

# Niche Modelling: An Introduction

Chris Yesson



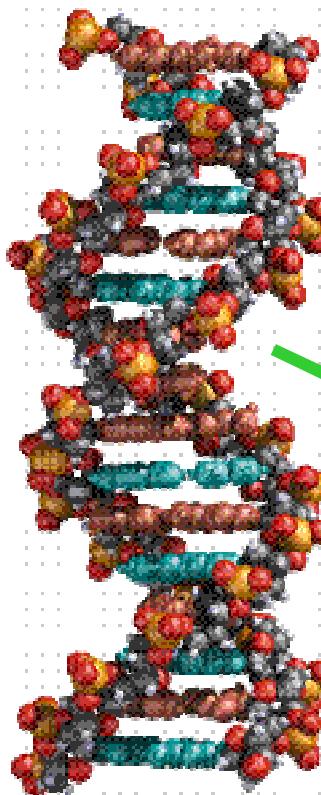
Institute of Zoology

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# Where does diversity come from?

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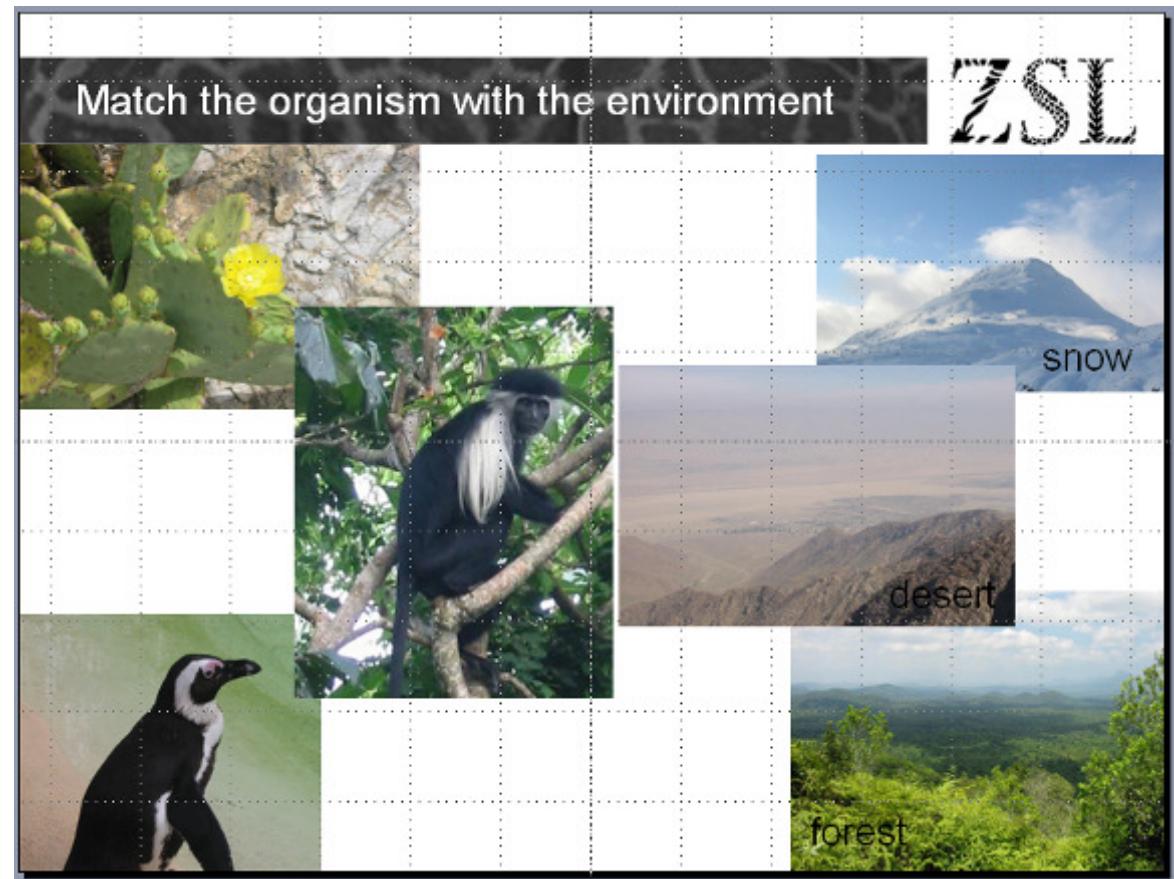
- Mutation
  - Selection
  - Chance
  - Time
- 
- **Geology**
  - **Geography**
  - **Climate**



# Niche concepts are innate

ZSL

- *A fish out of water*
- *He took to it like a duck to water*
- *As happy as a pig in ... clover*





# What is a niche?



- A niche is an ecological construct defining the optimum environment for growth, reproduction and survival of a species
- Niches are defined by three main factors:
  - Substrate
  - Microclimate
  - Competition

# What is a climatic niche?



- A bioclimatic niche is the area in which the climate is suitable for a species to succeed
- Consequentially it does not take into direct account other factors such as soil and competition
- Areas of a bioclimatic niche can be empty of the expected species because:
  - It can't get there
  - The substrate is wrong
  - It can't compete with other species

- One way to investigate species response to climate is through examination of climatic preferences by constructing bioclimatic niche models (=species distribution models = environmental niche models)
- Preferences of a given species, based on its known distribution
- Model using the climate parameters correlating with this

# Developing predictive ‘niche’ models

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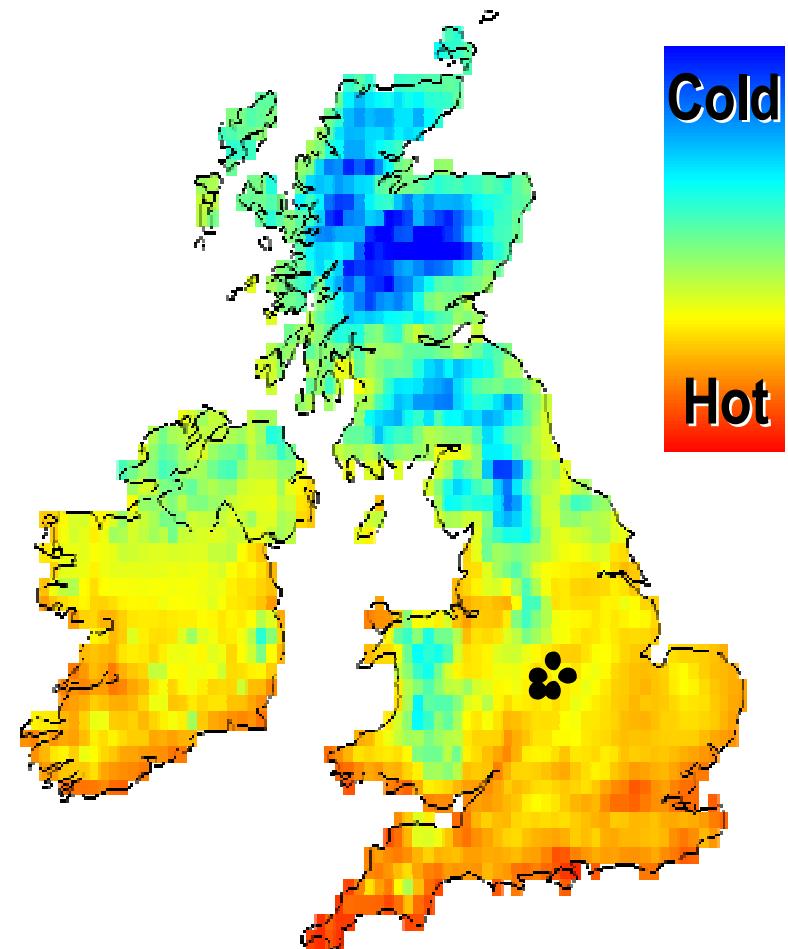
## 1. Distribution data



# Developing predictive ‘niche’ models

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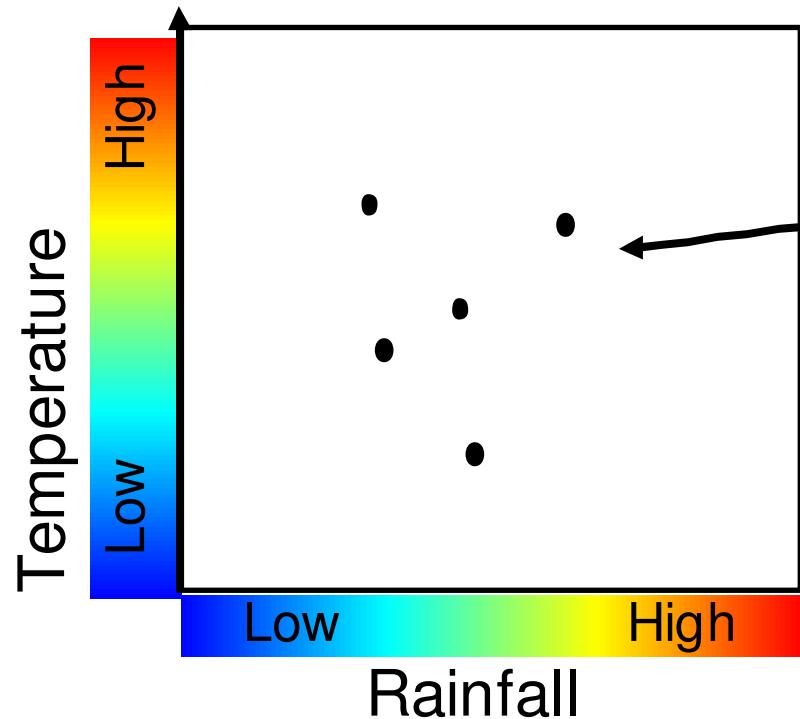
## 2. Environmental data



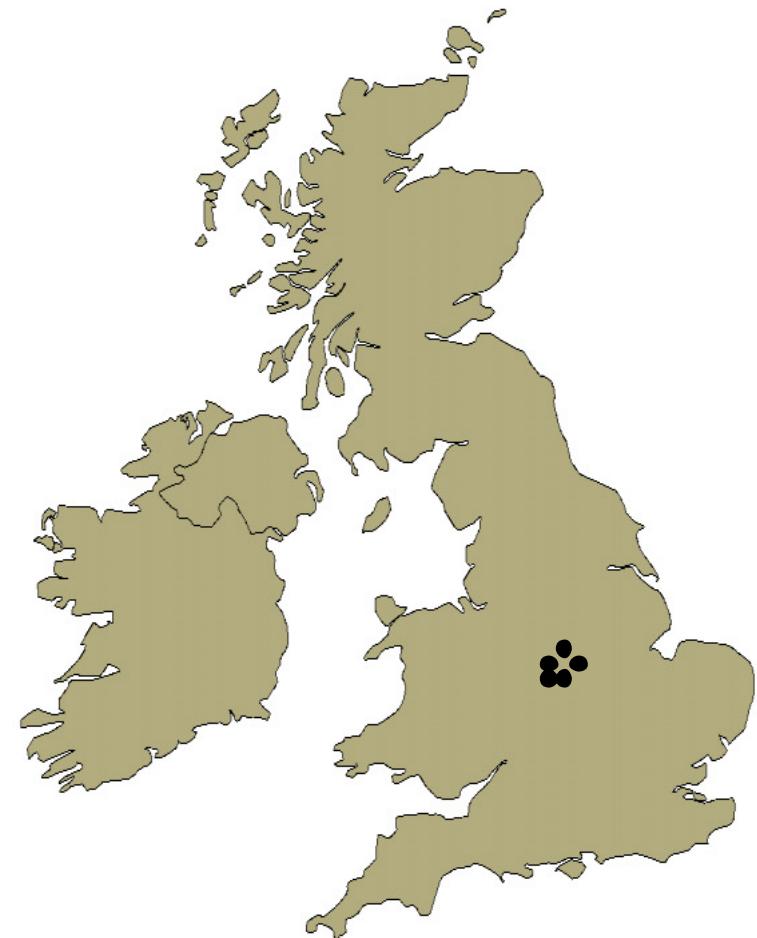
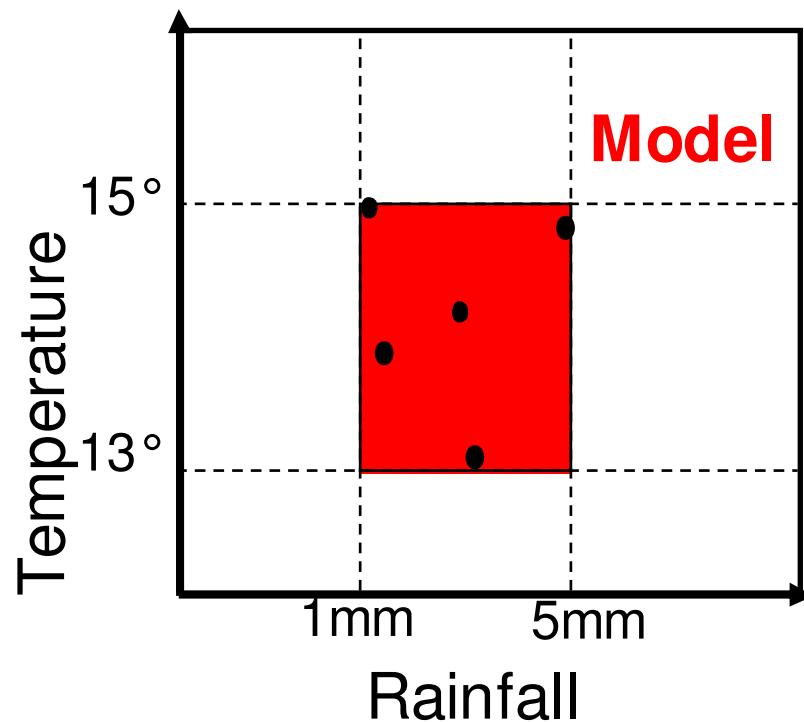
# Developing predictive ‘niche’ models

