

Individual-based modelling in RangeShiftR

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https://damariszurell.github.io

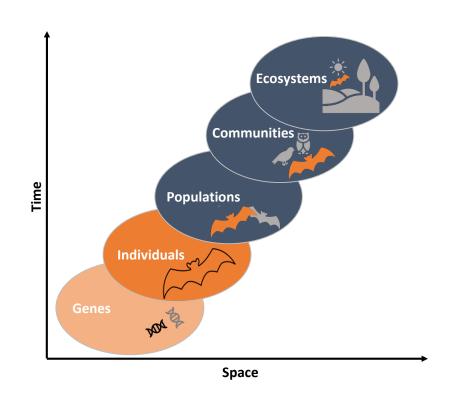




Ecology is hierarchical



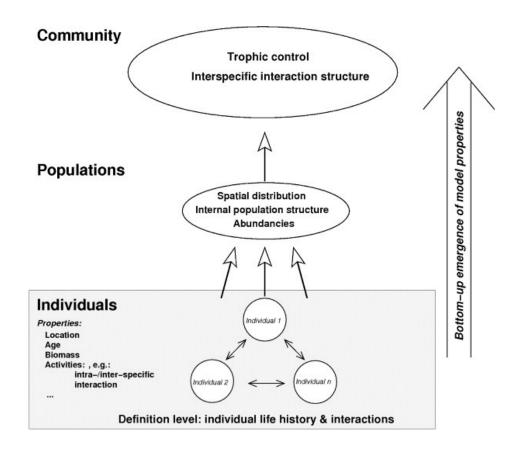
- Hierarchy of organisational levels from genes to ecosystems
- Each level operates and evolves on a characteristic spatial and temporal scale
- ➤ It is crucial to find the the appropriate modelling level for the given ecological problem or application



Emerging population dynamics

Universitär Torsdam

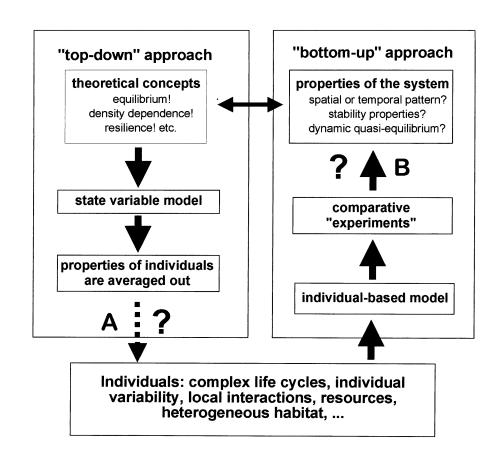
- "The whole is more than the sum of its parts." –
 Aristotle
- Bottom-up approach: use knowledge about behavior of single individuals to infer properties of the larger population or community



Individual-based models (IBM)



- Individuals/agents as unique discrete entity
 - Characterised by at least one state/property (additional to age) that changes through time
- Allow to account for individual variability, local interactions, individual space use, behaviour ...



IBM state variables and processes



Population

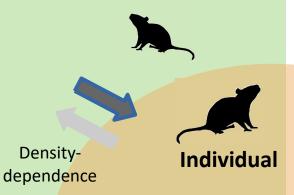
Growth rate, demographic structure, range dynamics, ...

Environment



Space is represented as a regular 2D grid, where each cell is characterised by:

- Habitat suitability index / Carrying capacity
- Land cover proportions
- Strength of abiotic drivers









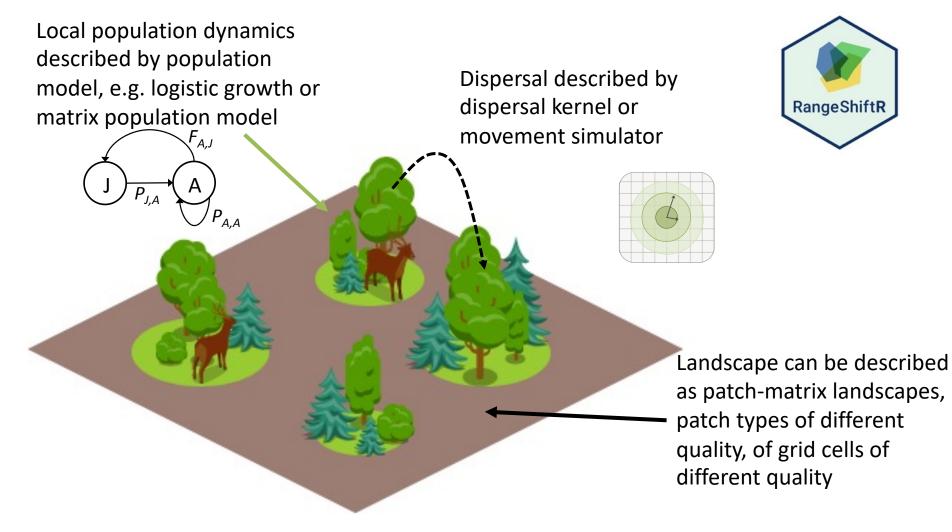


- State: sex, age/stage, position, traits, genes, etc.
 - can vary among individuals and/or across time
- **Behaviour/Processes**: reproduction, aging, maturing/developing, dispersal, mortality
 - can be dependent on current state and environment

