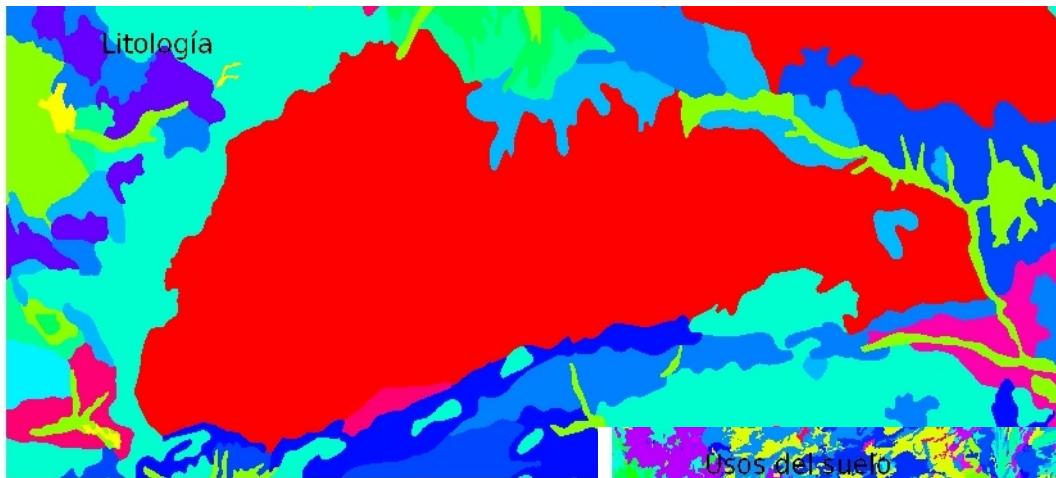
# HUMAN FOOTPRINT



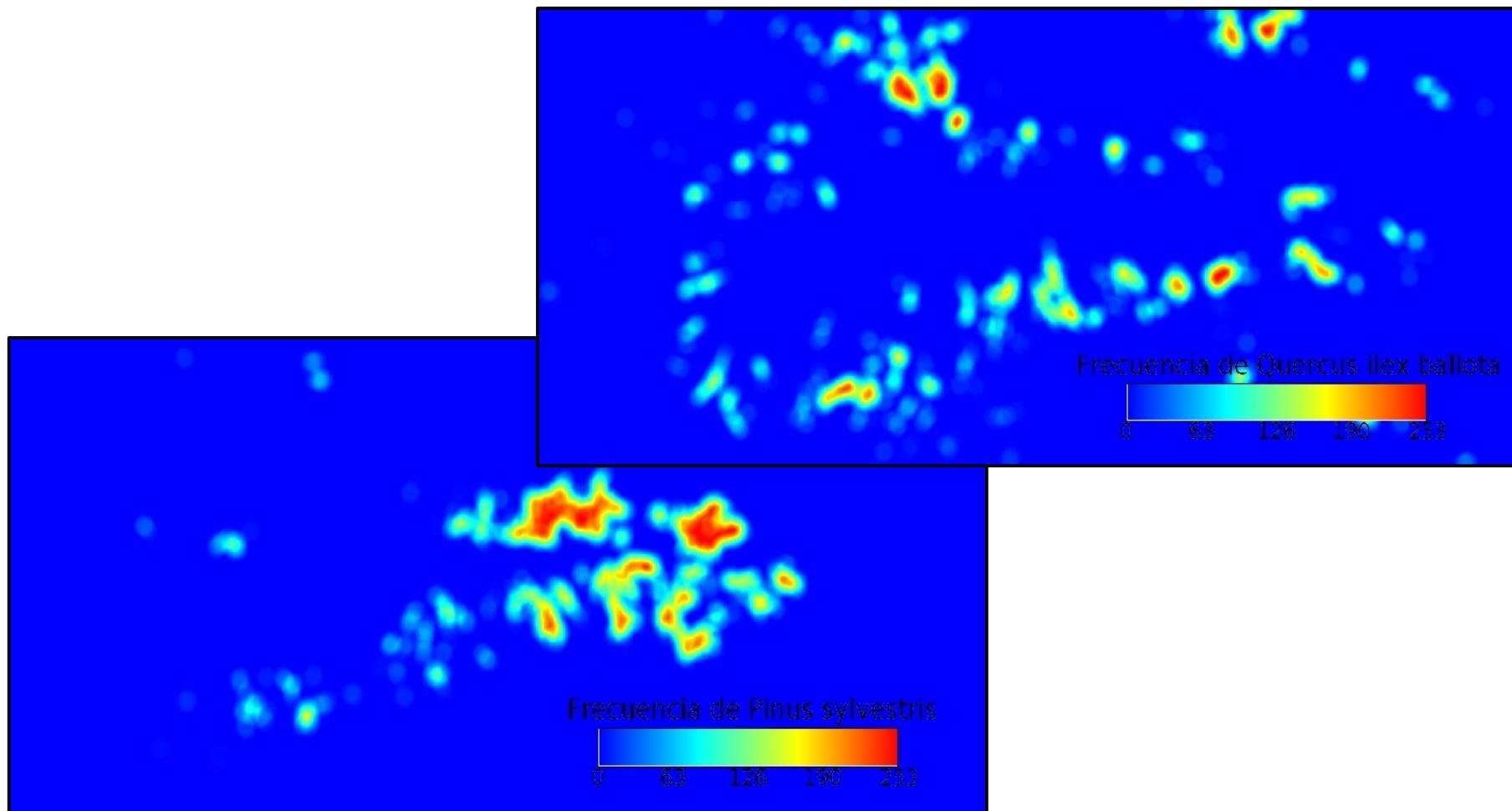
# REMOTE SENSING



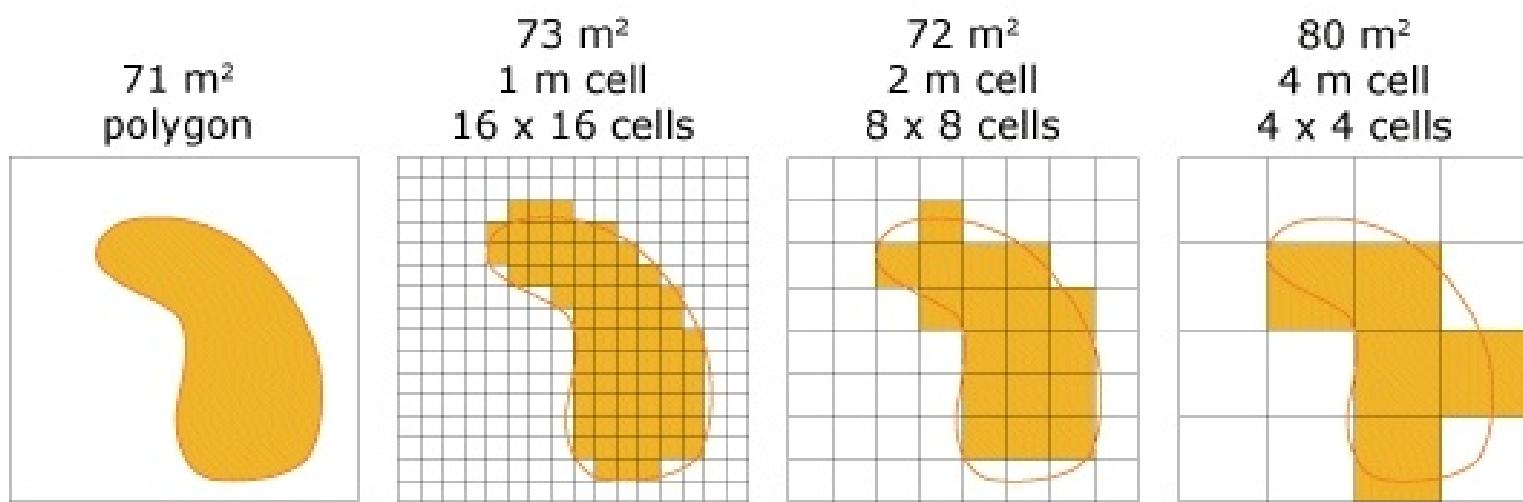
# QUALITATIVE PREDICTORS



# BIOLOGICAL PREDICTORS



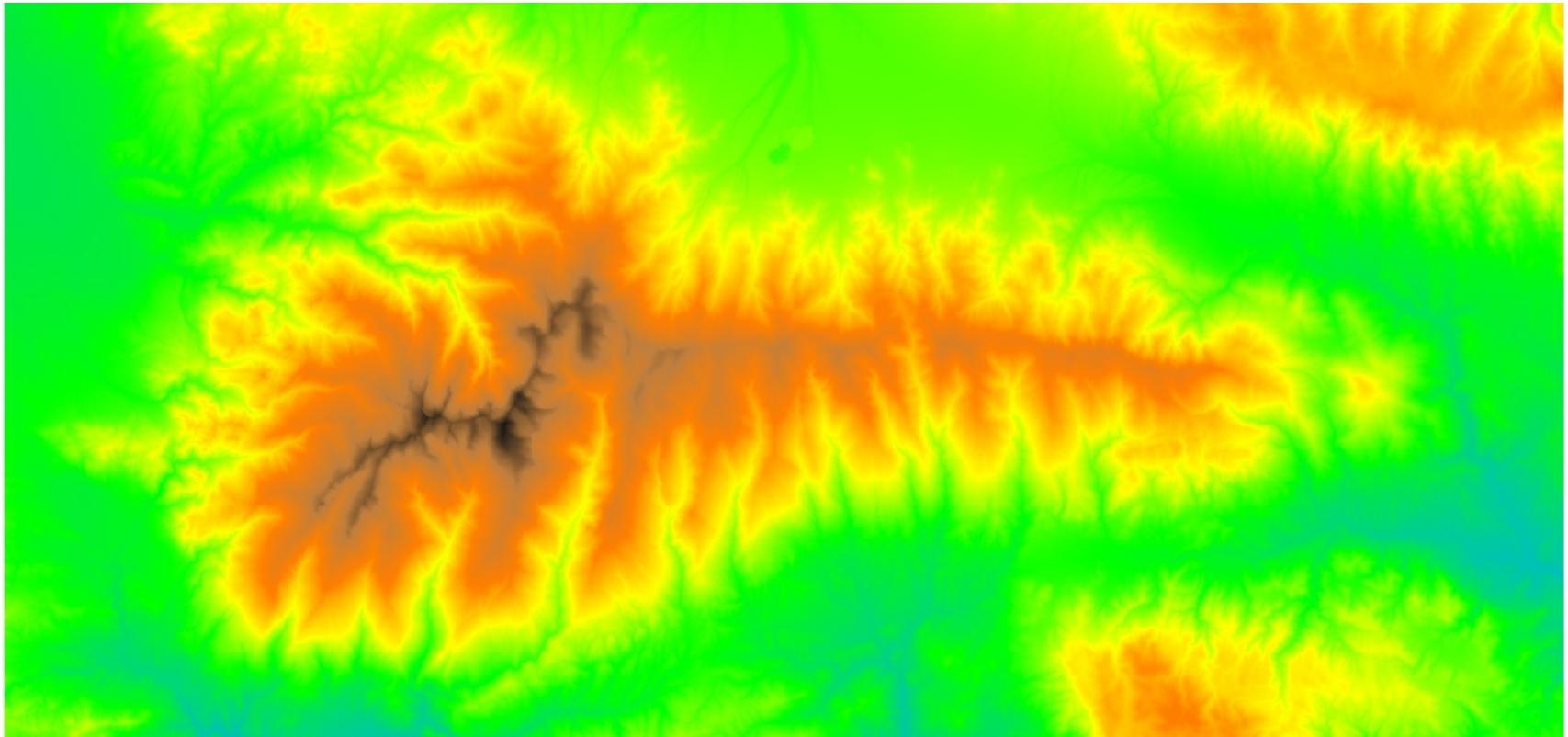
# RESOLUTION



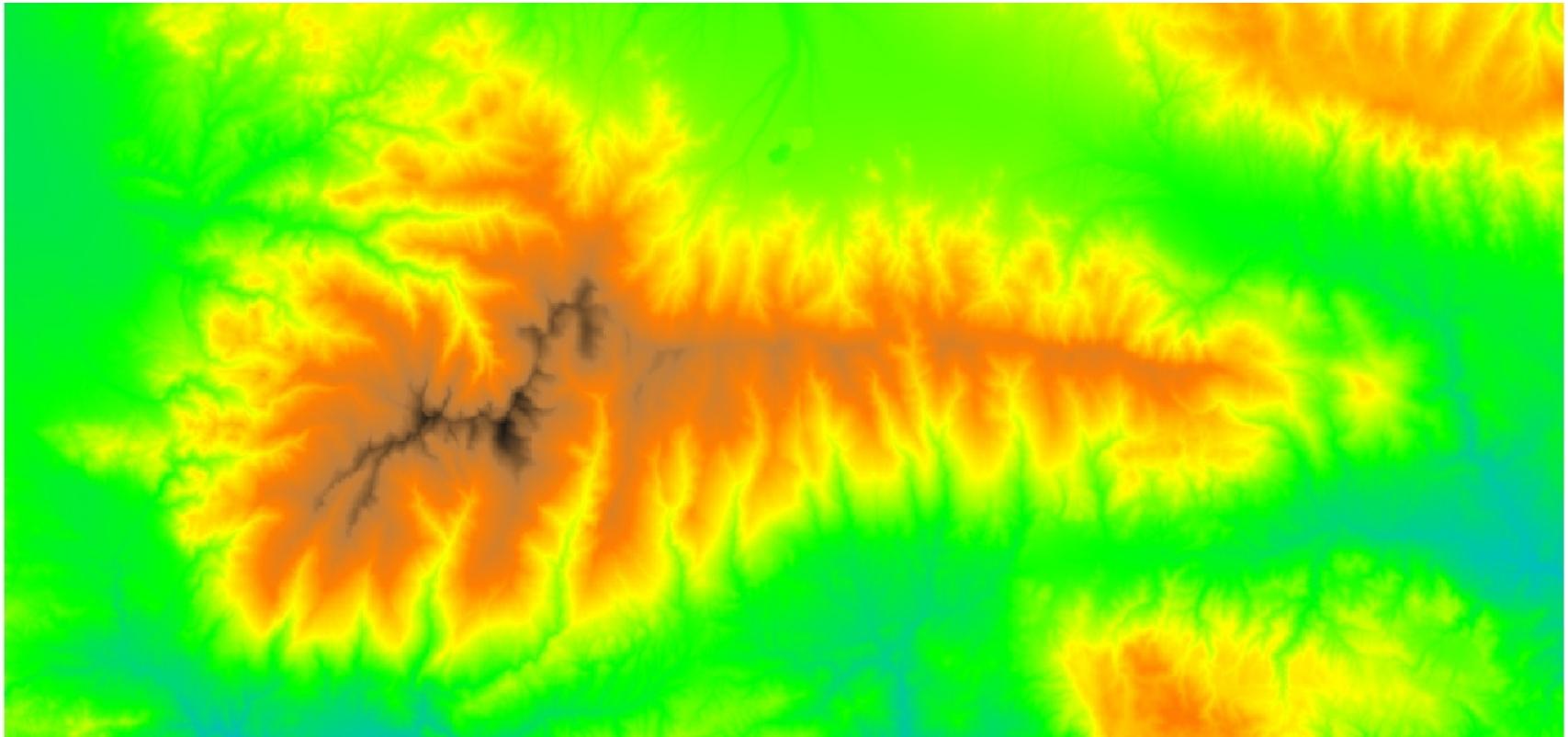
- Smaller cell size
- Higher resolution
- Higher feature spatial accuracy
- Slower display
- Slower processing
- Larger file size

- Larger cell size
- Lower resolution
