

# A multidisciplinary framework for biodiversity prediction in the Atlantic Rainforest hotspot

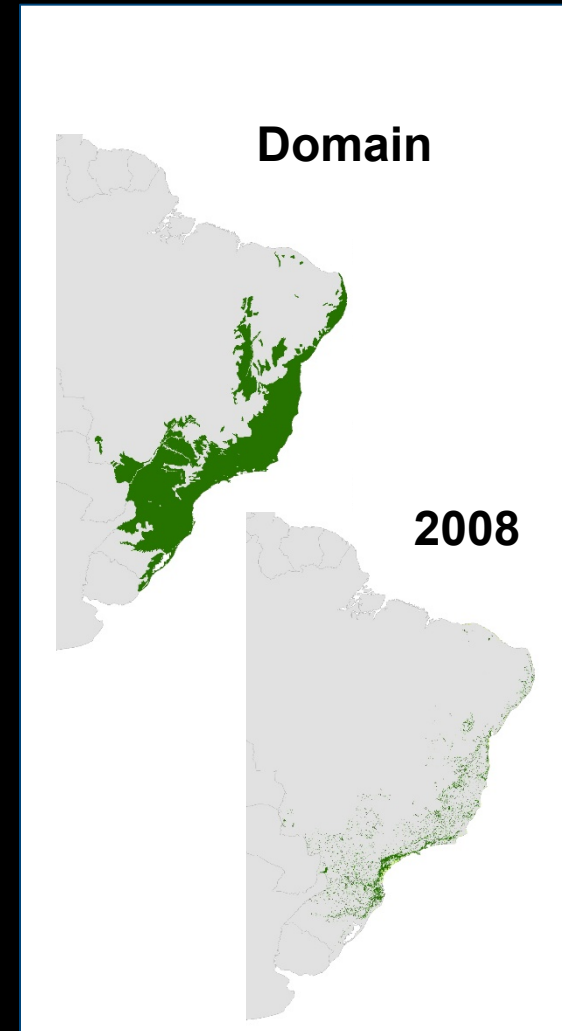


[www.afbiota.org](http://www.afbiota.org)

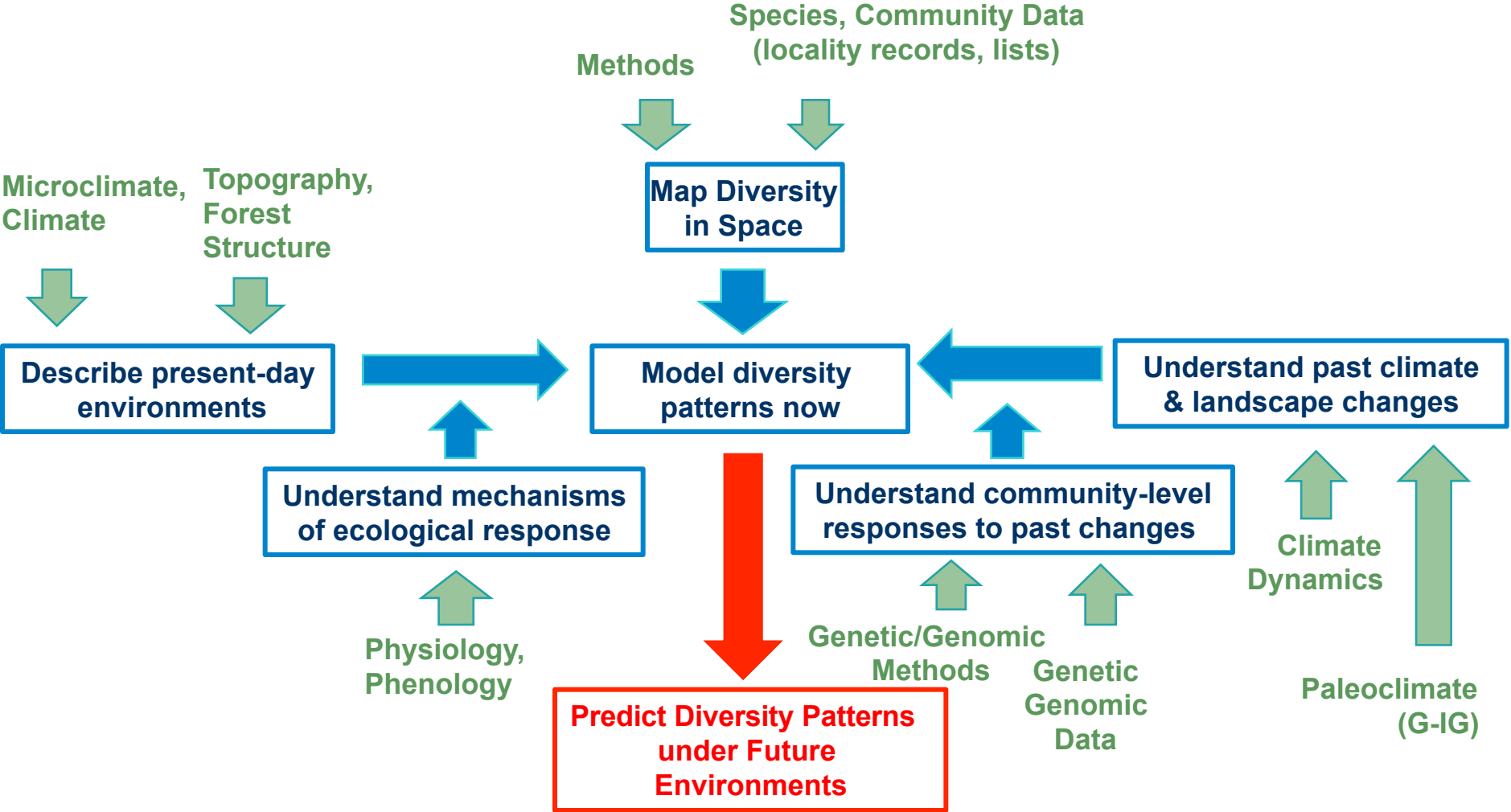




Our DoB project implements an integrative framework to improve biodiversity prediction in the Atlantic Forest hotspot

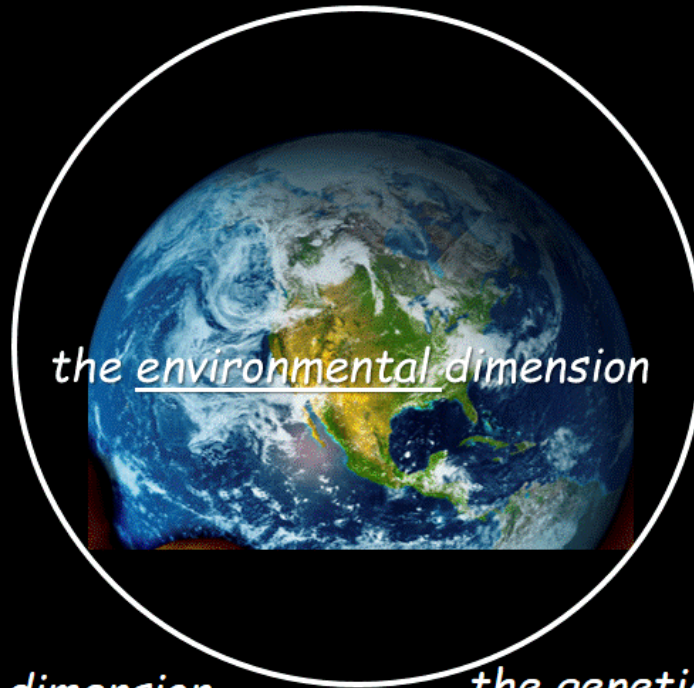


# Our DoB project implements an integrative framework to improve biodiversity prediction in the Atlantic Forest hotspot





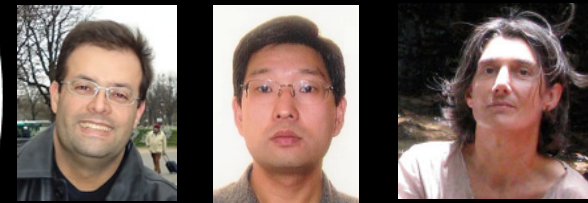
*the taxonomic dimension of biodiversity*



