A multidisciplinary framework for biodiversity prediction in the Brazilian Atlantic Forest hotspot









UNICAMP

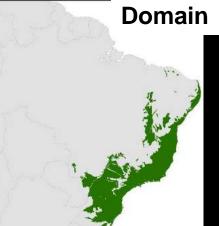


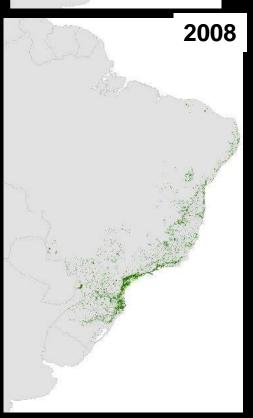


The Brazilian Atlantic Rainforest





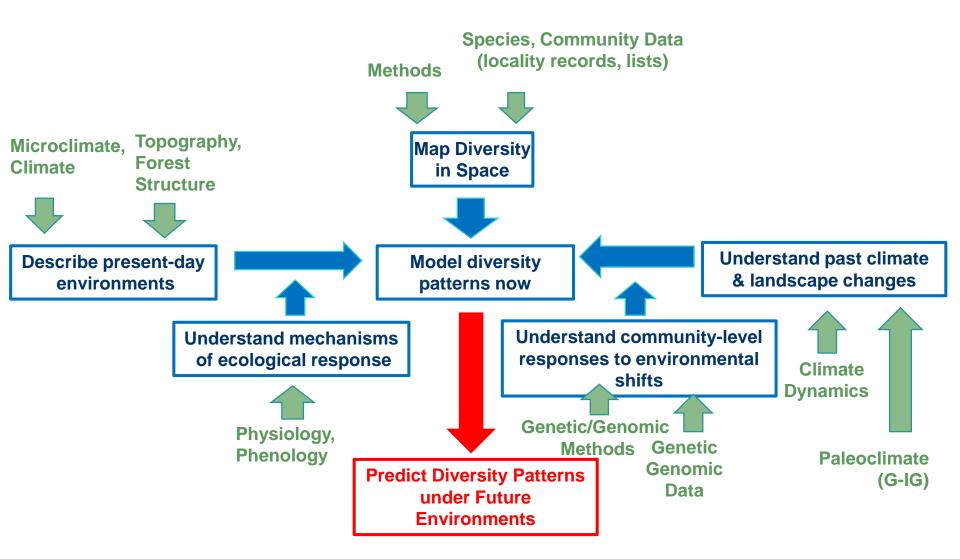




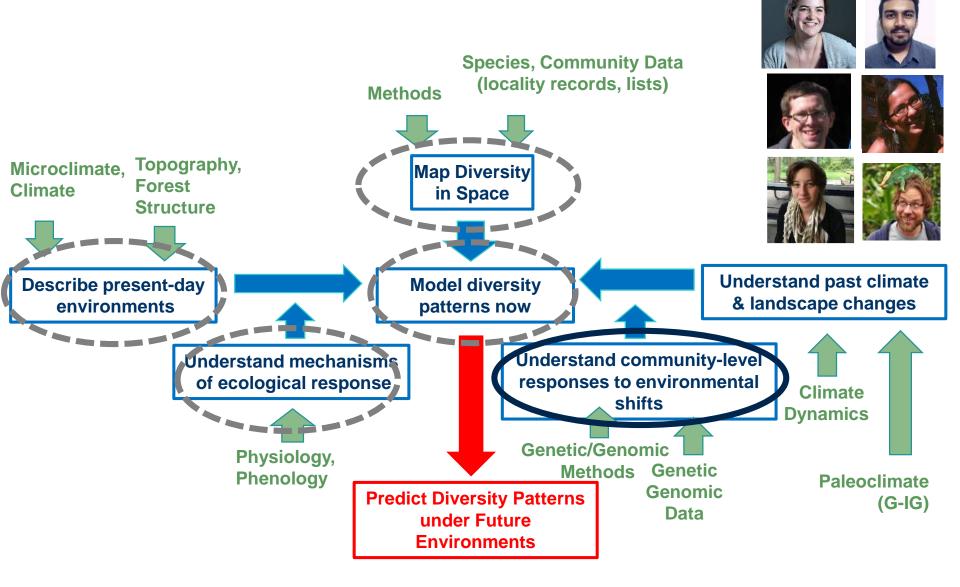




Framework

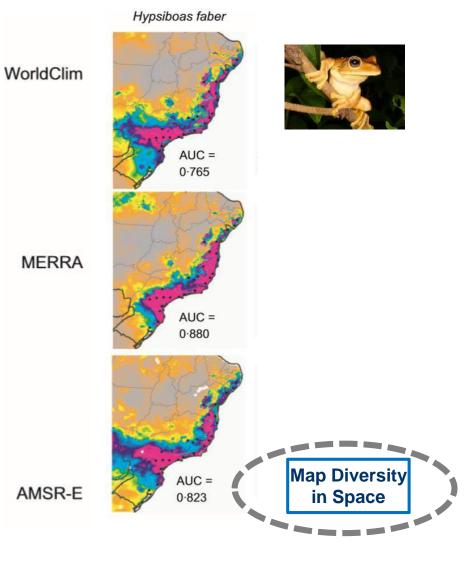


How can the incorporation of remote sensing components improve biodiversity prediction?



