

Spatially continuous identification of beta diversity hotspots using species distribution models

Gabriel Dansereau^{1,2,3} Timothée Poisot^{1,2,3,4} Pierre Legendre^{1,3}

¹Département de sciences biologiques, Université de Montréal

²BIOS²

³Quebec Center for Biodiversity Science

⁴Groupe de recherche interuniversitaire en limnologie et environnement aquatique

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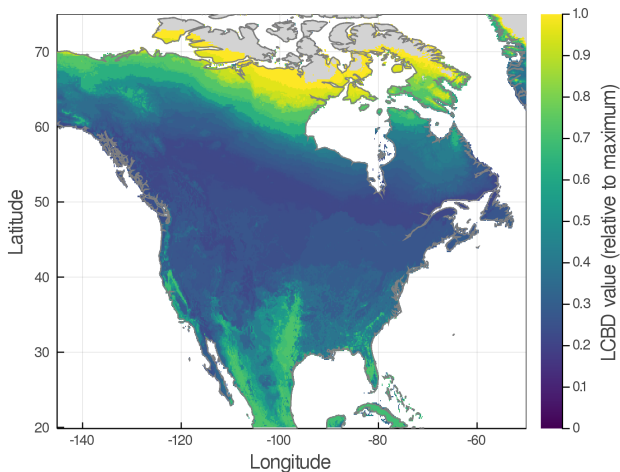
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Suggestion

Bring together 2 elements:

1. Identification of beta diversity hotspots
2. Species distribution modelling (SDM) on continuous scales



While we're at it...

Beta diversity

- ▶ Community composition, not turnover

"Variation in species composition among sites within a geographical region of interest" (Legendre et al. 2005)

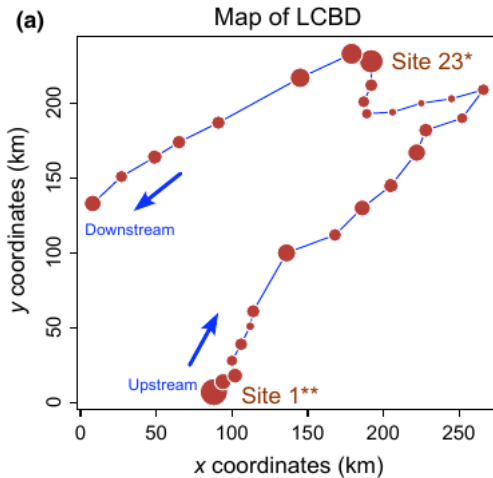
Local contribution to beta diversity (LCBD)

- ▶ Highlights exceptional species compositions

"Comparative indicators of the ecological uniqueness of sites in terms of community composition" (Legendre & De Caceres, 2013)



Why continuous scales?

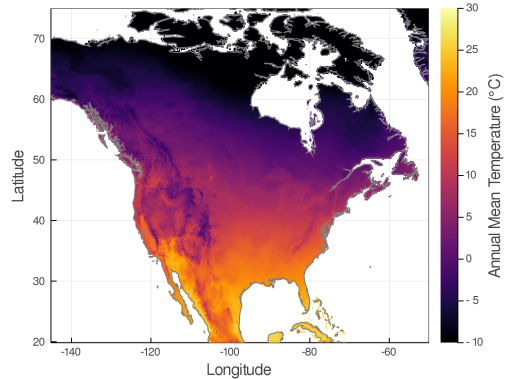
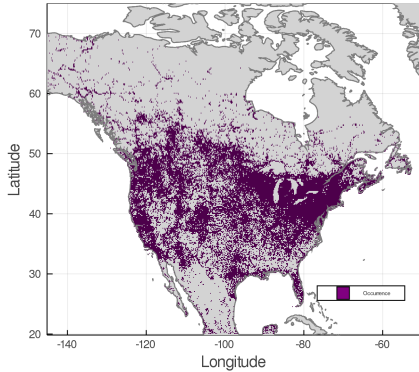


Original LCBD example (Legendre & De Caceres, 2013)



Why continuous scales?