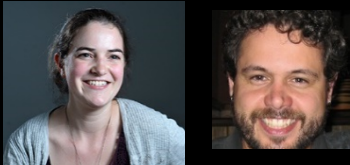
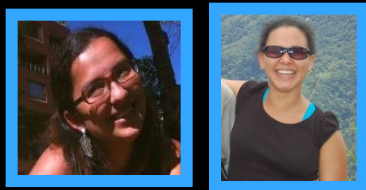
*the environmental dimension*



*the functional dimension*



*the genetic dimension*





Species, Community Data  
(locality records, lists)

Methods



Map Diversity  
in Space

Microclimate, Climate  
Topography, Forest  
Structure



Describe present-day  
environments

Model diversity  
patterns now

Understand past climate  
& landscape changes



Understand mechanisms  
of ecological response

Understand community-level  
responses to past changes

Physiology,  
Phenology

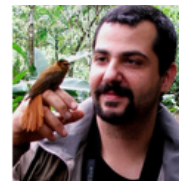
Genetic/Genomic  
Methods

Genetic  
Genomic  
Data

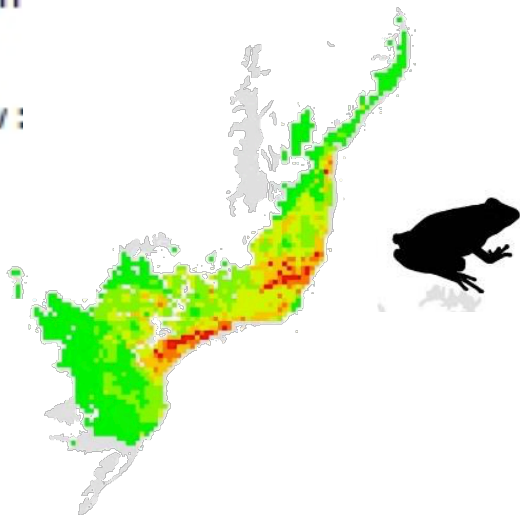
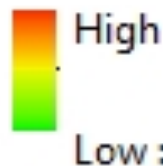
Climate  
Dynamics

Paleoclimate  
(G-IG)

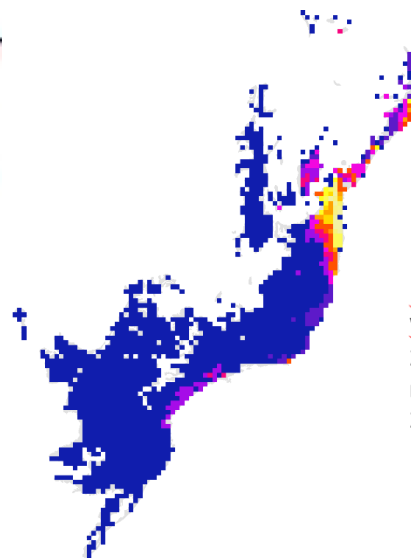
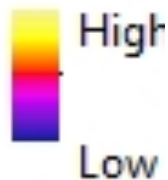
Predict Diversity Patterns  
under Future  
Environments



# Species Richness



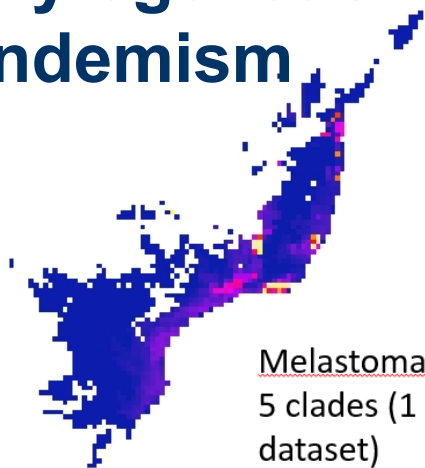
# Phylogenetic Diversity



Bromeliaceae  
Wittmackia clade  
24 spp. (ETS, g3pdh,  
rpb2, matk, rps, trn)  
228 points



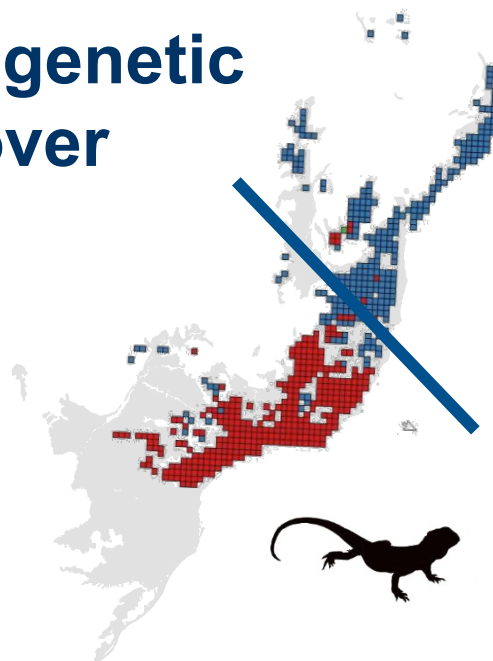
# Phylogenetic Endemism



Melastomataceae  
5 clades (1  
dataset)  
184 spp.  
25737 points



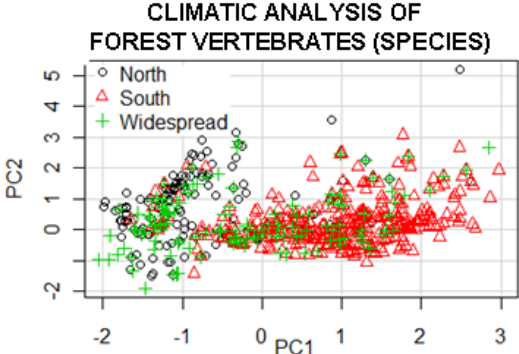
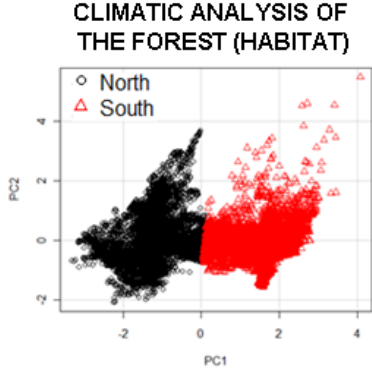
# Phylogenetic Turnover





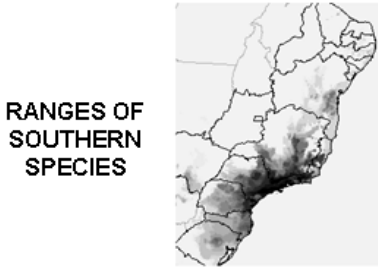
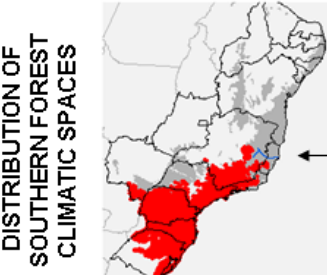
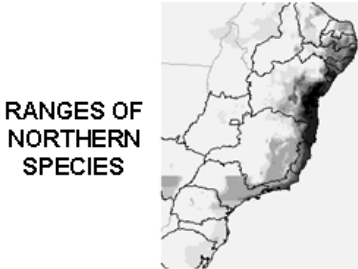
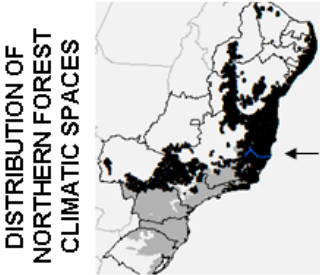
# Improving characterization of present-day environments for biodiversity modeling