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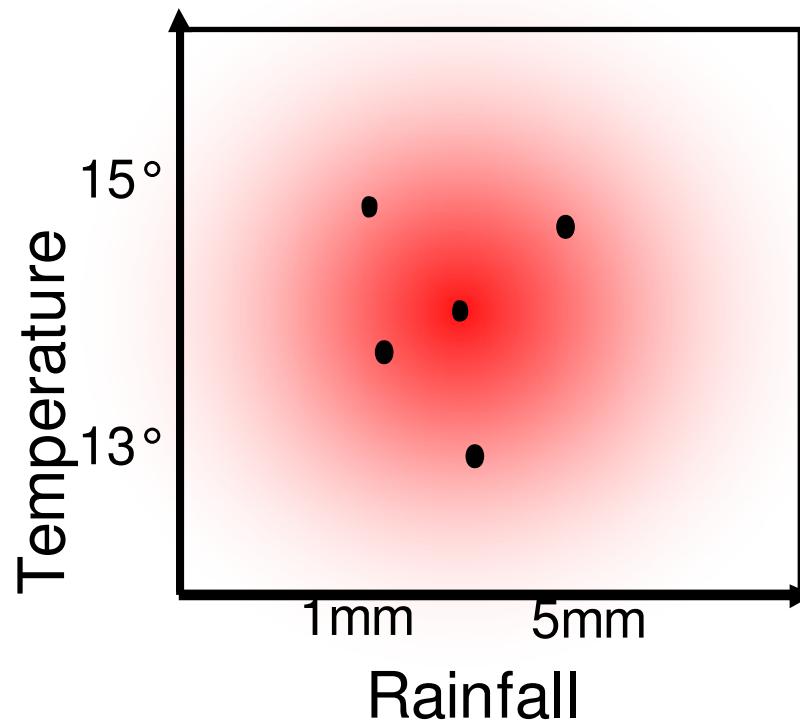
## 3. Profile species



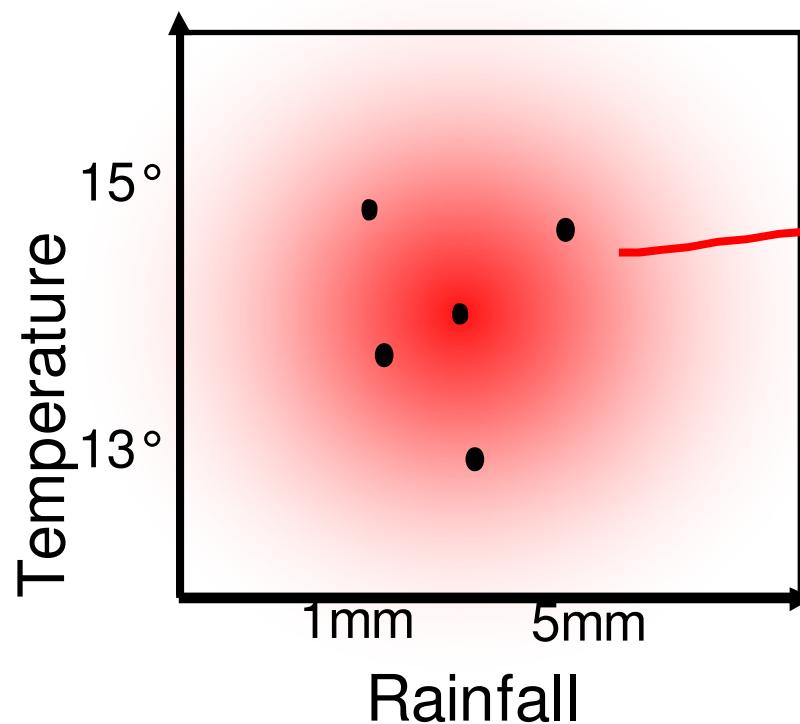
## 4. Develop model



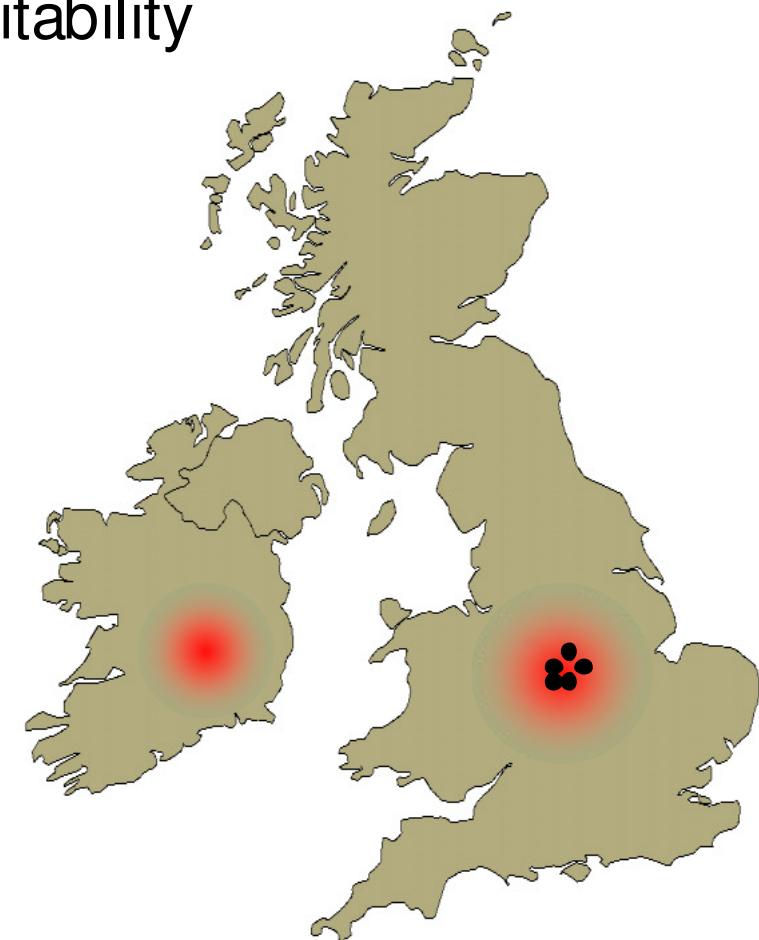
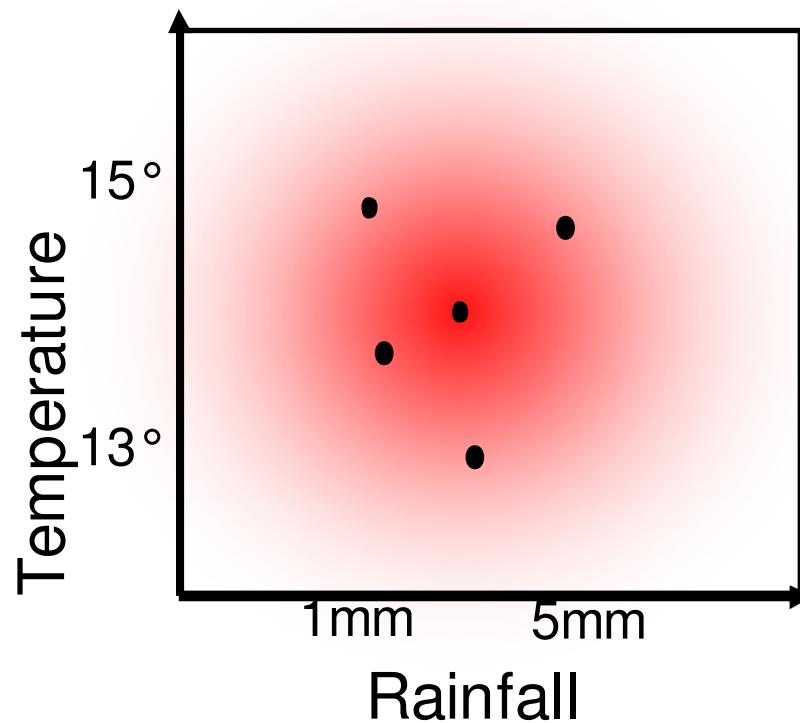
## 4. Develop model



## 5. Project model into environment



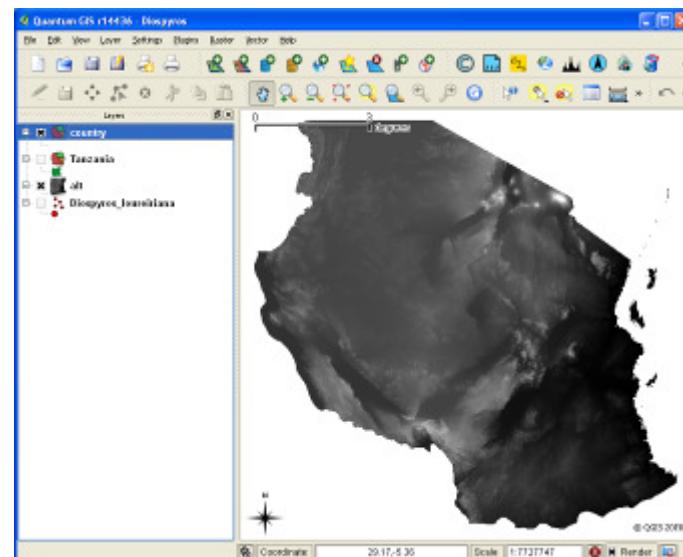
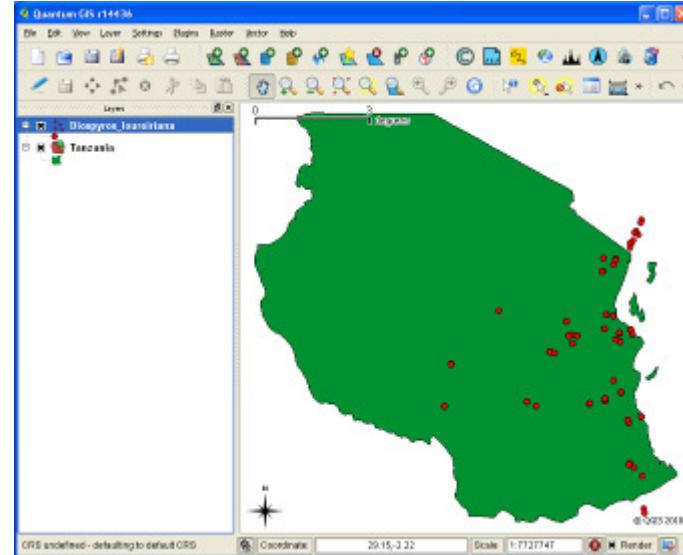
## 6. Discover areas of niche suitability



# Common data sources



- Species distribution
  - museums / herbaria  
(<http://data.gbif.org/>)
  - literature
- Environmental data
  - Global climate grids  
(<http://www.worldclim.org/>)
  - Global topographic grids  
(strm30)
- *More details this afternoon*



# Niche modelling software



- Biomapper
- Desktop GARP\*
- DIVA-GIS
- Maxent
- R packages
- Many others

... and

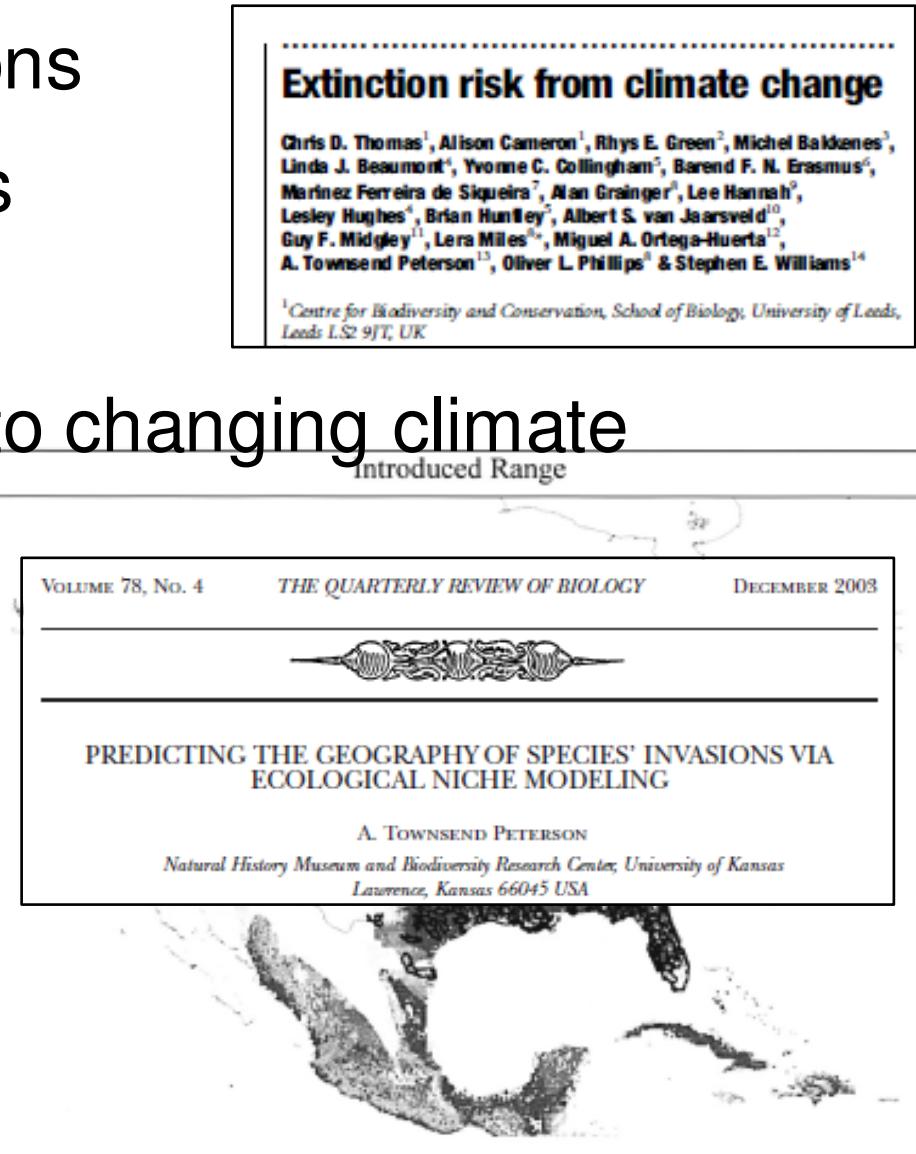
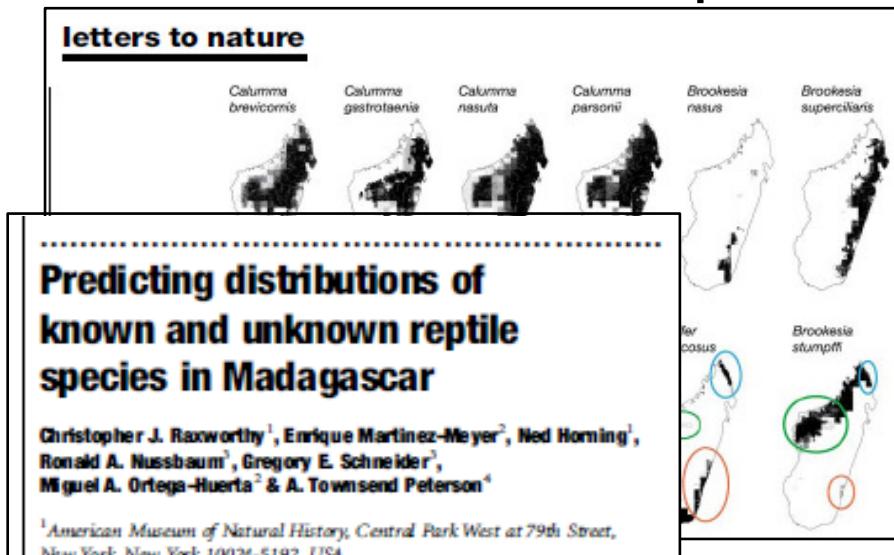
- **openModeller**