- ▶ Online data on extended scales is increasingly accessible
- ▶ Potential for novel ecological insights



Relevance

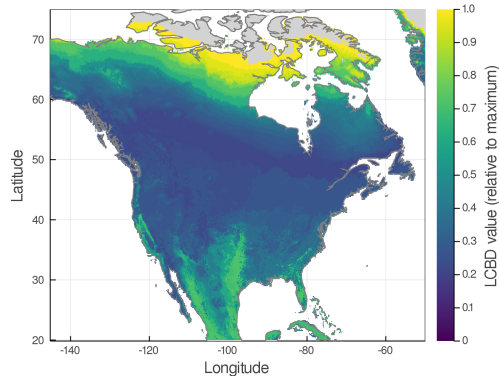
Novel ecological insights

- ▶ Tool for poorly sampled regions, or with sparse sampling
- ▶ Identification of conservation targets

Combination with IPCC climate change scenarios

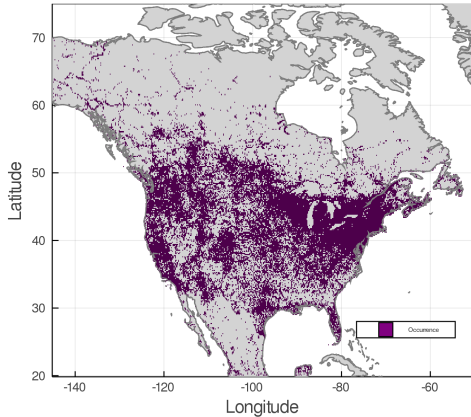
- ▶ Model beta diversity changes
- ▶ Identify sites with significant changes

⇒ Insight-oriented approach, exploratory analyses



Occurrence data

- ▶ Data from the eBird Basic Dataset
- ▶ All species of Warblers (*Parulidae* family) in North America

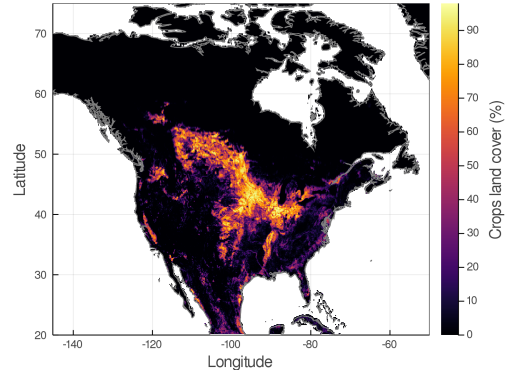
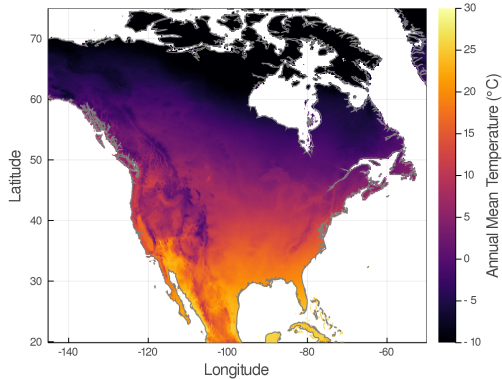


eBird

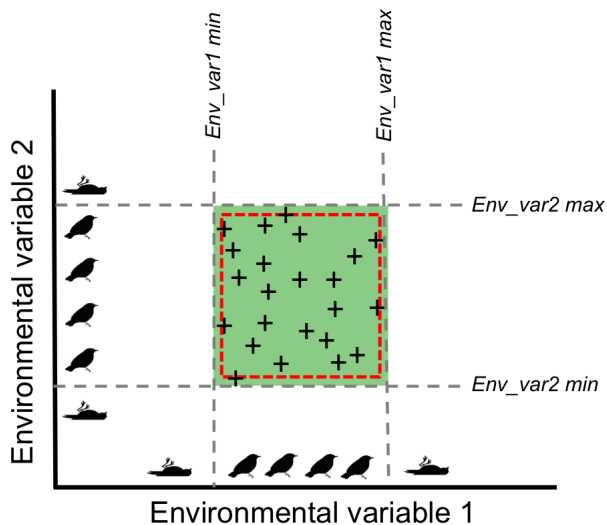


Environmental data

- ▶ 2 climates variables : mean annual temperature, mean annual precipitation
- ▶ 10 land cover variables : bare, crops, grass, moss, shrub, snow, tree, urban, permanent water, seasonal water