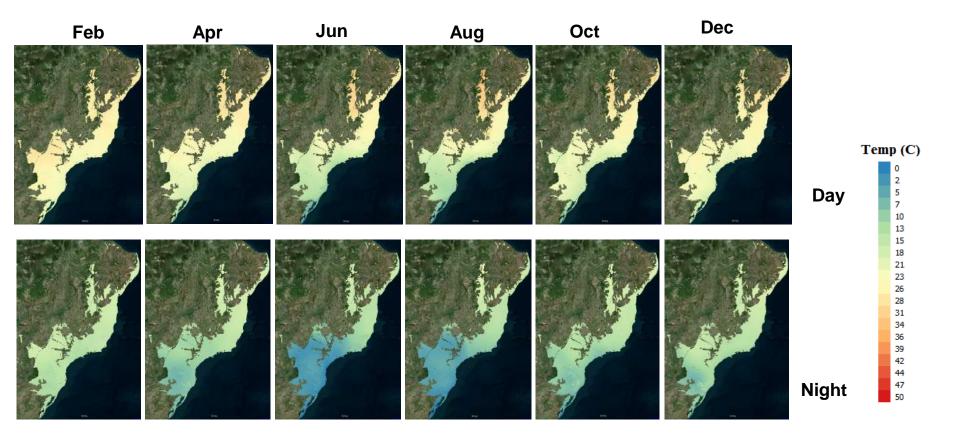
Microwave data enable characterization of present-day environments for biodiversity modeling

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





AMSR-E Climatology enables description of day/night temperature shifts (here, LST Min; 2003-2011)

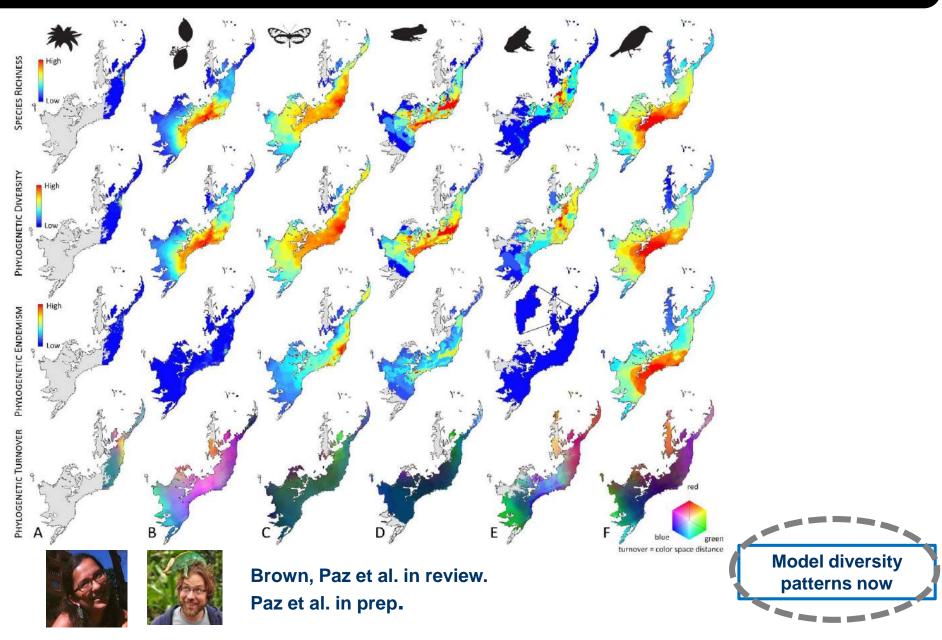








Remote sensing products predict diversity patterns at larger ecological and geographical scales