The screenshot shows the homepage of the openModeller project. At the top, the word "open" is written in a small, italicized font above "Modeller" in a large, bold, blue sans-serif font. Below the title, a blue banner contains the text: "openModeller is a fundamental niche modelling library, providing a uniform method for modelling distribution patterns using a variety of modelling algorithms." To the right of the banner, there is a diagram illustrating the process: a map of South America with a red dot representing "Species Occurrence" is combined with a stack of three environmental layers (represented by white rectangles) using a plus sign, resulting in a map of "Probability Distribution". On the left side of the page, there is a sidebar with links to "Home", "News", "Overview", "Developers", "Documentation", "View Source Code", "Environmental Data", "Screenshots", "oM Desktop", "Algorithms", "Publications", "Resources", "Download", and "Bugs!". Below the sidebar, there is a "SOURCEFORGE.NET" logo and a "FAPESP" logo. A banner at the bottom of the page announces "openModeller 1.1 released". The main content area contains a brief description of the project's purpose and development, mentioning the Centro de Referência em Informação Ambiental (CRIA), Escola Politécnica da USP (Poli), and Instituto Nacional de Pesquisas Espaciais (INPE) as partners, and funding from Fundação de Amparo à Pesquisa do Estado de São Paulo (FAPESP) and the Ineofish project.

# Niche Modelling studies



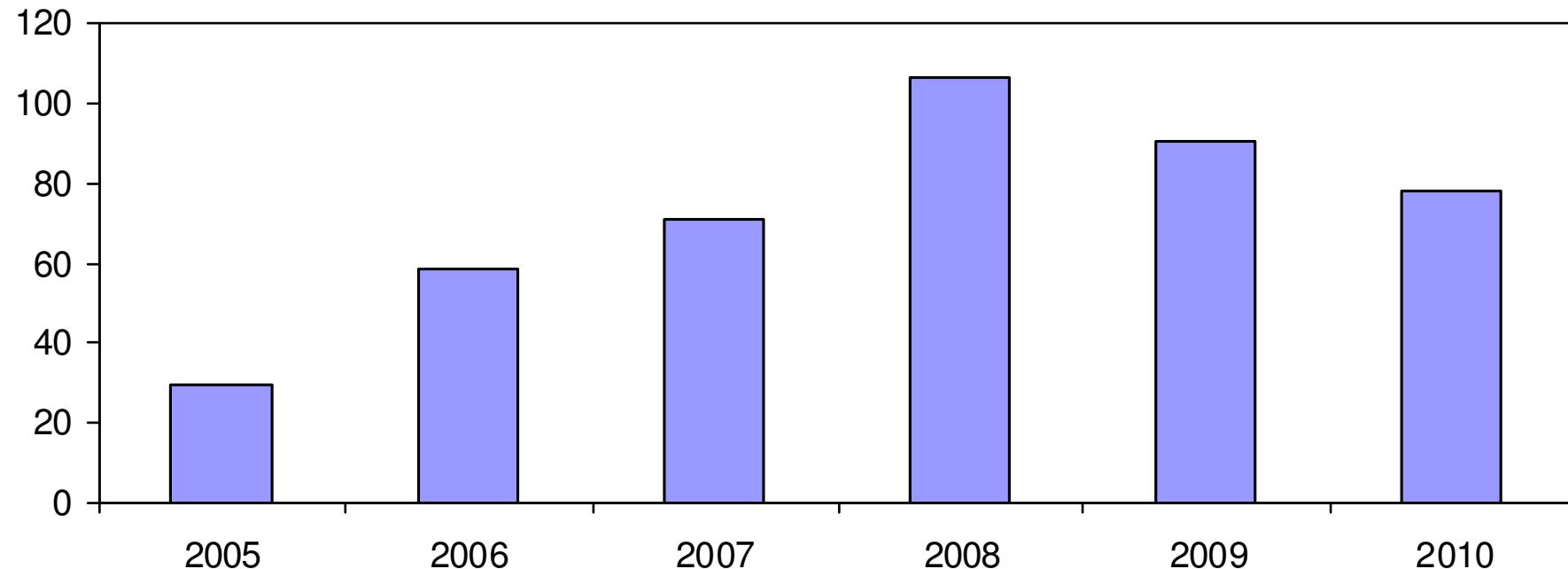
- Used to predict distributions
  - Closely related species
  - Invasive species
  - Distribution shifts due to changing climate
  - Undiscovered species!



# Niche modelling studies

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WOK citation search for "Niche model" OR "Species distribution model" or  
"habitat suitability model" or "bioclimatic model"



Thomas et al. (2004) Extinction risk from climate change. Nature 427: 145-148



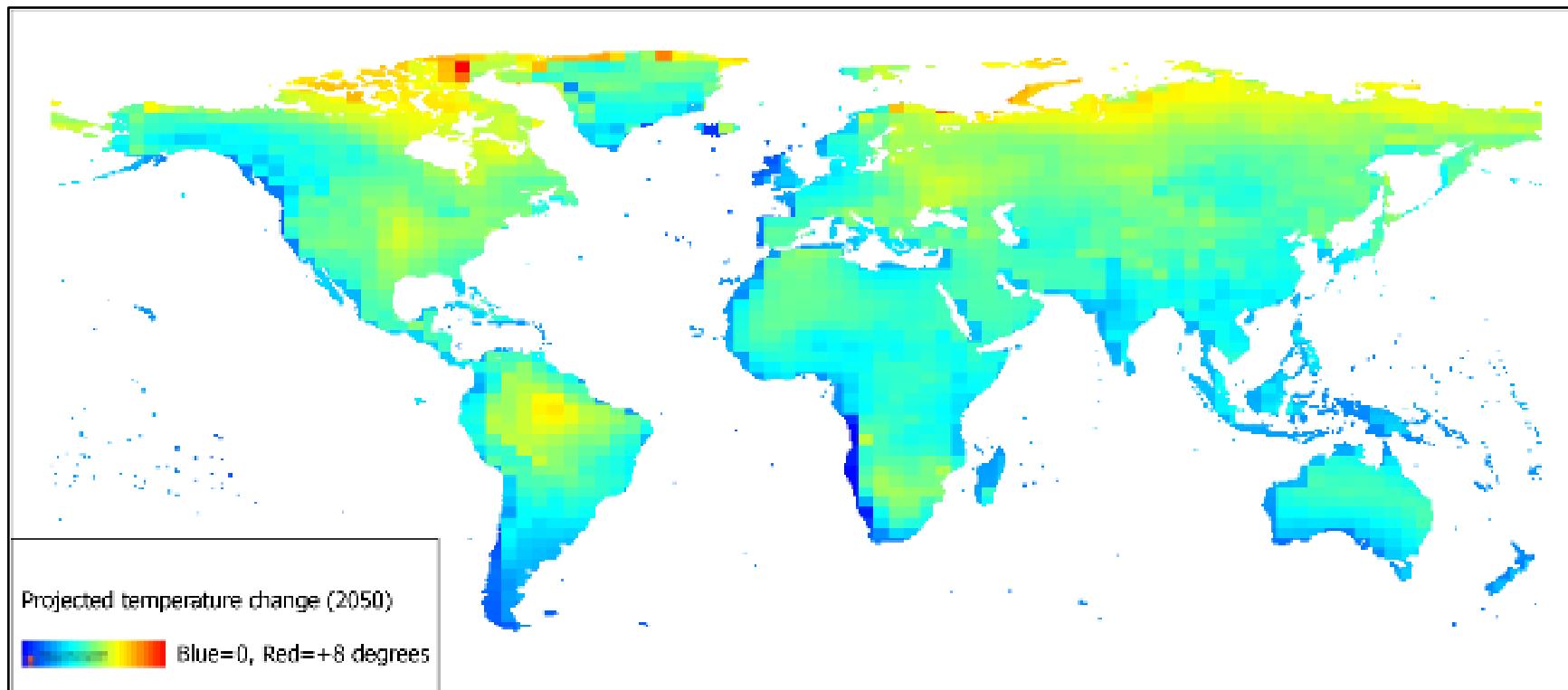
- Meta analysis of niche modelling studies using many different methodologies
- Modelled thousands of terrestrial plants & animals
- Projected models into a mid-range future scenario
- 15 - 37% species “will be ‘committed to extinction’”
- Lots of publicity, lots of criticism, lots of citations (>1,500)



# Practical applications – climate change



- How will climate change affect species distribution?



# Legumes

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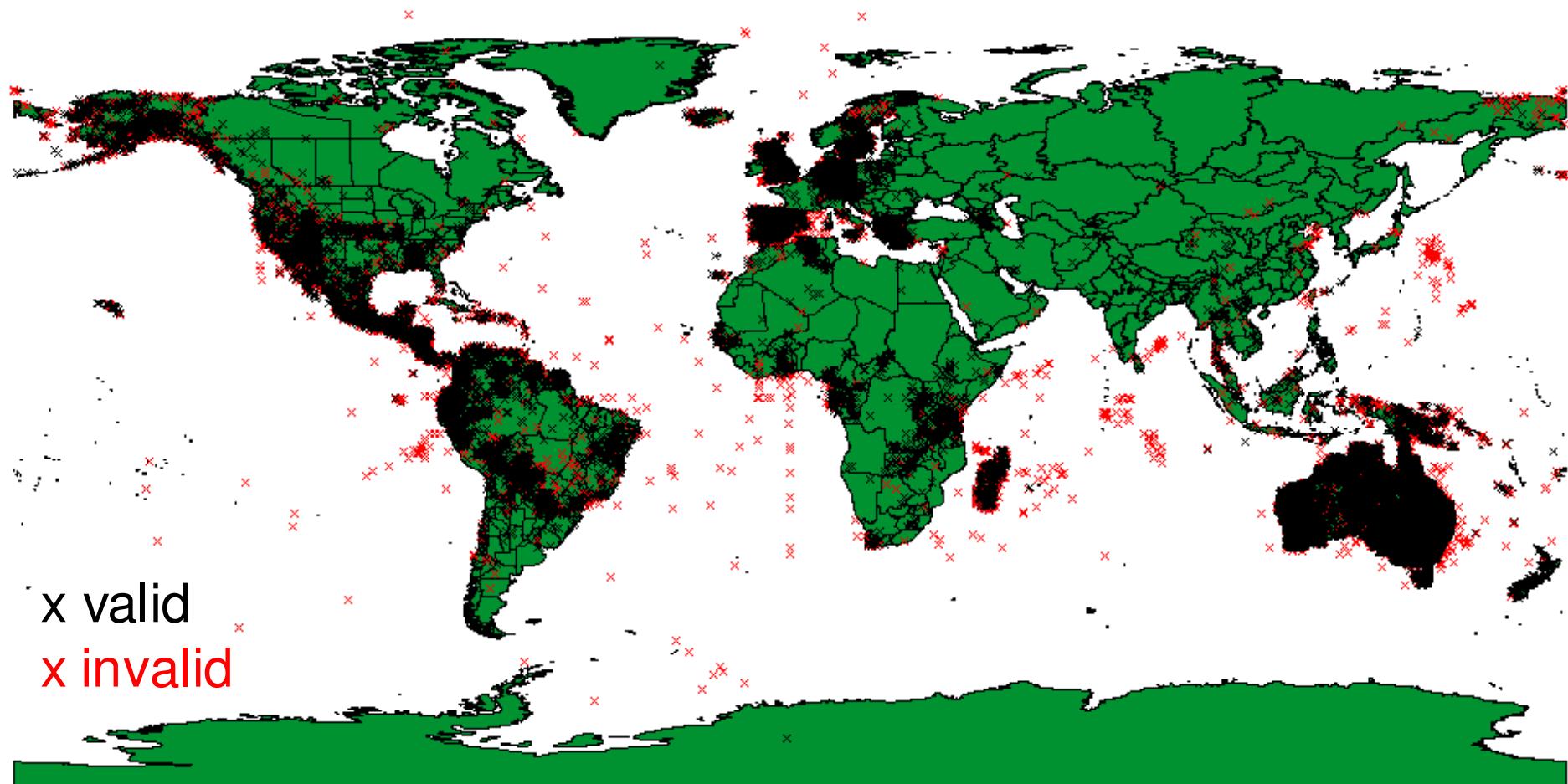
- Legumes (Pea family) include ~20,000 species
- Many are economically important for food (beans & peanuts), wood (*Acacia*) and other uses (e.g. dyes, forage)
- How will climate change affect global diversity?