- Lower feature spatial accuracy
- Faster display
- Faster processing
- Smaller file size

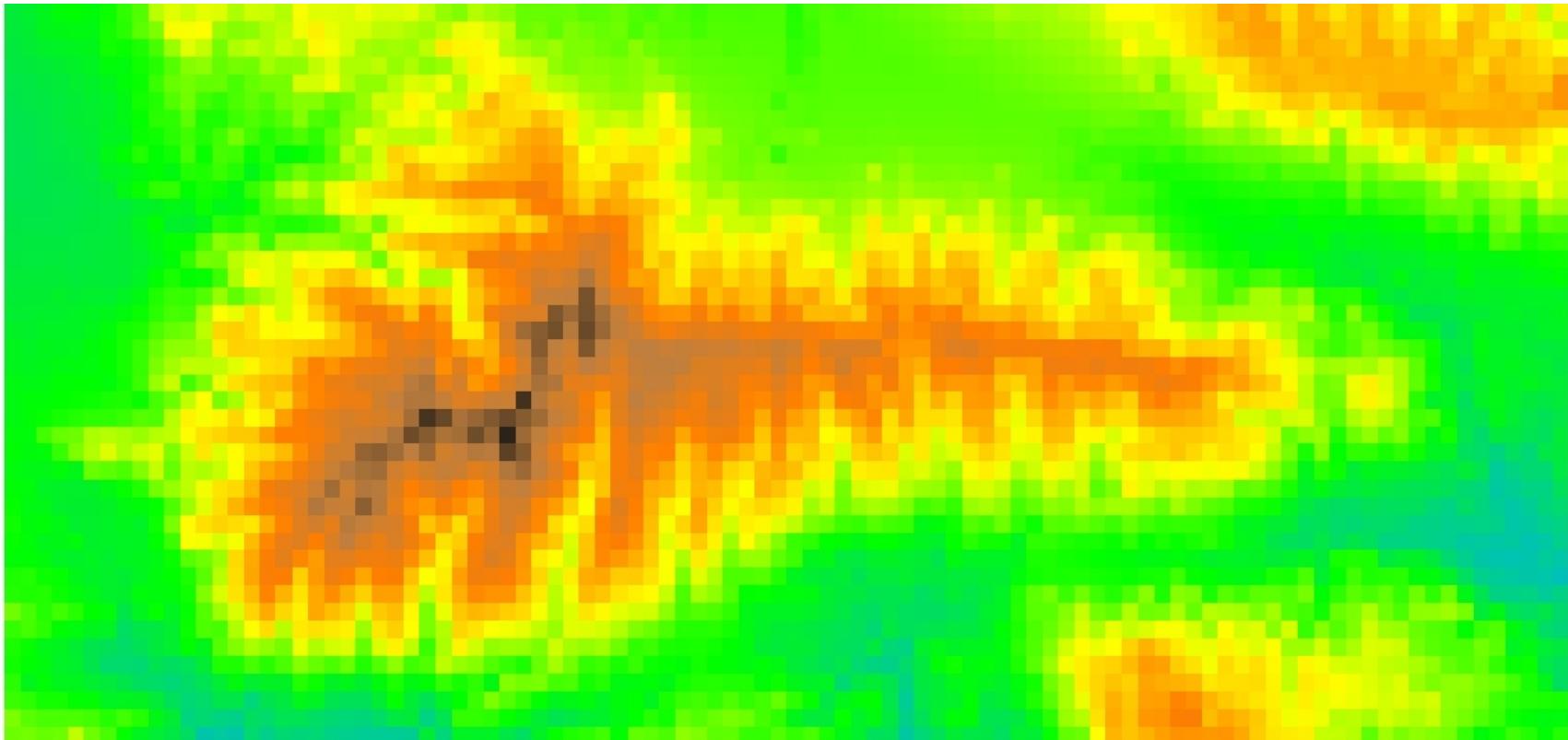
# RESOLUTION (10 m)



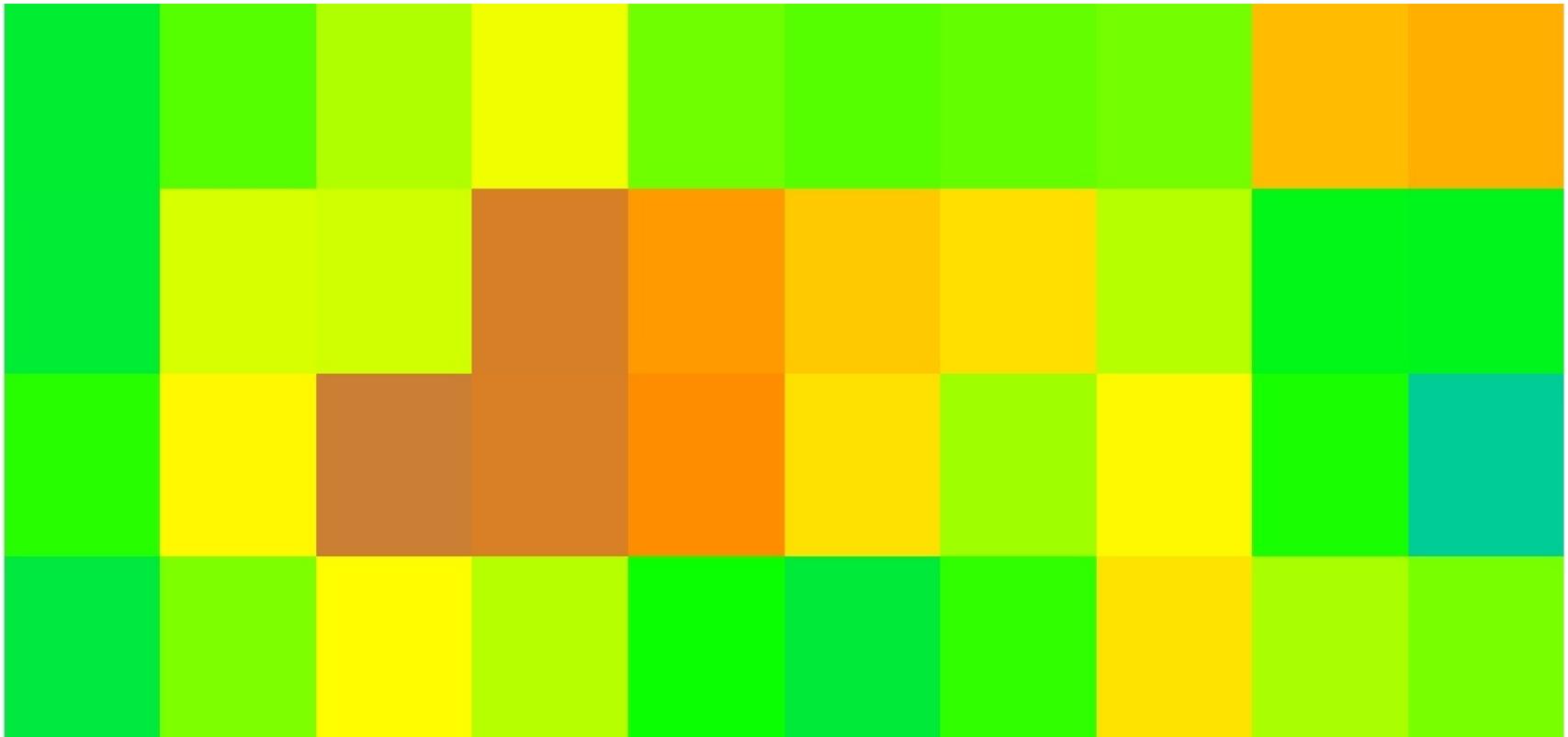
# RESOLUTION (100 m)



# RESOLUTION (1000 m)

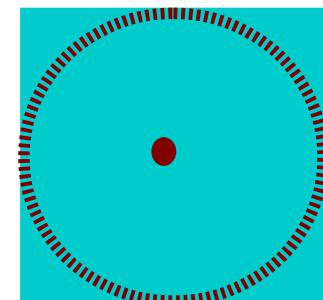
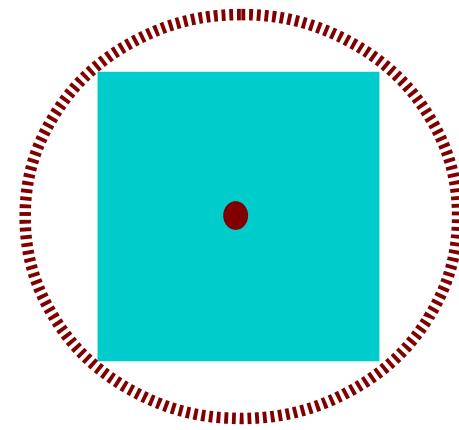
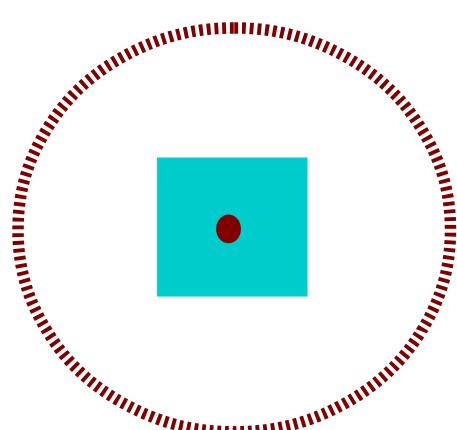


# RESOLUTION (10000 m)



# IMPORTANT

Resolution of predictors should match the coordinates precision of the presence records!