Methods - BIOCLIM model

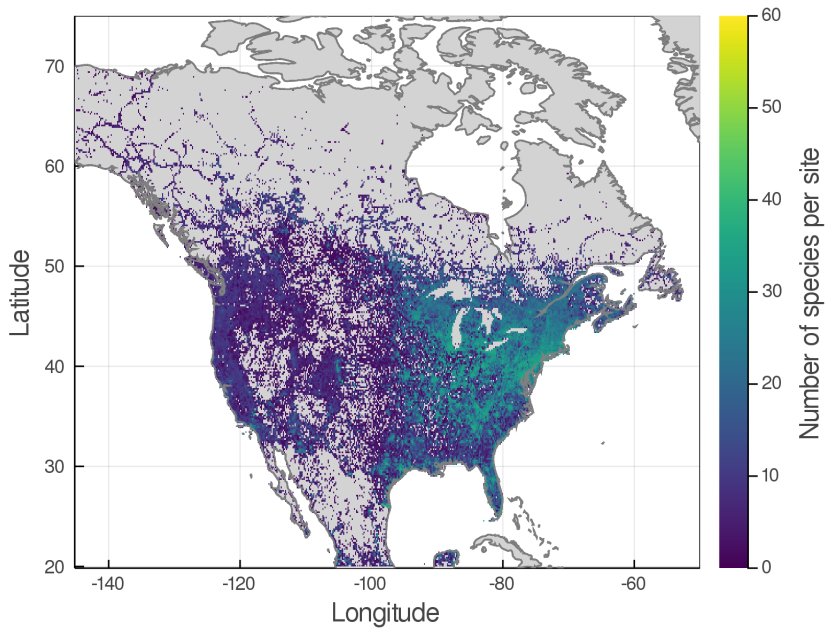


Climate envelope in the BIOCLIM model¹

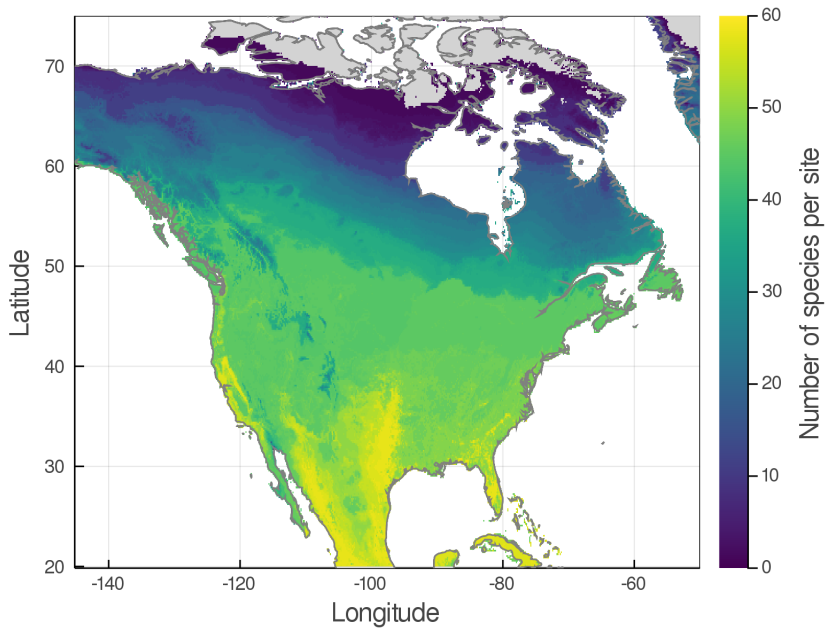
¹<https://support.bccvl.org.au/support/solutions/articles/6000083201-bioclim>