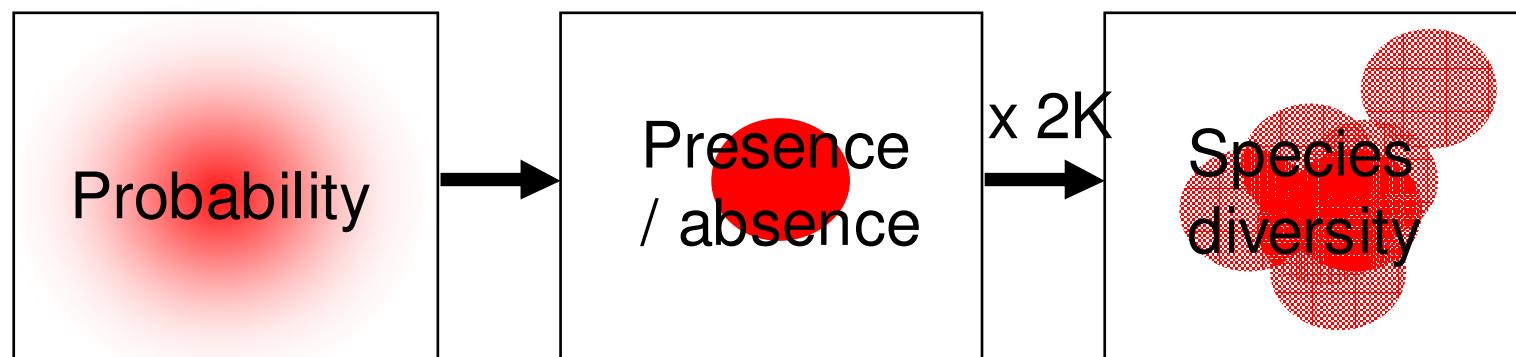
# Legumes – assessing global diversity



- 20K species
- Only 2K species with valid data!

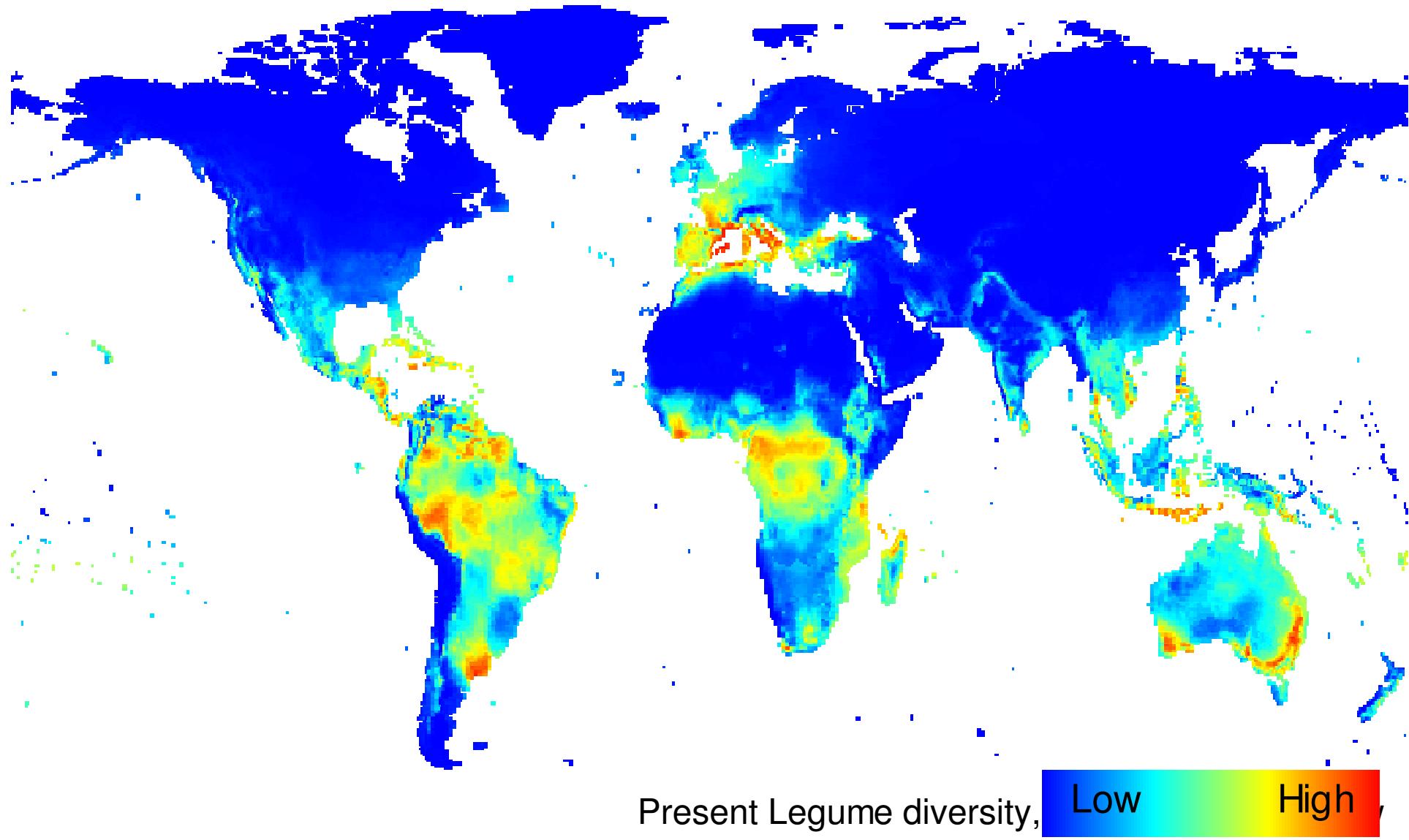


- Model species with data
- Choose threshold to estimate presence-absence
- Overlay models to produce map of diversity



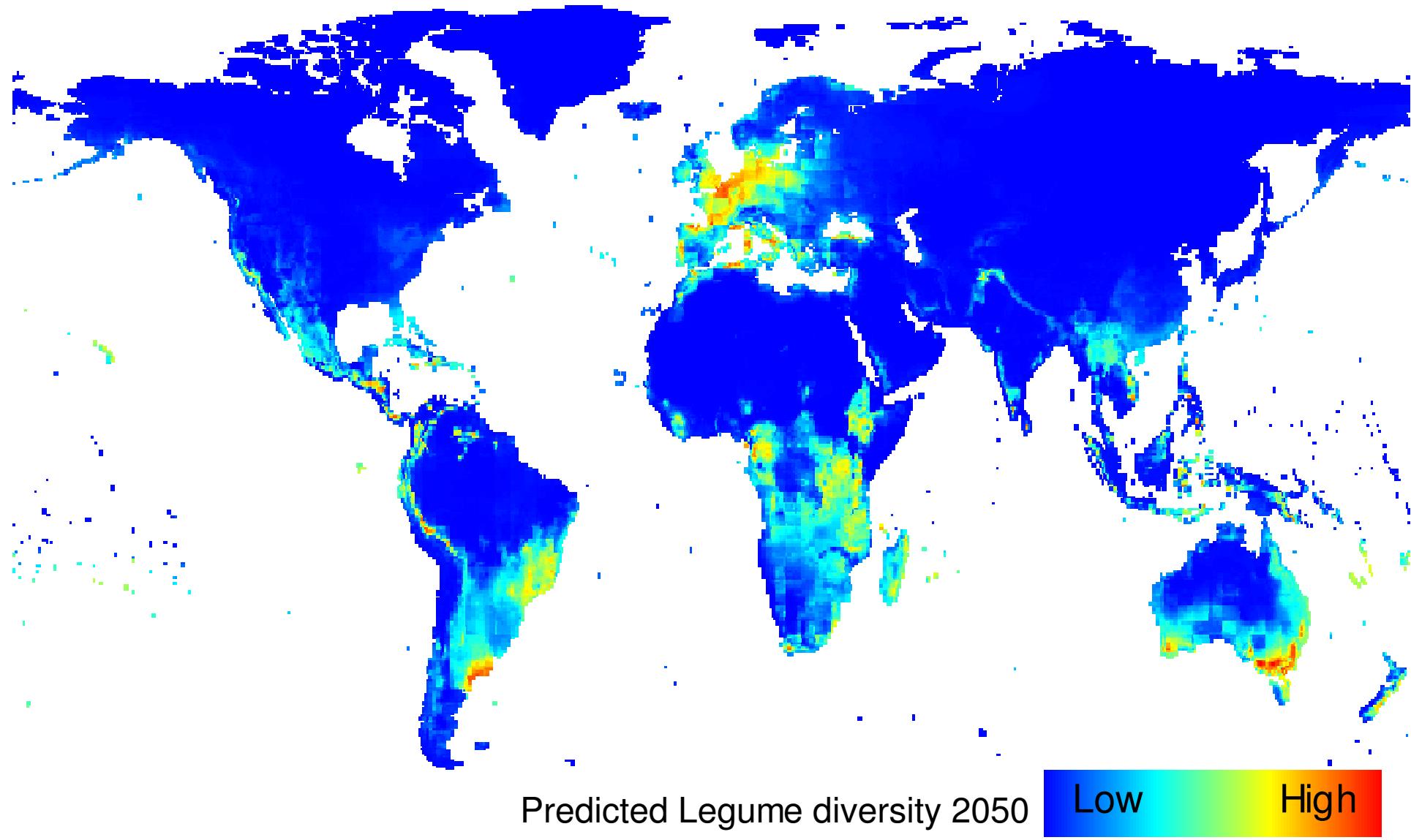
# Estimating change in diversity

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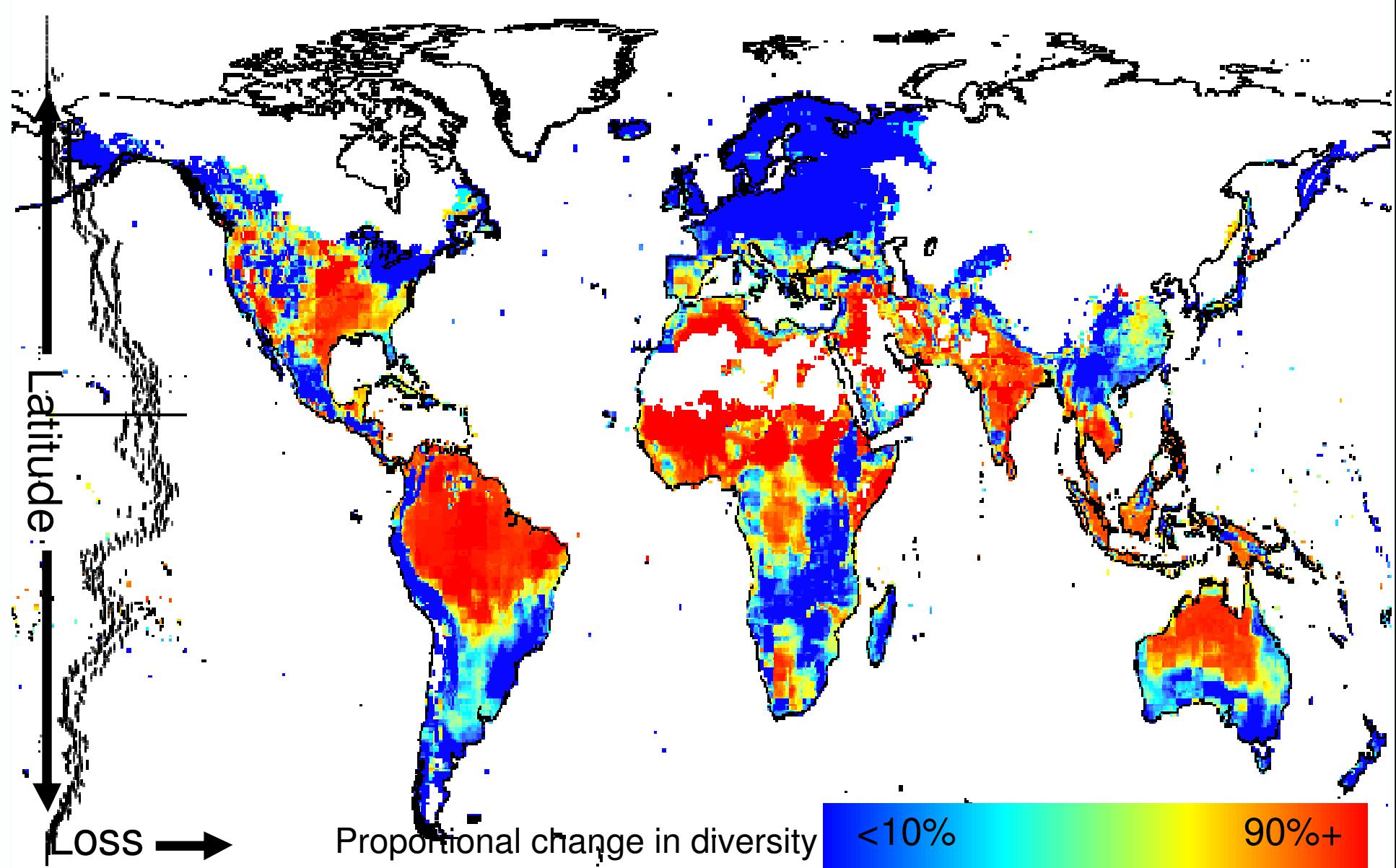
# Estimating change in diversity