## Many forests and forest communities in the AF

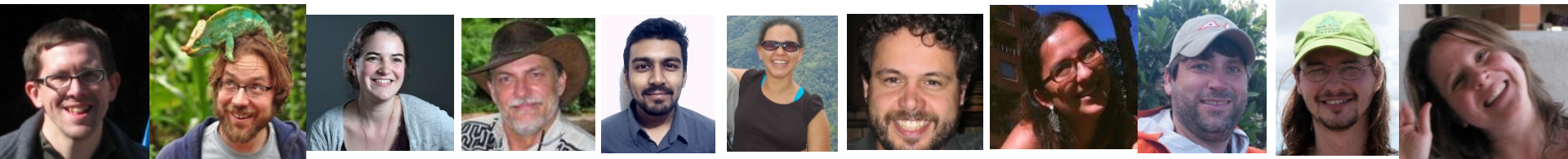
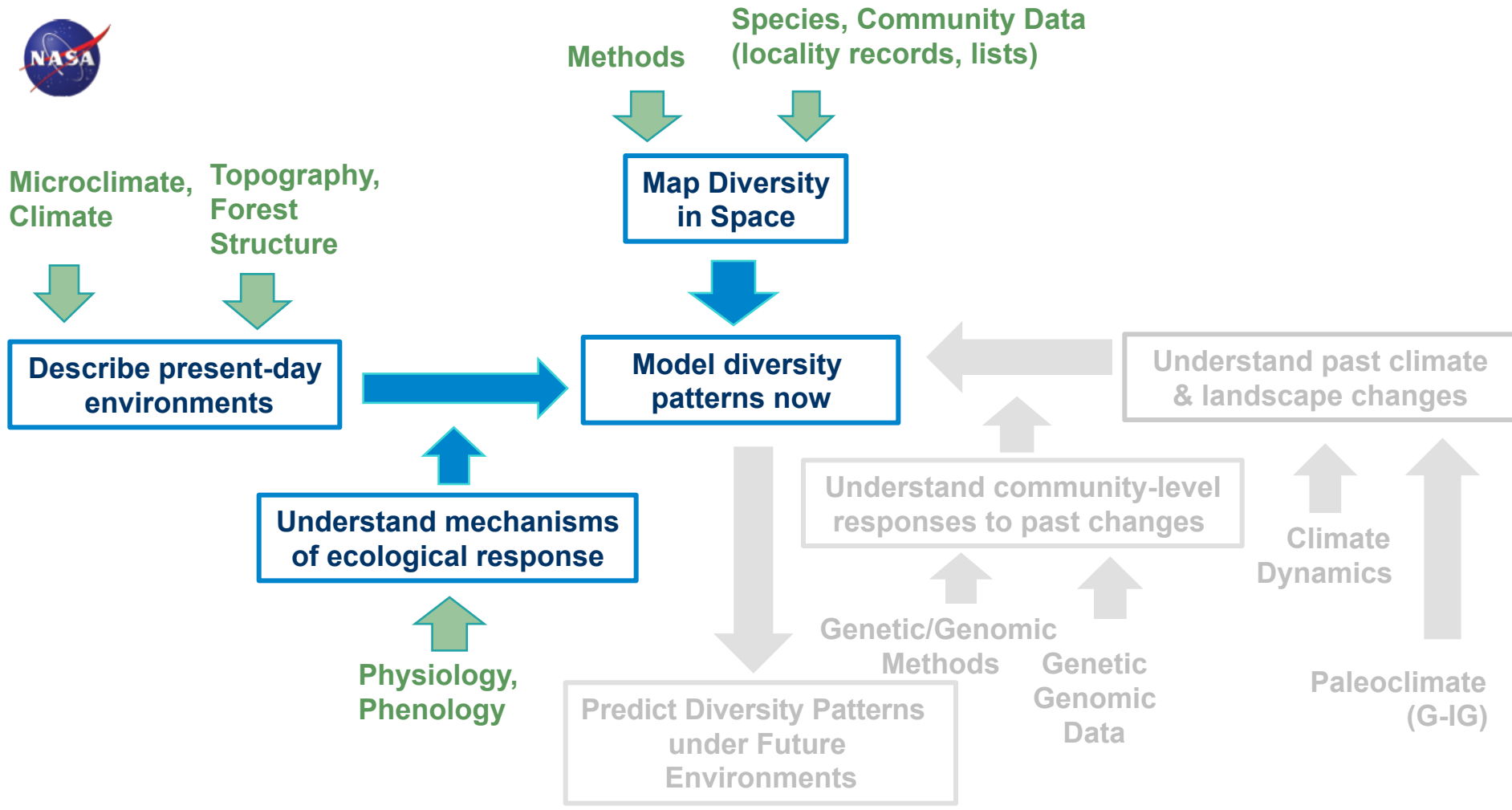


**Left:** PCA-based identification of two climatically distinct spaces within the Atlantic rainforest (top), and their respective northern (black) and southern (red) geographical ranges (bottom).

**Right:** PCA-based identification of species assemblages along the climatic axes of the Atlantic forest (top) depicting their northern (black), southern (red), or widespread (green) distributions.



Carnaval et al. 2014.  
Proc Roy Soc.





# Improving characterization of present-day environments for biodiversity modeling



## Methods in Ecology and Evolution



*Methods in Ecology and Evolution* 2014

doi: 10.1111/2041-210X.12264

### **Bioclimatic variables derived from remote sensing: assessment and application for species distribution modelling**

**Eric Waltari<sup>1\*</sup>, Ronny Schroeder<sup>1,2</sup>, Kyle McDonald<sup>1,3</sup>, Robert P. Anderson<sup>1,3,4,5</sup> and Ana Carnaval<sup>1,3,4</sup>**

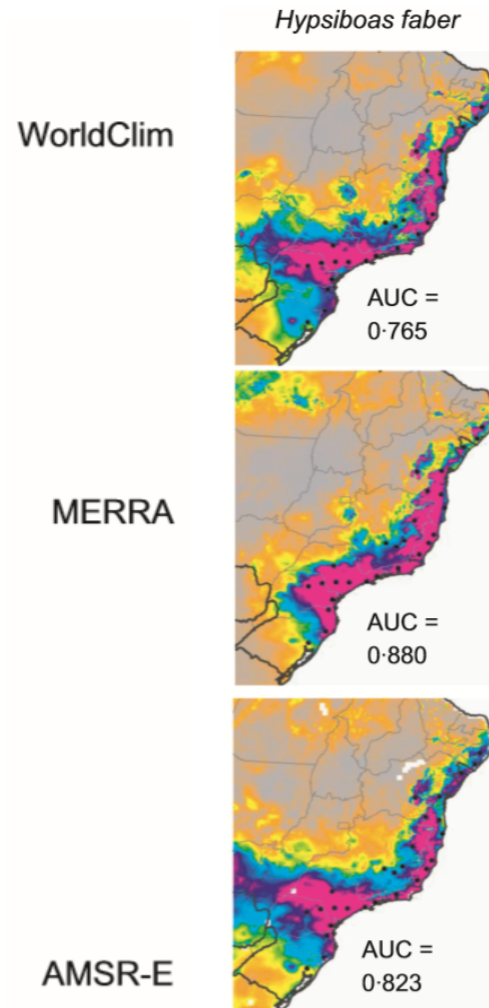
#### NASA products

- the Modern-Era Retrospective Analysis (MERRA), which incorporates remote sensing information and a subset of the station-based data
- pure remote sensing information (Advanced Microwave Scanning Radiometer-Earth Observations, AMSR-E).

Coarse spatial resolution (MERRA 55 x 75 km. AMSR-E 25 km), high temporal frequency of data collection (near-daily at low latitudes).

# Improving characterization of present-day environments for use in correlative modeling

- Species Distribution Models derived from MERRA-derived layers performed better than models built with WorldClim data.
- Models constructed with AMSR-E-based layers had similar performance to models built with WorldClim.