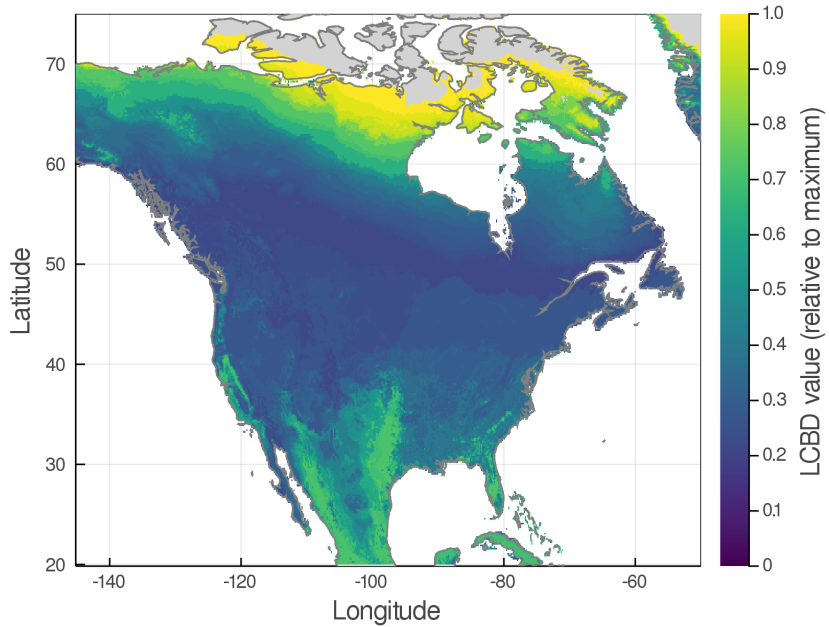
Species richness - Raw data



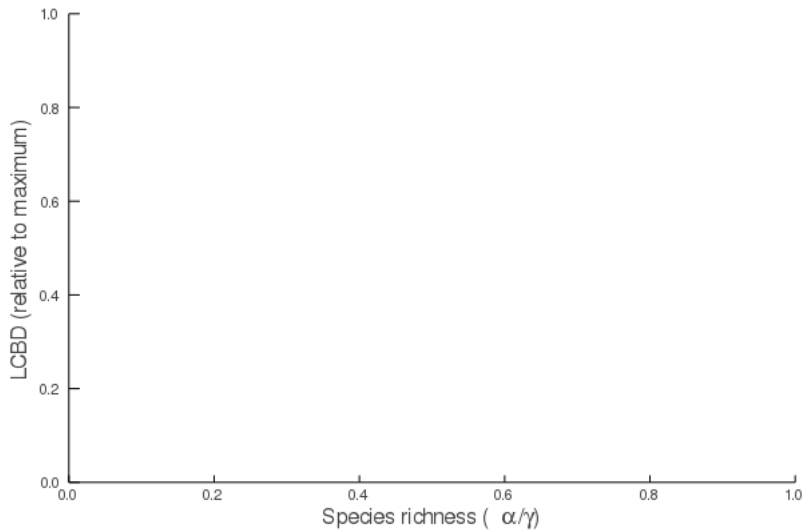
Species richness - SDM results



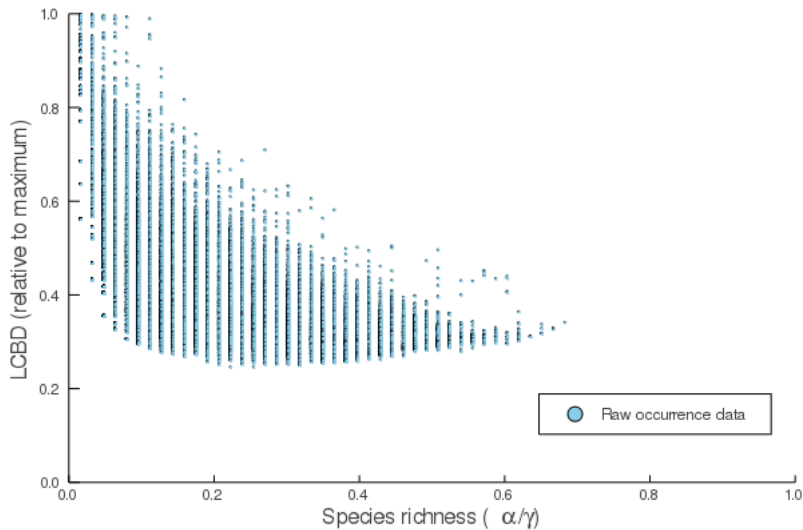
LCBD values - SDM results



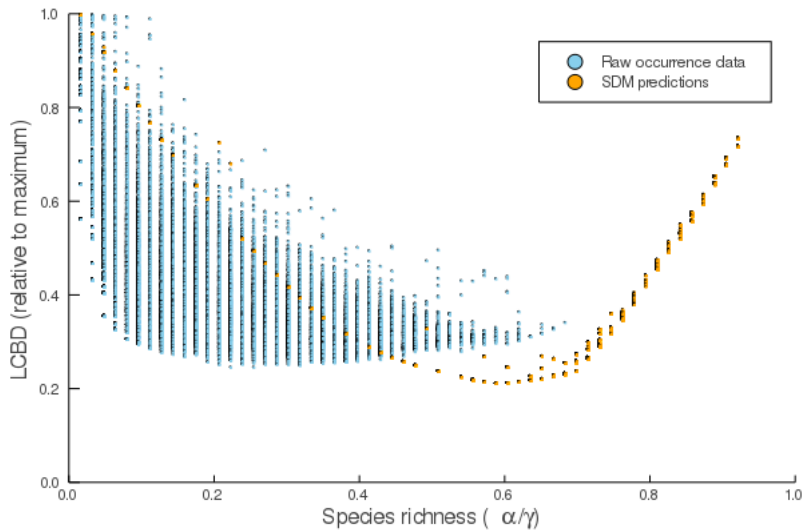