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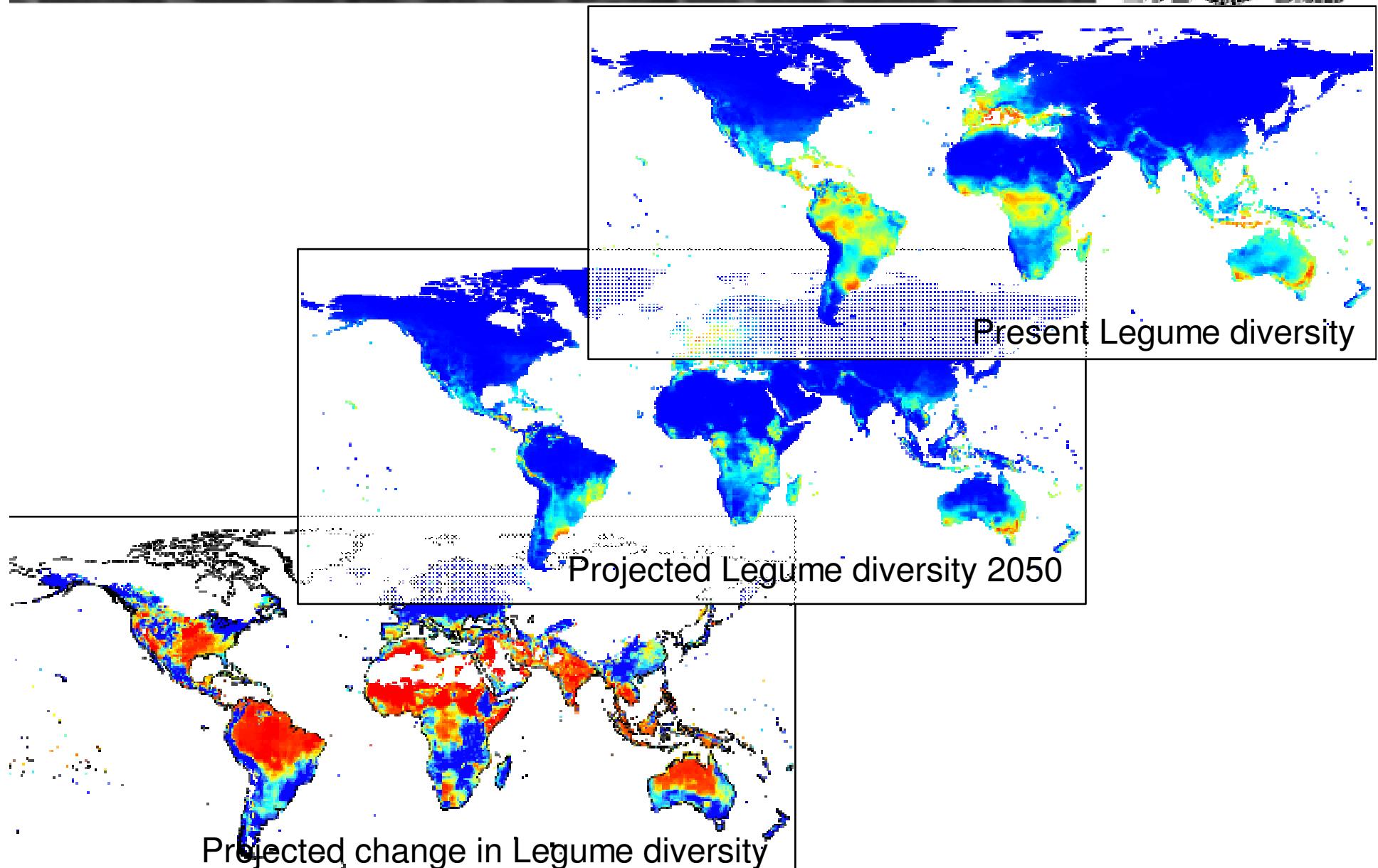
# Estimating change in diversity

ZSL



# Estimating change in diversity

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- Practical
  - Invasive species (weeds)
  - Disease
  - Data deficient species
  - Marine protected areas
- Theoretical
  - Diversity through time
  - Evolutionary patterns

# Practical applications - weeds

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- *Cyperus rotundus* is *the world's worst weed*
- It can dramatically reduce crop production
- Are some areas at risk from the spread of this weed?

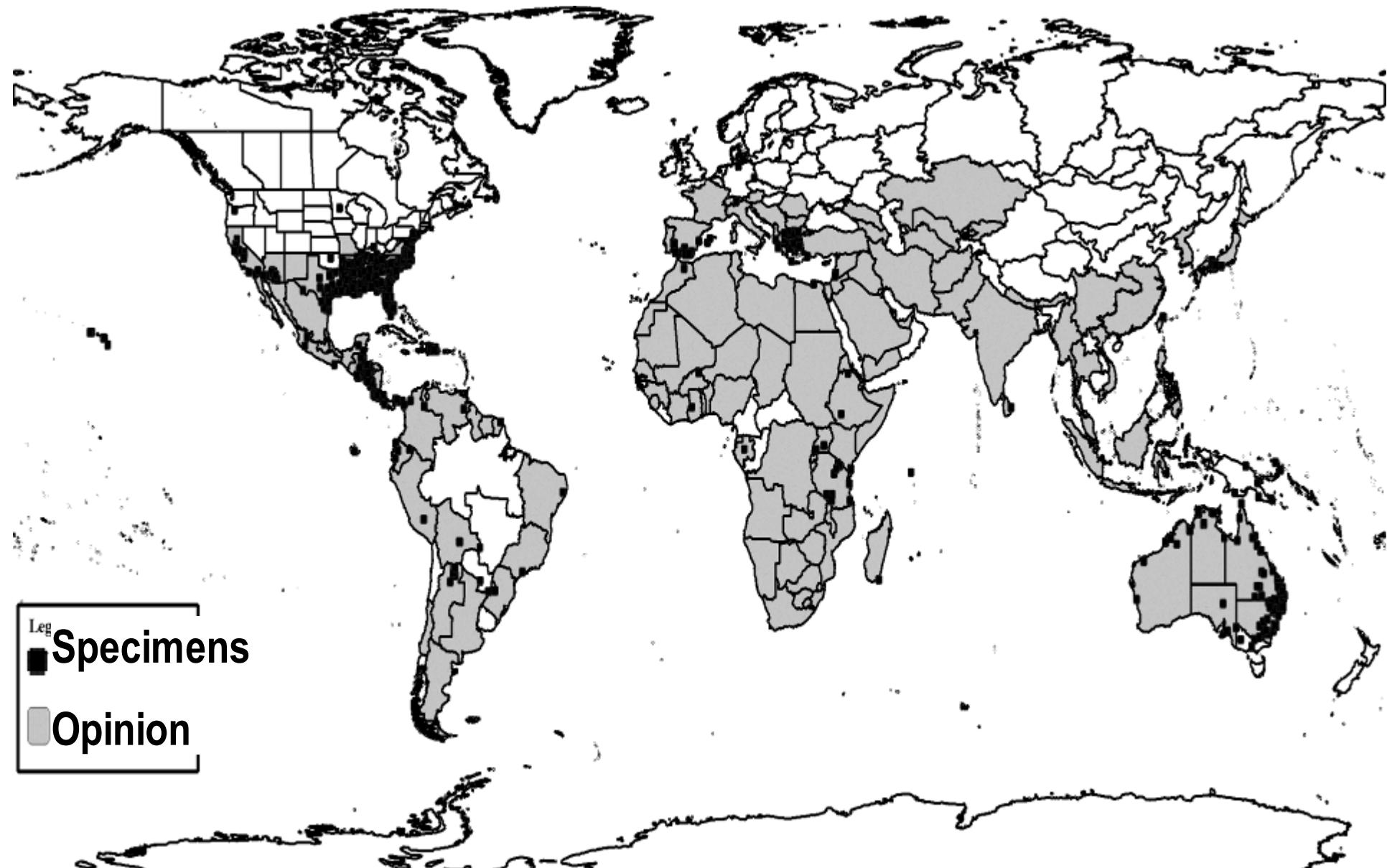


UGA1624072



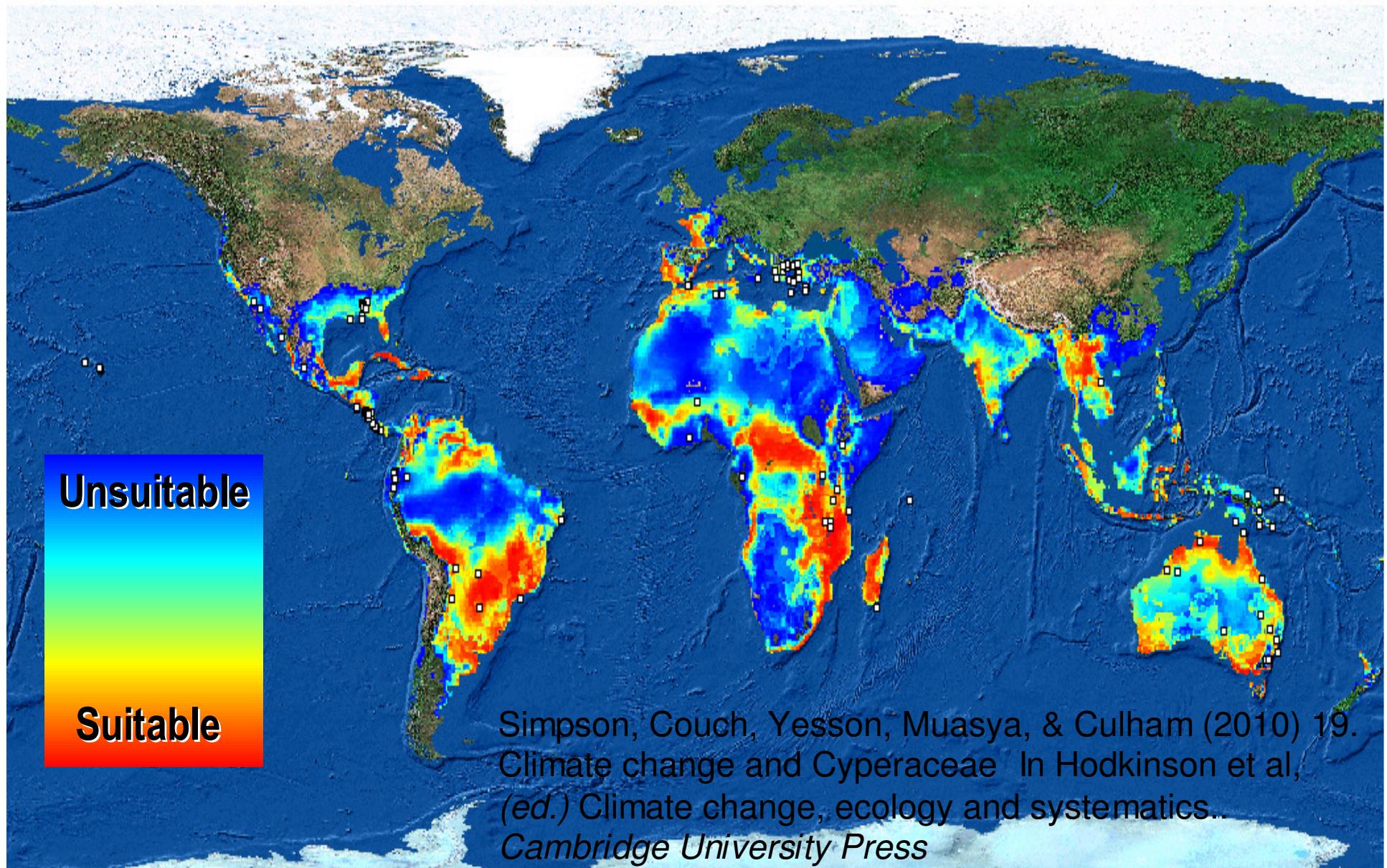
# Practical applications - weeds

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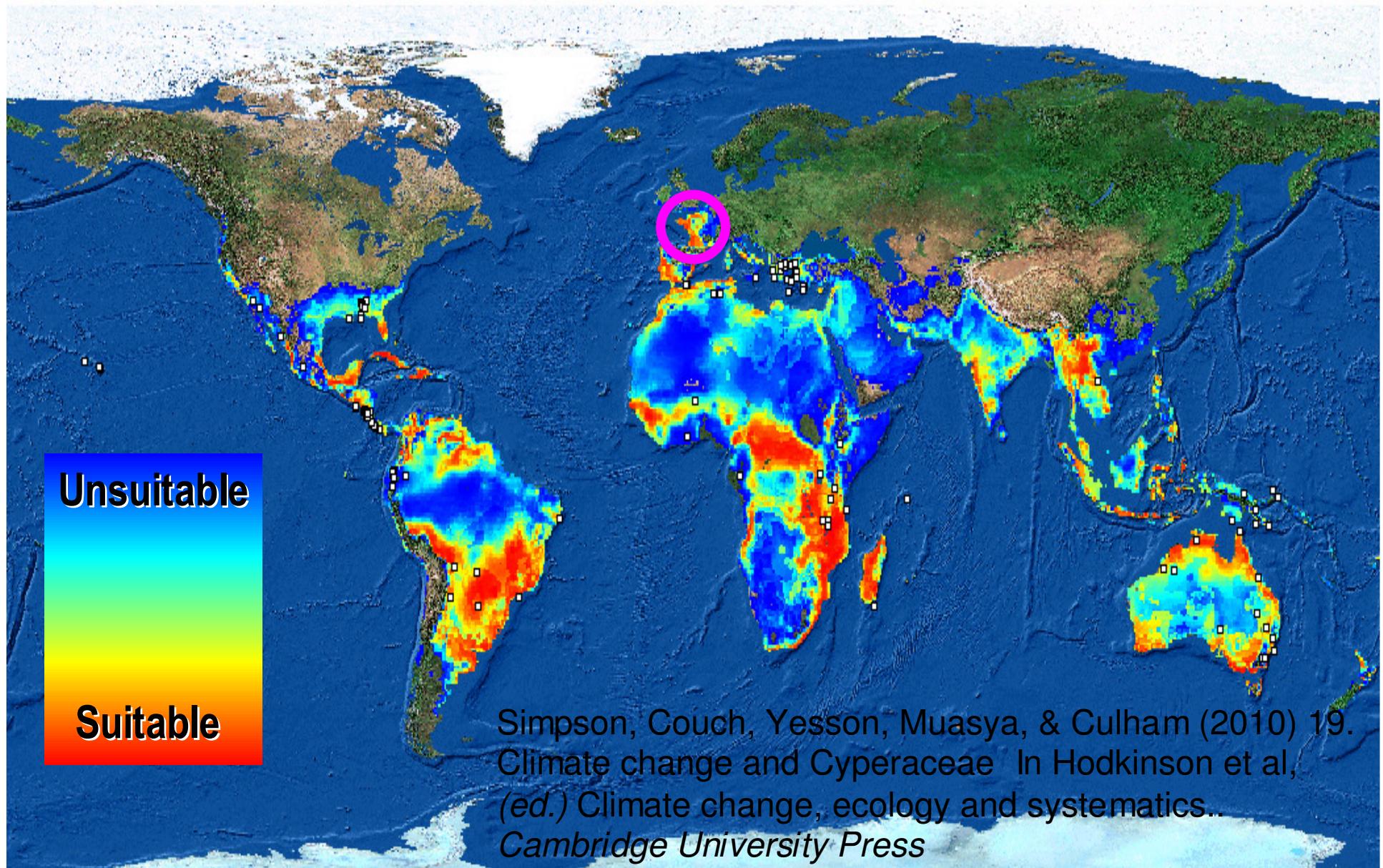
# Practical applications - weeds