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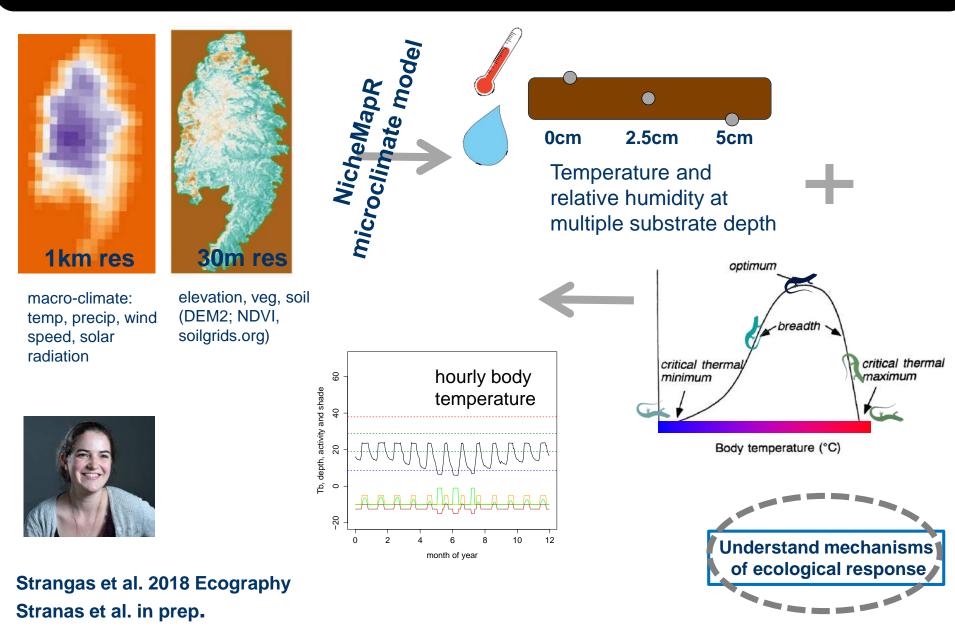
R ² Phylogenetic Diversity		R ² Phylogenetic Endemism	PRELIMINARY
8 6 89	0.92	0.77	ANALYSES BASED ON MODIS AND CHIRPS DATA
	0.95	0.51	Conditional Autoregressive Models used to address spatial
	0.94	0.46	autocorrelation; probability of values at any given location is conditional on
*	0.78	0.27	neighboring values.
	0.89	0.05	
	0.86	0.05	Model diversity patterns now



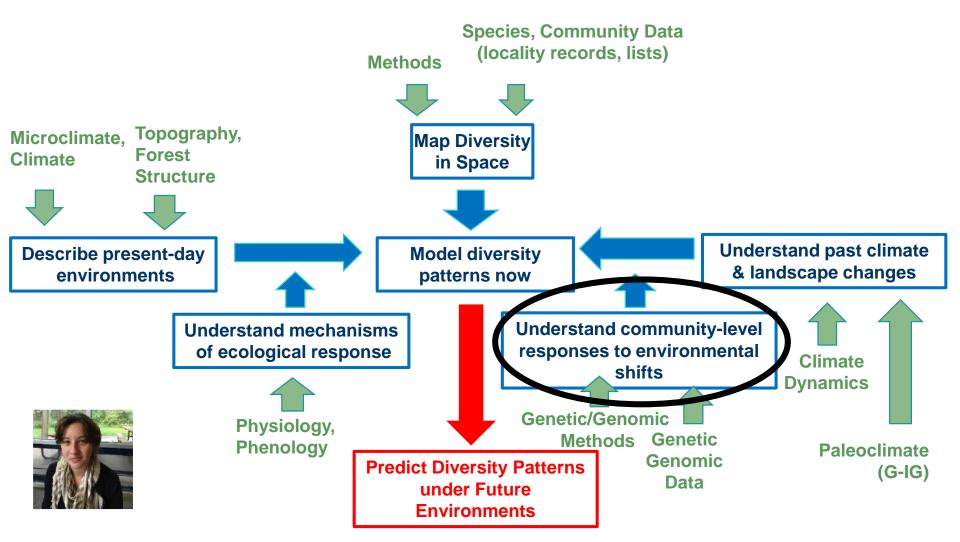


Paz et al. in prep.