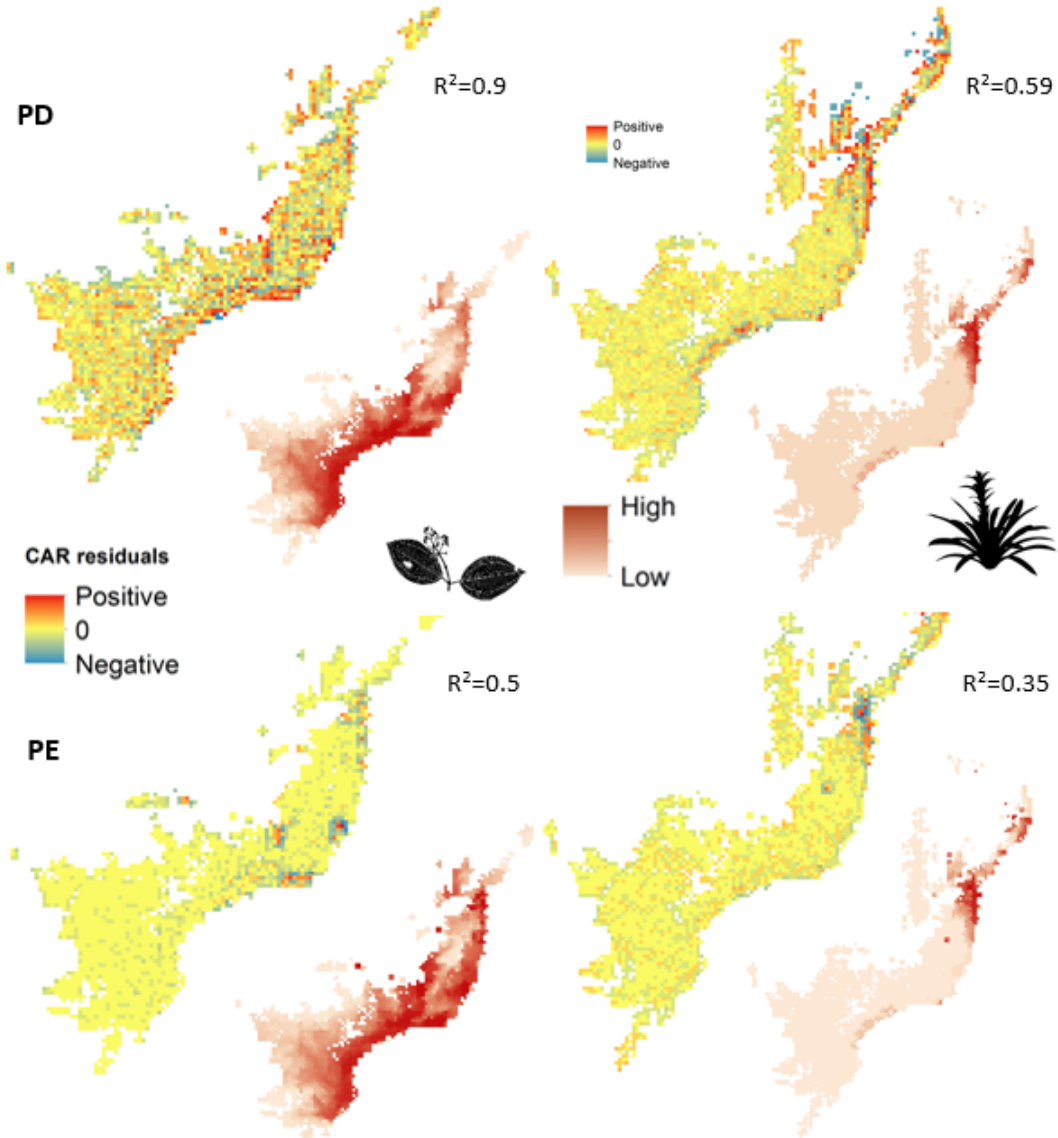
# Improving characterization of present-day environments for biodiversity modeling



**Conditional autoregressive (CAR) model analysis:** using bioclimatic variables (from WorldClim vs. MODIS/CHIRPS\*) as predictors of PD and PE.

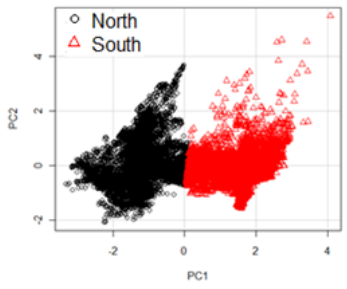
Response variable	AIC (CAR with MODIS/CHIRPS vs. CAR with Worldclim)
Bromeliads PD	61.79
Bromeliads PE	<b>-42.52</b>
<u>Melastomataceae</u> PD	<b>-103.75</b>
<u>Melastomataceae</u> PE	<b>-39.88</b>

\*Deblawe et al. 2016. Glob Change Biol

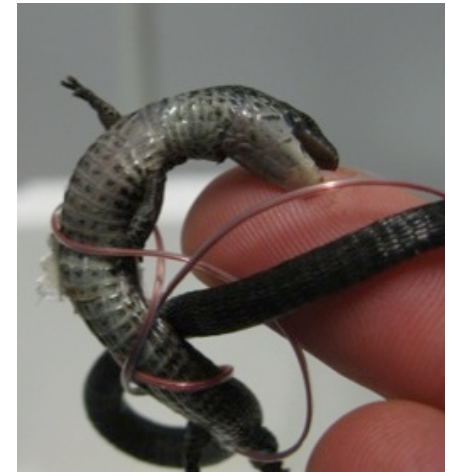
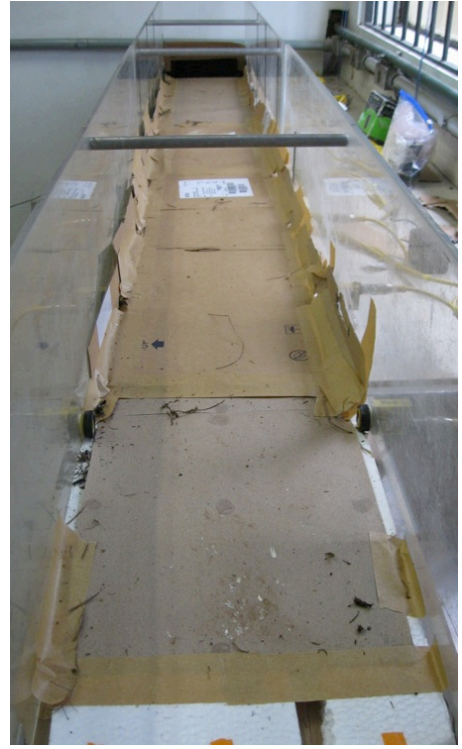
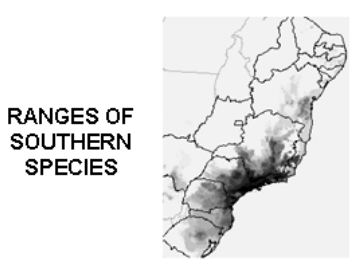
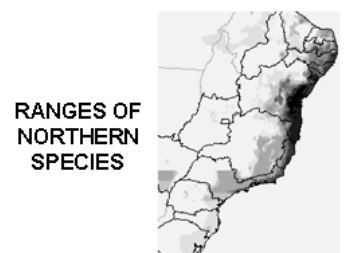
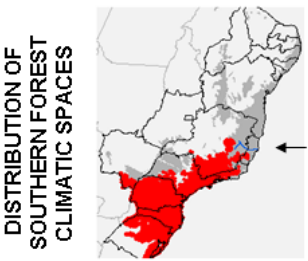
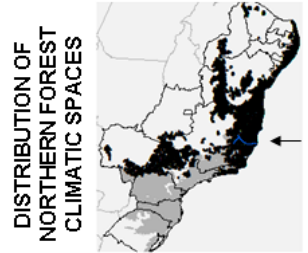
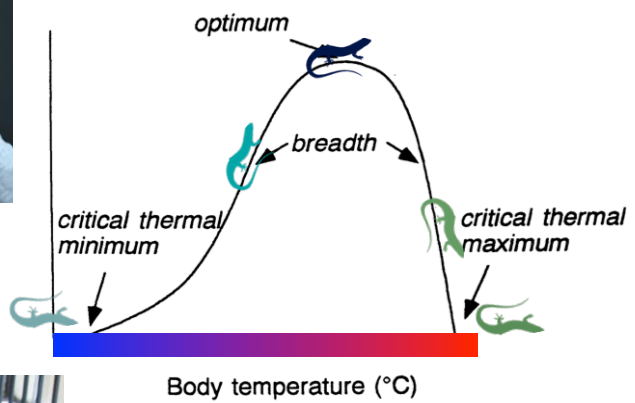
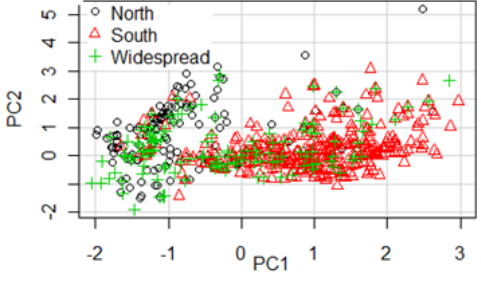