ZSL



# Practical applications - weeds

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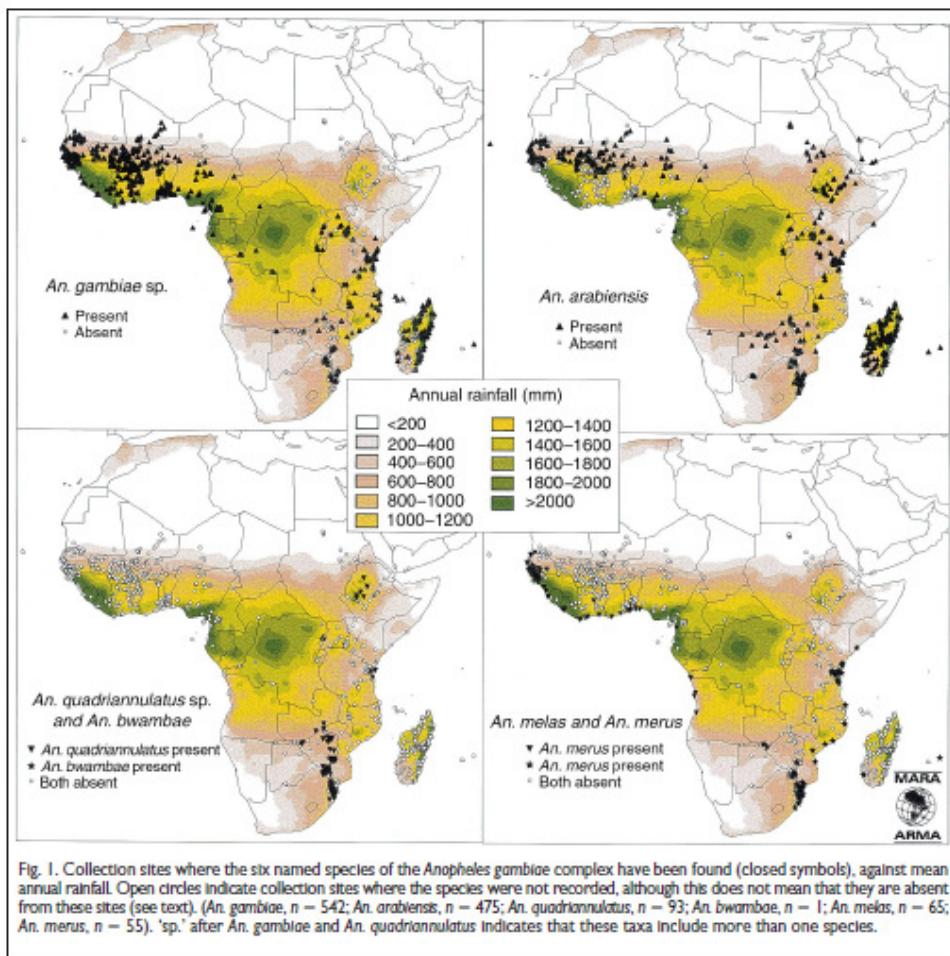


# Practical applications - disease



## Distribution of African Malaria Mosquitoes Belonging to the *Anopheles gambiae* Complex

M. Coetzee, M. Craig and D. le Sueur



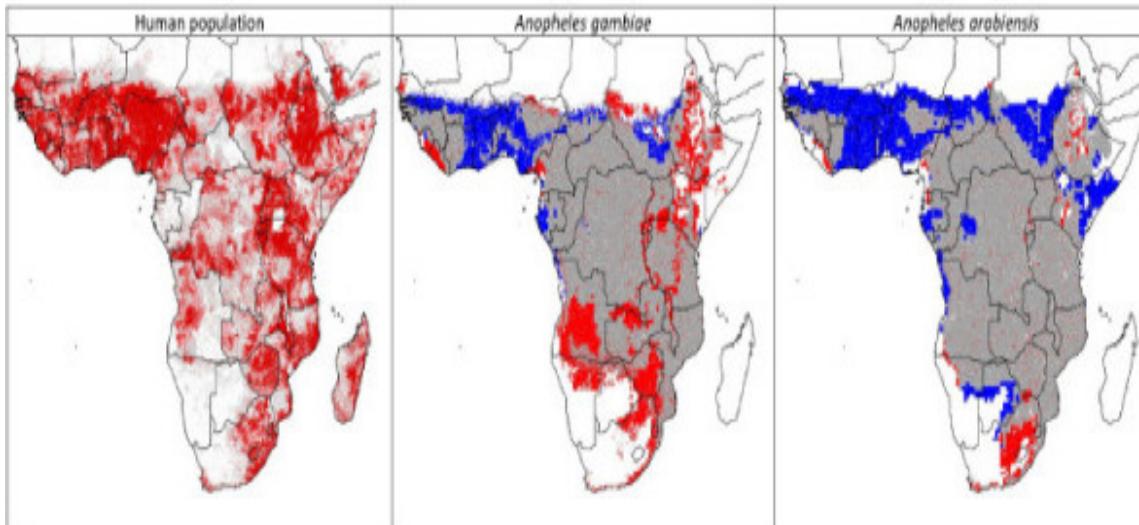
- Understanding habitat requirements helps estimate the distribution of disease vectors

Coetzee & le Sueur (2000)  
Parasitology Today, vol. 16,  
no. 2, 2000

# Practical applications - disease

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Human population    A. gambiae    A. arabiensis



**Figure 1 Human distributions and modeled present and future malaria vector distributions in Africa**

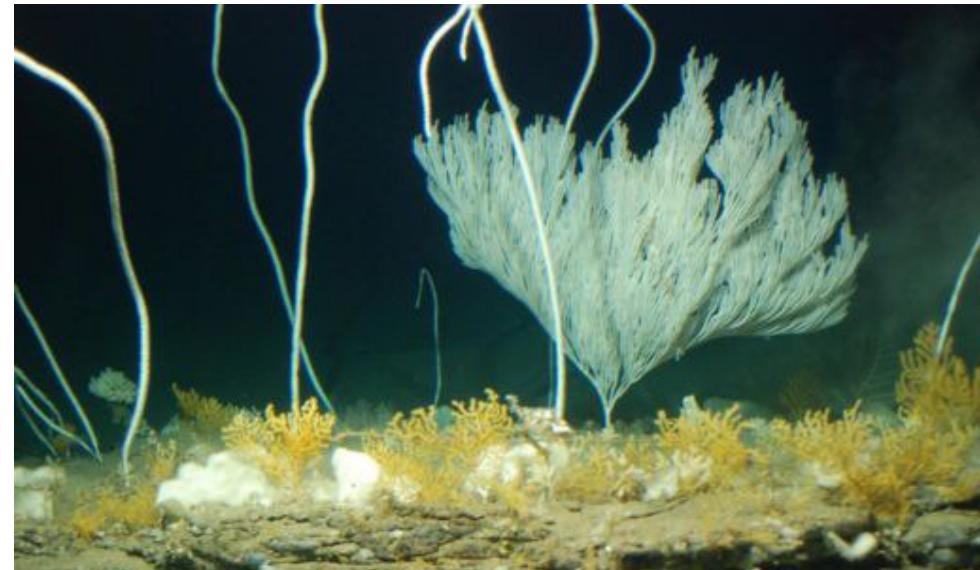
- Understanding disease vector habitat requirements and population dynamics helps understand the spread of disease

Peterson et al (2009) BMC  
Infectious Diseases 9:59

# Practical applications – deep sea

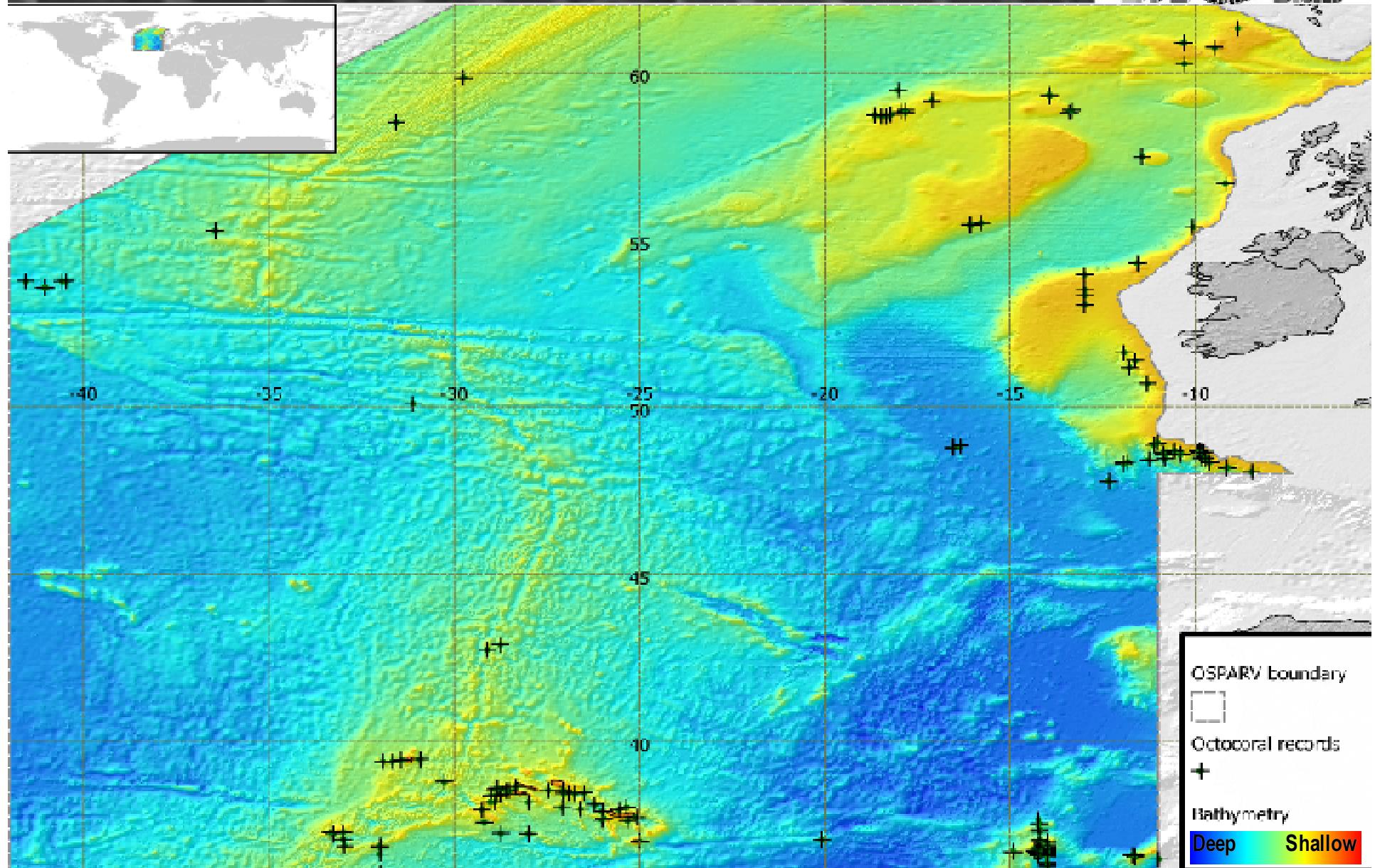
ZSL

- Where do corals live?
  - Some are thousands of metres below the surface
  - Very expensive and difficult to observe
- 
- Trawling can devastate coral gardens
  - Which areas should we protect from trawling?



