

# Reconstructing the Patterns of Alien Plant Species Habitat Niche Expansion

**Gilles Colling**

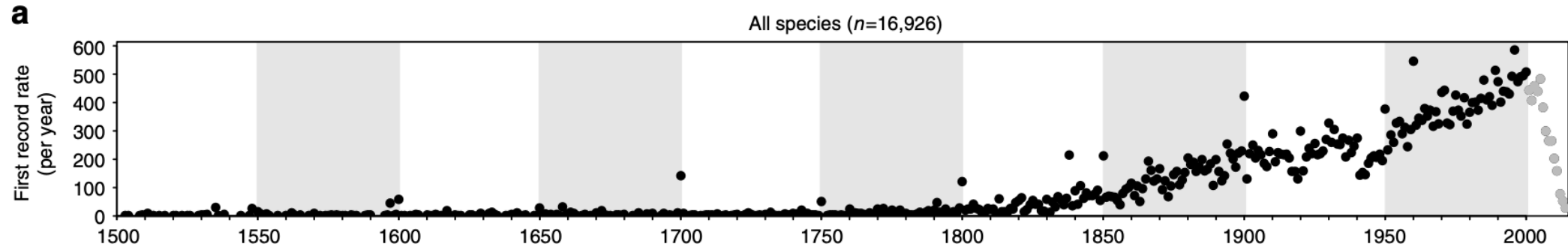
Division of BioInvasions, Global Change & Macroecology

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**TAC members:** Dr. Bernd Lenzner, Dr. Hanno Seebens, Dr. Michael Glaser

# No Signs of Saturation



## ARTICLE

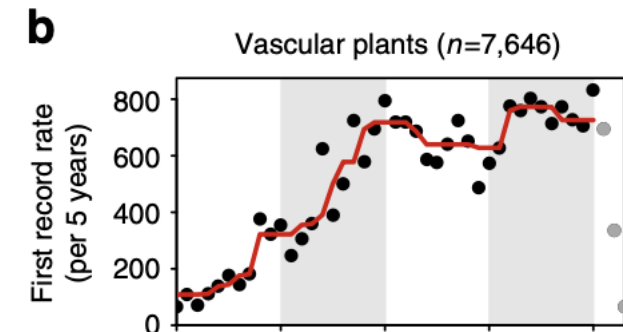
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DOI: [10.1038/ncomms14435](https://doi.org/10.1038/ncomms14435)

OPEN

## No saturation in the accumulation of alien species worldwide

Hanno Seebens *et al.*<sup>#</sup>

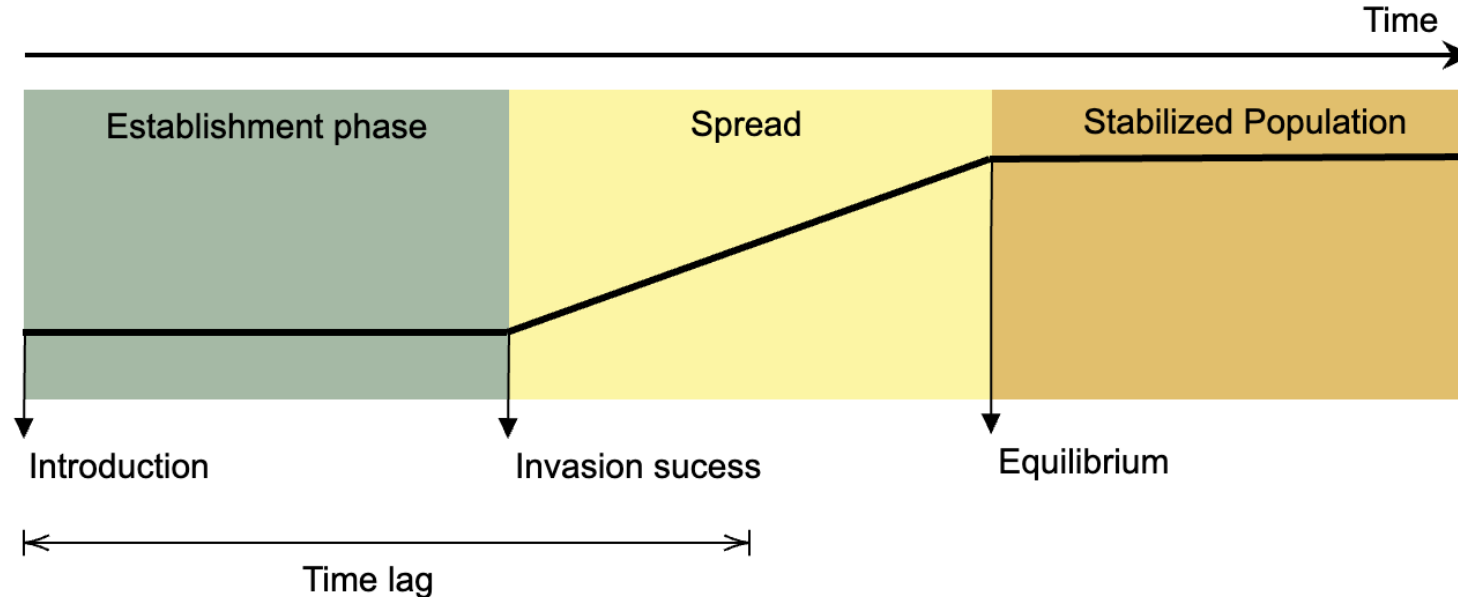


# Corollary

- **Regional accumulation** of alien species well-documented (Juozaitienė et al., 2023; Seebens et al., 2017) but **local-scale** effects **less explored**

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- **Regional accumulation** of alien species well-documented (Juozaitienė et al., 2023; Seebens et al., 2017) but **local-scale** effects **less explored**
- **High numbers** of aliens in **regional** species pools, but **low representation** in **local** communities, mostly a **time-lag phenomenon**



# European Perspective - Habitat Origins

- **7'300 alien plants** are recorded across **55 European territories**, most = **neophytes** (Kalusová et al. 2024)

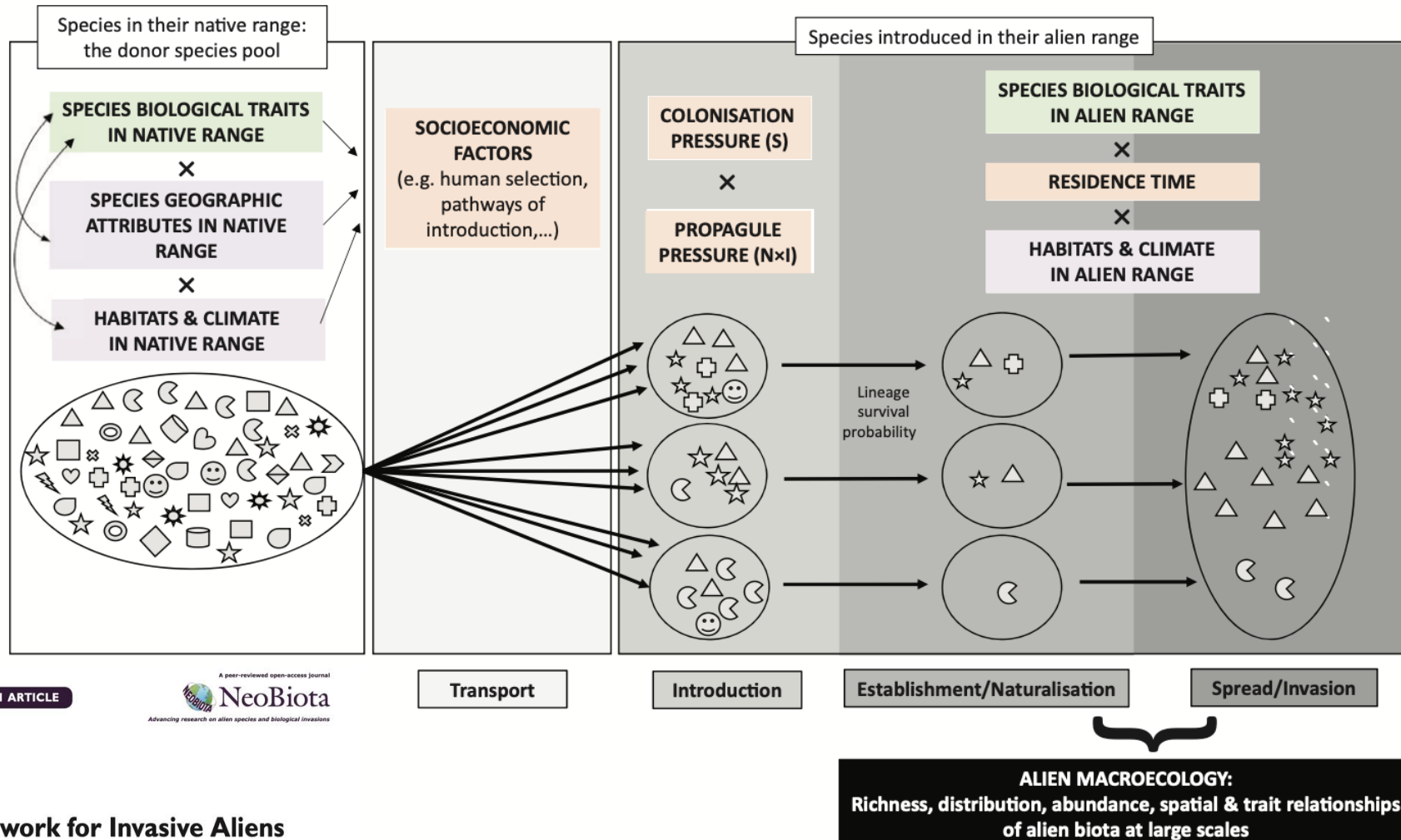
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- **Habitats shaped by disturbance** and environmental **variability**, (coasts, floodplains, and urban margins) → **main entry points** for alien species. (Kalusová et al. 2013)
- Interconnected habitats **likely facilitate expansion** of alien species **into new habitat types**