# We are studying the physiological limits of AF species to understand the MECHANISMS that underscore the distribution of diversity

CLIMATIC ANALYSIS OF THE FOREST (HABITAT)

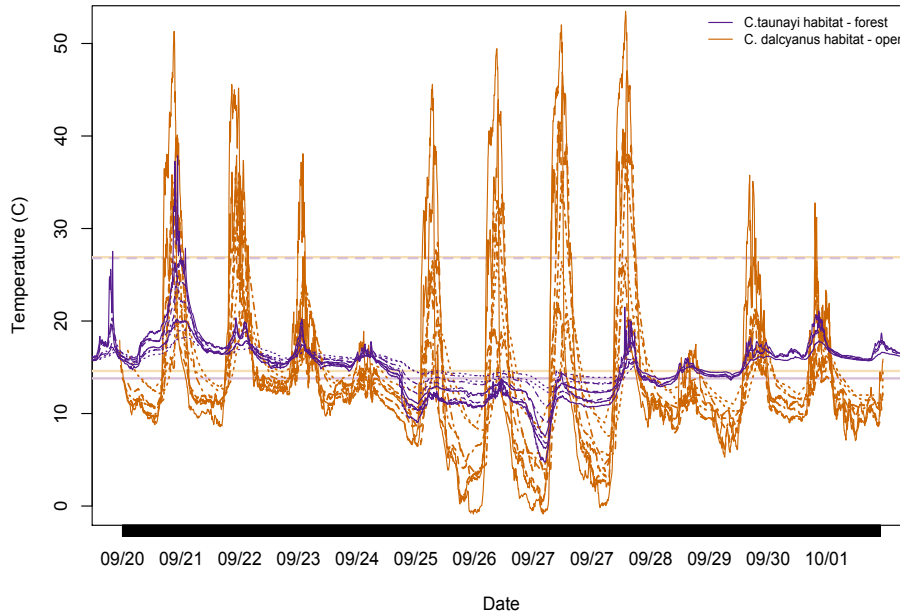
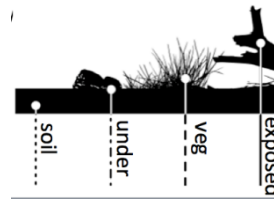
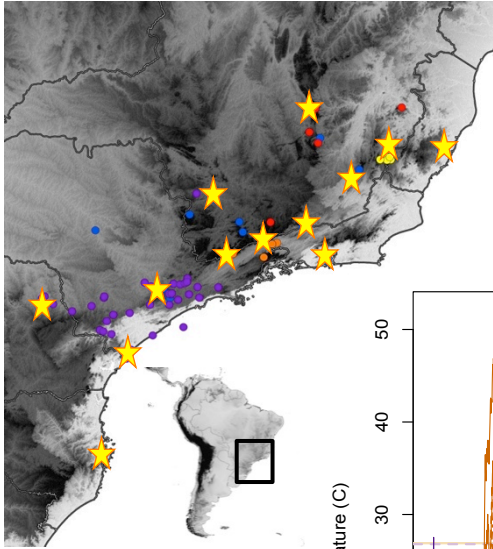


CLIMATIC ANALYSIS OF FOREST VERTEBRATES (SPECIES)