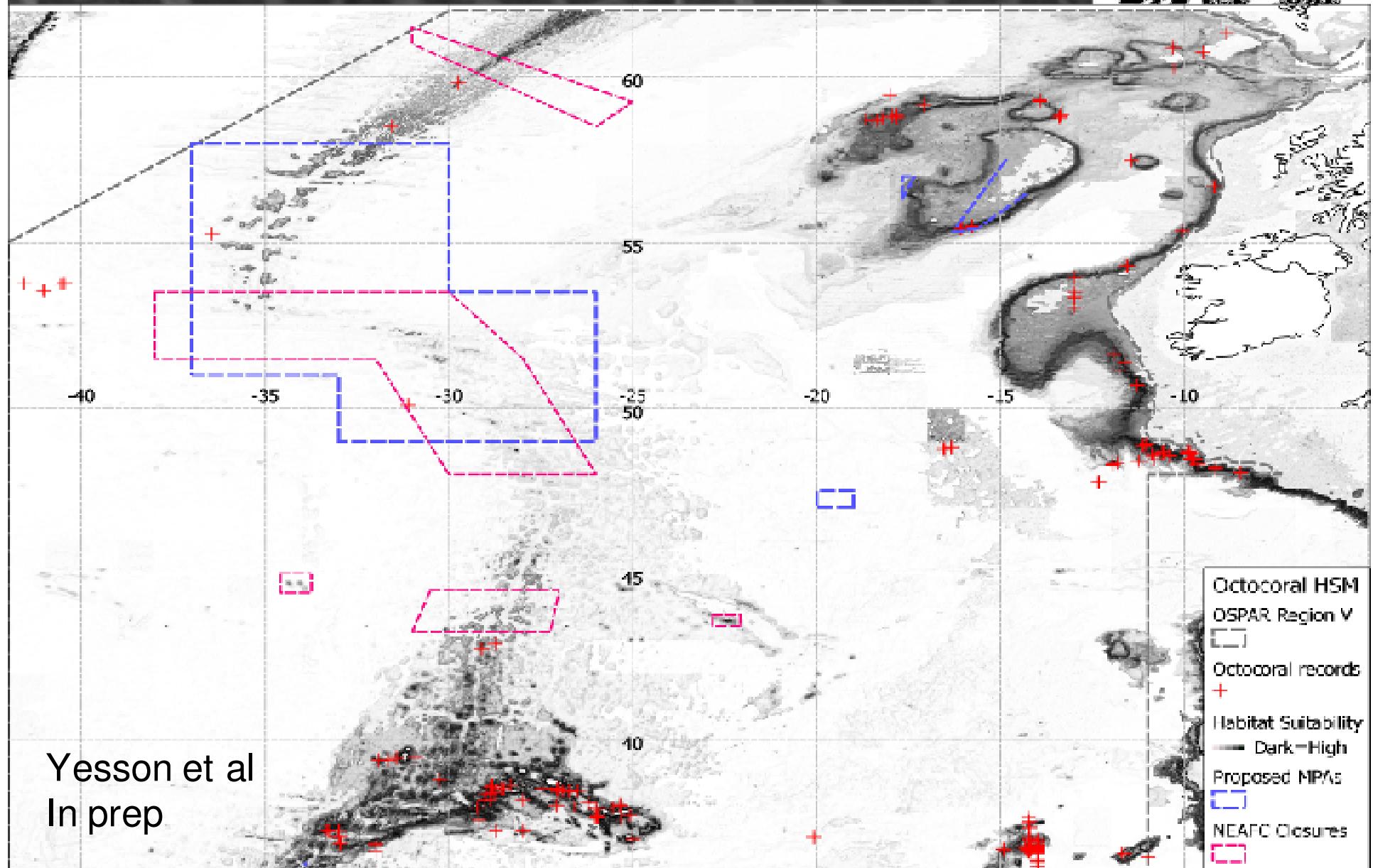
# Octocoral records from N. Atlantic

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# Estimated distribution based on niche

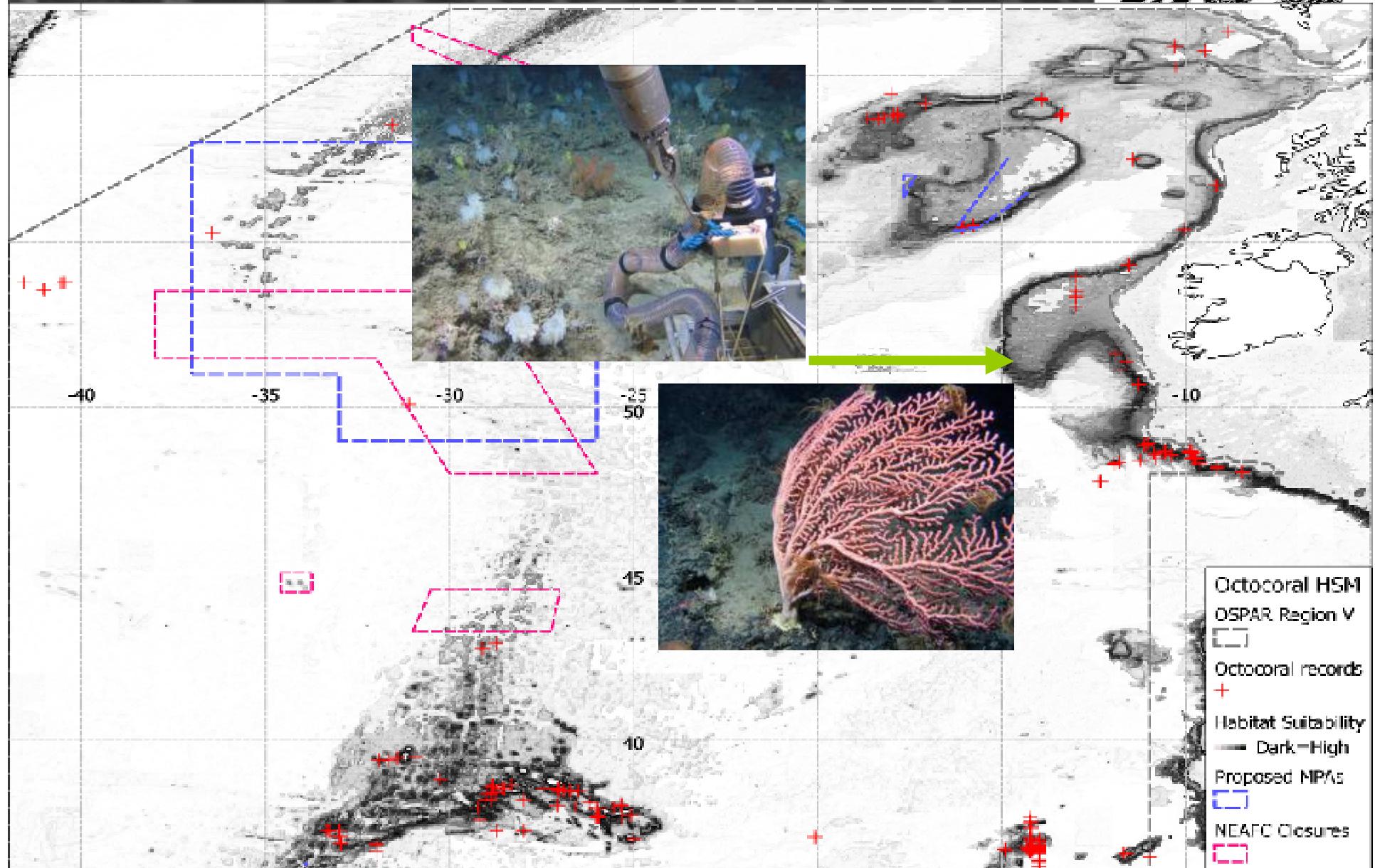
ZSL



Yesson et al  
In prep

# Estimated distribution based on niche

ZSL



# Conclusion

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- Habitat suitability modelling of corals is potentially useful due to a lack of distribution data
- Habitat suitability modelling of corals is difficult (see left)



© Ifremer 2003

# Niche model applications

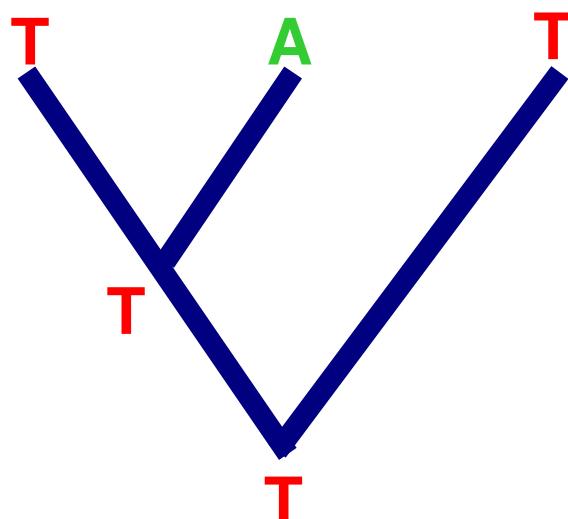


- Practical
  - Invasive species (weeds)
  - Disease
  - Data deficient species
  - Marine protected areas
- Theoretical
  - Diversity through time
  - Evolutionary patterns

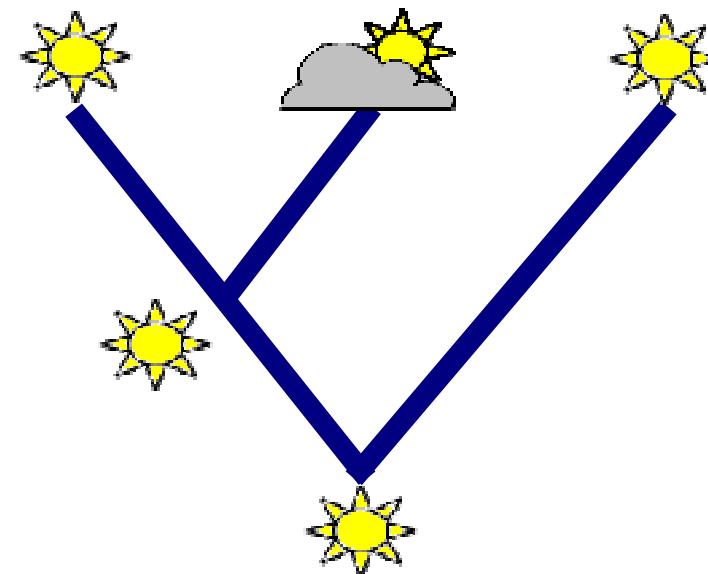
# Theoretical work

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- Using palaeoclimate reconstructions to estimate historical, palaeohistorical and ancestral niches



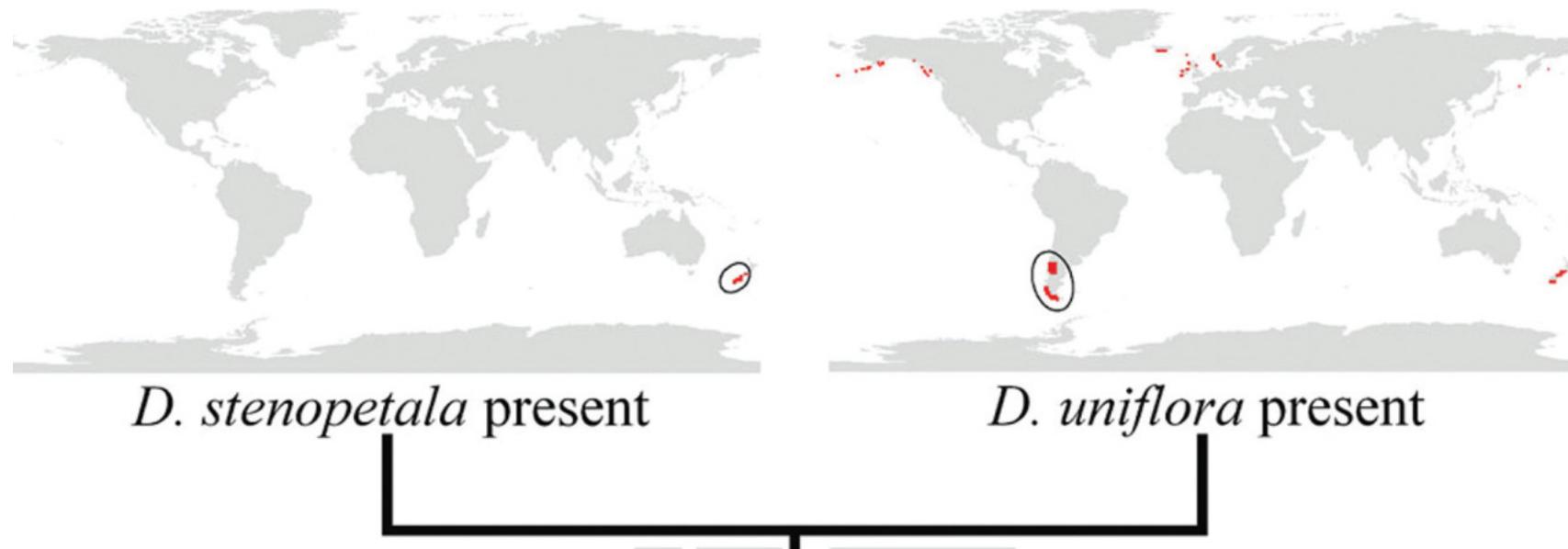
*Parsimony optimisation on  
a phylogenetic tree*



*Parsimony optimisation of  
environmental character*

# Phyloclimatic modelling – an example

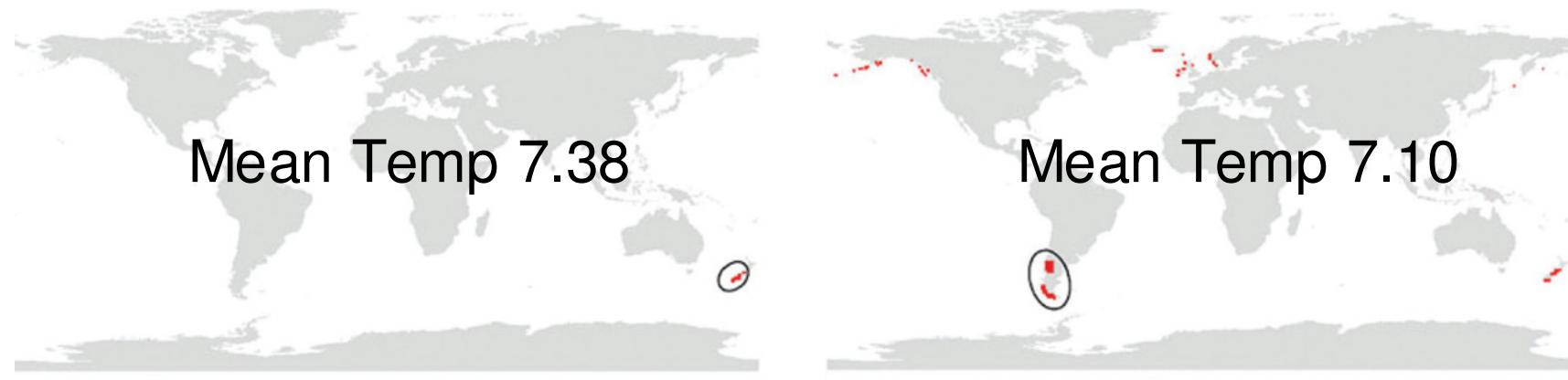
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- *D. stenopetala* and *D. uniflora* are sister taxa
- One is from New Zealand, the other is from South America
- We estimate an ancestral area?

