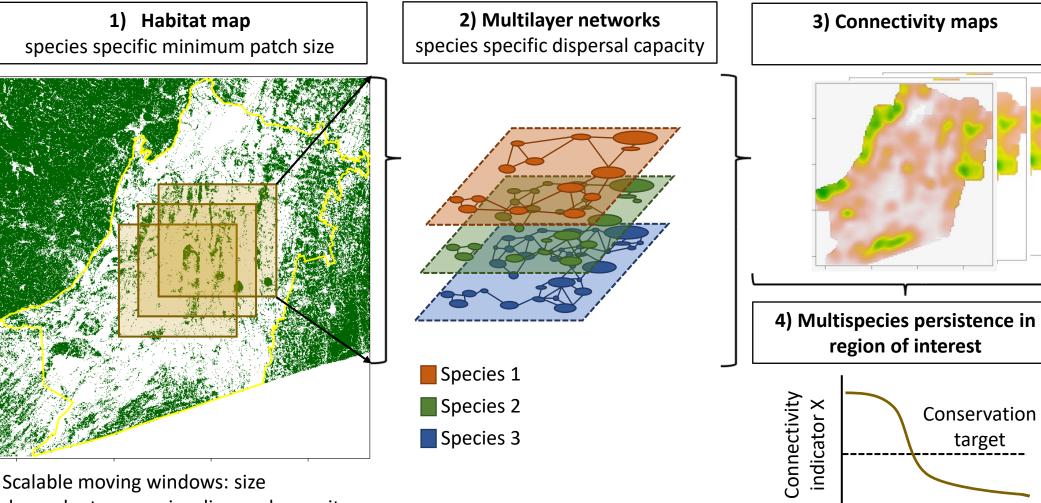
# Rapid evaluation of multispecies connectivity (REMC)

R-tool to efficiently compute multiple connectivity indicators for multiple species needs and across large regions of interest

> by Jacqueline Oehri

## Rapid evaluation of multispecies connectivity (REMC)



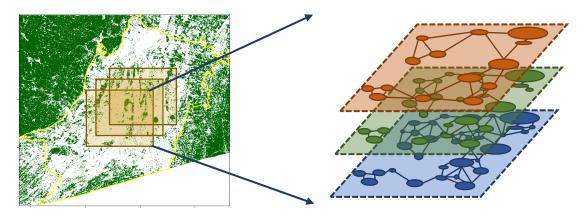
Species rank

dependent on species dispersal capacity

## **REMC R-tool core functions**

#### 1) REMC\_wrap()

- Implement moving windows, scales and resolution of interest
- Read inifile (a), apply connectivity functions in moving windows using REMC\_core (b)



#### 1a) REMC\_inifile.xlsx

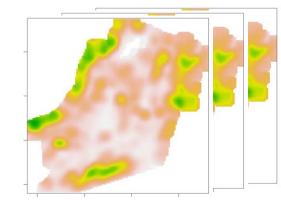
- define connectivity functions
- define species needs (habitat, dispersal)

## 1b) REMC\_core()

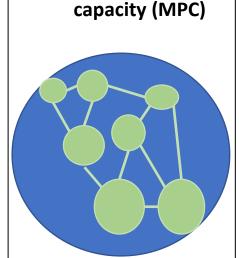
• Apply connectivity functions in moving windows

#### 2) REMC\_summary()

- Summarize moving window outputs into seamless maps
- At pixel, patch and landscape-level



## **Connectivity indicator – functions**



**Metapopulation** 

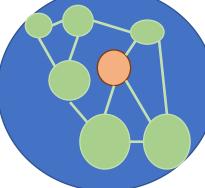
1.

Highlights:potential longterm species persistence. Metapopulation carrying capacity, based on area and connectance of habitat (Hanski & Ovaskainen 2000, Schnell et al. 2013). 2. Equivalent Connected Area (ECA)

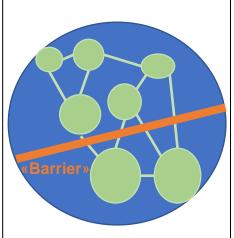
Highlights: size (area) of connected habitat. The size of a single habitat patch providing the same probability of connectivity than the actual habitat pattern in the landscape (Saura et al. 2011). **3. Fraction of connected** habitat (ECA<sub>AP</sub>, ECA<sub>AL</sub>)



<u>Highlights: underused</u> <u>connectivity potential.</u> Fraction of habitat that is connected: ECA divided by the total habitat area (AP) or landscape area (AL; cf. ProtConn index, Saura et al. 2017). 4. Betweenness centrality (BC), node degree (ND)



Highlights: stepping <u>stones.</u> BC (Brandes 2001): nr. of shortest paths between pairs of habitat patches passing through a focal patch. ND (Minor & Urban 2008): nr. of habitat patches connected to a focal patch. 5. Inverse cumulative resistance (invCR)



<u>Highlights: ease of</u> <u>landscape traversability.</u> Omnidirectional inverse cumulative resistance (Albert et al. 2017).

