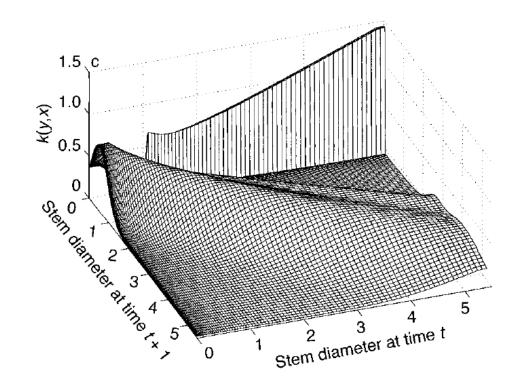
# Basics of plant population modelling and its application





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### Programme

Monday: Matrix models

Tuesday: Integral Projection Models: construction
Plus first paper discussion

Wednesday: Integral Projection Models: output

Plus: second paper discussion

Thursday: Integral Projection Models: more applications

Plus: preparing presentations

Friday: **Presentations** 

## Programme

## Thursday February 13<sup>th</sup>: Integral Projection Models: output & applications

9-9.30 Lecture: IPM output & applications

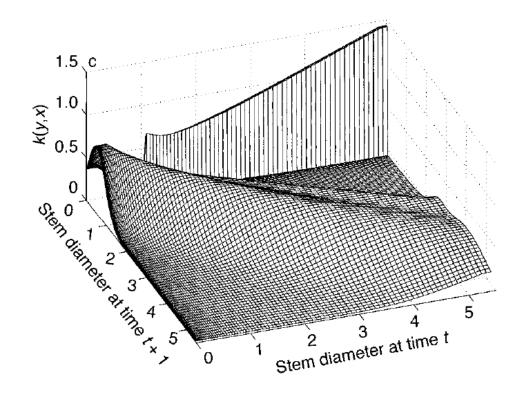
9.30-12 Exercises: IPM output & applications

Lunch

2-5 Work on your own data & prepare your presentation

## Lecture: Running & applying IPMs

- My tips for IPMs
- Rpadrino package



- An interesting question
  - On population viability
  - On population expansion (invasive species)
  - On harvesting effects
  - On climate/environmental change

- The question determines:
  - What you measure and study in the field: where, how long, what classes, how many plots
  - What stages your IPM includes: seeds, seedlings
  - What structure your IPM has: time varying, habitat differences, treatments, landscape level?

#### Sound data

- Make sure you sample sufficient individuals (minimum ~ 200; ideally 500)
- Ensure sufficient coverage of all classes
- Use statistical logic to determine sampling and experimental design
- Pool data across habitats, treatments, etc.
- All vital rates: survival, growth, reproduction & recruitment
- From permanent sample plots, tree-ring analysis, mark-recapture
- From individuals outside plots that you remeasure

- Sound statistical analyses
  - Use multiple regressions to include effects of both habitat and size
  - Only include fixed effects that you would like to include in your IPM: habitats, years
  - Use mixed effect models if you're interested in variability across years (or plots). In ipmr you can add random year factors to produce stochastic models
  - Some vital rates will not respond to treatment/habitat/etc. In that case they need to be the same ascross your different treatment/habitat/year IPMs

- Vital rate regressions: survival
  - Use logistic regressions (glm binomial): survival ~
     size or

```
survival ~size + size^2
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- Add habitat or year if applicable
- If not significant, then survival ~ 1

- Vital rate regressions: growth
  - For organisms with large size changes compared to maximum size, you can use size\_next ~ size
  - For long-lived organisms with small size changes compared to maximum growth, better to use size\_next ~ size + growth\_function
  - Growth function can be linear (a\* size) or more complex (a\*size + b\*size^2) or Hossefeld or something else
  - In exercises we used curve fitting for this (nls function)

- Vital rate regressions: sexual reproduction
  - Most complex
  - You need at least: (1) info on which individuals are reproduction, (2) the number of new recruits per reproductive individuals.
  - But can also be: (1) repro probability, (2) number of inflorescences or seeds produced per repro individual, (3) number of new recruits per inflorescence or per seed
  - Make sure that the recruits are expressed per unit at which you have information about the reproduction.

- Vital rate regressions: clonal reproduction
  - You need: (1) info on which individuals are producing new shoots, (2) the number of new shoots per clonally reproducing individual.
  - Add a new kernel in ipmr, see example code

#### IPM construction

- Build your kernel carefully, step by step and check the transition rates and lambda values carefully
- Start simple to test your model

#### Mesh size checks

- Mesh size influence lambda, ages & growth variability
- Make cross-cuts in IPM to ensure growth variability is described by at least 5 points
- Check effect of mesh size on lambda
- Check effect of mesh size on age estimates

- IPMs for multiple treatments/habitats
  - build one IPM for all of them together, based on your statistical models
  - Extract each habitat-specific IPM transition
     matrices and conduct further analyses with these
  - In this way, you have used the strength of all data to build your IPMs and you can use the separate IPMs

- Example code!
  - Please feel free to use or share the code we used in the course
  - This is yours and you can use it and adapt it

- Assistance?
  - Feel free to ask me for help, involve me in a study,
     visit us in Wageningen
  - I'm happy to help an collaborate