LCBD-richness relationship



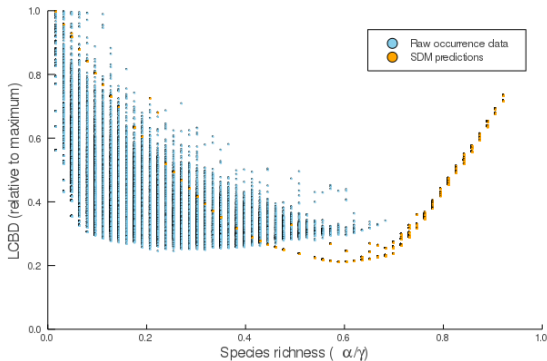
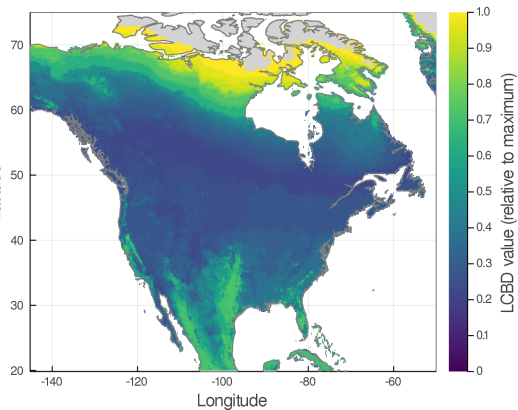
LCBD-richness relationship



LCBD-richness relationship



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Gabriel Dansereau
gabriel.dansereau@umontreal.ca