# SOME DATA SOURCES

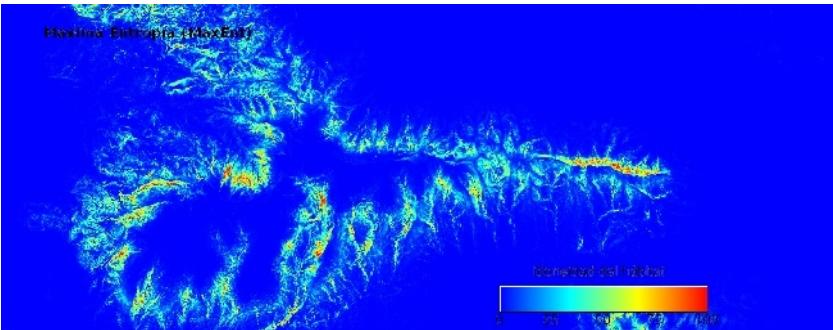
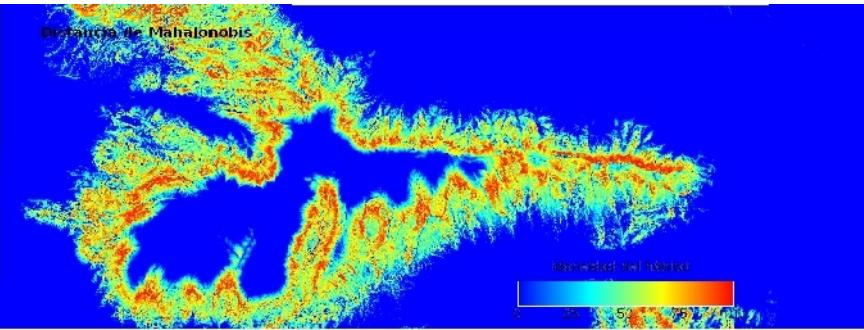
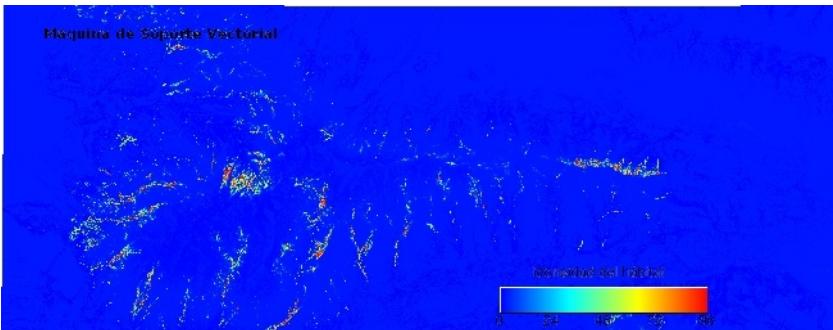
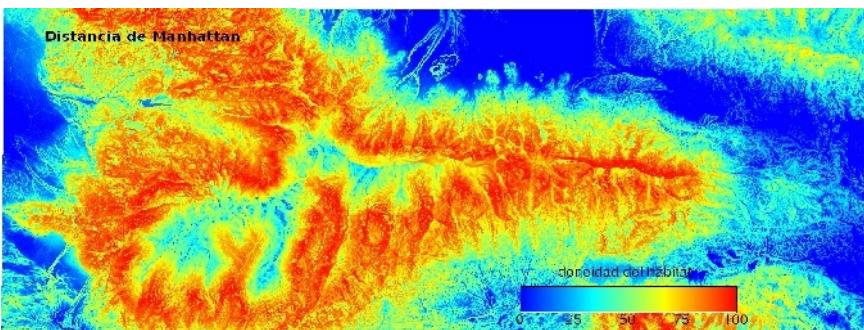
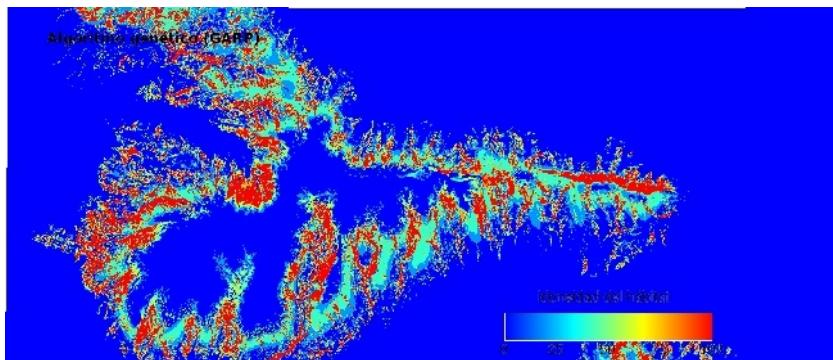
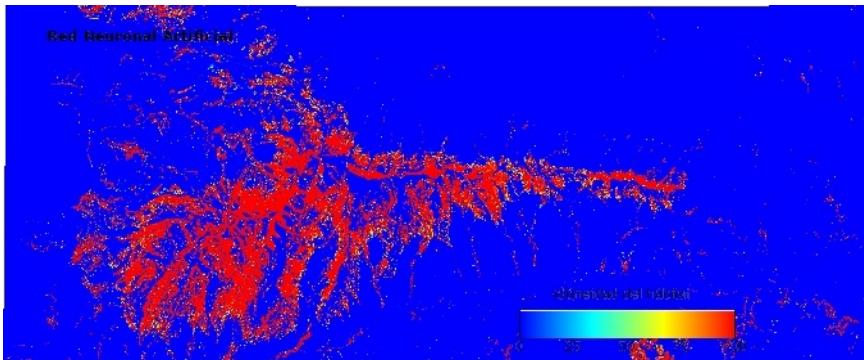
<b>Tipo</b>	<b>Dataset</b>	<b>Sitio web</b>
Climate	WORLDCLIM	
Topography	SRTM	
NDVI	GIMMS	
Vegetation	MODIS VCF	
Human footpr.	Human Footprint	
Land uses	GLOBCOVER	

# ALGORITHMS

Main families of methods

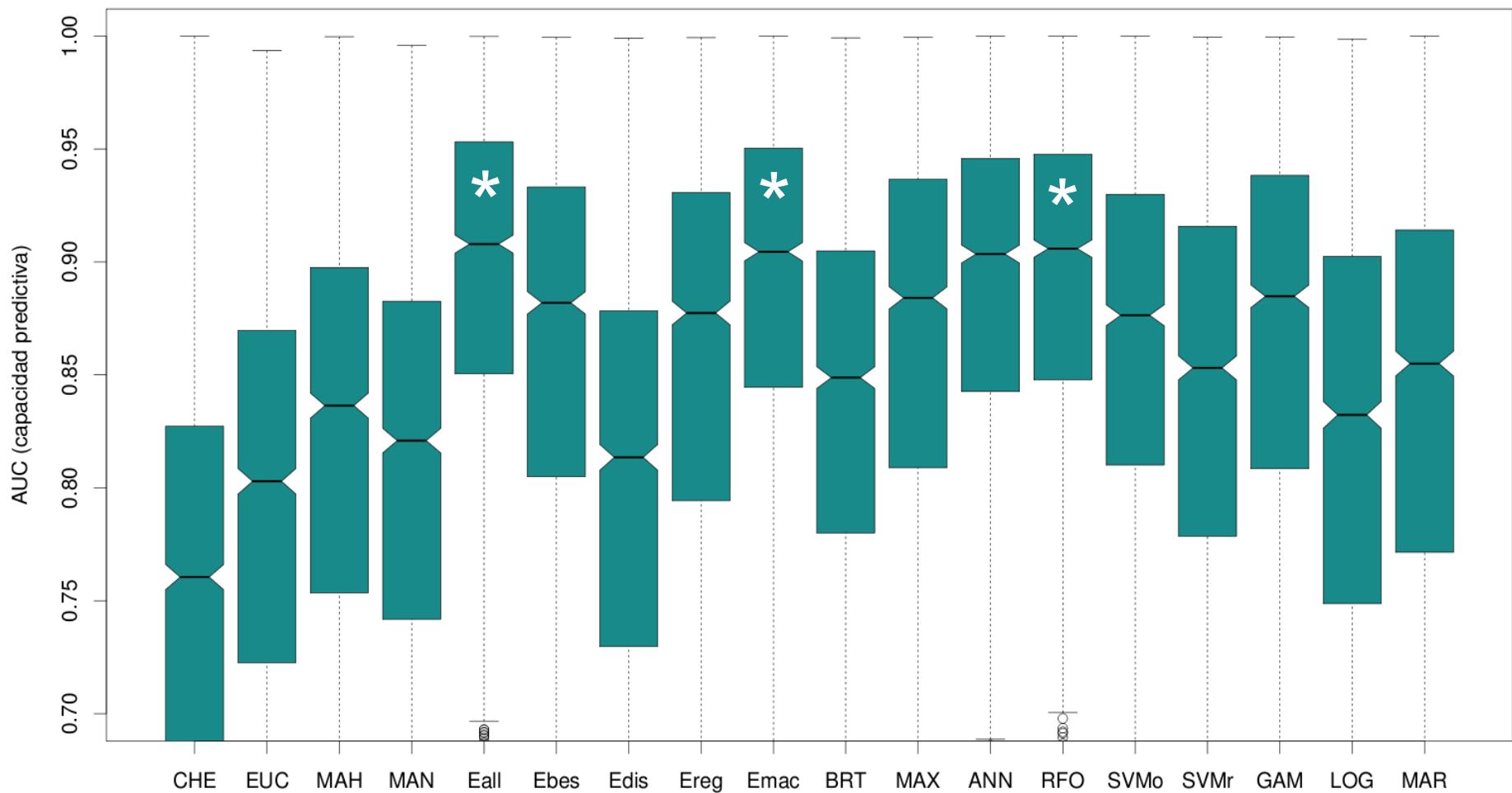
- Bioclimatic envelopes
- Ecological similarity
- Regression
- Machine learning

# DIFF METHODS = DIFF RESULTS

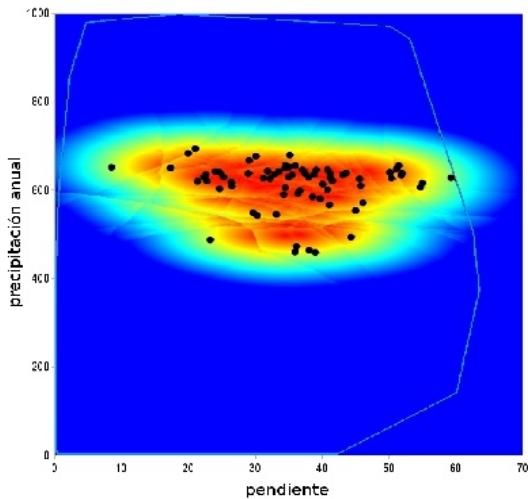


# DIFFERENT PREDICTIVE ABILITIES

Evaluación de modelos de 1700 especies de árboles en Mesoamérica

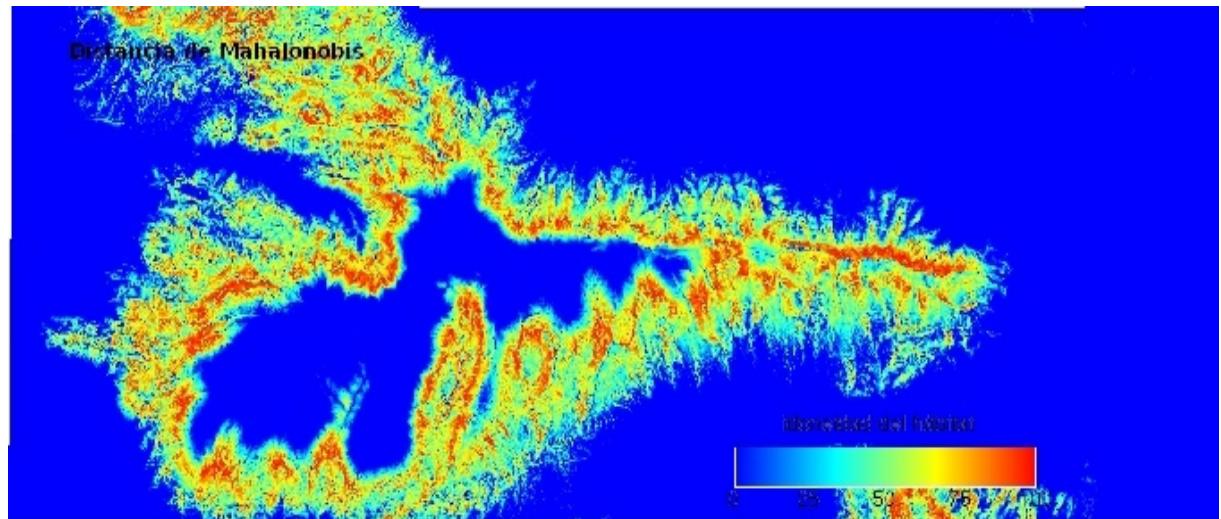


# MODEL



Ecological space

Geographical space

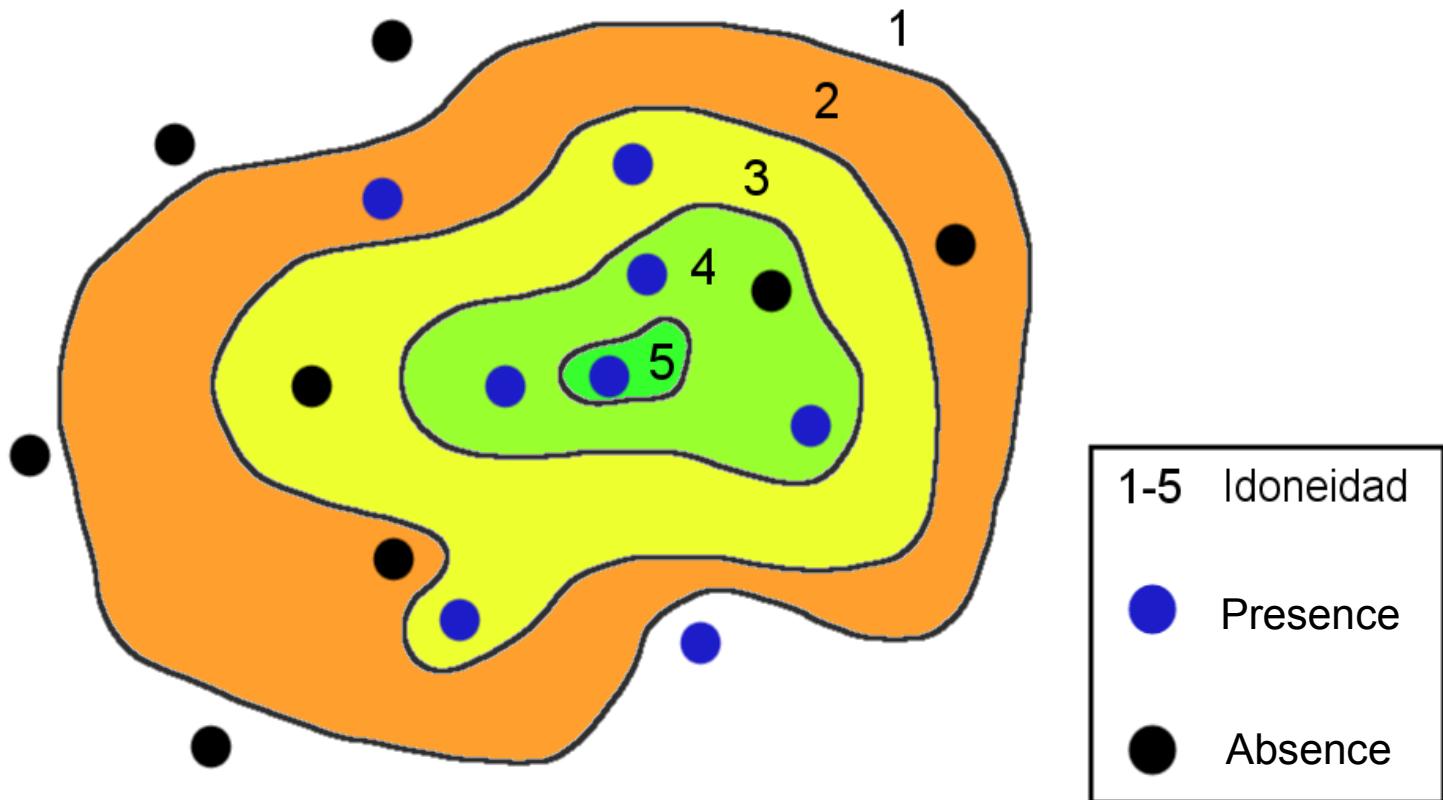


# EVALUATION

Key paper:

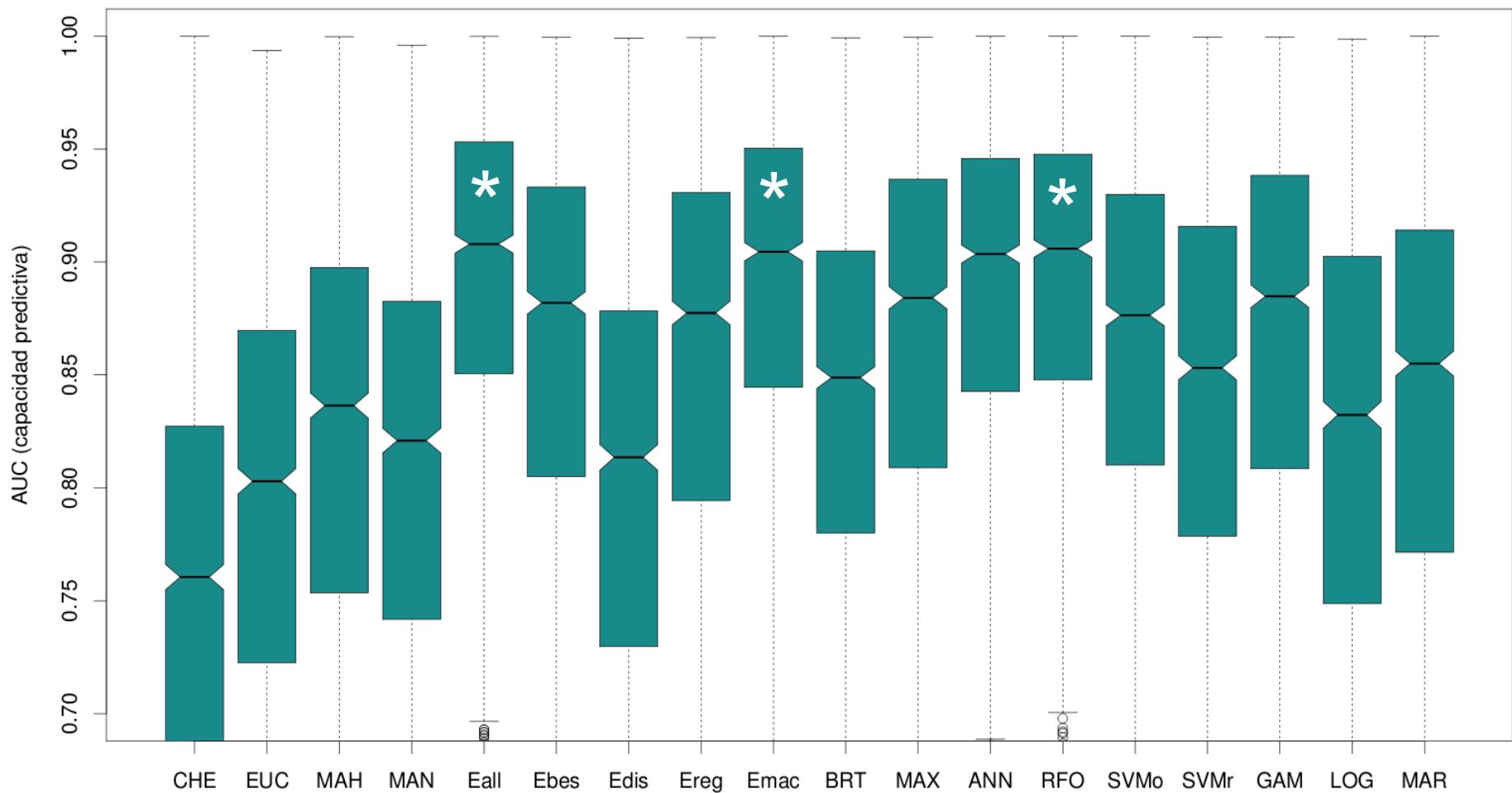
Fielding AH y Bell JF 1997. A review of methods for the assessment of prediction errors in conservation presence/absence models.  
*Environmental Conservation* 24(1), 38-49

# EVALUATION



# DIFFERENT PREDICTIVE ABILITIES

Evaluación de modelos de 1700 especies de árboles en Mesoamérica



# CONFUSION MATRIX

**A** → correctly predicted presences

**D** → correctly predicted absences

**B** → missed absences (false positives or commission error)

**C** → missed presences (false negatives or omission error)

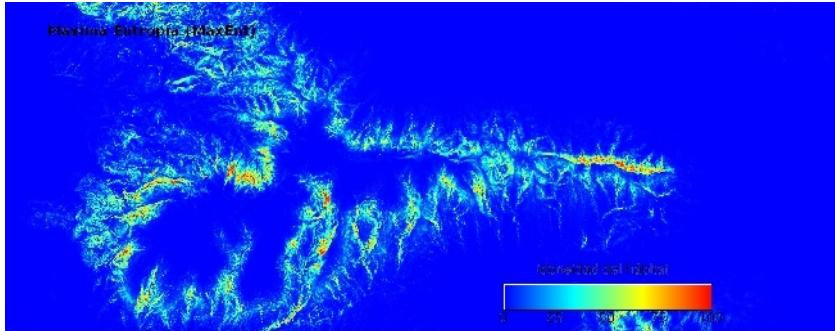
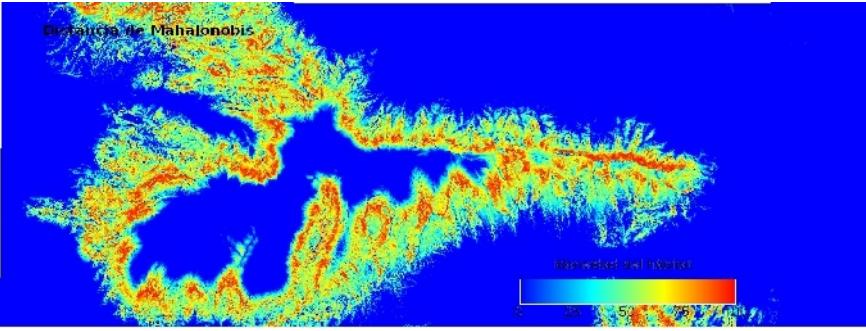
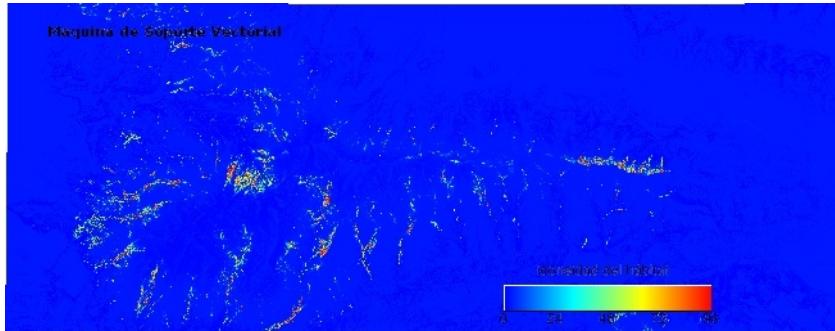
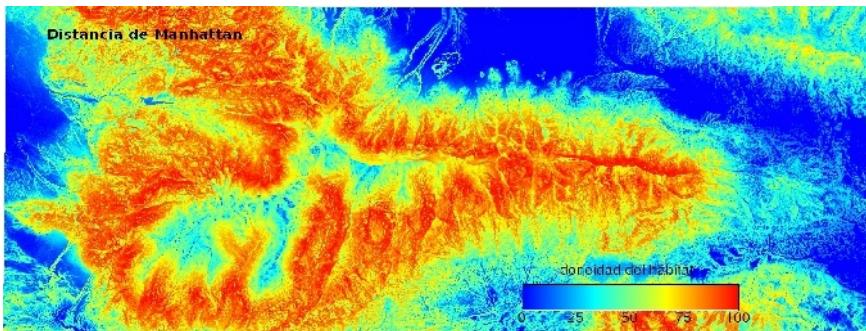
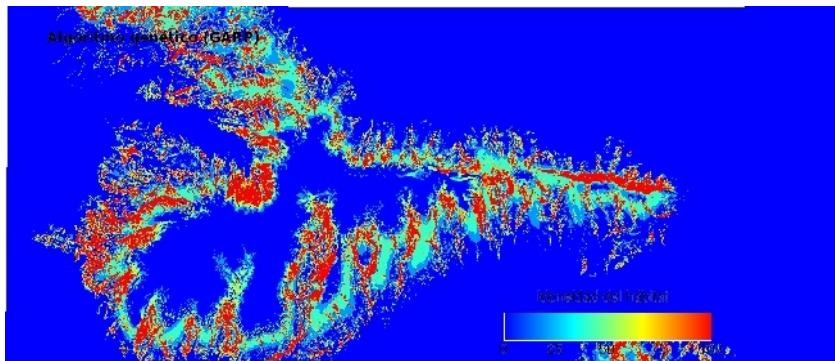
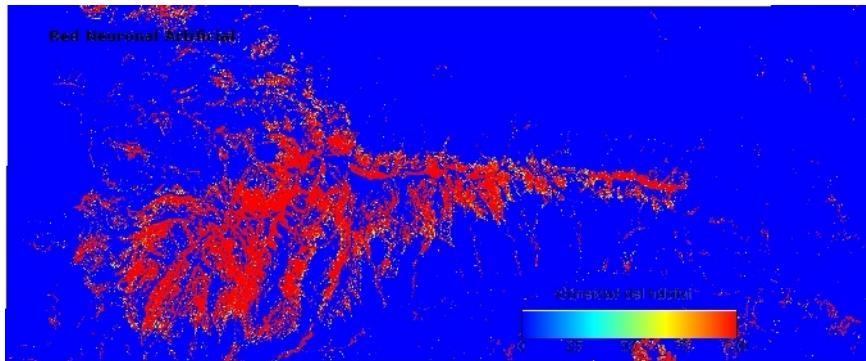
		PRESENCE DATA	
		Presence	Absence
MODEL PREDICTION	Presence	<b>A</b>	<b>B</b>
	Absence	<b>C</b>	<b>D</b>

sensitivity →  $S = A/(A+C)$

specificity →  $E = D/(B+D)$

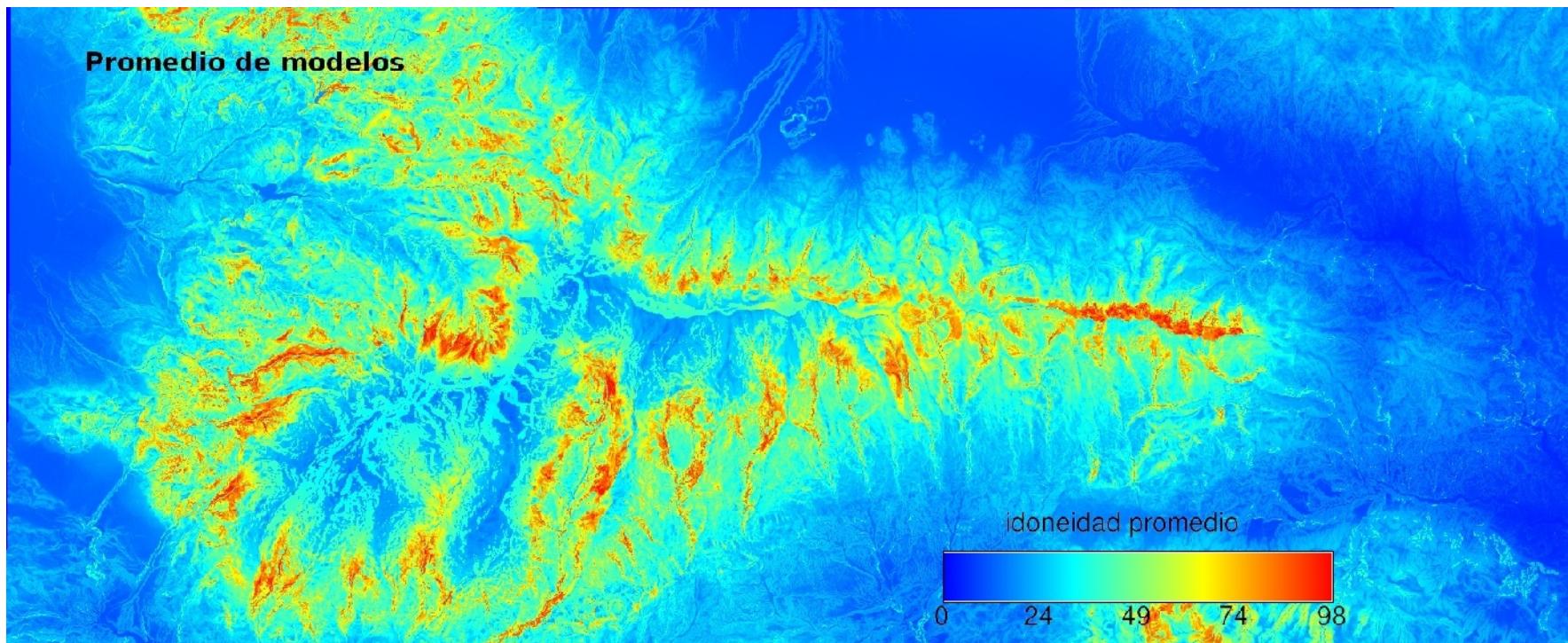
true skill statistic →  $TTS = S + E - 1$