RangeShiftR



Simulating local population dynamics and dispersal



RangeShiftR – local population dynamics



- Option 1: Non-overlapping generations and no stage structure
 - Discrete generations
 - In each generation life cycle: reproduction, death of adults, offspring dispersal
 - Described by discrete form of logistic growth

$$Poisson\left(\frac{R_{i,t}}{1+|R_{i,t}-1|*\left(\frac{N_{i,t}}{K_{i,t}}\right)^{b_c}}\right)$$

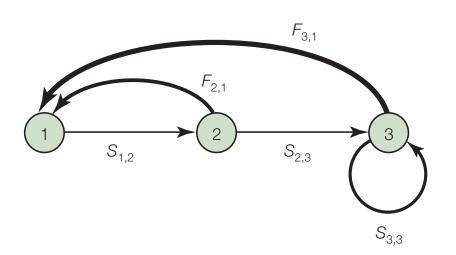
 Important: individual-based formulation, meaning the number of offspring produced by adult is drawn from random distribution

RangeShiftR – local population dynamics



- Option 2: overlapping generations and stage structure
 - Parameters provided through classical transition matrices (but applied stochastically)
 - Transition matrices can be age-structured (Leslie matrix) or stage-structured (Lefkovitch matrix)

Life-cycle graph:



Stages:

- 1 Juveniles / first-year birds do not breed
- Second-year birds are inexperienced breeders
- 3 Mature adults are experience breeders

Transition rates:

- $F_{i,1}$ Fecundity
- $S_{i,j}$ Development rate (transition to next stage)
- $S_{i,i}$ Survival probability

RangeShiftR – local population dynamics

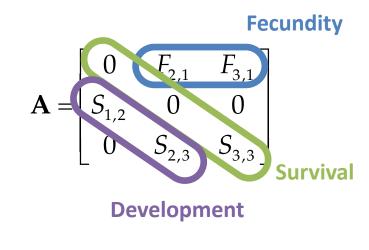


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Life-cycle graph:

 $F_{3,1}$ $F_{2,1}$ $S_{1,2}$ $S_{2,3}$ $S_{3,3}$

(Lefkovitch) Transition matrix:



Dispersal



- Natal/juvenile dispersal: spatial process by which individuals move away from birth location
 - avoid kin competition, exploit new resources
- Critical for range expansion and spatial gene flow
- Associated with various costs and trade-offs that determine the eventual displacement distance
 - has impacts on both population dynamics and genetics

Dispersal



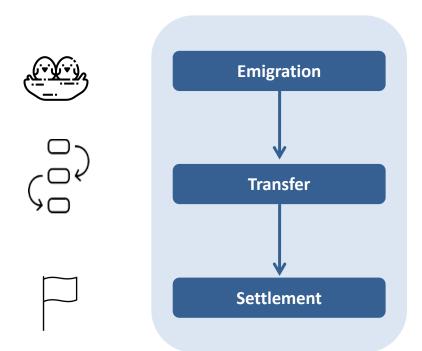


Fig. 1. Dispersing individuals are a nonrandom subset of the population.

(**Left**) Dispersing females of the Glanville fritillary, Melitaea cinxia, have a higher flight metabolic rate and are more fecund than sedentary ones.

(**Right**) In Siberian jays, Perisoreus infaustus, the subordinate individuals disperse, whereas the heavier and more dominant remain in their natal territories.