# Improving characterization of present-day environments for biodiversity modeling

## Characterizing microclimates with hygrobuttons



Precise measurements, limited coverage



# Improving characterization of present-day environments for biodiversity modeling

▼  
**Ibutton day  
& Ibutton night  
(8-day and monthly)**

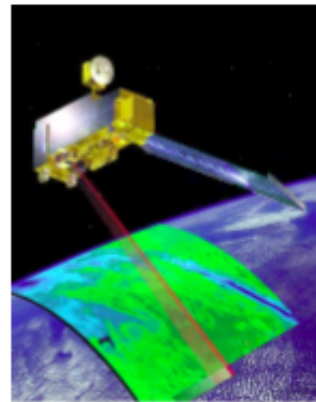
**Why Ibutton?**

Represents environment at  
the scale of the organisms



**MODIS day  
& MODIS night  
(8-day and monthly)**

**Why remote sensing  
and MODIS?**



Broad and even  
coverage with  
high spatial  
resolution

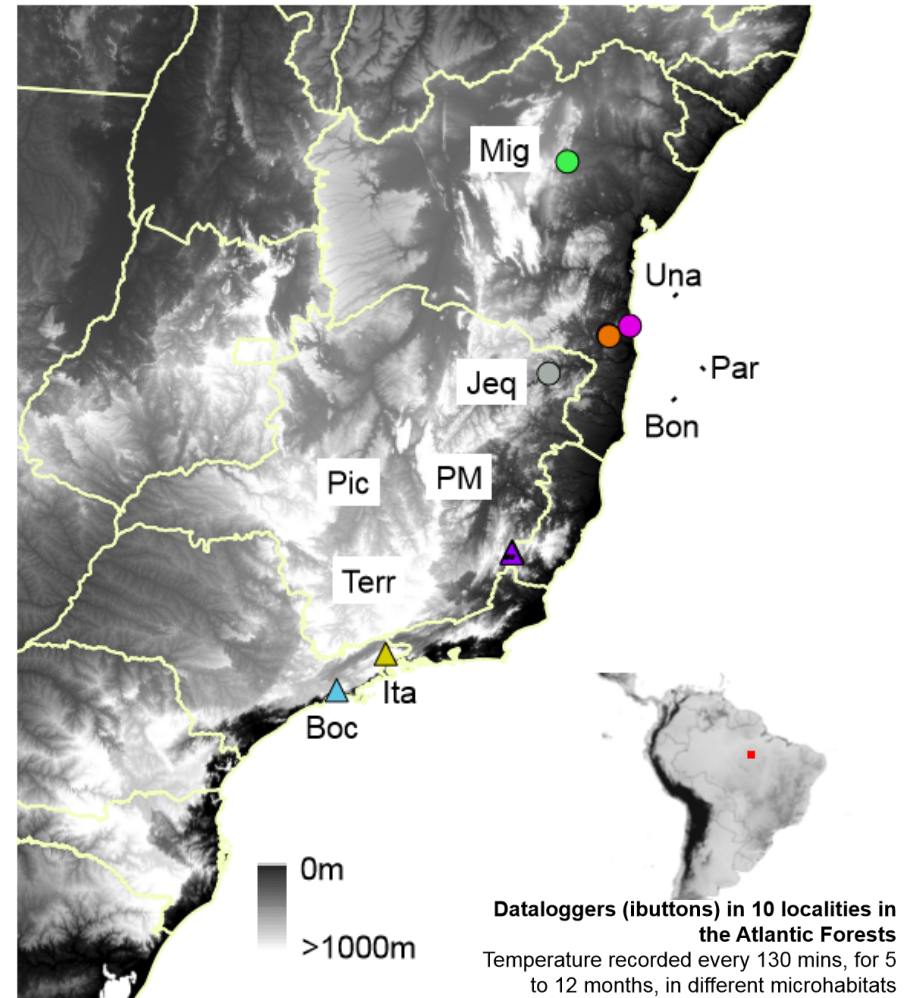
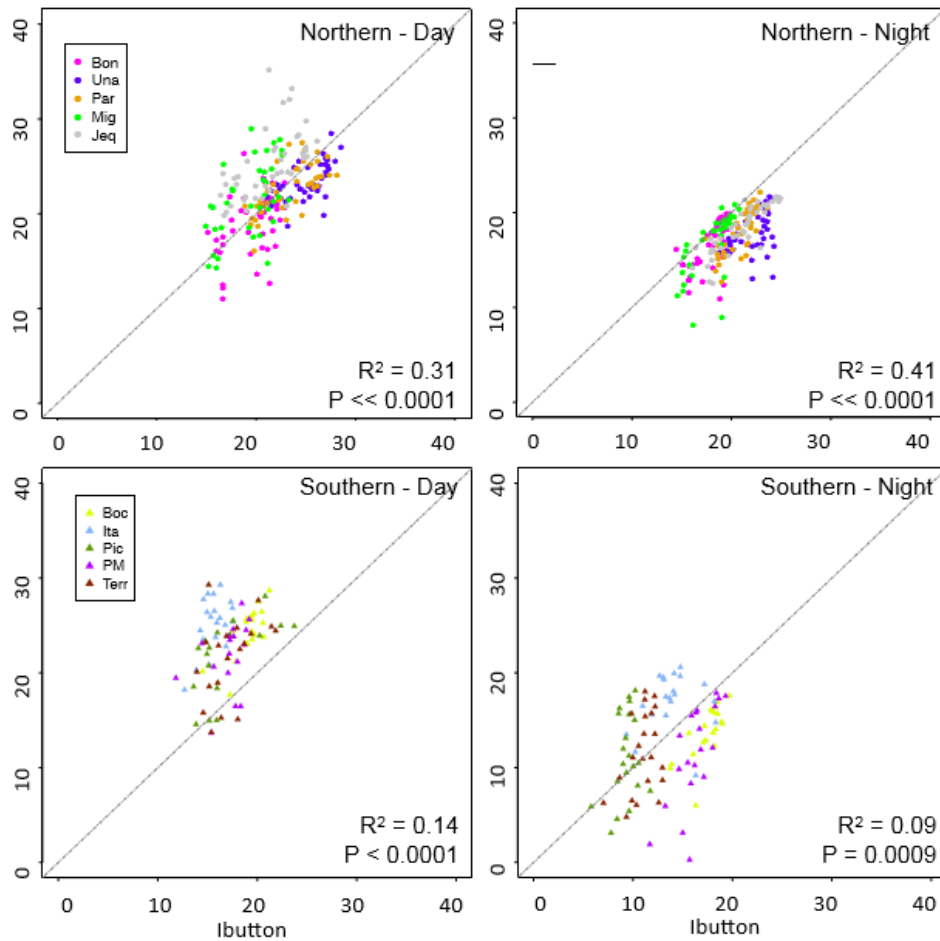
# Microclimate (ibutton) vs. MODIS (8-day comparison)



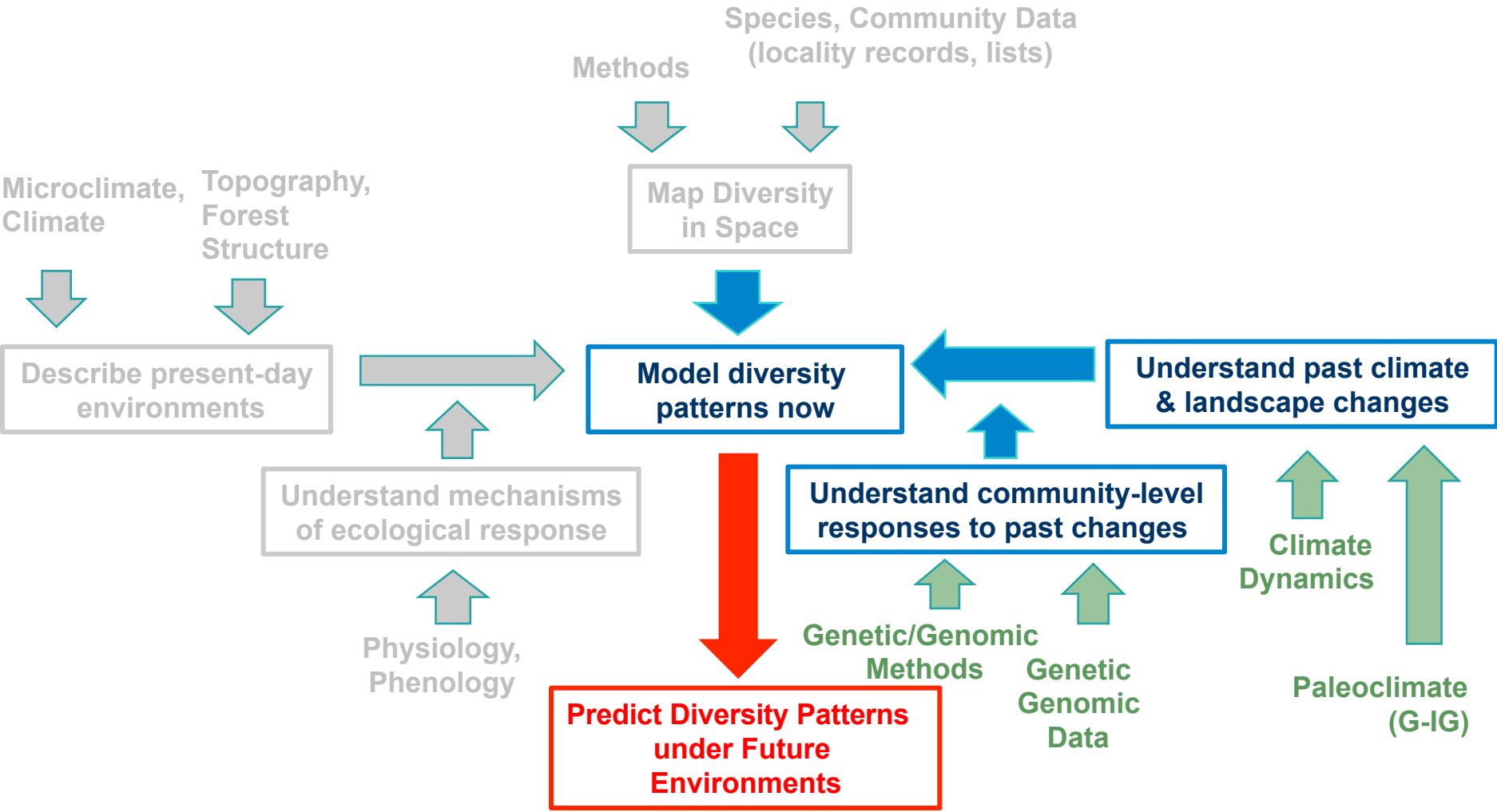
Stronger correlations in North (effect of S topography)

MODIS: higher diurnal temperatures, lower nocturnal temperatures (buffering effect of vegetation)