# Residence Time and Habitat Expansion



NeoBiota 62: 407–461 (2020)  
doi: 10.3897/neobiota.62.52787  
http://neobiota.pensoft.net

RESEARCH ARTICLE

A peer-reviewed open-access journal  
**NeoBiota**  
Advancing research on alien species and biological invasions

**MAcroecological Framework for Invasive Aliens (MAFIA): disentangling large-scale context dependence in biological invasions**



# Hypotheses

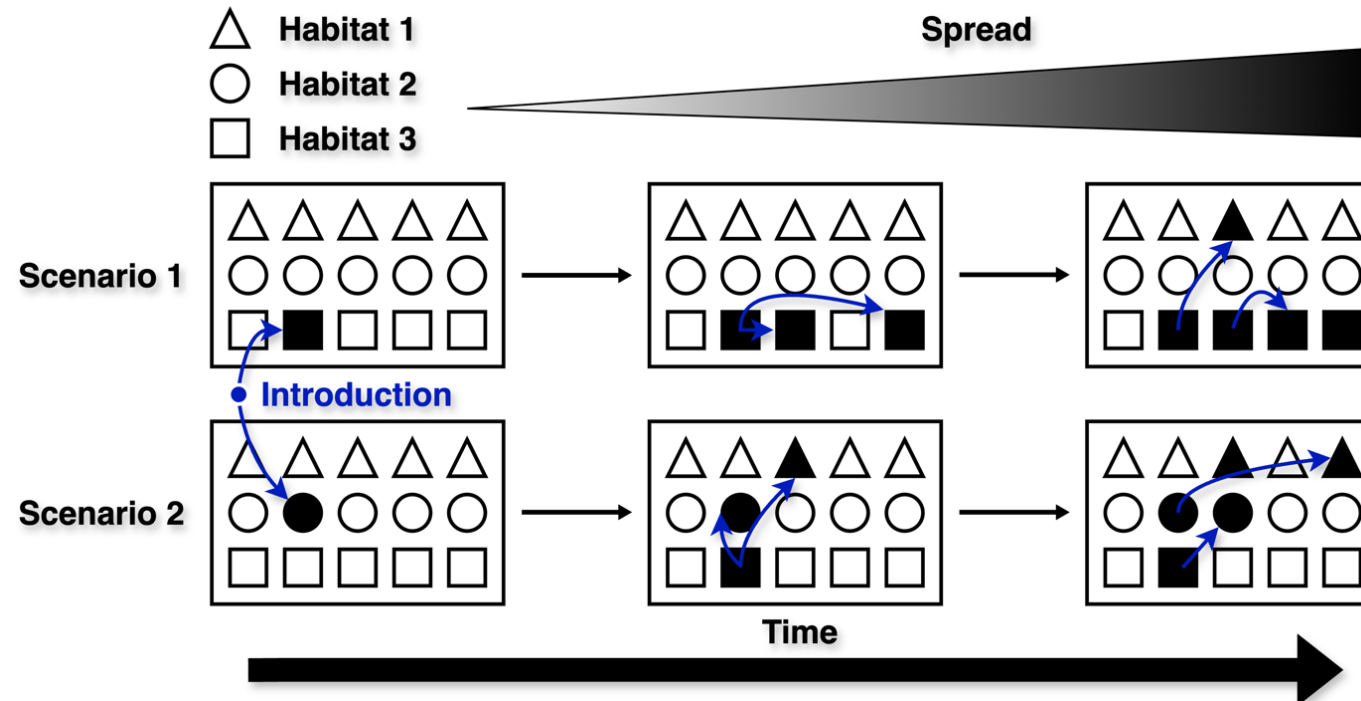
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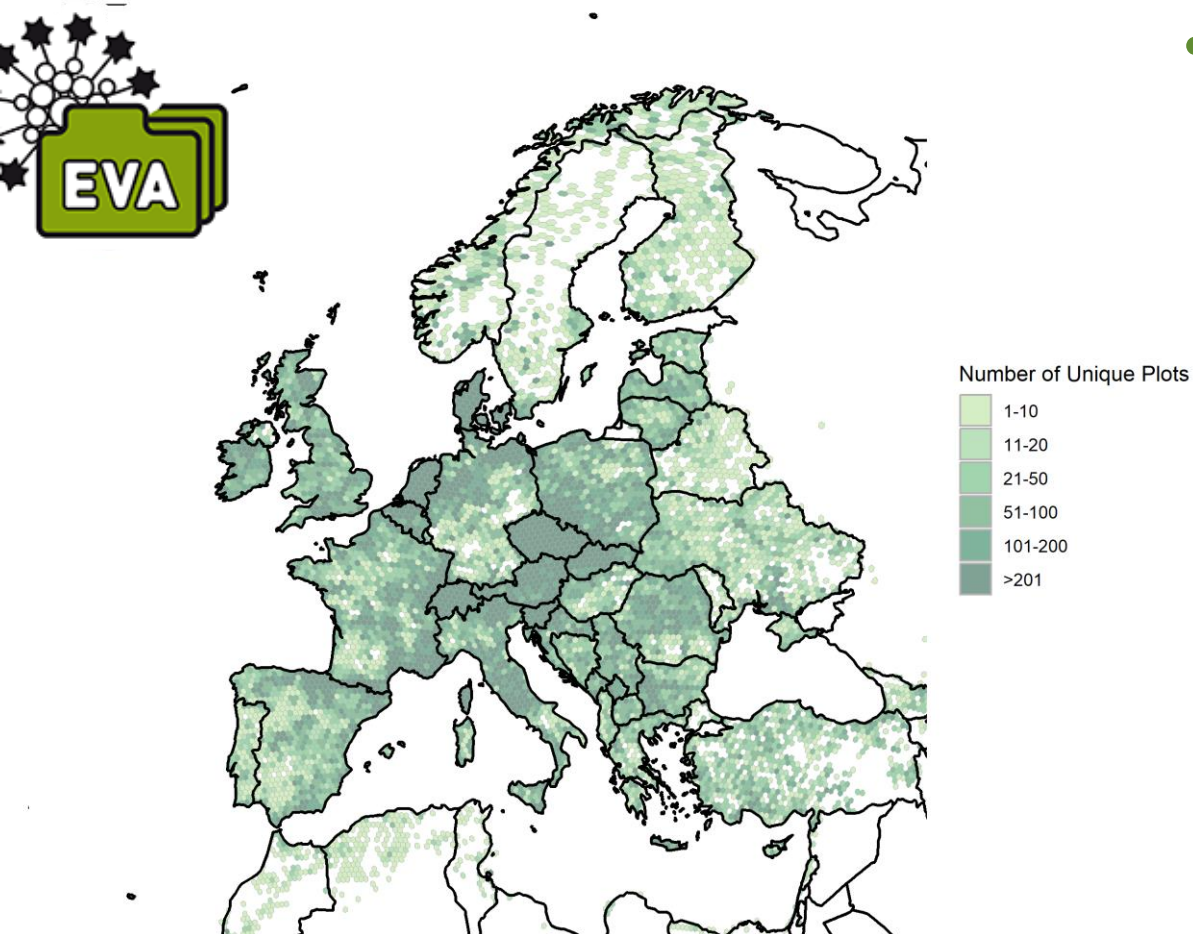
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  - (b) **more local communities of different habitat types**.