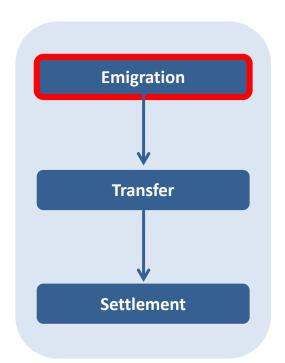
Leaving the natal site (area where born)

Movement through the landscape

Establish in a new habitat site /area



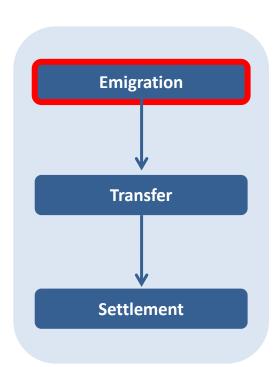




Leaving the natal site (area where born)

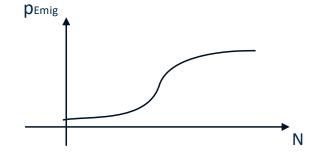






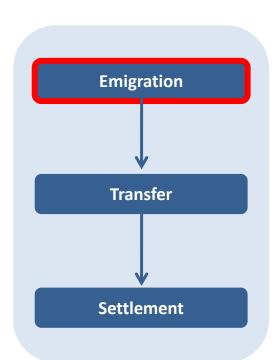
Emigration probability:

- the probability that an individual will leave its natal patch during the present year
- > can be density-dependent



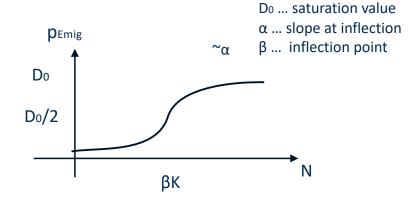






Emigration probability:

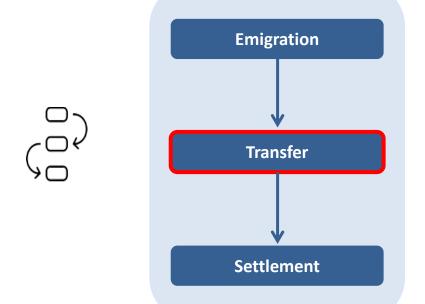
- the probability that an individual will leave its natal patch during the present year
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stage	sex	D ₀	α	β
0	0	0.4	10	1
0	1	0.9	10	1

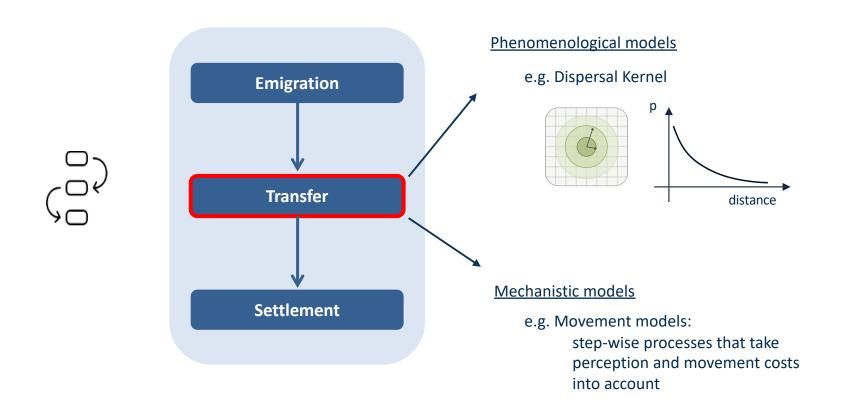
sex = 0 ... female sex = 1 ... male





Movement through the landscape







1.) Stochastic Movement Simulator

step-wise process

3	4	1	0	0
1	1	3	3	5
2	3	1	1	4
1	1	0	0	0
5	4	1	0	0

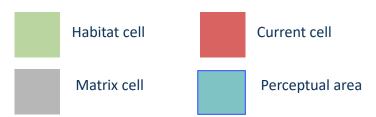




1.) Stochastic Movement Simulator

- step-wise process
- takes perception into account
 - perceptual range (e.g. 1 cell)

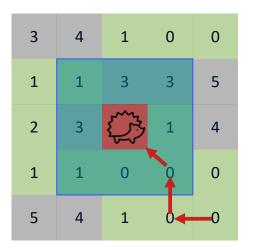
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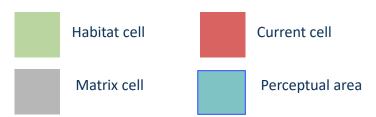




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- decision of next direction based on:
 - lowest average cost
 - preceeding steps in memory (e.g. 3 steps)
 - directional persistence

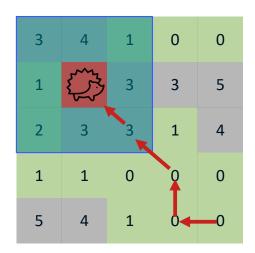


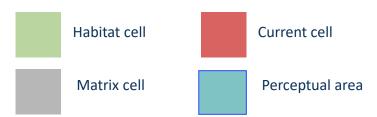




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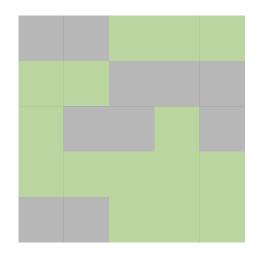