Remote sensing improves description of microclimates, which are needed to understand physiological responses to climate

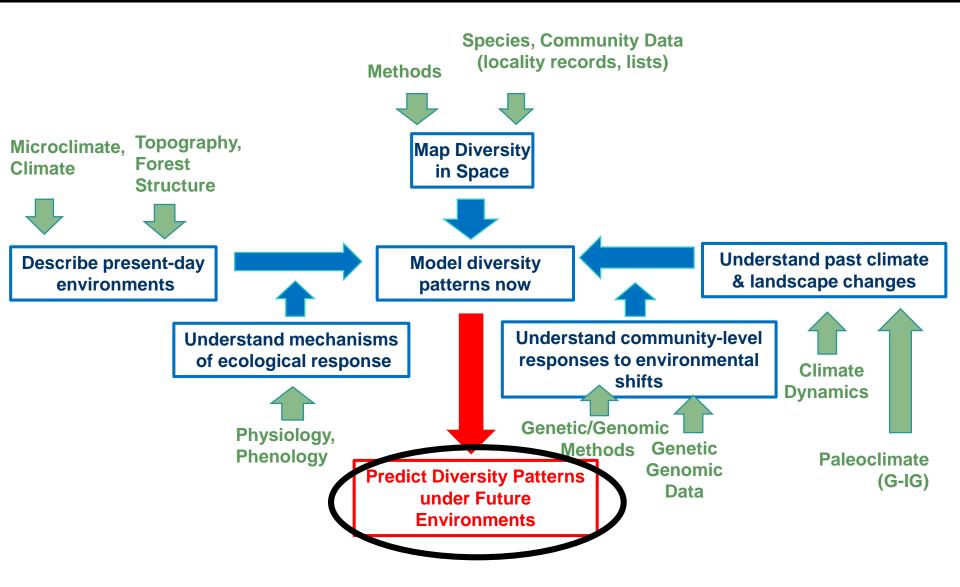


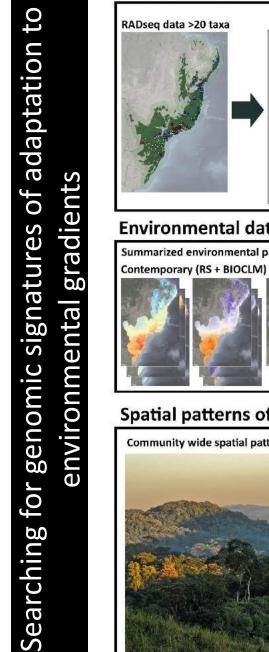
Searching for genomic signatures of adaptation to environmental gradients

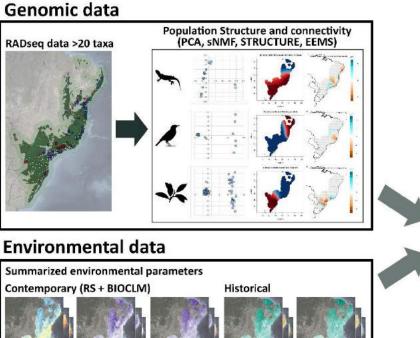


Bertola et al. in prep.

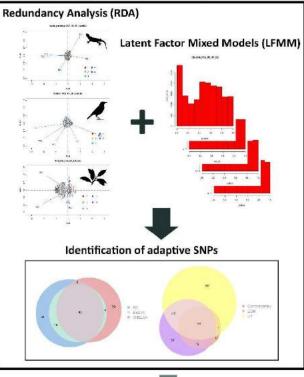
Stay tuned for next year's meeting







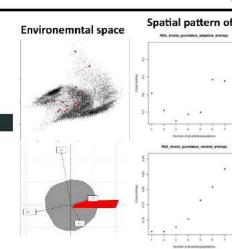
Adaptive genomics analyses

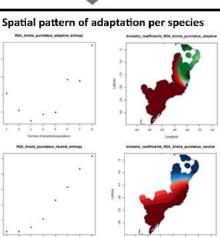


Spatial patterns of adaptation

Community wide spatial pattern of adaptation







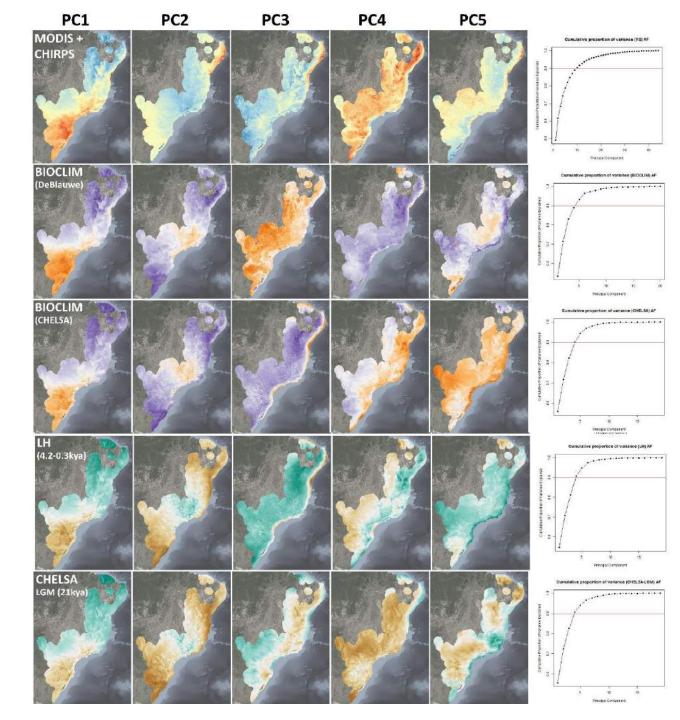
Environmental data:

Contemporay:

- Remote sensing
- BIOCLIM (DeBlauwe)
- BIOCLIM (CHELSA)

Historical:

- BIOCLIM LH
- BIOCLIM LGM



Genomic data:

- RADseq data
- >20 taxa
- Different ranges/ environmental space

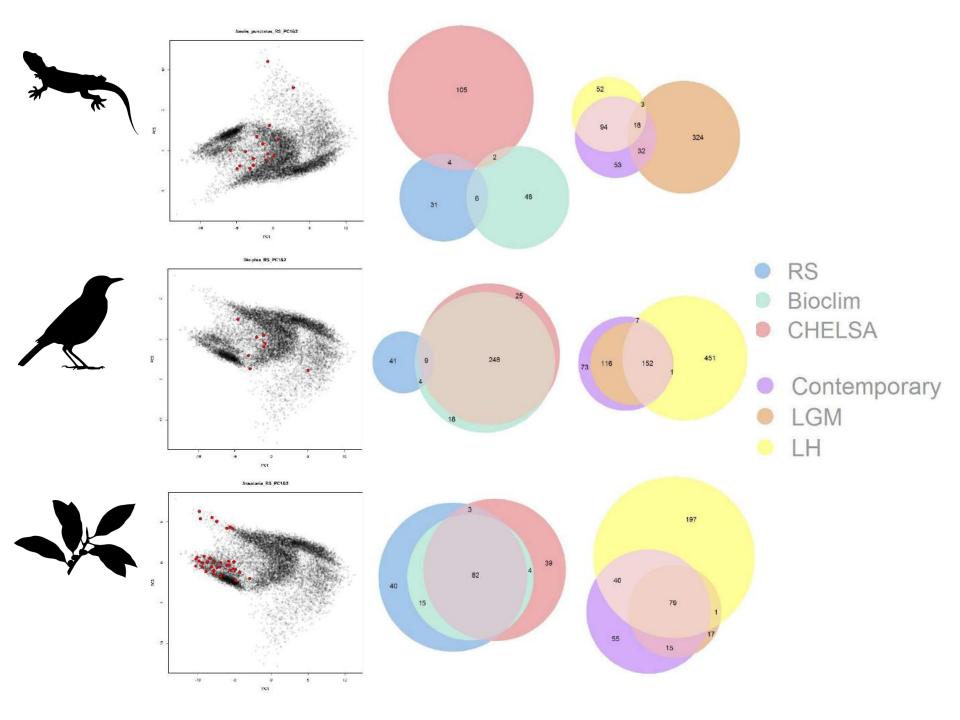












Ancestry_coefficients_RDA_Anolis_punctatus_neutral

40 19

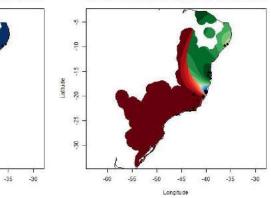
10

-52

-30

-20

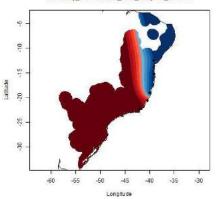




Longitude Ancestry_coefficients_RDA_Dixiphia_neutral

-55 -60

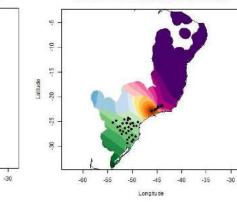
-50 -45 -40



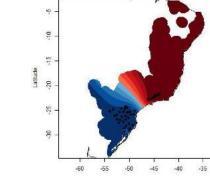




Ancestry_coefficients_RDA_Araucaria_neutral Ancestry_coefficients_RDA_Araucaria_adaptive

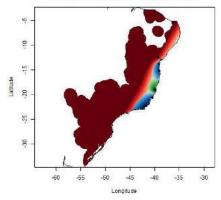






Longitude

Ancestry_coefficients_RDA_Dixiphia_adaptive



Acknowledgements

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