# Dataset

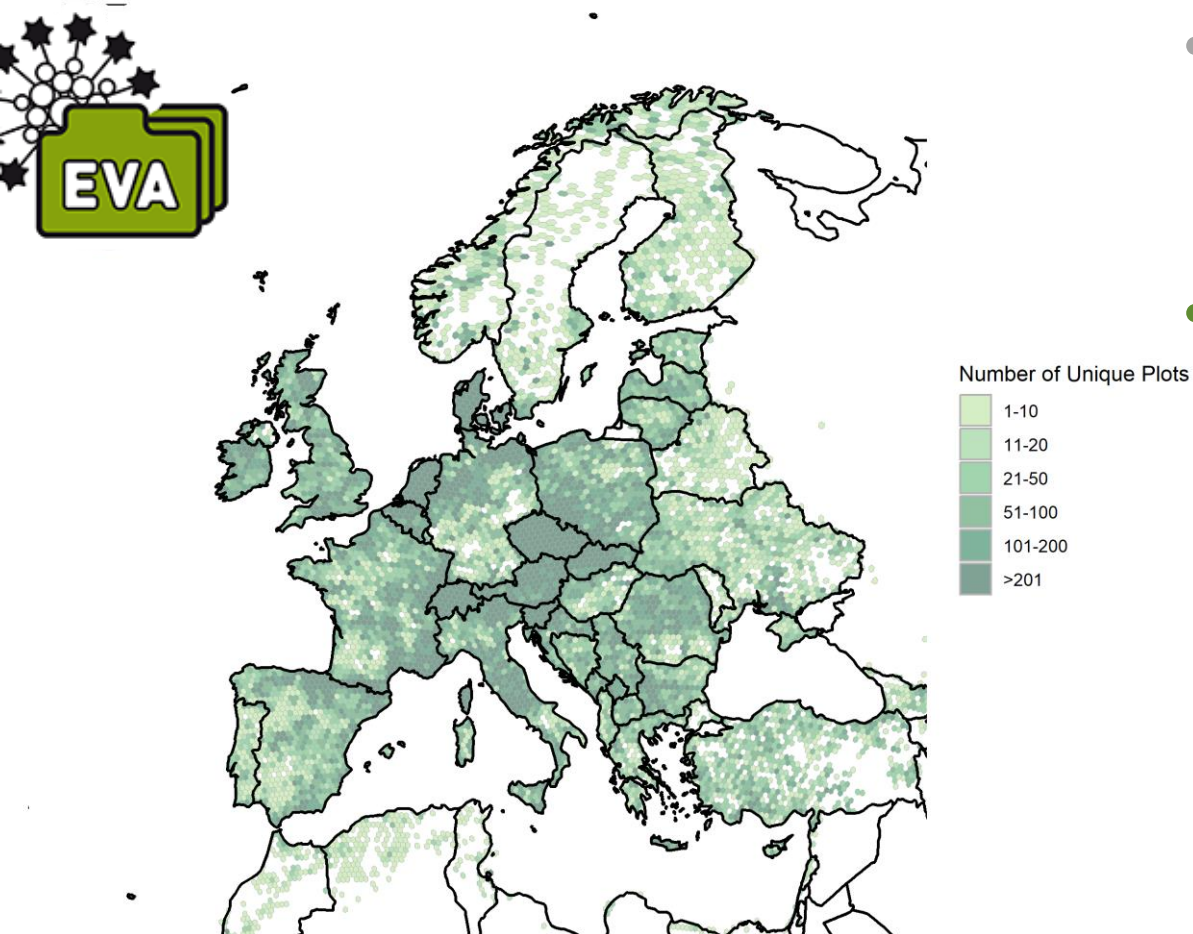
- **EVA** (The European Vegetation Archive)



- Data for **ASAAS** compiles **1.9 million vegetation plots** from more than 100 regional databases **across Europe**

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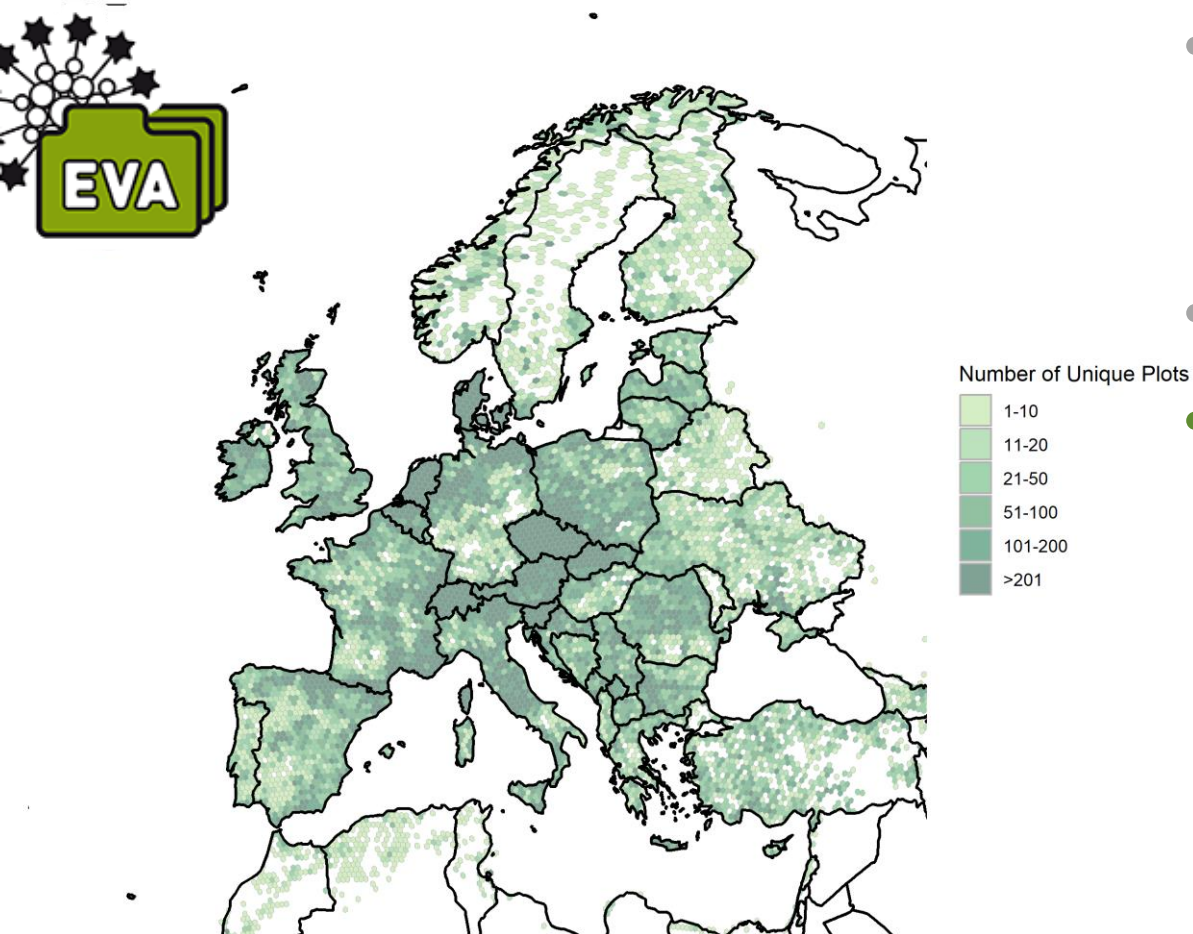
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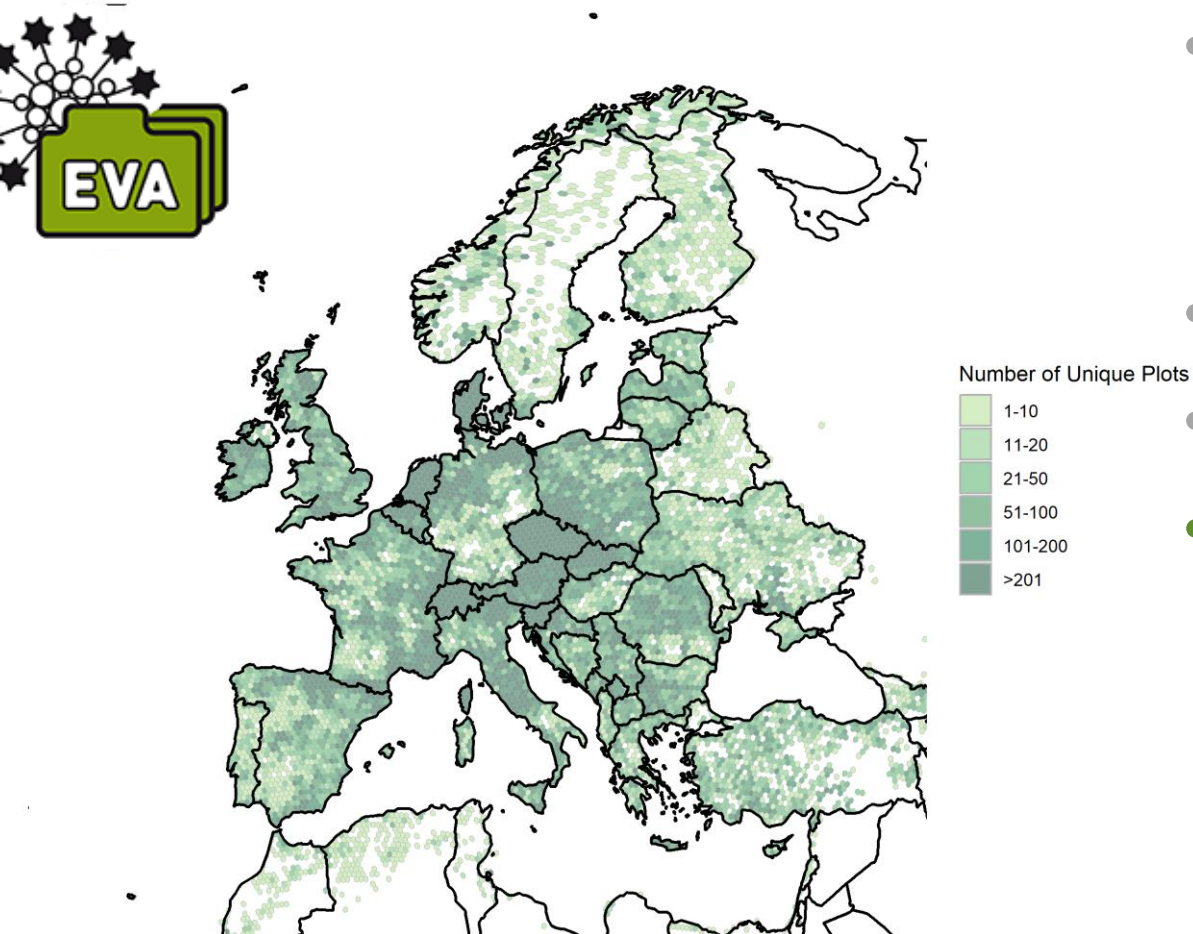