# ENSEMBLE MODEL FORECASTING



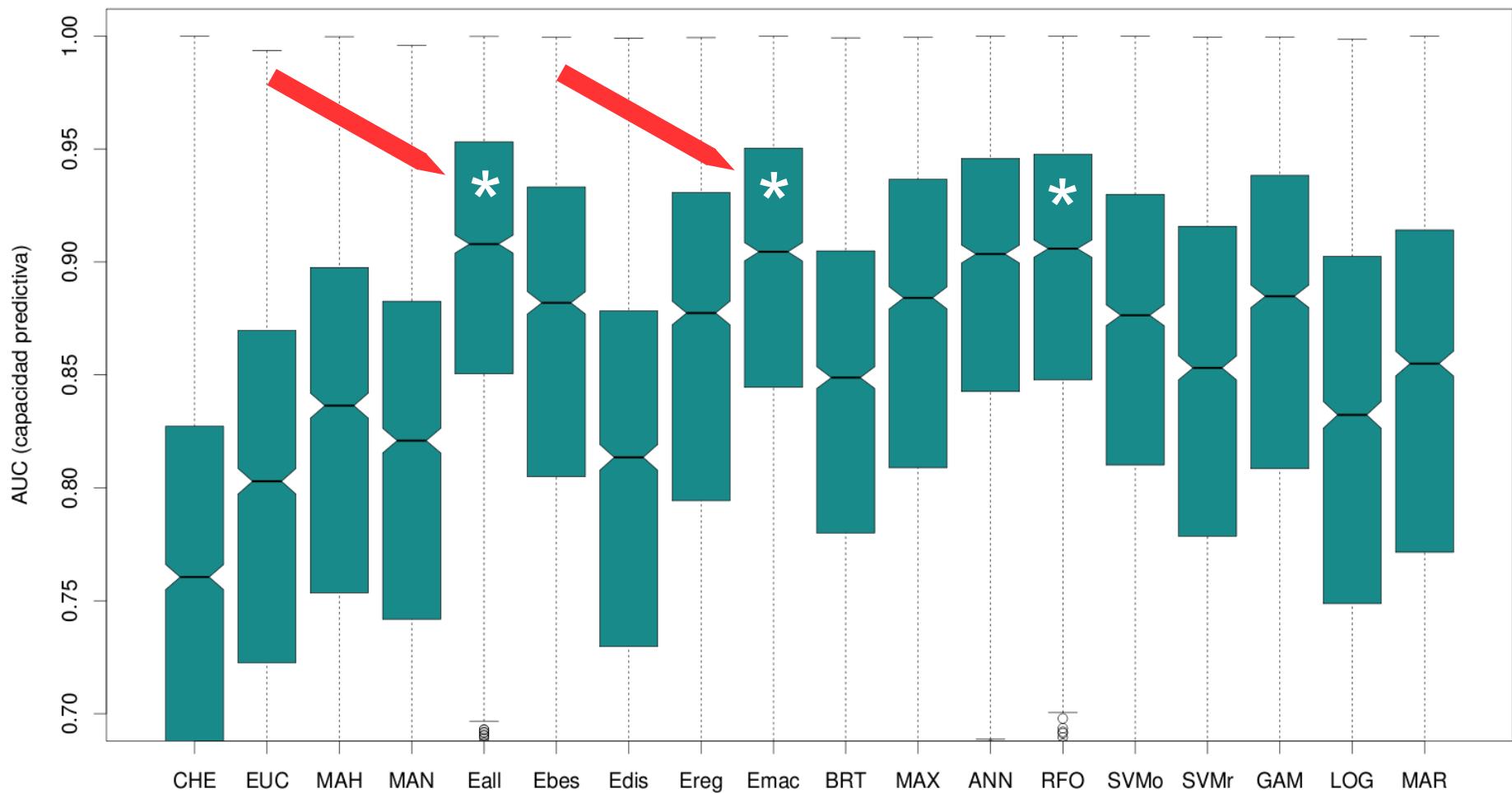
# ENSEMBLE MODEL FORECASTING

AVERAGE OF THE PREVIOUS MODELS

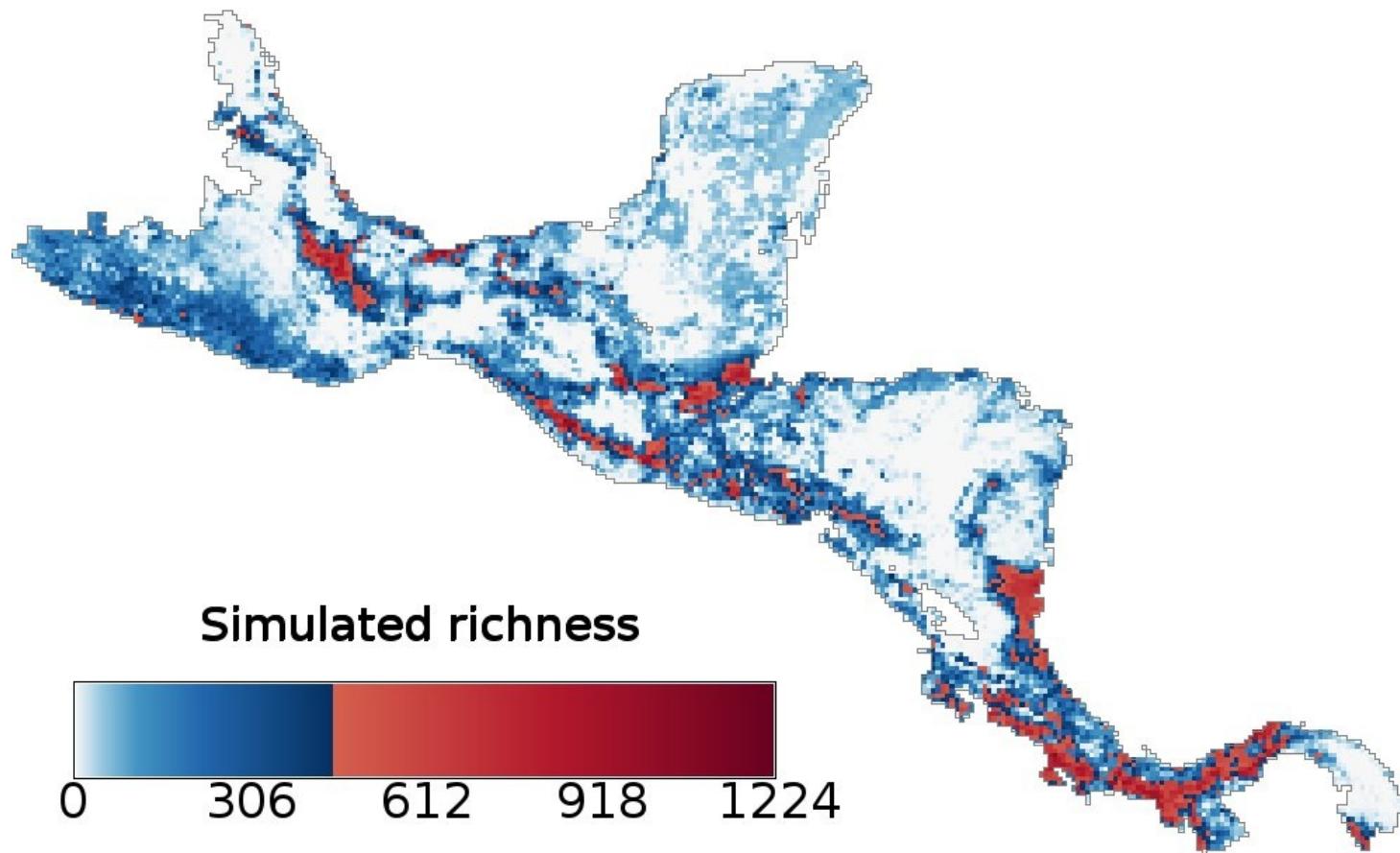


# ENSEMBLES ACTUALLY WORK!

Evaluación de modelos de 1700 especies de árboles en Mesoamérica

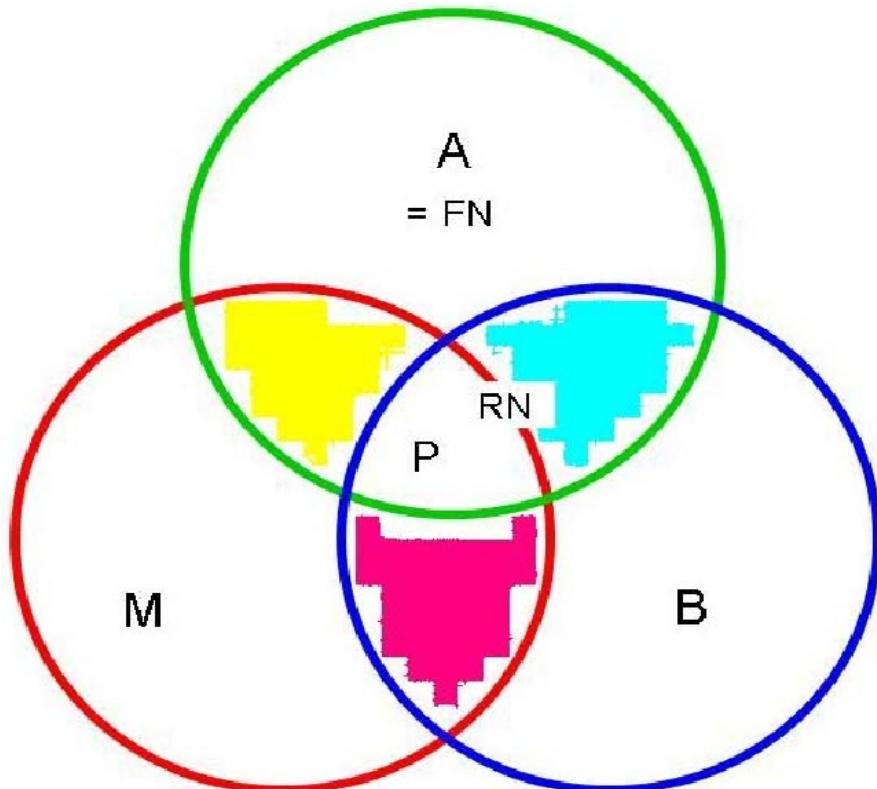


# RICHNESS MODELS



Benito et al. 2013 MEE

# INTERPRETATION



BAM diagram

- **A** – abiotic environment = **FN** (fundamental niche)
- **B** – biotic environment
- **M** – accessible habitat
- **RN** – *realized niche*
- **P** - *presence*

Soberón 2005 Biodiversity Informatics

# INTERPRETATION

Soberón (2005):

*“Whether the result is interpreted as the species' distribution, the spatial extent of its fundamental niche, or some other phraseology, these algorithms **only find regions that 'resemble' those where occurrence points are located.**”*