2.) Correlated Random Walk

- step-wise process
- considers continuous space, not grid cells

no consideration of costs

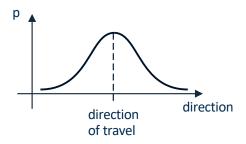




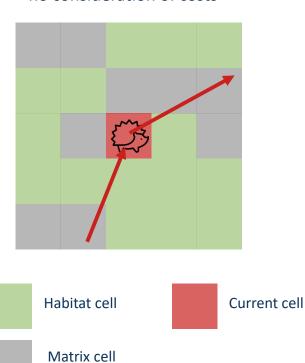


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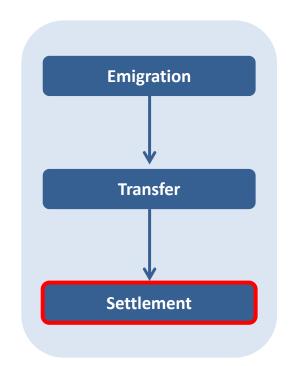
- step-wise process
- considers continuous space, not grid cells
- draws direction from a distribution:



no consideration of costs

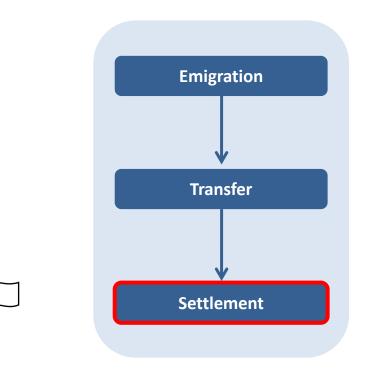






Establish in a new habitat site /area





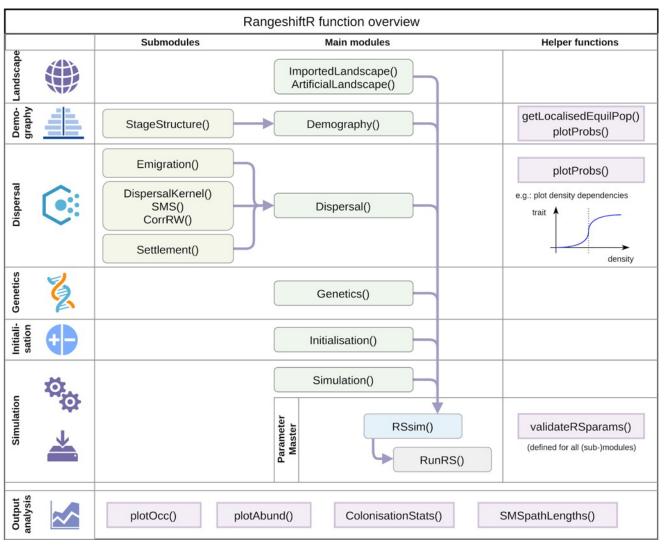
Settlement probability:

- the probability that an individual will stay in a new habitat patch
- can be density-dependent (analogous to emigration probability)

RangeShiftR overview









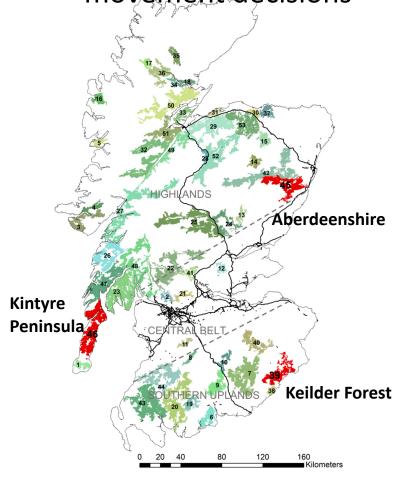
RangeShiftR case study

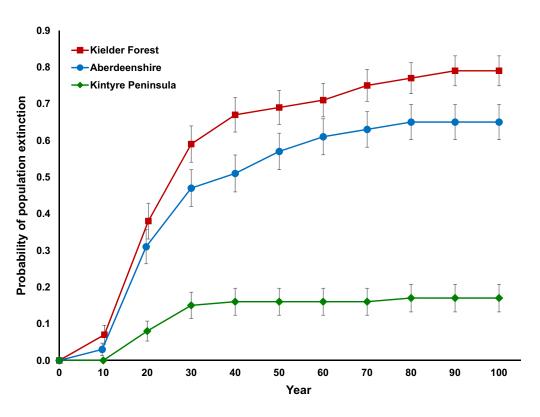


Example: Eurasian lynx, reintroduction Scotland

Stage-structured model, individual-based movement decisions







Thank you for your interest



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