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- **9 Decades (1930-2020)**



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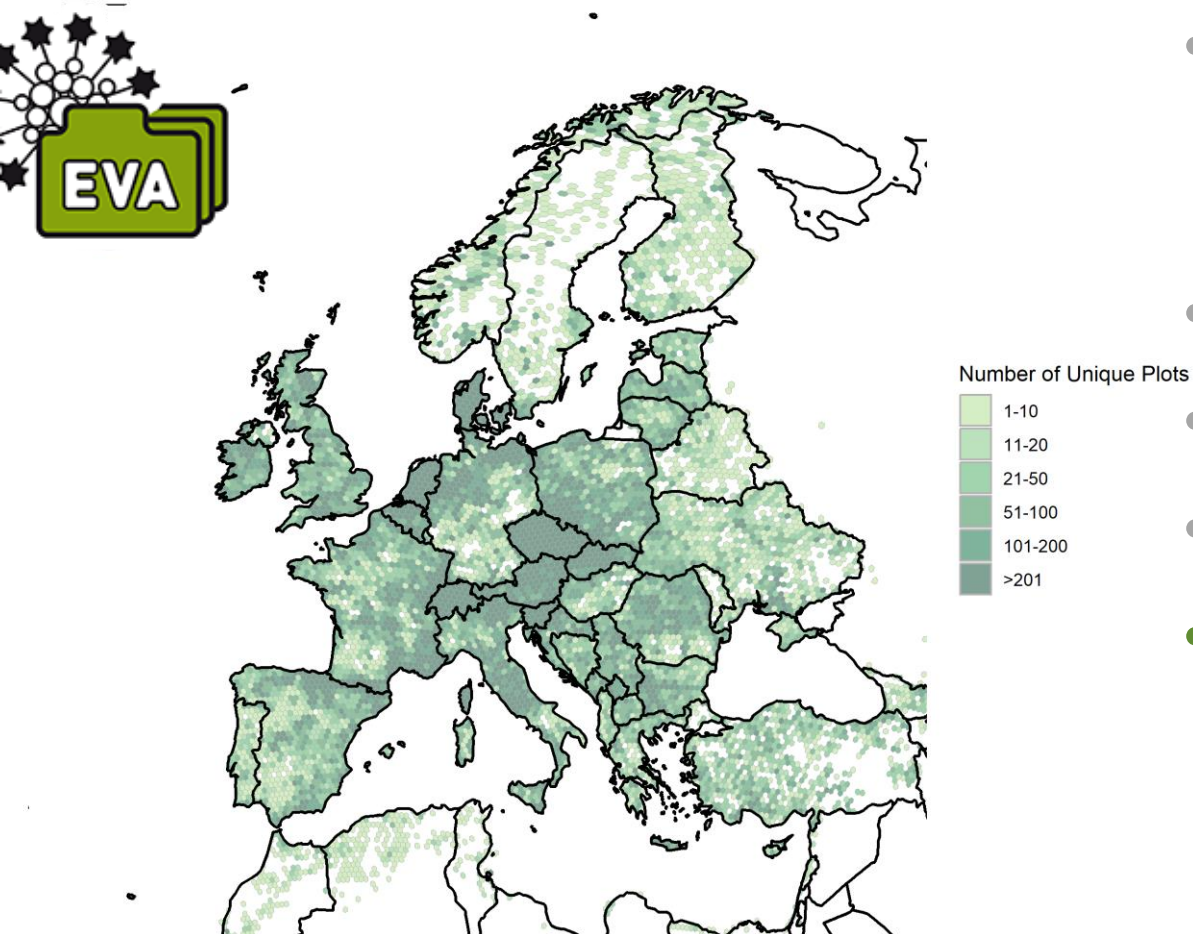
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- **EUNIS classification**



# Habitat Classification

- EUNIS Level 2 aggregated into 18 habitats

Group	Examples
Man-made & ruderal	Arable, gardens, artificial grasslands
Broadleaved deciduous	Temperate deciduous forests
Mesic & wet grasslands	Hay meadows, wet grasslands
Dry grasslands	Xerophytic grasslands
Alpine & subalpine	High-altitude grasslands
Bogs & poor mires	Peat-accumulating wetlands

**Intermediate resolution:** Level 1 too coarse, Level 2 too detailed

# The Problem with Counting Habitats

- If we *observe a species* in **grasslands and wetlands**...
  - a) is it: **Habitat preference**, or **sampling** where it is common?
  - b) Raw **occurrence** conflates distribution with **habitat availability**
  - c) EVA is **sampled opportunistically**: some habitats heavily, some sparse

**Need: Compare species to *local* habitat availability**

# How to measure Habitat profiles?

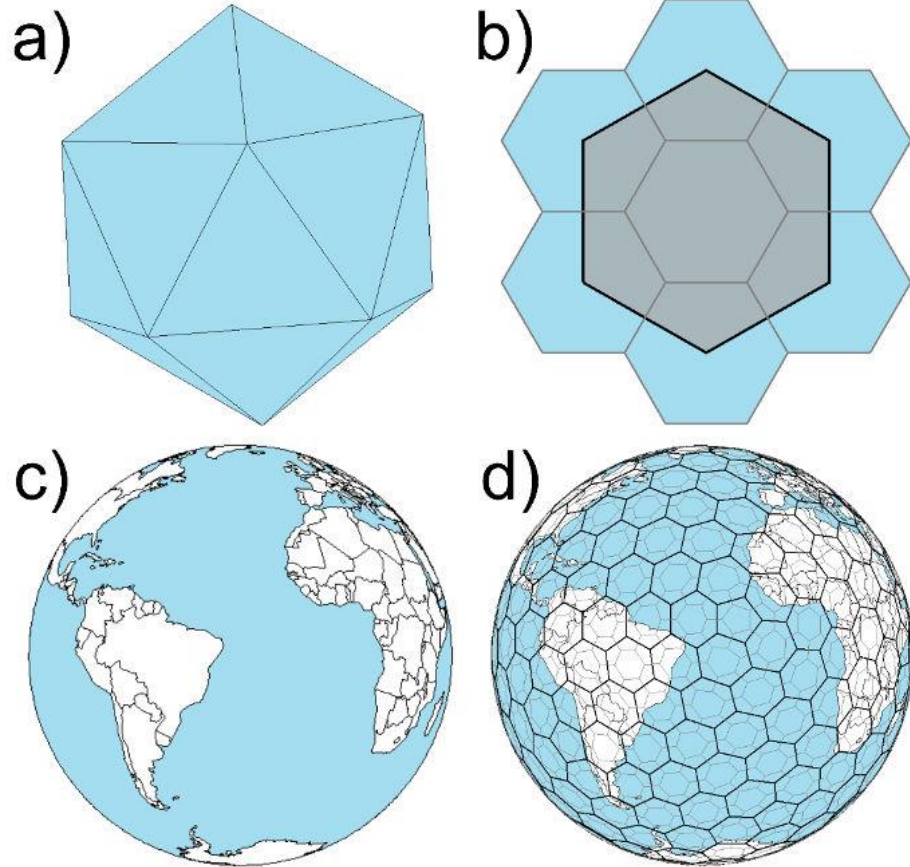
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# How to measure Habitat profiles?

- **Fidelity, IndVal, diversity indices** = designed for **systematic** surveys
- **Problems** within EVA data:
  - **Denominators unstable** when habitats **sampled unevenly**
  - **Shannon diversity increases** with **sample size** regardless of true distribution and isn't well defined for "habitat diversity"
  - Does **not answer**: "Is the **species more common than random chance** given local availability?"