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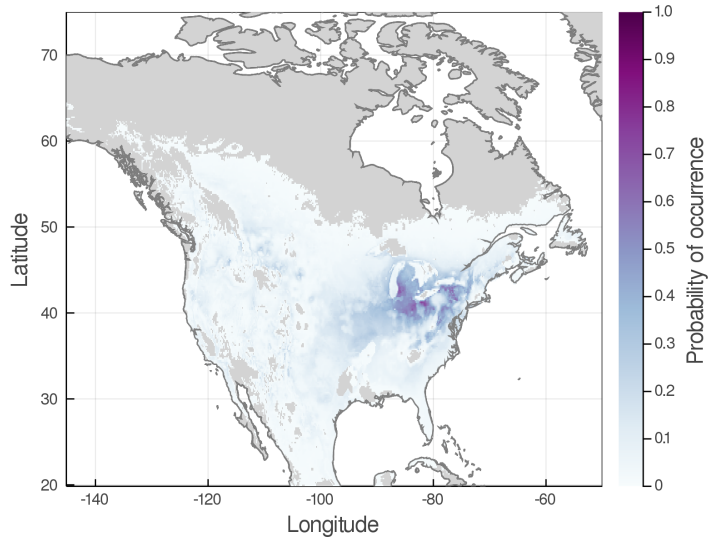
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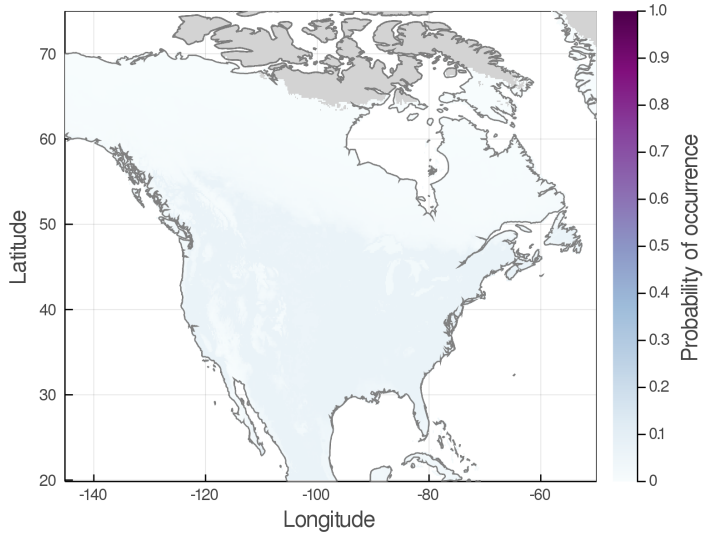
Appendix



Single species example - SDM with threshold



Single species example - SDM without threshold



LCBD - Raw data (with Hellinger transformation)

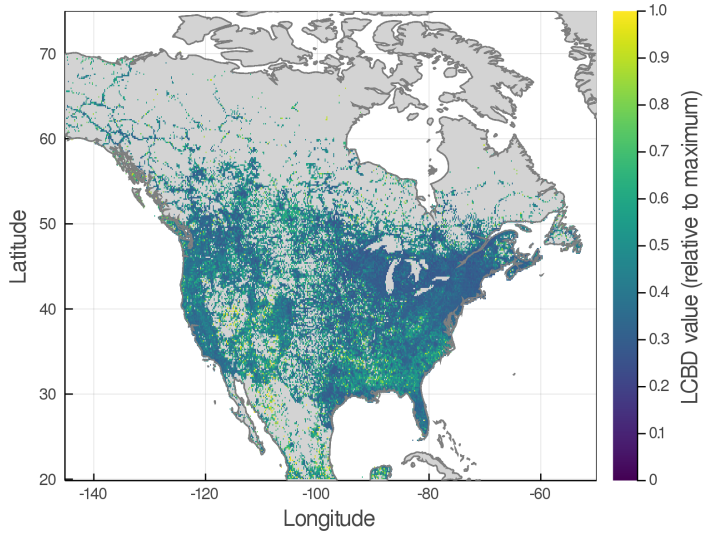


Figure 1: LCBD values relative to maximum value based on the raw data after Hellinger transformation