# **APPLICATIONS OF SPECIES DISTRIBUTION MODELS**

# VEGETATION MAPPING

**Box et al. (1981)** Predicting physiognomic vegetation types with climate variables.

*Vegetatio* 45, 127-139

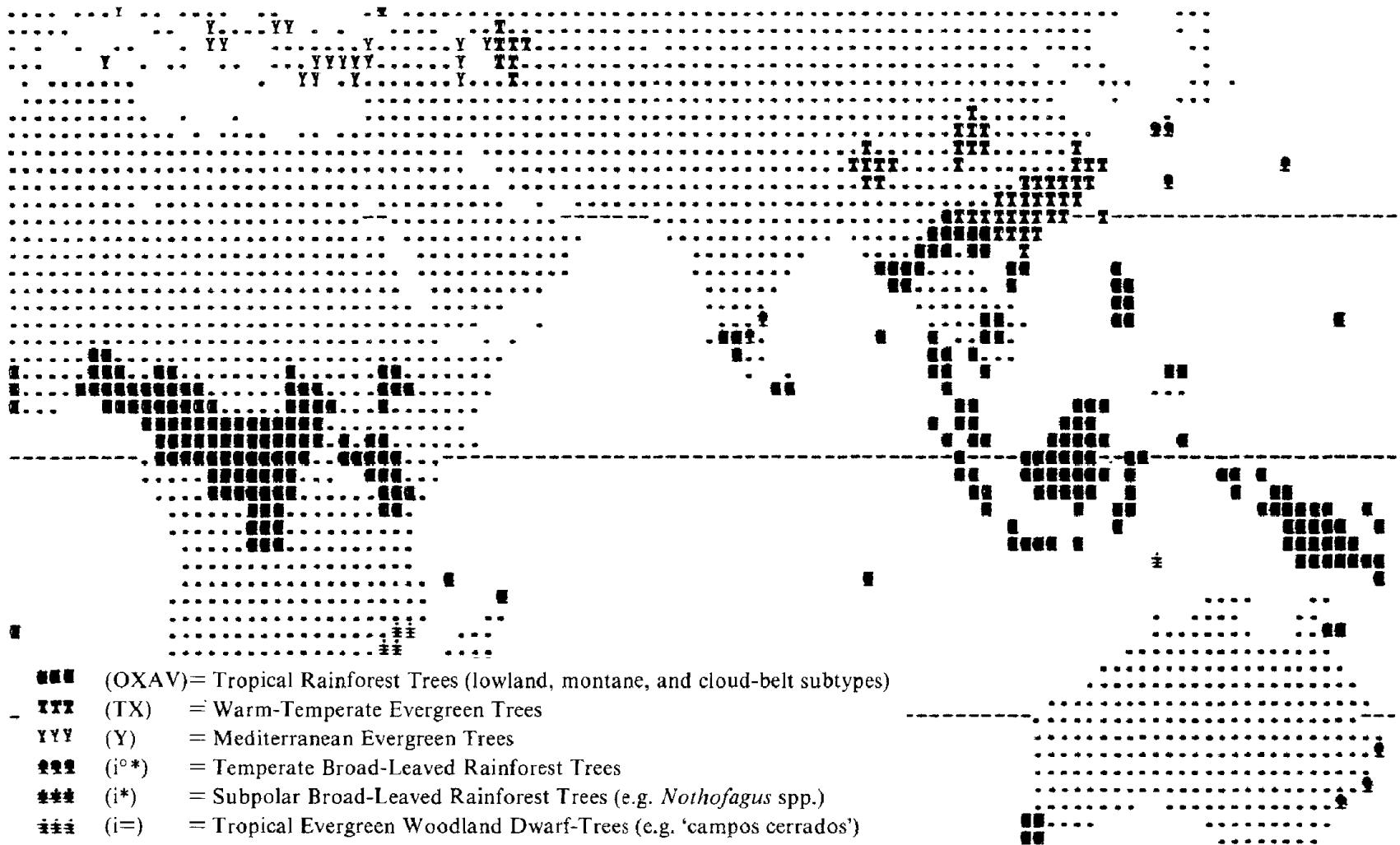
- **Objective:** mapping of broad vegetation types
- **Methods:** bioclimatic envelopes
- **Results:** 85 % success in predicting plant types and 50 % in predicting vegetation structure.
- **Conclusion:** Simple models can predict vegetation patterns at continental to global scales with remarkable accuracy.

# Box et al. (1981)

## Bioclimatic envelopes

	TMAX	TMIN	DTY	PRCP	MI	PMAX	PMIN	PMTMAX
Tropical rainforest								
Max.	30.0	28.0	12.0	*****	***	*****	***	*****
Min.	20.0	18.0	0.0	1 400.	1.00	150.	5.	10.
Raingreen monsoon forest								
Max.	35.0	28.0	15.0	*****	3.00	*****	30.	*****
Min.	23.0	14.0	0.0	800.	0.90	100.	0.	20.
Summergreen broad-leaved trees								
Max.	32.0	10.0	50.0	*****	***	*****	***	*****
Min.	15.0	-20.0	8.0	300.	0.80	40.	2.	30.
Short-needed boreal/ montane trees								
Max.	22.0	3.0	60.0	*****	***	*****	***	*****
Min.	10.0	-25.0	10.0	100.	0.60	25.	0.	10.

# Box et al. (1981)

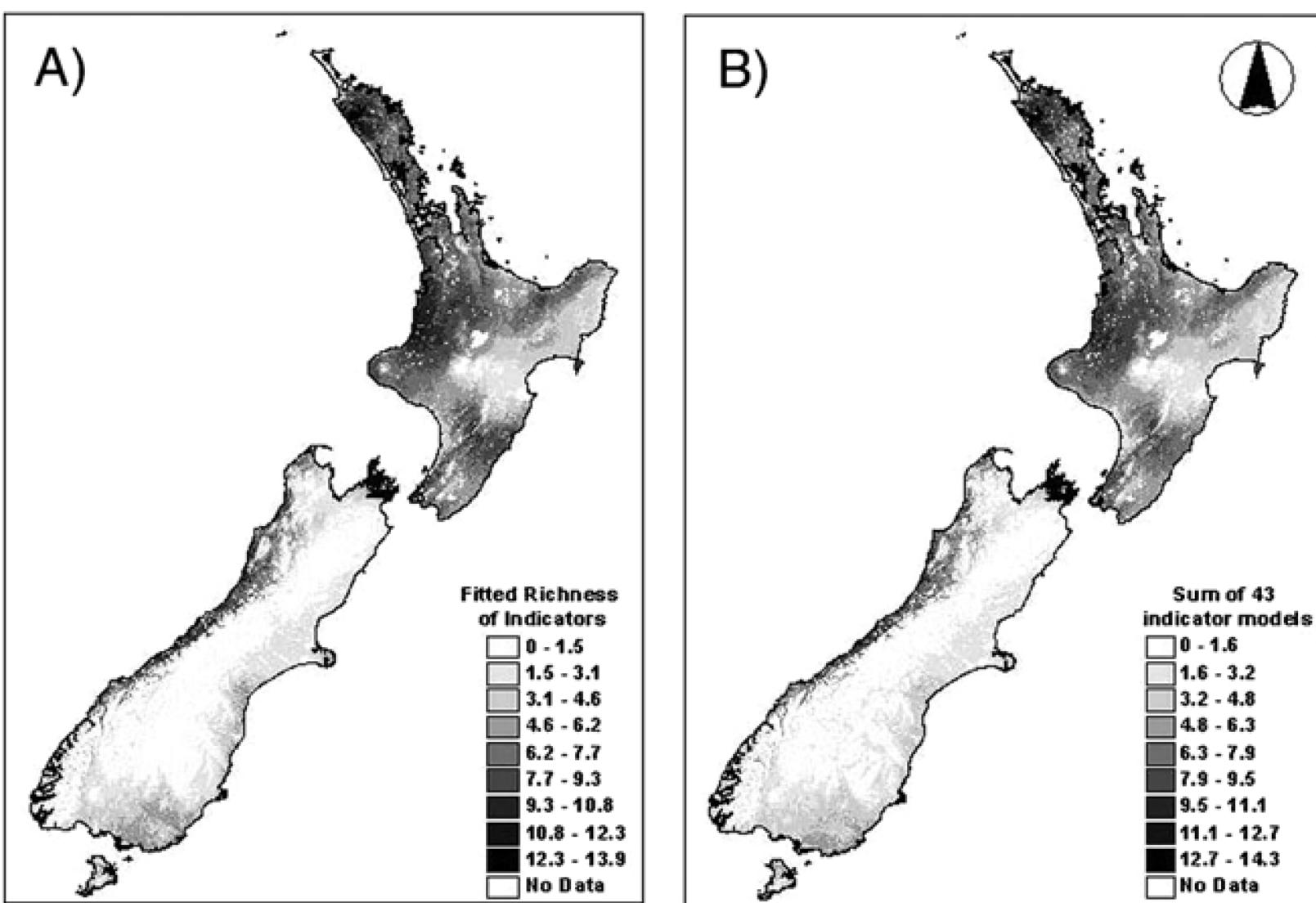


# RICHNESS MODELLING

**Lehmann et al. 2002.** Assessing New Zealand fern diversity from spatial predictions of species assemblages. *Biodiversity and Conservation* 11, 2217-2238

- **Objective:** Biodiversity modelling.
- **Methods:** Stacked SDMs compared with a classic richness-modelling method.
- **Results:** High agreement between models.
- **Conclusion:** Stacked SDMs allow to identify biodiversity hotspots, useful in conservation planning.

# Lehmann et al. 2002

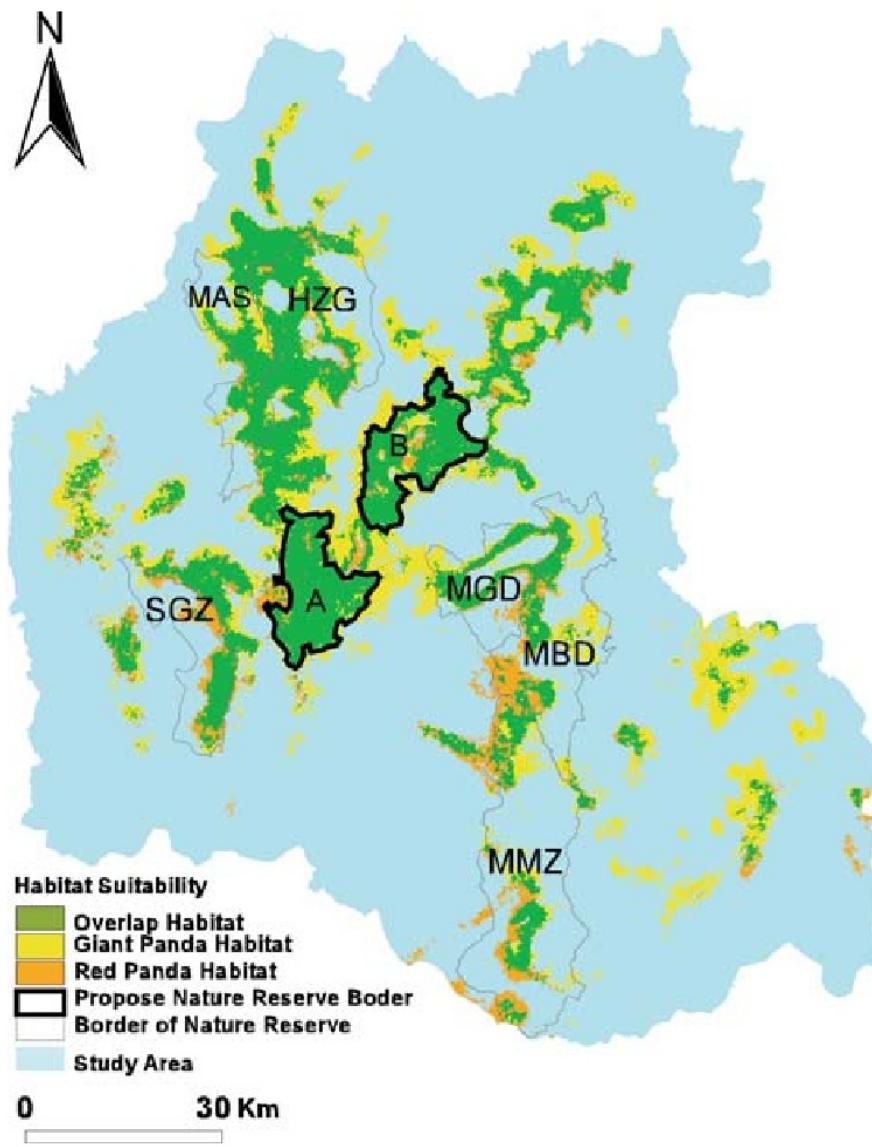


# CONSERVATION PLANNING

**Dunwu et al. 2002.** Ecological niche modeling of the sympatric giant and red pandas on a mountain-range scale. *Biodiversity and Conservation* 18, 2127-2141

- **Objective:** Find suitable conservation areas for Red and Giant pandas.
- **Methods:** Overlap of SDMs of each species.
- **Results:** Current reserves have non-optimal designs, SDMs can help to make them more efficient.
- **Conclusions:** SDMs can be useful for conservation planning!

# Dunwu et al. 2002

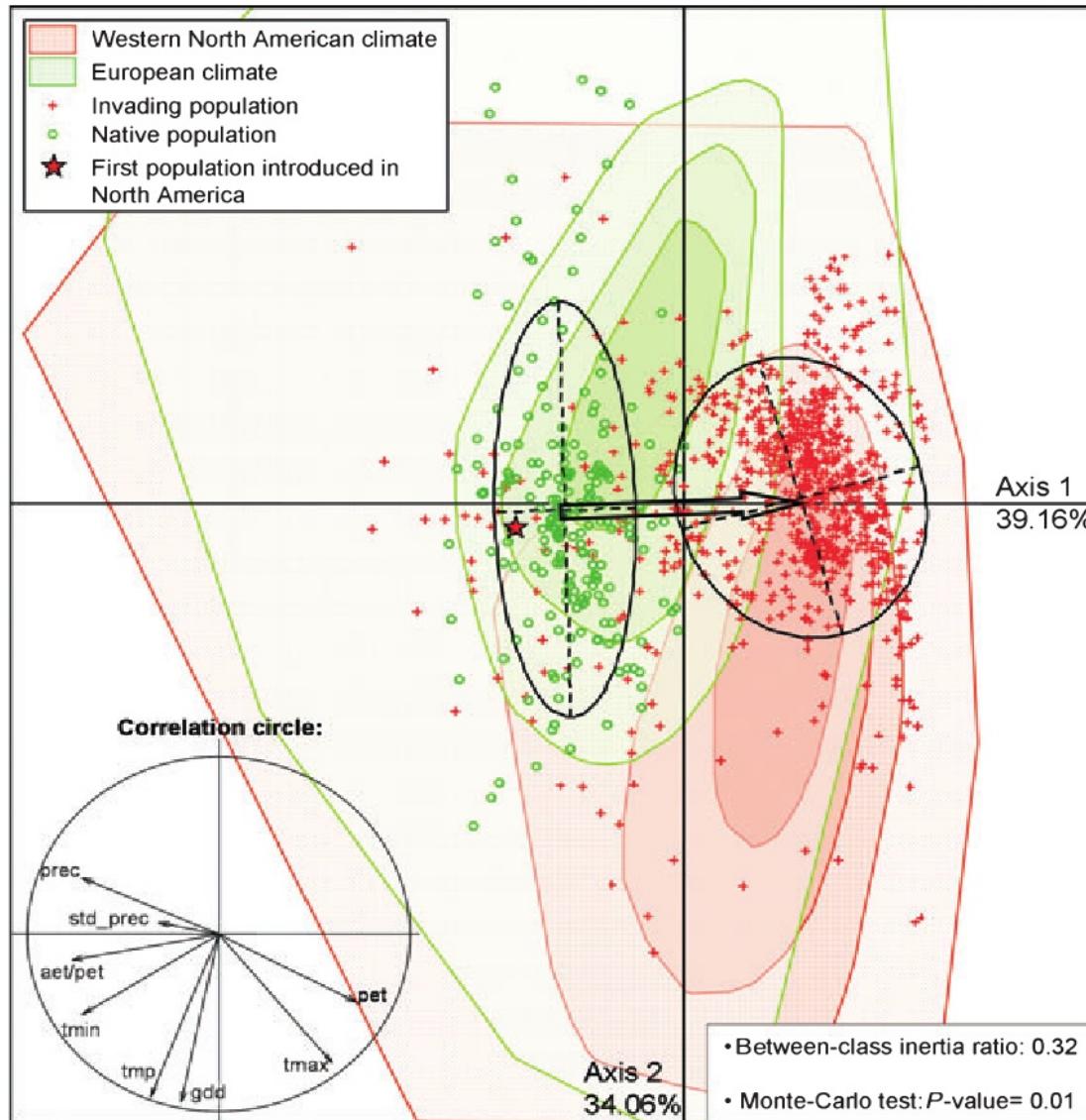


# UNDERSTANDING BIOLOGICAL INVASIONS

**Broenninmann et al. 2007.** Evidence of climatic niche shift during biological invasions.  
*Ecology Letters* 10, 701-709

- **Objective:** Evaluate how niche dynamics promotes invasions.
- **Methods:** SDMs of an invasive species in native and invaded ranges.
- **Results:** The target species expanded its niche after invasion.
- **Conclusion:** SDMs can predict introduction areas, but can have issues to forecast invaded ranges due to niche expansion.