# Methods

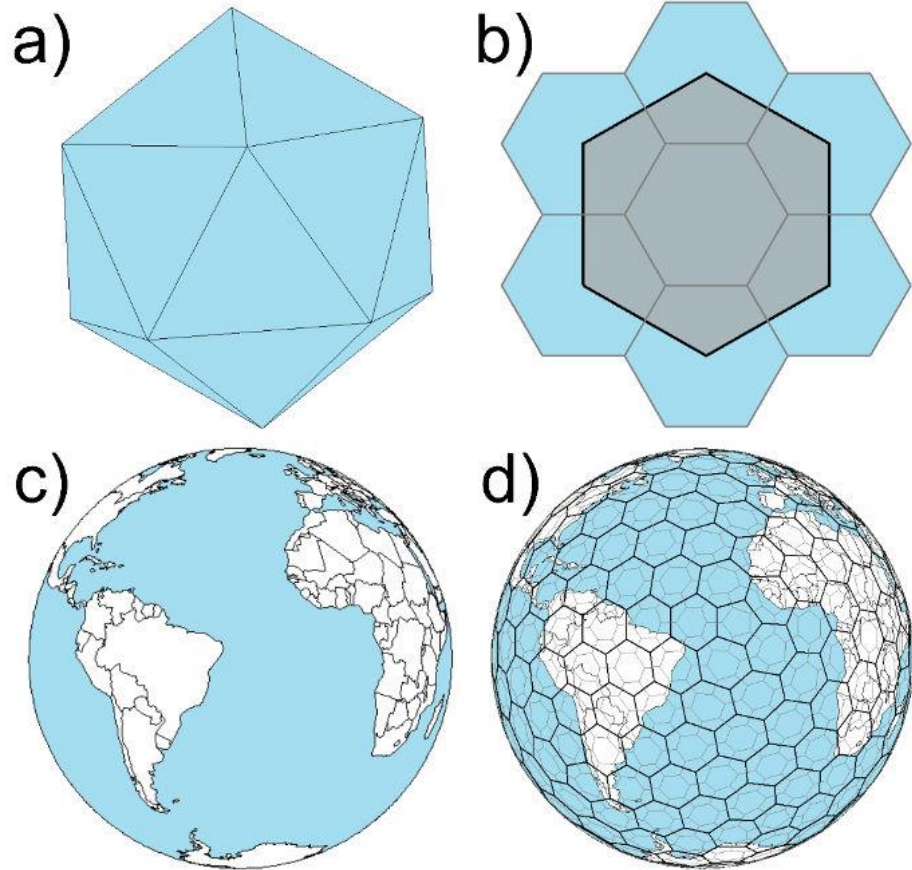
## Comparing Habitat use across 863 km<sup>2</sup> x 10 years hexcells



- **Accounting for sampling:**  
→ Reorganize data into hexagonal grid cells

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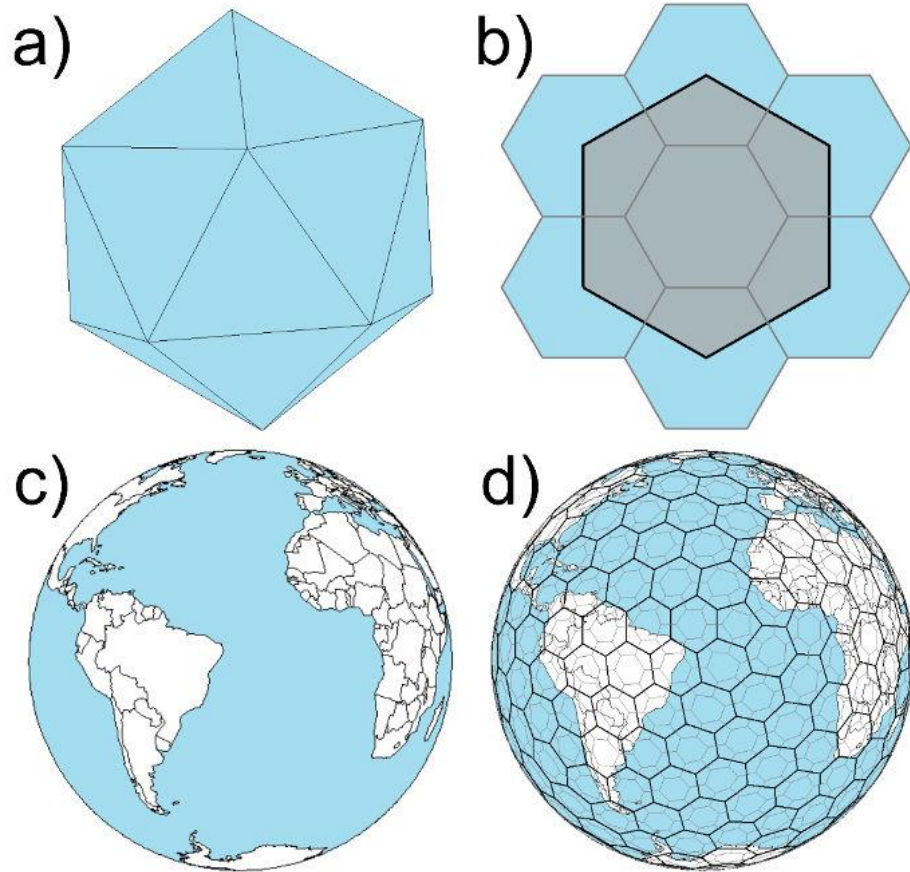
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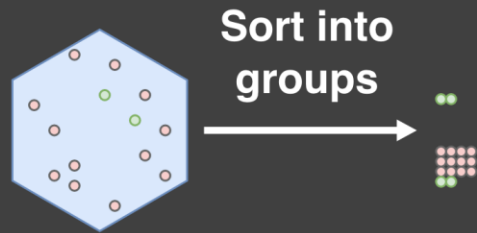
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- Within each **hexcell** we collect:
  - **Neophyte** habitat distribution (**p**)
  - **Background** habitat distribution (**q**)
- Within each habitat (per decade):
  - Null-model:  **$p \approx q$**
  - Overrepresented:  **$p > q$**

# Methods

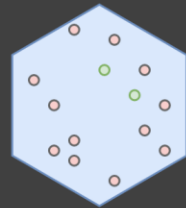
- Cell with species
- Cell without species
- Plot with species
- Plot without species