MODIS 8-day composite seem to perform reasonably well in cloud-covered areas



# Our DoB project implements an integrative framework to improve biodiversity prediction in the Atlantic Forest hotspot



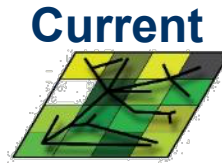
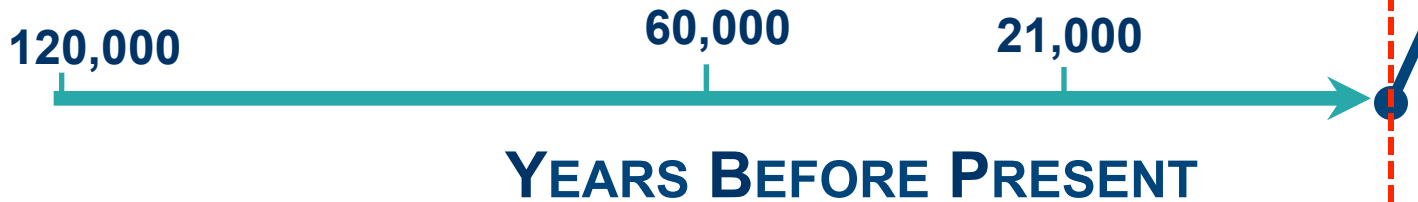




# A framework with two phases

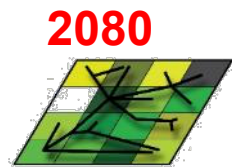
- Phase 1: Learning from the past

Use empirical genetic data and SDMs under paleoclimate to estimate demographic parameters



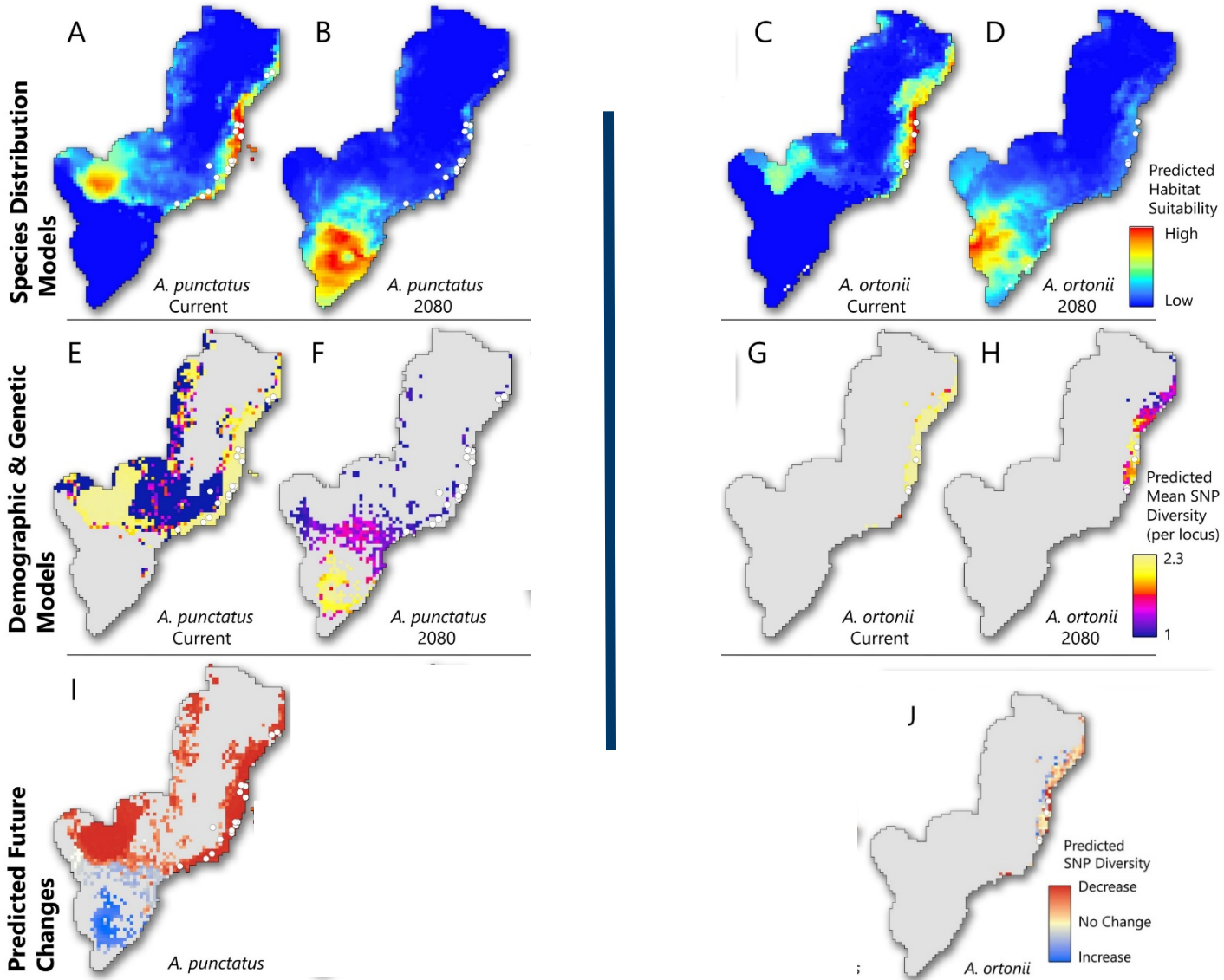
- Phase 2: Using inferred parameters to infer plausible distribution of diversity in the future

Estimate genetic composition of landscape: Past to Present and Future




Spatially explicit genetic effects of climate change

# Plausible scenarios of the spatial distribution of genetic diversity of two anole species differ extensively



## Aims for years 3+

- Continue to explore MODIS, AMSR-E, improve environmental characterization 
- Develop models to describe microclimatic conditions, link with physiological data
- Link predictions of changes in diversity with near-real time monitoring of the forest





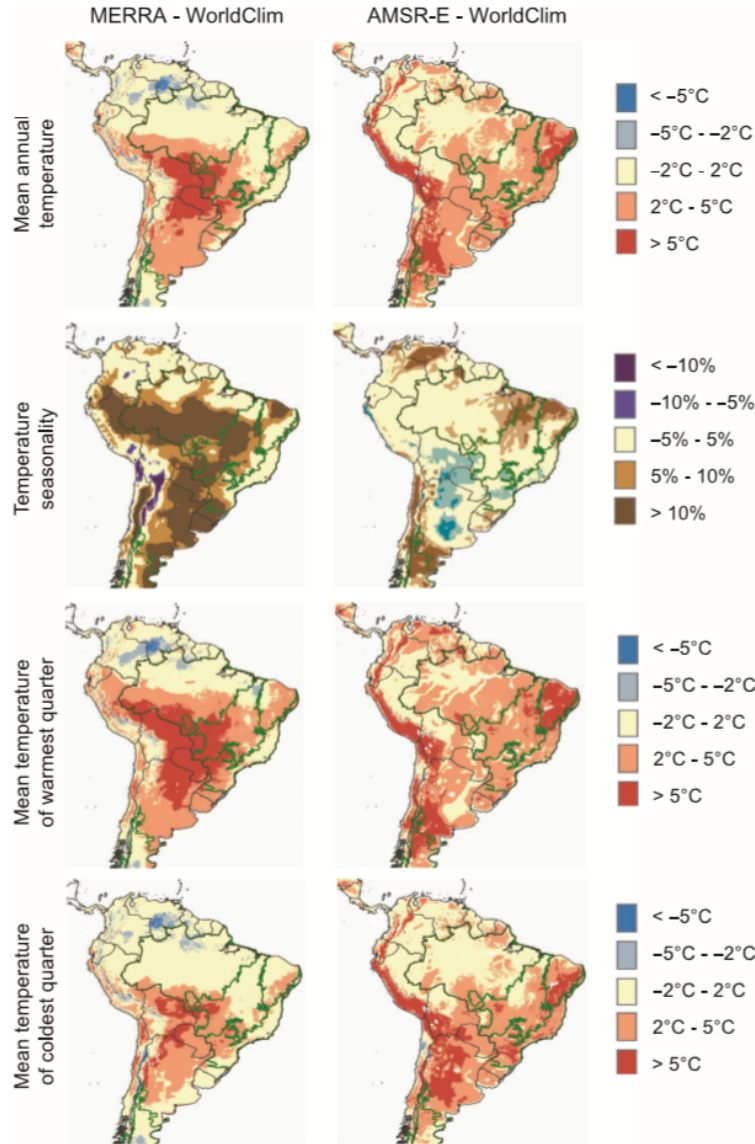


# New Bioclims - MERRA

- Used time-averaged hourly MERRA single-level temperature at 2 m above the displacement height.
- Created monthly maximum and minimum temperatures, which we converted to four bioclimatic temperature fields matching those available through WorldClim.
- Reanalysis of temperature provided at the native spatial resolution of MERRA. Using ancillary information on elevation and geopotential height from the Global Land One-km Base Elevation Project (GLOBE) and MERRA, the native resolution temperature fields from MERRA were downscaled to match the 1km spatial resolution of GLOBE using cubic convolution interpolation
- Used MERRA data from its inception in 1979 to 2000 to focus on the overlap with the existing WorldClim data set, which employed weather station data primarily from 1960 to 2000.