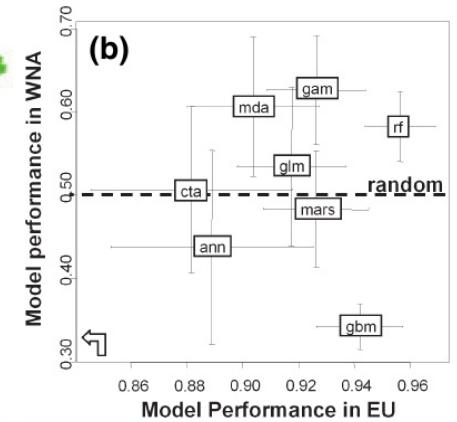
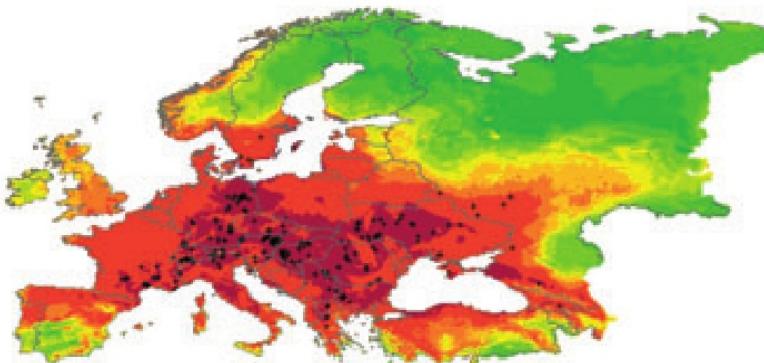
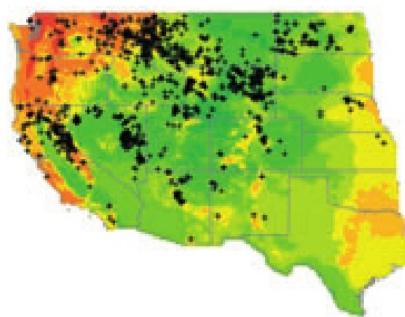
# Broenninmann et al. 2007



# Broenninmann et al. 2007

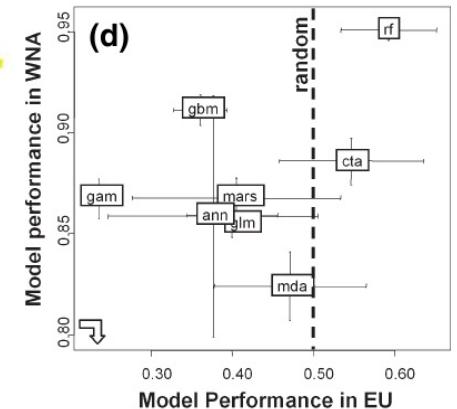
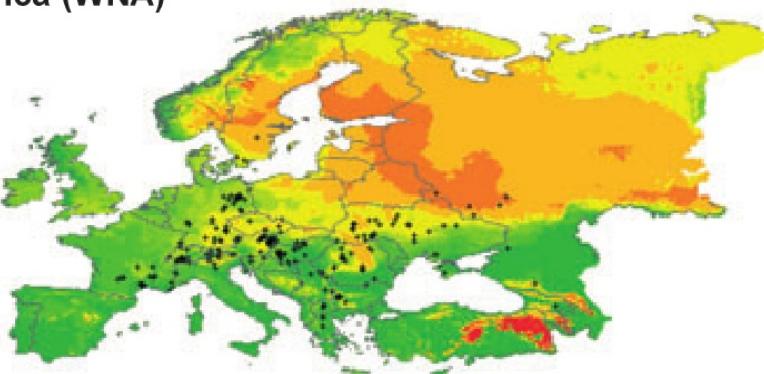
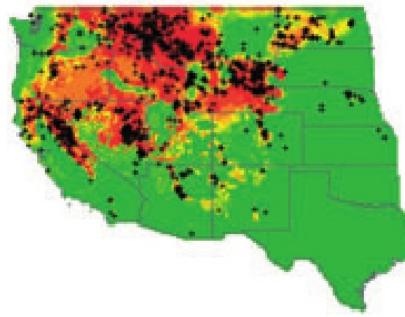
## Calibration in Europe (EU)

(a)



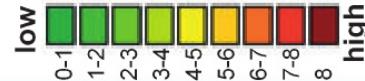
## Calibration in Western North America (WNA)

(c)

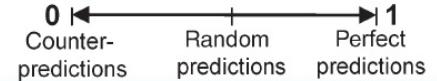


Occurrences of  
Centaurea maculosa : +

Predicted climatic  
suitability :



Model  
performance :

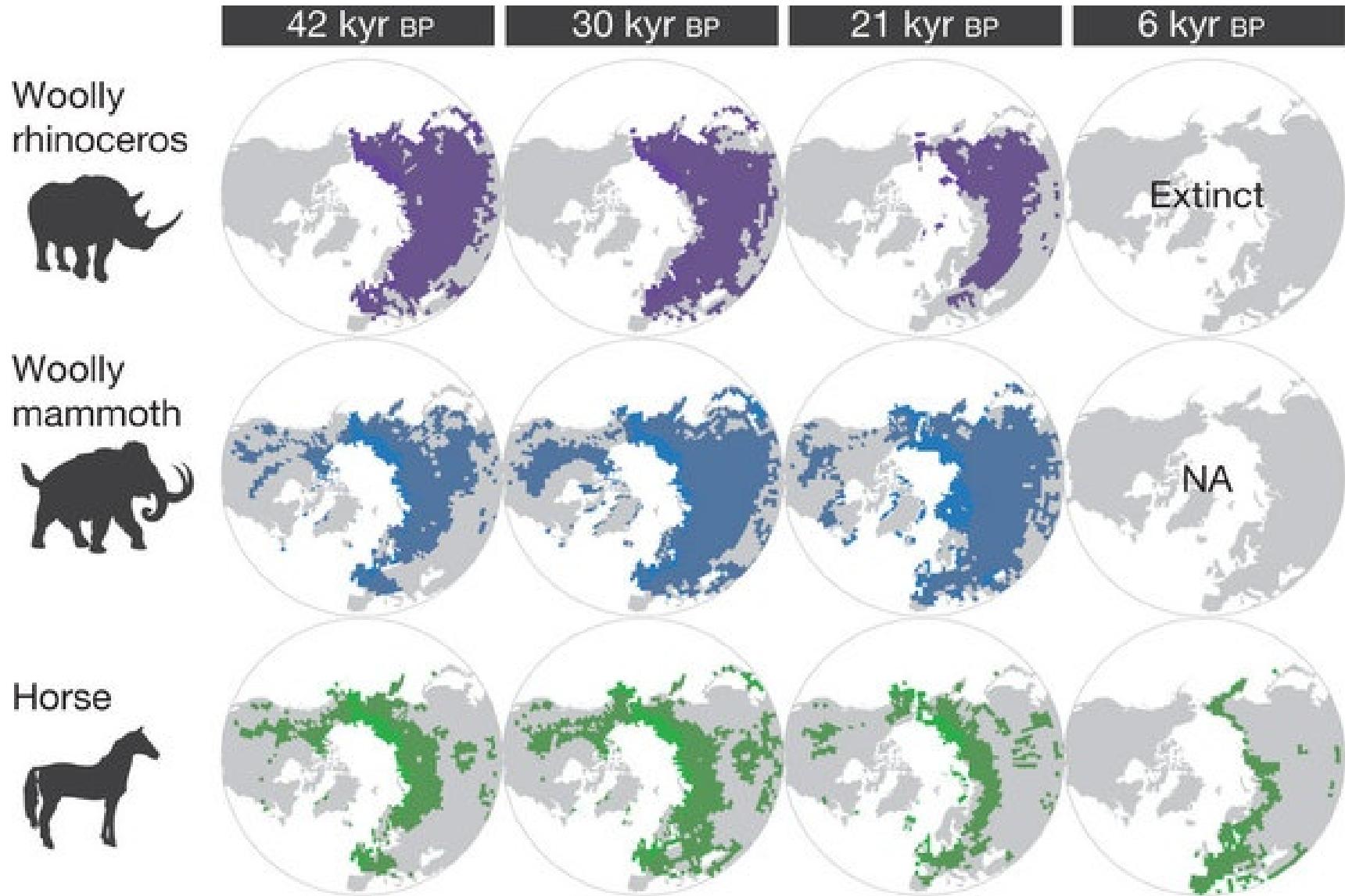


# PALAEODISTRIBUTION ANALYSIS

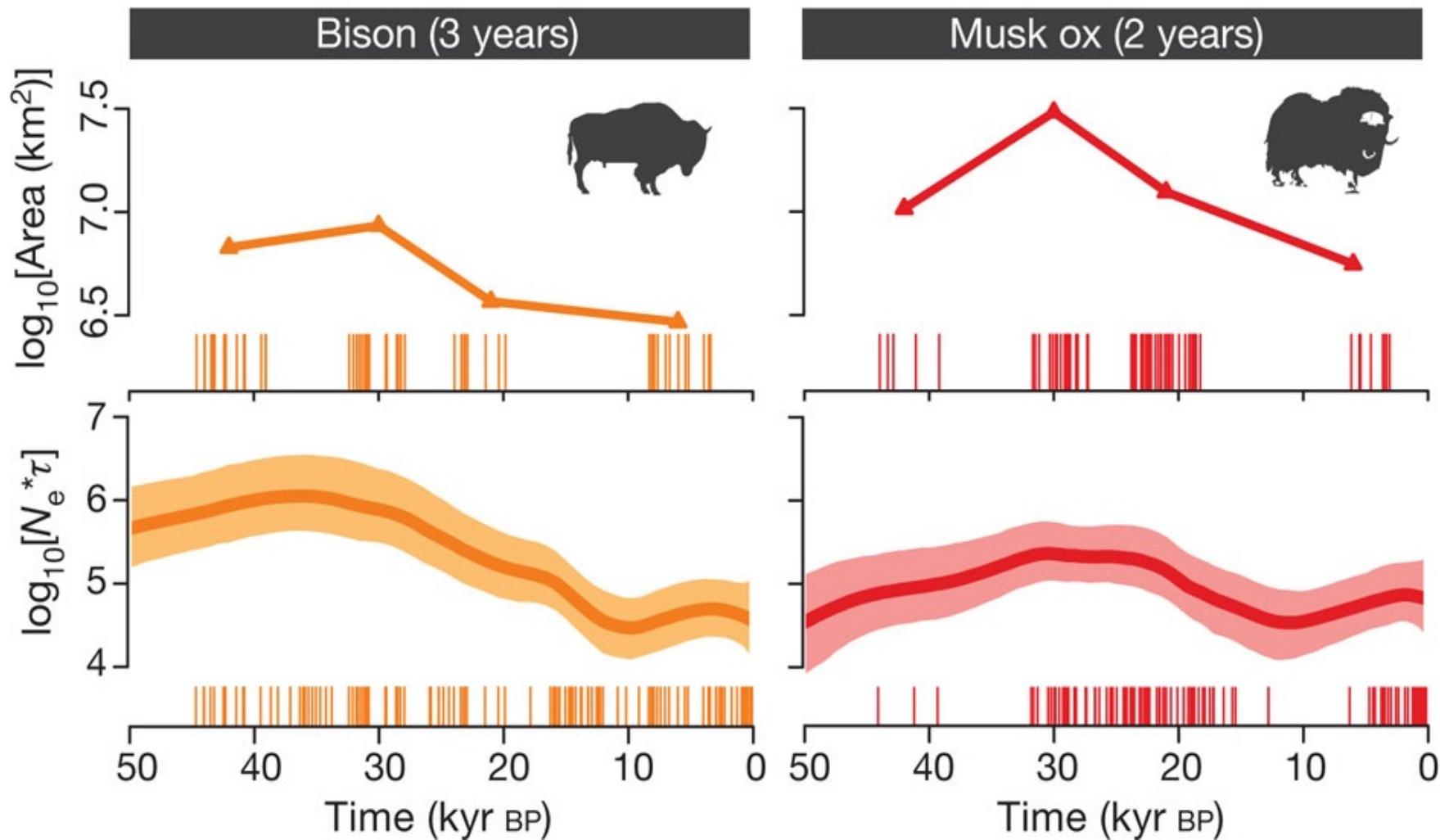
**Lorenzen et al. 2011.** Species-specific responses of Late Quaternary megafauna to climate and humans. *Nature* 479, 359-364

- **Objective:** Test the effect of climate and humans on the distribution of Quaternary megafauna.
- **Methods:** SDMs, ancient DNA, fossil record.
- **Results:** Different species showed different responses to climate change and huma presence.
- **Conclusiones:** SDMs coupled with other methods can provide new insights on palaeoecological dynamics.