# Methods

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Sort into  
groups

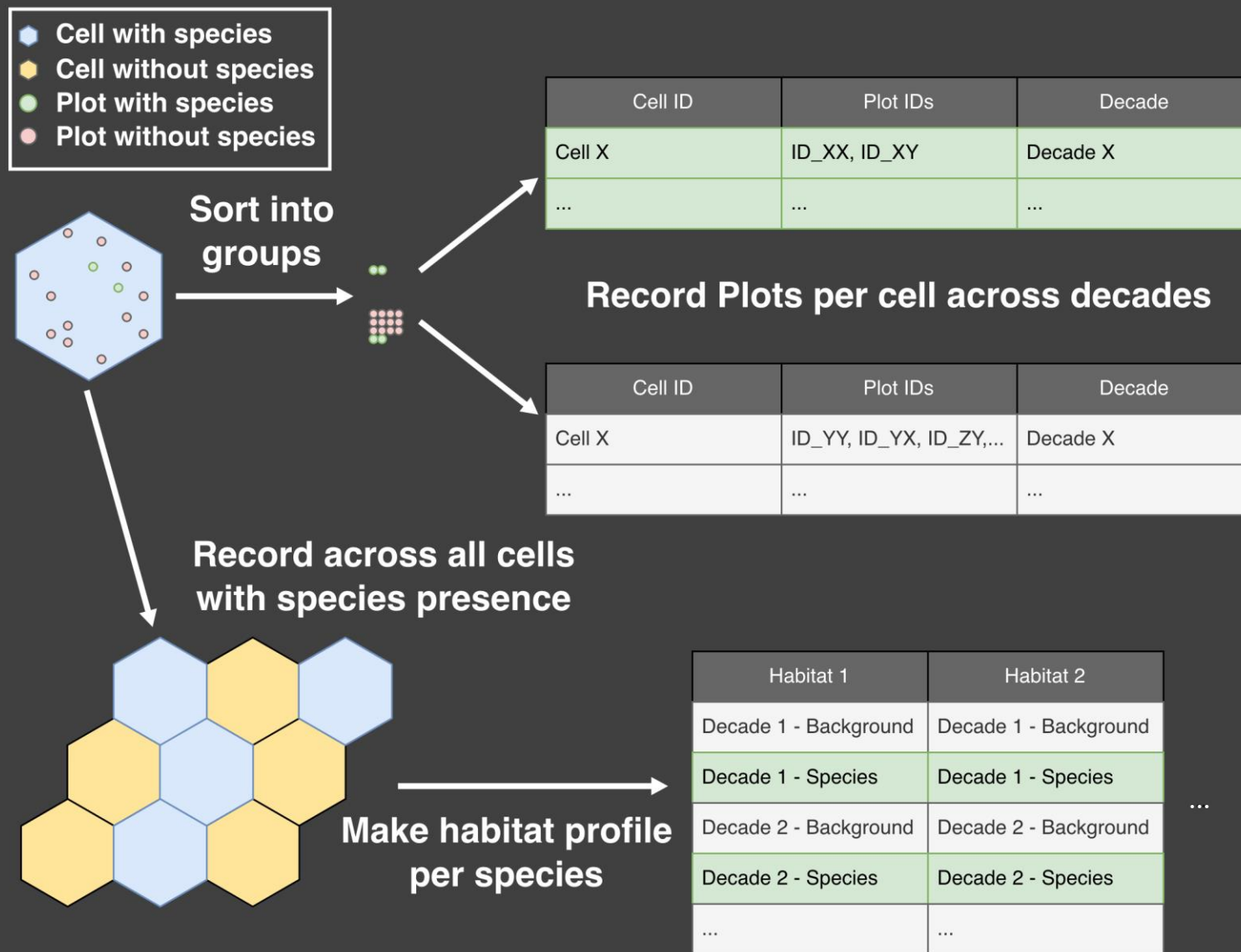


Cell ID	Plot IDs	Decade
Cell X	ID_XX, ID_XY	Decade X
...	...	...

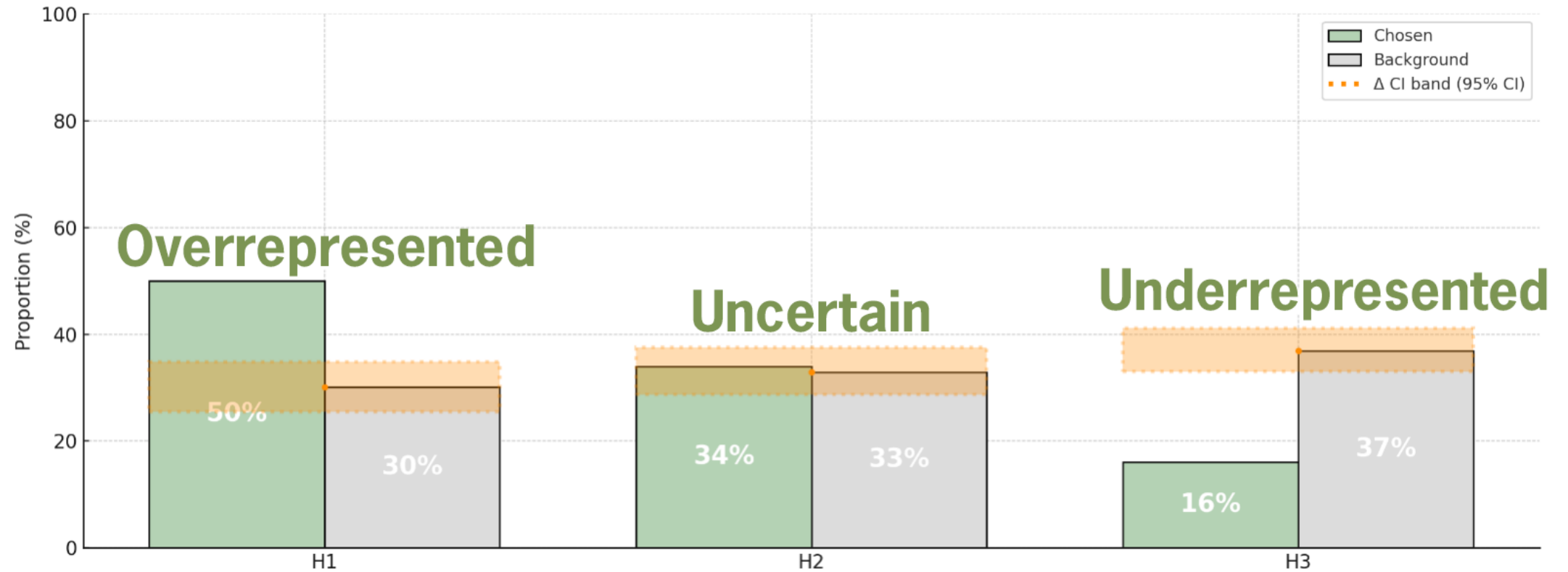
Record Plots per cell across decades

Cell ID	Plot IDs	Decade
Cell X	ID_YY, ID_YX, ID_ZY,...	Decade X
...	...	...

# Methods



# Methods



**Species (green) vs background (grey) with 95% CI band**

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  - Each hexcell x decade is independent of global sampling
  - Oversampled regions don't dominate the signal
- **Categorical output**
  - Species is overrepresented **(1) or not (0)** per habitat
  - Directly **usable as response**

# Methods

**Biologically, two distinct processes are involved:**

- 1. Can the species **extend beyond one habitat?** (generalist potential vs. specialist constraint)**

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1. **Can the species extend beyond one habitat?** (generalist potential vs. specialist constraint)
  - Many species remain constraint
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2. **How many additional habitats** does it occupy once it expands?
  - Given expansion, what determines breath?
  - Outcome: Counts (number of additional habitats)



# Methods

$$Y_i \sim \text{TruncHurdleNegBin}(\pi_i, \mu_i, \phi)$$

- **Hurdle** component (logit link)

$$\text{logit}(\pi_i) = \alpha^{(\text{hur.})} + \mathbf{X}_i \boldsymbol{\beta}^{(\text{hur.})} + u_{\text{county}[i]}^{(\text{hur.})}$$

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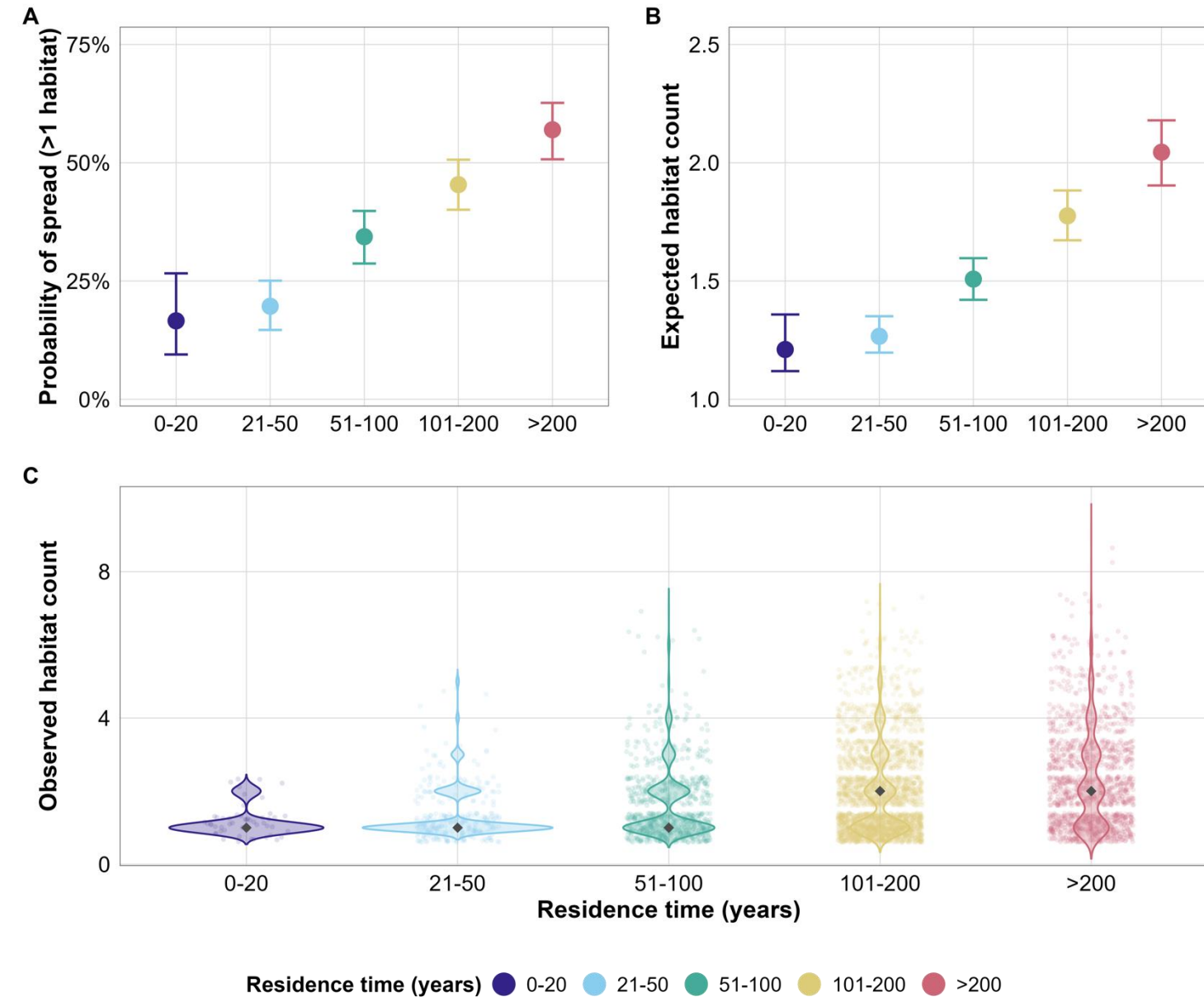
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- **Random effects:** country intercepts