#### **Connectivity indicator – functions**

Metapopulation capacity (MPC)

Equivalent Connected Area (ECA)

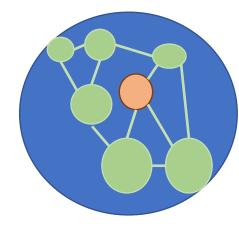
Highlights potential longterm species persistence. Highlights size (area) of connected habitat.

Fraction of connected habitat ( $ECA_{AP}$ ,  $ECA_{AL}$ )



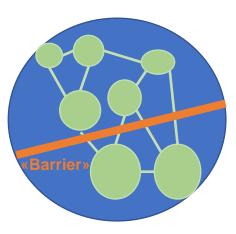
Highlights underused connectivity potential.

Betweenness centrality (BC), node degree (ND)



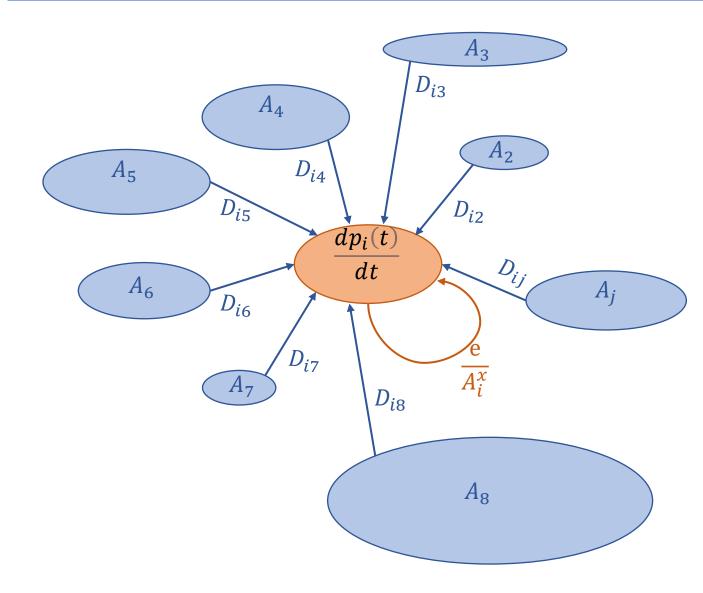
Highlights stepping stones.

Inverse cumulative resistance (invCR)



Highlights: ease of landscape traversability.

## Metapopulation capacity indicator – based functions



Occupancy-based, spatially explicit metapopulation model (SEM)

$$\frac{dp_i(t)}{dt} = C_i[1 - p_i(t)] - E_i p_i(t)$$
$$C_i = c \sum_{j \neq i} f(D_{ij}) A_j p_j(t)$$
$$E_i = \frac{e}{A_i^x}$$

Metapopulation capacity (MPC) as leading eigenvalue of 'landscape matrix' (M)

$$M_{ij} = \begin{cases} f(D_{ij})A_jA_i^x & i \neq j \\ A_jA_i^x & i = j \end{cases}$$
$$MPC = \lambda_M$$

## Neutral landscape models

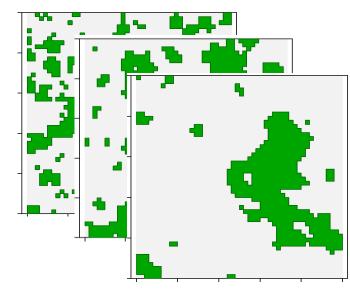
Generate simulated landscapes «libraries» covering a gradient in habitat amount & fragmentation (clumping)Functions for two types available:

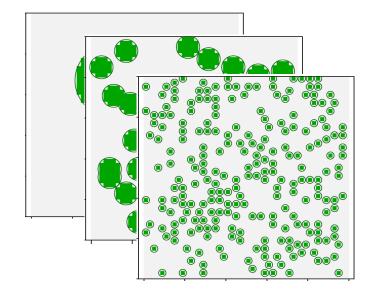
## **Random cluster**

Based on the algorithm by Saura & Martínez-Millán 2000. and the NLMR R-package (Sciaini et al. 2018)

## Simple-circle

Simple indication of habitat amount and number of patches.





## Introduction

Ecological connectivity: The 'unimpeded movement of species and the flow of natural processes that sustain life on Earth'

(Convention on Migratory Species, UN, 2019)

#### Important for

- Persistence of biodiversity
- Landscape ecosystem functioning & resilience
- Access to Nature's Contributions to People