# Phyloclimatic modelling – an example

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*D. stenopetala* present

*D. uniflora* present

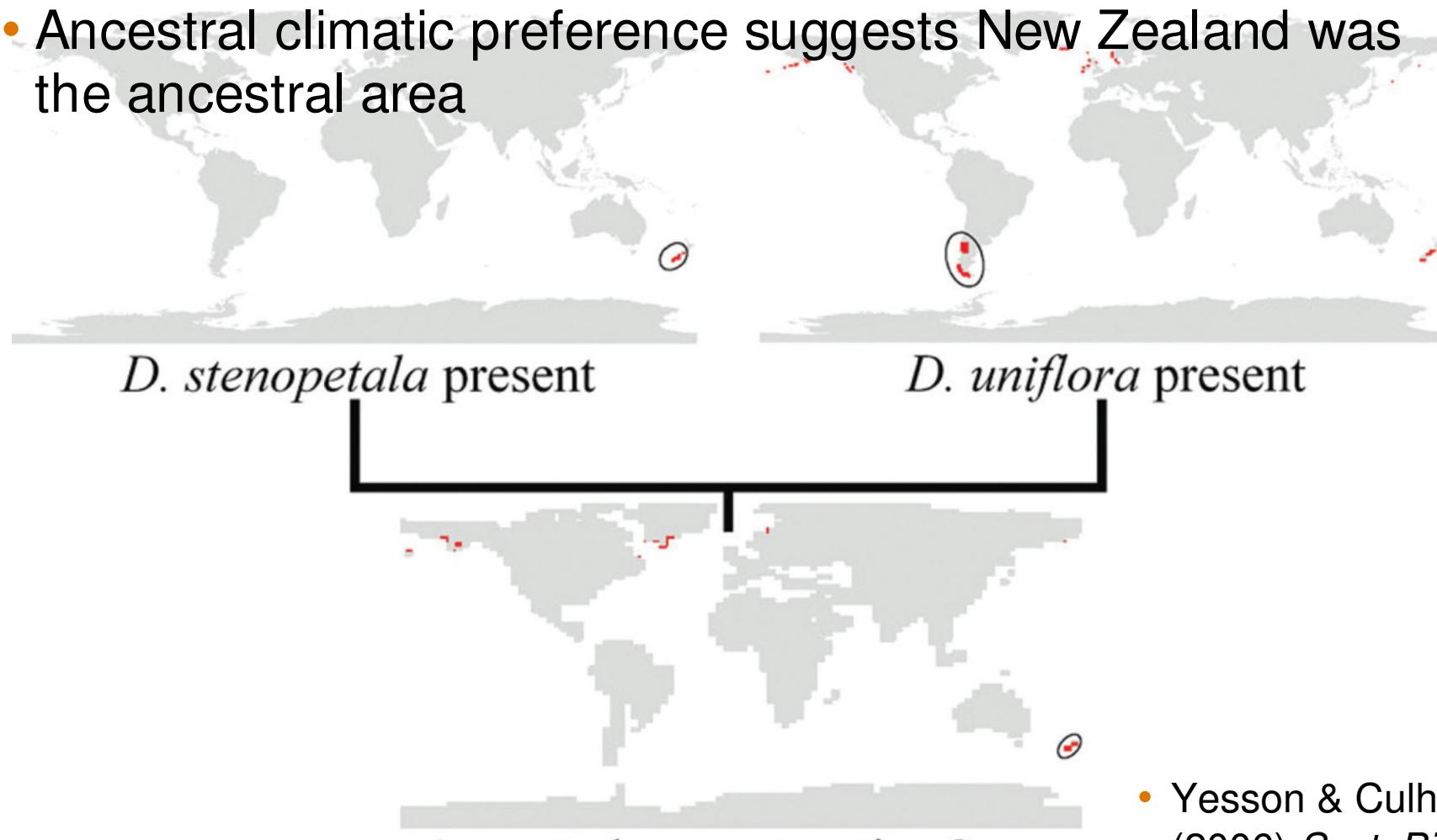
Ancestral node  
Mean Temp 9.87  
Age ~8.40-7.95 Ma

Project ancestral niche into palaeoclimate for ~8Ma

# Phyloclimatic modelling – an example

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- Ancestral climatic preference suggests New Zealand was the ancestral area

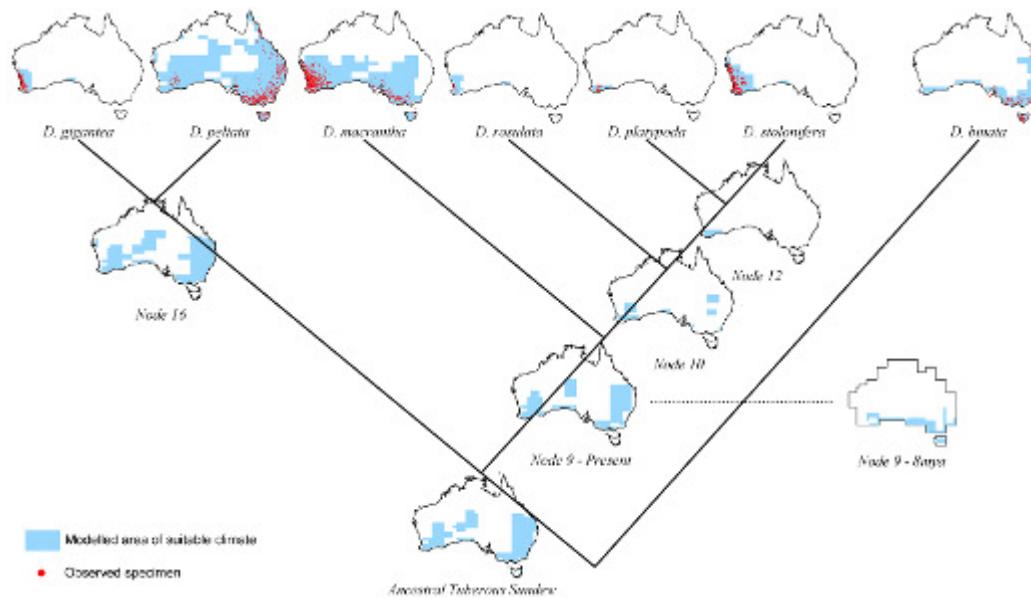


- Yesson & Culham (2006) *Syst. Biol.* 55(5): 785-802.

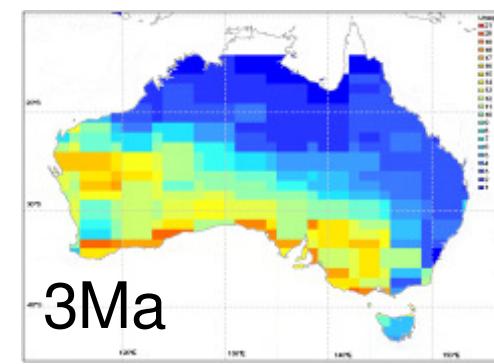
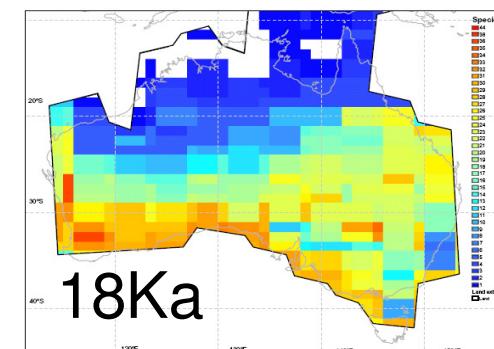
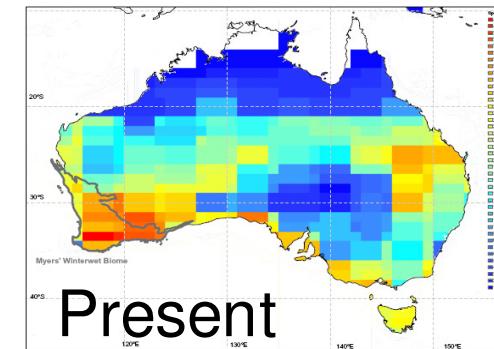
# Diversity through time

ZSL

## Evolutionary & ecological diversity of sundews



- Yesson & Culham  
(2006) *Syst. Biol.*  
55(5): 785-802.



- Yesson & Culham In prep

Gastrolobium diversity

# Practical – run your own niche model



## openModeller Practical



### Obtaining specimen data

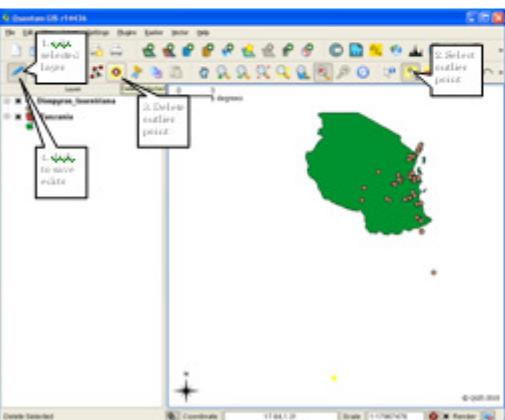
Before you can do any species distribution modelling you need some specimen data with associated point localities. For this practical we have prepared a sample file containing specimens from Tanzania.

Typically you would search for data by inspecting museum catalogues or searching through literature. The GBIF portal is a good place to start (<http://data.gbif.org>). openModeller has a search utility that allows you to search GBIF for data and feed the results directly into your model. Instructions for this are provided as an appendix.

I have prepared an example using the star apple *Diospyros kuteirana*, an East African endemic shrub, using data from GBIF (including Tanzanian). Data for this practical is in the Data directory in the handouts.

### Modelling setup

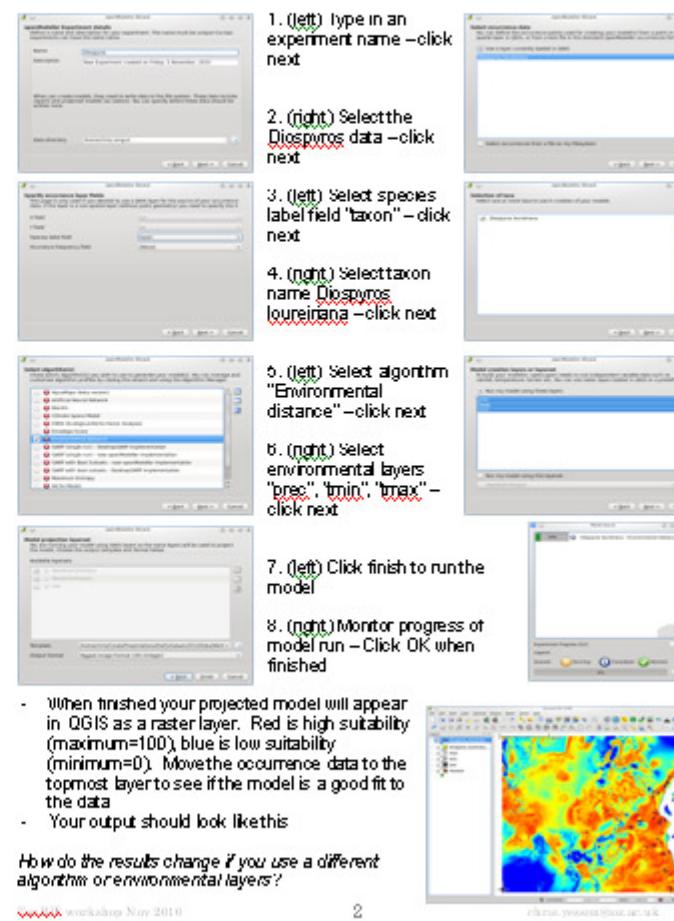
- Start by opening a reference map (.shp)
- open file Specimens\Diopsporus\_kuteirana.shp in qgis
- check the distribution – some records are outside the reference map
- use the qgis edit function to remove the outliers
- Note that removing outliers is not essential as these will be automatically excluded



1  
www.zsl.ac.uk/workshop-Nov-2010  
r.lucas.yousouf@zsl.ac.uk

### Modelling

- open raster environmental layers for modelling, use present day climate data minimum & maximum temperature and precipitation (tmin, tmax, precip in Worldclim/Present)
- select new experiment in the openModeller plugin menu
- follow the wizard instructions



1. (left) Type in an experiment name – click next  
2. (right) Select the *Diospyros* data – click next  
3. (left) Select species label field "taxon" – click next  
4. (right) Select taxon name *Diospyros kuteirana* – click next  
5. (left) Select algorithm "Environmental distance" – click next  
6. (right) Select environmental layers "prec", "tmin", "tmax" – click next  
7. (left) Click finish to run the model  
8. (right) Monitor progress of model run – Click OK when finished  
When finished your projected model will appear in QGIS as a raster layer. Red is high suitability (maximum=100), blue is low suitability (minimum=0). Move the occurrence data to the topmost layer to see if the model is a good fit to the data  
Your output should look like this  
How do the results change if you use a different algorithm or environmental layers?  
www.zsl.ac.uk/workshop-Nov-2010  
r.lucas.yousouf@zsl.ac.uk