# Results

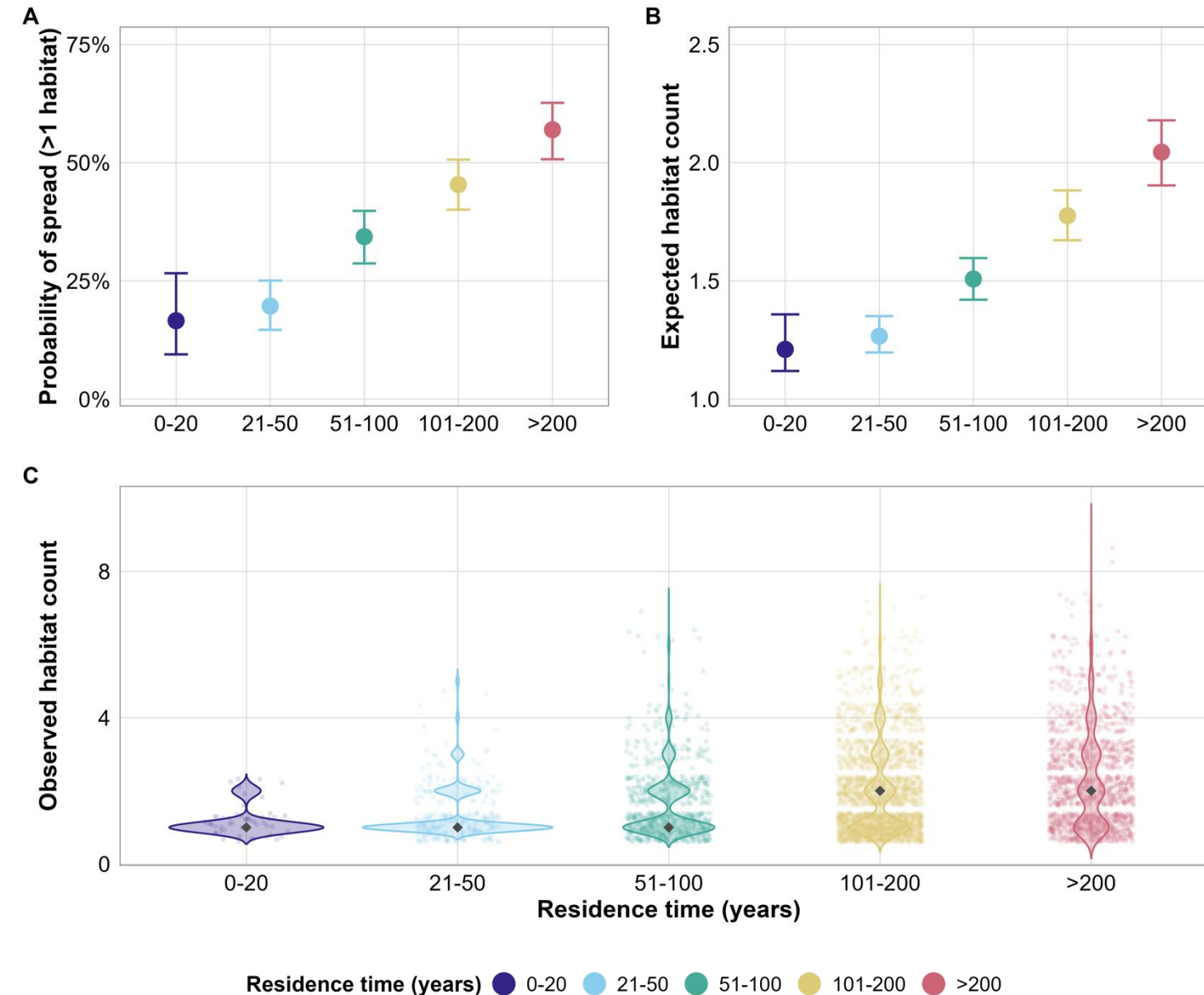


- **Probability of > 1 habitat:**

- 0-20 years: 17%

- >200 years: 57%

# Results



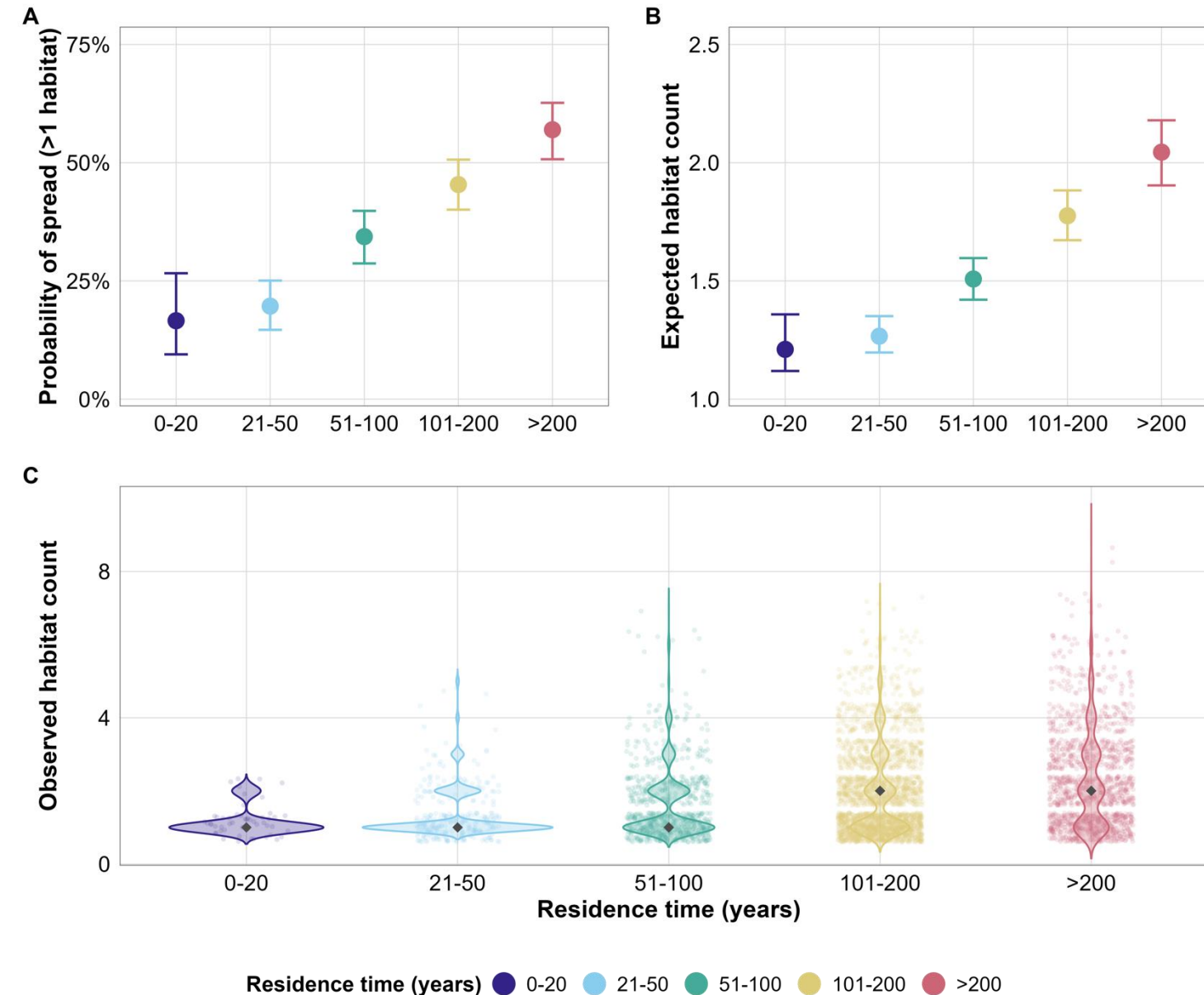
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# Results

- **Hurdle** component (P of > 1 habitat )
  - Trees, vines: reduced prob. (OR 0.57-0.69)
  - Long-lived perennials: reduced prob. (OR 0.55)
  - Human-mediated dispersal: reduced prob. (OR 0.73)