Kendra Hoff, CLLC

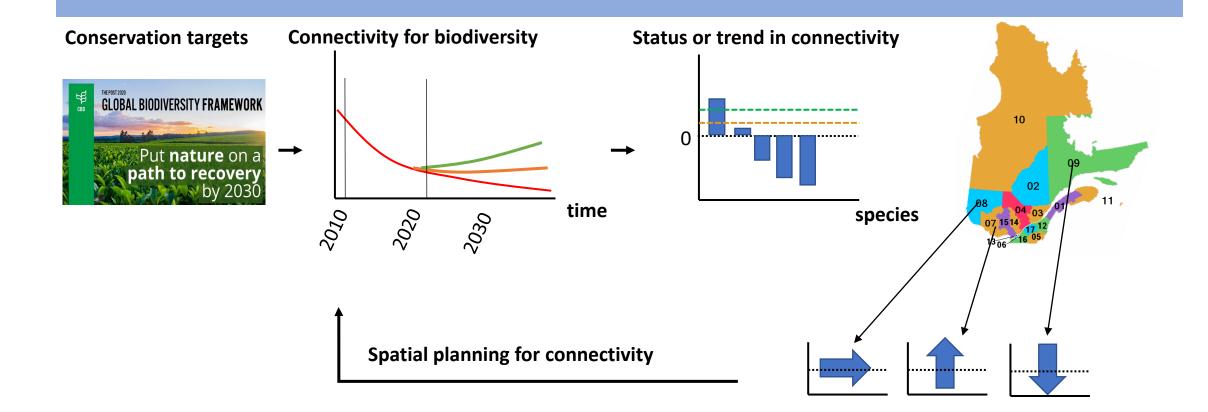
#### **Central to**

 the targets of the Kunming-Montreal Global Biodiversity Framework



## Motivation

#### Safeguarding connectivity for biodiversity (many species!)



Need for tools that enable <u>monitoring</u> of <u>multiple aspects of connectivity</u> for <u>multiple species</u> that are <u>efficient</u>, <u>scalable</u>, <u>validated</u> & that <u>allow evaluation</u> regarding conservation targets!

## Rapid assessment of multiple connectivity indicators for multiple species and large regions of interest

#### **REMC data input**

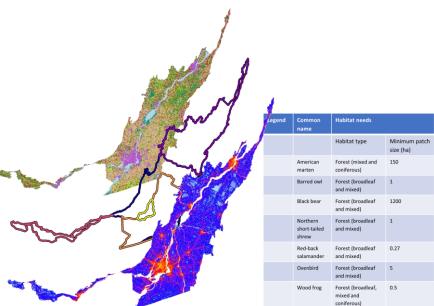
- 1. Land cover map
- 2. Species habitat needs and dispersal capacity
- 3. Region of interest (optional)
- 4. Resistance map (optional)

#### **REMC** settings

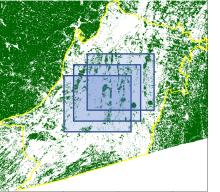
- 1. Moving window size
- 2. Spatial overlap
- 3. Connectivity functions
- 4. Landscape, patch and/or pixel level?

#### Runtime dependent on..

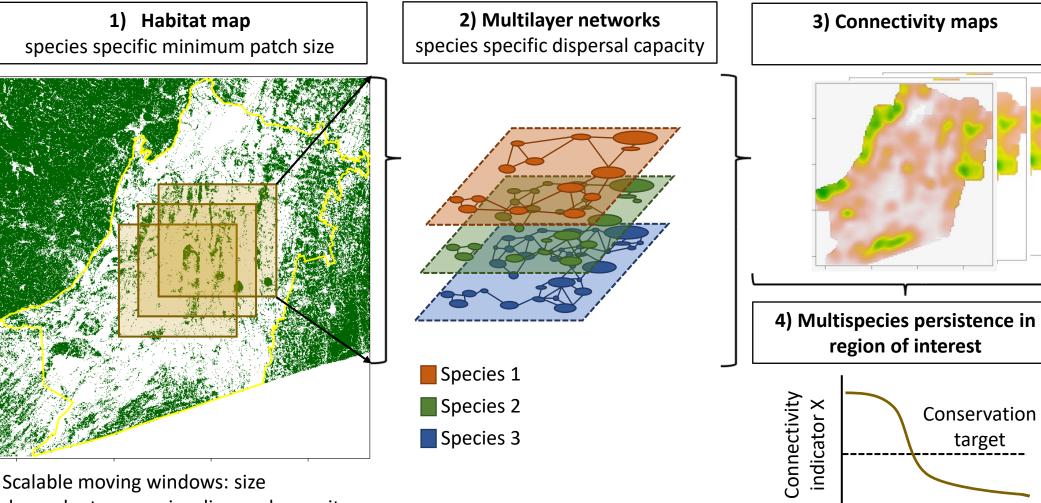
- Resolution & extent of Data input
- Number of species
  - Moving window size and overlap
  - Number and complexity of connectivity functions...



| Habitat needs                    |                            | Dispersal<br>capacity        |
|----------------------------------|----------------------------|------------------------------|
| Habitat type                     | Minimum patch<br>size (ha) | Gap crossing<br>distance (m) |
| Forest (mixed and<br>coniferous) | 150                        | 220                          |
| Forest (broadleaf<br>and mixed)  | 1                          | 209                          |
| Forest (broadleaf<br>and mixed)  | 1200                       | 236                          |
| Forest (broadleaf<br>and mixed)  | 1                          | 39                           |
| Forest (broadleaf<br>and mixed)  | 0.27                       | 10                           |
| Forest (broadleaf<br>and mixed)  | 5                          | 54                           |
| Forest (broadleaf,<br>mixed and  | 0.5                        | 39                           |



## Rapid evaluation of multispecies connectivity (REMC)



Species rank

dependent on species dispersal capacity

#### **REMC R-tool core functions**