# Lorenzen et al. 2011



# Lorenzen et al. 2011

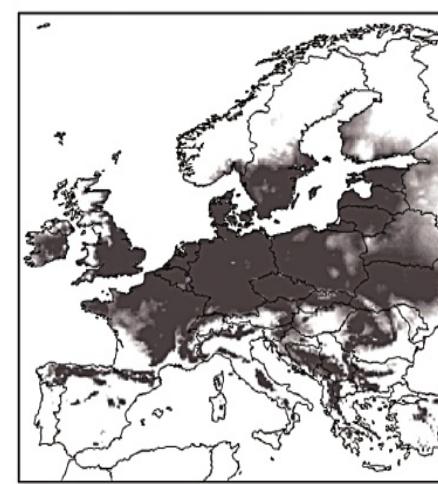
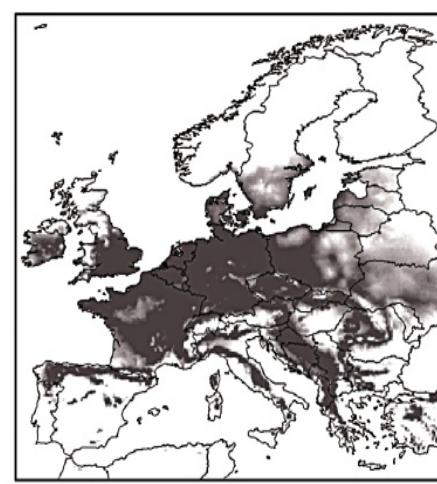
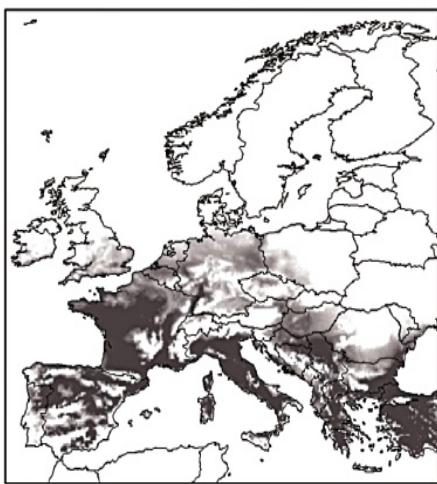
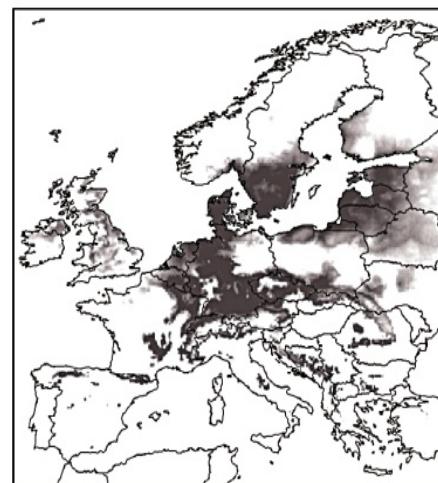
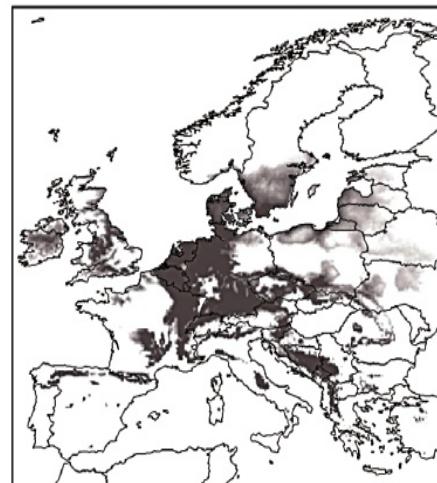
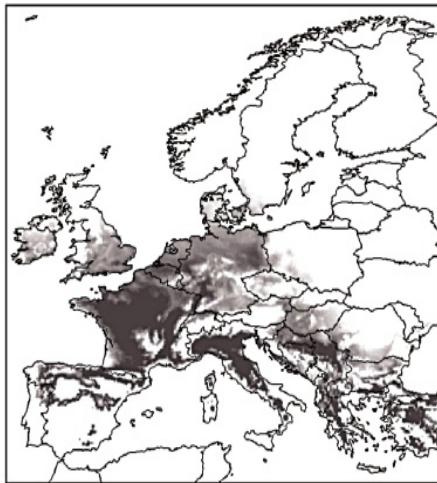
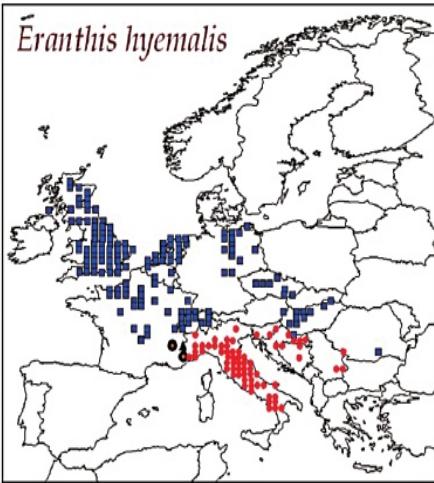


# RANGE SHIFT UNDER CLIMATE CHANGE

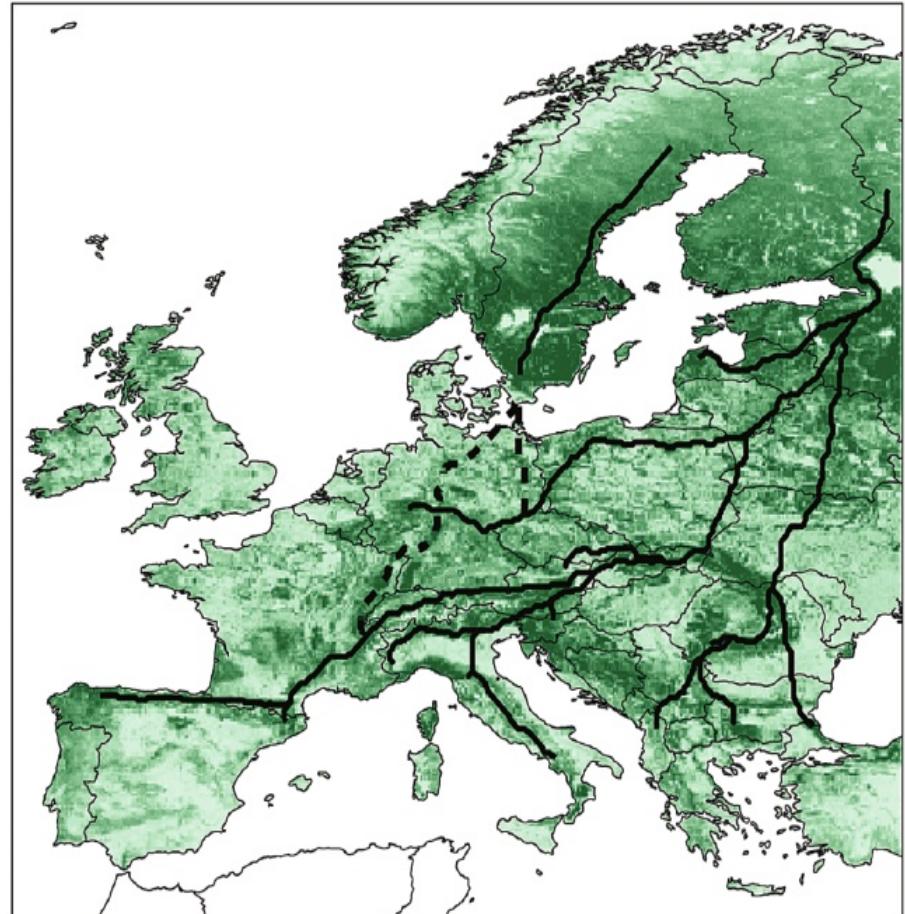
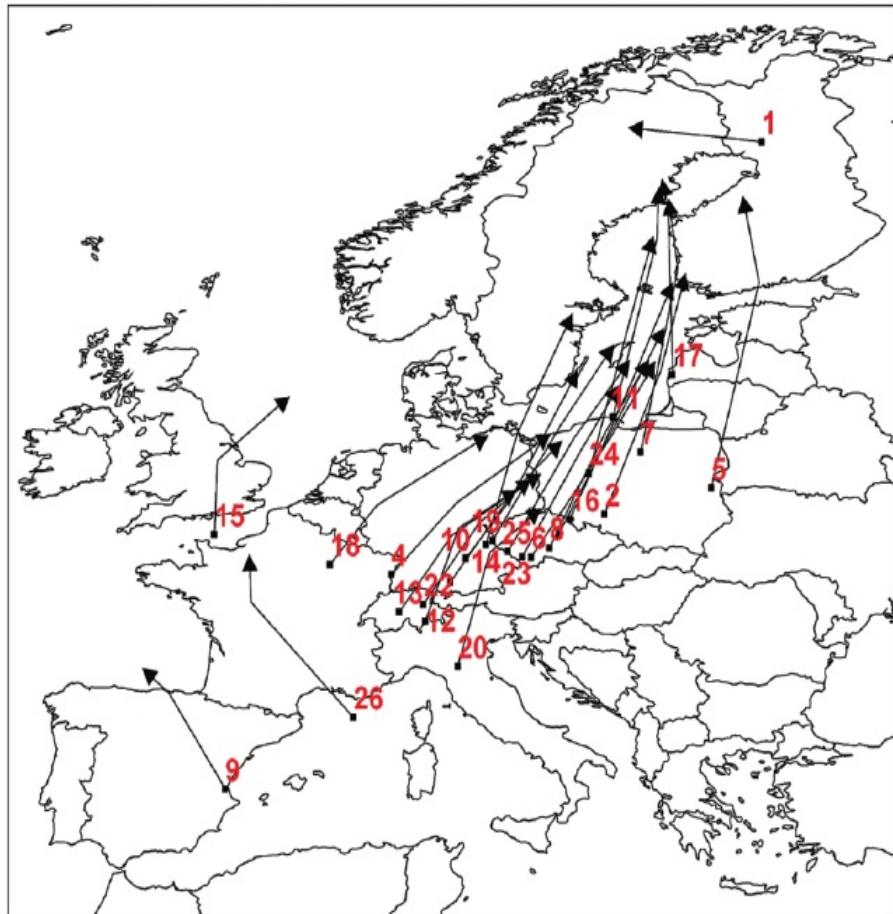
**Skov y Svenning 2004.** Potential impact of climatic change on the distribution of forest herbs in Europe. *Ecography* 27, 366-380

- **Objective:** Forecast range shift under climate change in Europe.
- **Methods:** Fuzzy bioclimatic envelopes.
- **Results:** Range reduction for all studied species.
- **Conclusions:** Low dispersal rates can hinder the ability of species to track their bioclimatic niche.

# Skov y Svenning 2004



# Skov y Svenning 2004





# PRIMEROS PASOS CON R y Rstudio

- Tools>Global Options> General> untick “Restore .Rdata”
- Tools>Global options>Pane Layout, tick “Files” (upper right panel)
- Tools> Global options> Packages> CRAN mirror> Tick a local mirror.
- Menú Session> Set Working Directory>  
c:/taller1/intro\_R/datos
- Open .../**taller1/intro\_R/introduccion.R**

# **PREPARING ENVIRONMENTAL PREDICTORS AND PRESENCE RECORDS**

# MAIN R SCRIPT

**1\_prepara\_variables\_y\_presence.R**

# COLLINEARITY ANALYSIS

- Two steps process:
  - Correlation matrix visualized as dendrogram to select meaningful predictors.
  - VIF (variance inflation analysis) to make sure that remaining predictors are not a linear combination of the other remaining predictors.

# PREPARING PRESENCES

- Adjusting presences to the study area
- Checking taxonomic coherence
- Removing fossil record
- Remove duplicated records
- Check accuracy of coordinates
- Reduce spatial autocorrelation

# PREPARING BACKGROUND, PSEUDO-ABSENCE AND ABSENCE DATA

- **Background:** comprehensive sampling of available ecological conditions (overlaps presences).
- **Pseudo-absence:** random points not overlapping presences.
- **Absences:** we don't have real absences, but we will create them following a realistic criteria (just for pedagogic purposes!)

# PRESENCE TABLES

species	x	y	var_1	var_2	var_n	presence
Ursus	26.48	63.01	456	856	...	1
Ursus	23.60	50.20	546	452	...	1
Ursus	30.20	57.34	569	145	...	1
Ursus	32.10	62.35	412	456	...	0
Ursus	28.45	27.41	158	658	...	0
Ursus	30.24	31.22	123	456	...	0
...	...	...	...	...	...	...

That's all Folks!