\*OR = Odds ratio

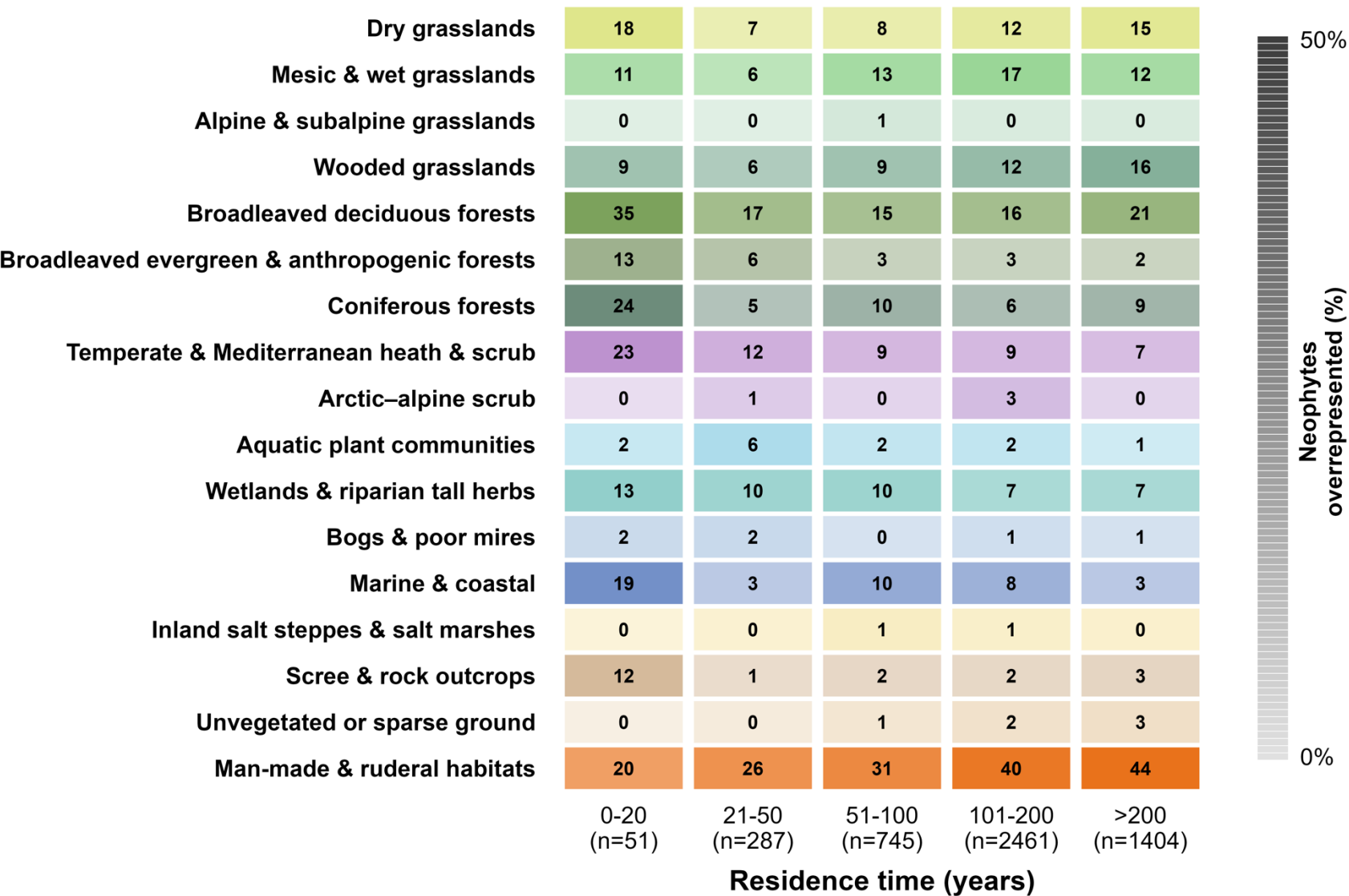


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- **Count** component (expected habitats given  $> 1$ )
  - Long-lived perennials: +43% more habitats (IRR 1.43)
  - Medium-lived perennials: +20% more habitats (IRR 1.20)

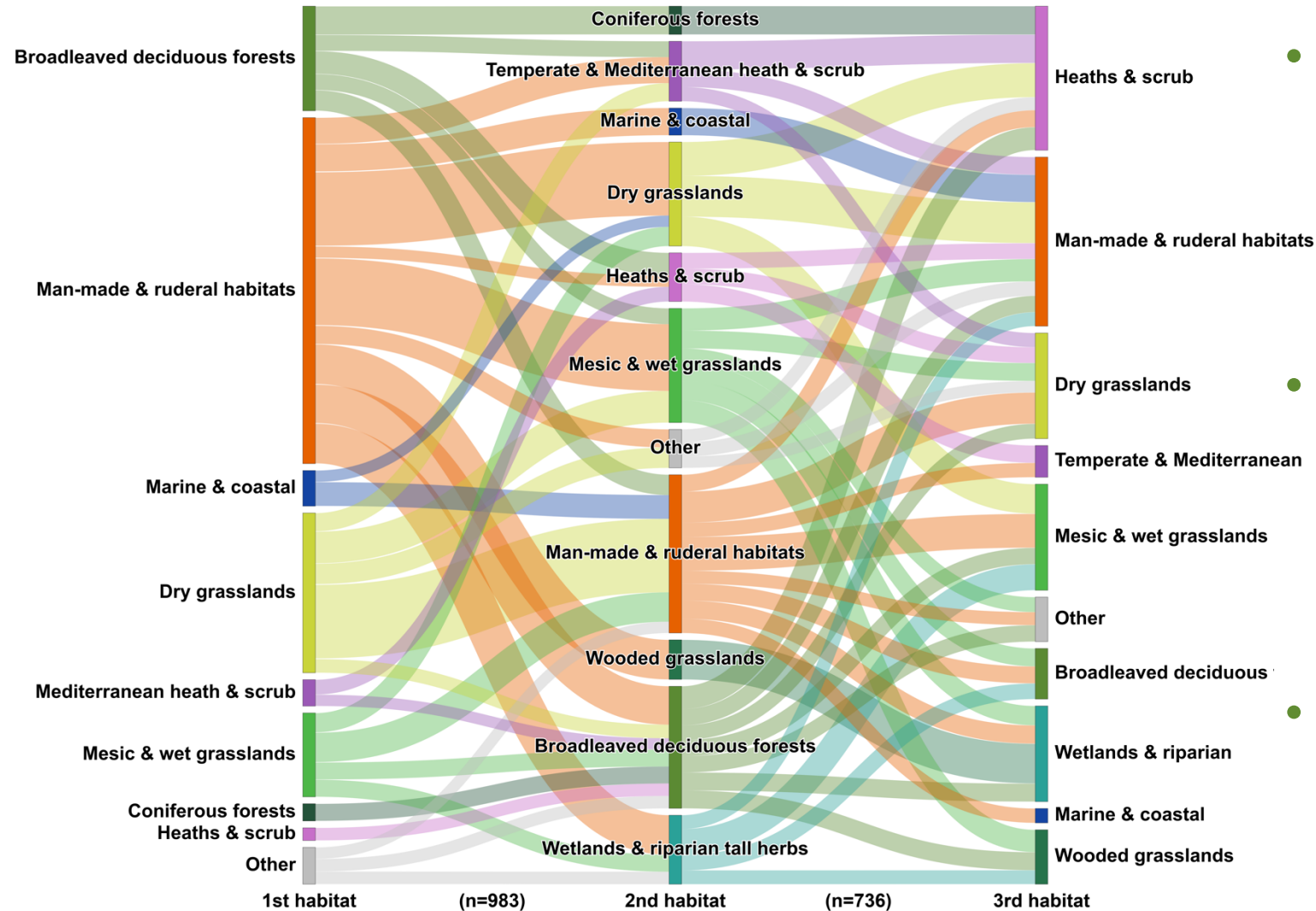
\*OR = Odds ratio, IRR = Incidence Rate Ratio

# Results



- **High Invasion:**
  - Man made & ruderal
- **High Resistance:**
  - Alpine grasslands
  - Bogs & poor mires
- **Intermediate**
  - Mesic grasslands
  - Deciduous forests

# Results



- **First habitat:**

- Man-made & ruderal (31%)
- Dry grasslands (16%)
- Broadleaved dec. (12%)

- **Common Establishment Patterns:**

- Dry grasslands ↔ man-made
- Man-made → mesic
- Man-made → wetlands

- **Third habitat**

- Mesic & wet grasslands = common tertiary site

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- **Transition analysis**
  - Continental aggregation, no local colonization pathways
  - Species are introduced independent across regions

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- **Ecological**
  - Habitat breadth grows with residence time → strong evidence of ongoing spread
- **Invasion debt**
  - The residence time effect continues up to >200 years → no sign of equilibrium

A low-angle, upward-looking photograph of a dense bamboo forest. The image is filled with numerous vertical bamboo stalks that create a strong sense of height and depth. Sunlight filters through the dense canopy of green leaves at the top, creating a dappled light effect on the bamboo stalks. The overall color palette is dominated by the dark greens of the bamboo and the bright highlights from the sun.

**Thank you**