# New Bioclimes – AMSR-E

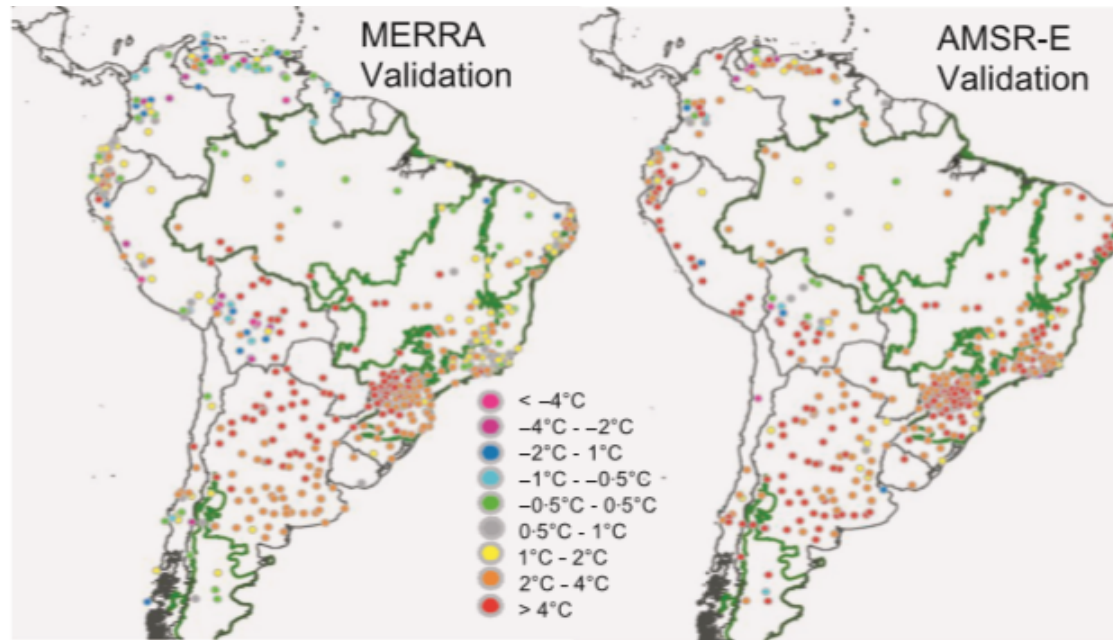
- Near-daily temperature minima and maxima obtained by inversion of a simplified semi-physical radiometric model that uses morning and evening brightness temperature observations. The temperature dataset provides global temperature retrievals over land for snow and ice-free non-frozen conditions for periods of no precipitation.
- Descending (morning) and ascending (evening) orbital nodes from AMSR-E's temperature retrieval provide respective minima and maxima for temperature at approximately 2 m height (Jones et al. 2010).
- We used the temperature observation from the morning and evening satellite overpasses, converted these temperatures to average monthly values, and then derived the four bioclimatic temperature fields.
- The grid resolution of the AMSR-E temperature fields is approximately 25 km; we downscaled the fields to 1 Km using cubic convolution interpolation. Downscaling followed that of the MERRA data set, except that the geopotential height was replaced by the 25-km EASE grid GLOBE DEM.
- Data generated from the 2003–2010 AMSR-E observation period.



**Fig. 3.** Differences across values of bioclimatic variables for temperature estimated from MERRA and AMSR-E data sets relative to WorldClim values. Left: difference between MERRA-derived and WorldClim values; right: difference between AMSR-E-derived and WorldClim values. Four bioclimatic variables are illustrated: Bioclim 1 (mean annual temperature), Bioclim 4 (temperature seasonality), Bioclim 10 (mean temperature of warmest quarter) and Bioclim 11 (mean temperature of coldest quarter; Hijmans *et al.* 2005). Values for Bioclim 1, 10 and 11 are given in degrees Celsius, whereas Bioclim 4 values are percentages.



# Exploring new ways to improve characterization of present-day environments



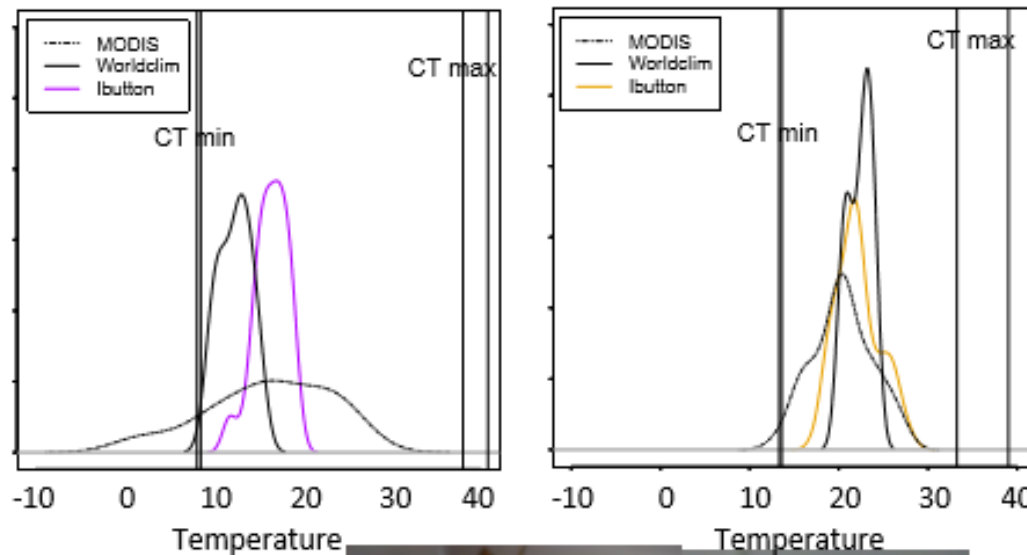
Surface temperature estimates showed warm temperature biases relative to in situ data fields; reliability of these datasets varied in space.

**Fig. 2.** Ground-validation of MERRA and AMSR-E data relative to mean ground temperatures recorded at 1006 weather stations (years 1950–2000) by the Global Historical Climatology Network and the World Meteorological Organization. Each dot represents a weather station; colours in a) and b) depict the net difference between respective MERRA and AMSR-E estimated temperatures and recorded ground temperature and those in c) indicate the difference in mean temperature recorded at surface weather stations between the more heavily sampled 1950–2000 period and the 2003–2008 period. Green boundaries outline the South American regions examined in the study.

Waltari et al. 2014.  
Methods in Ecol. & Evol.

## Validating environmental temperatures with thermal physiology

MODIS LSTs fall outside CT's more often than Worldclim and ibutton temperatures, especially in the topographically complex southern AF.



Critical temperatures (CT min and CT max) represent tolerance limits, with CT max potentially being a harder boundary than CT min.



CT min and max were estimated experimentally in 5 species of lizards, by heating up or cooling down lizards until they lose coordination.

