

# (Spatial) Ecology and Evolution: Integrating Theory and Data

Otso Ovaskainen

## OVERVIEW

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- . L1. Approaches to ecological modelling
- . L2. Model parameterization and validation (stat)
- . L3. Stochastic models of population dynamics (math)
- . L4. Animal movement (math + stat)
- . L5. Quantitative population genetics (math + stat)
- . L6. Community ecology (stat)

# mathematical biology group

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movements, populations, communities, genetics, evolution, bioinformatics

# L1. Approaches to Ecological Modelling

## OVERVIEW

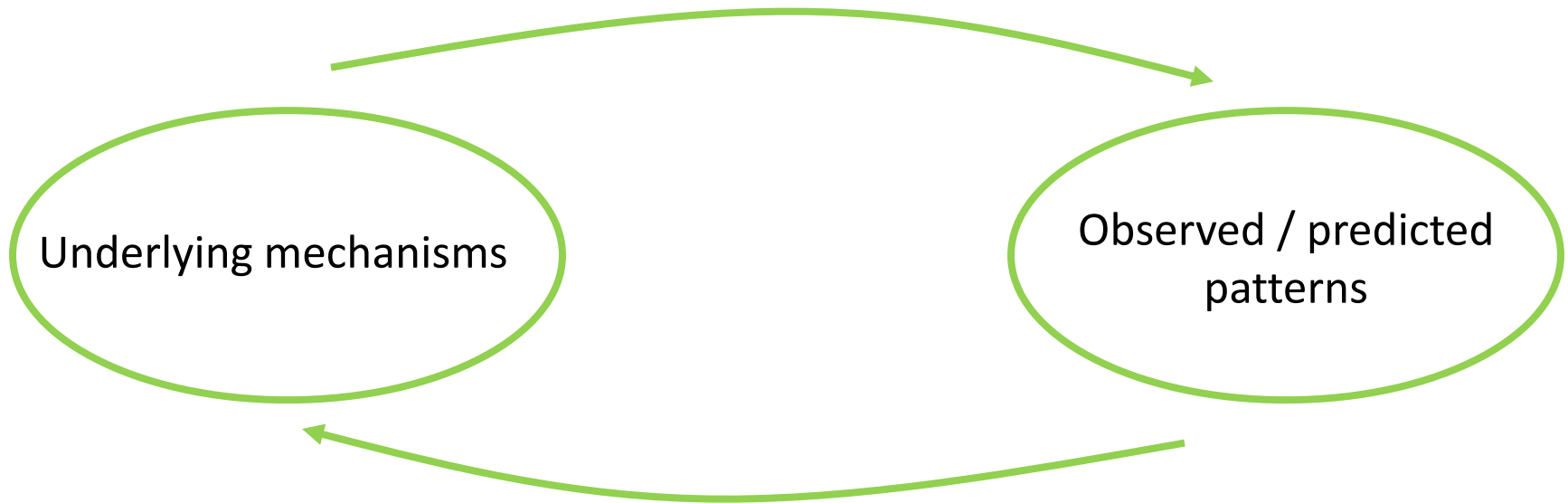
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# Approaches to ecological modelling

**The forward approach: mathematical (mechanistic) modelling.**

Aim: to understand causal relationships at the general level

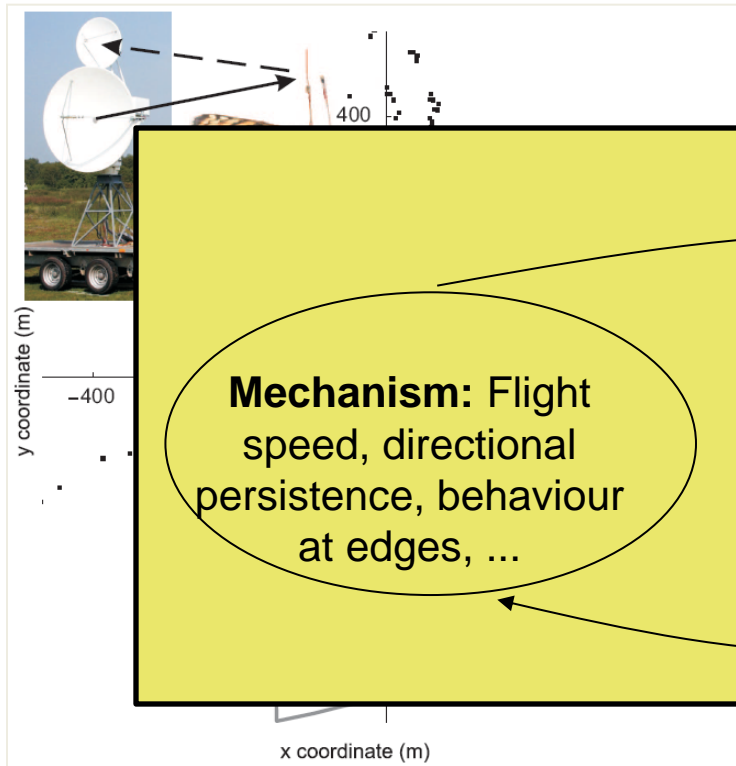


**The inverse approach: statistical (phenomenological) modelling.**

Aim: to find out factors shaping empirical data

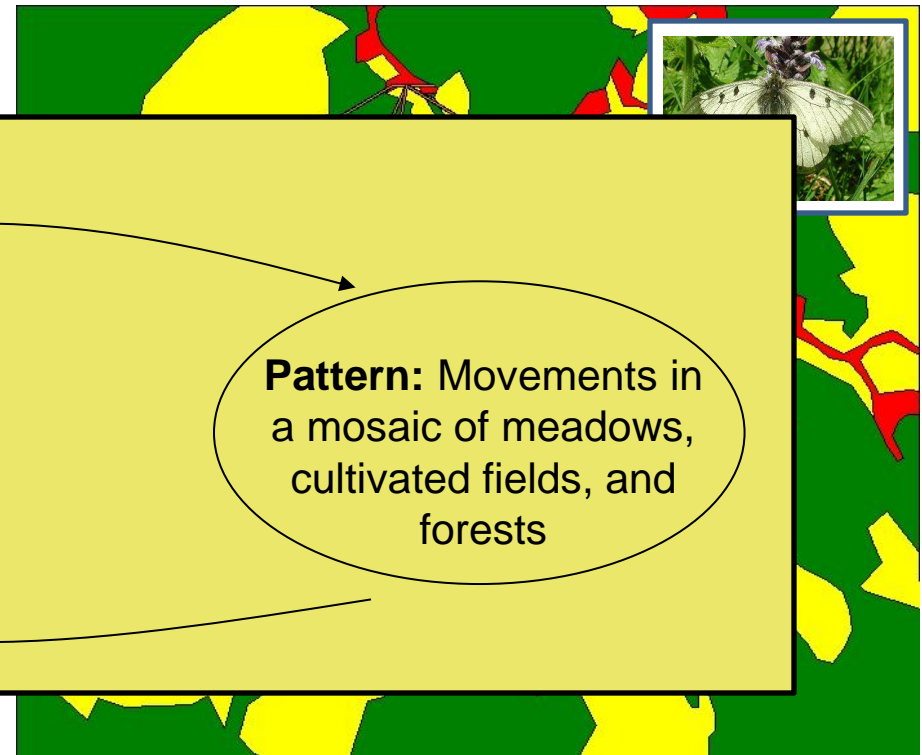
# Empirical approaches for studying butterfly movements

## Harmonic radar



**Mechanism:** Flight speed, directional persistence, behaviour at edges, ...

## Spatial mark-recapture

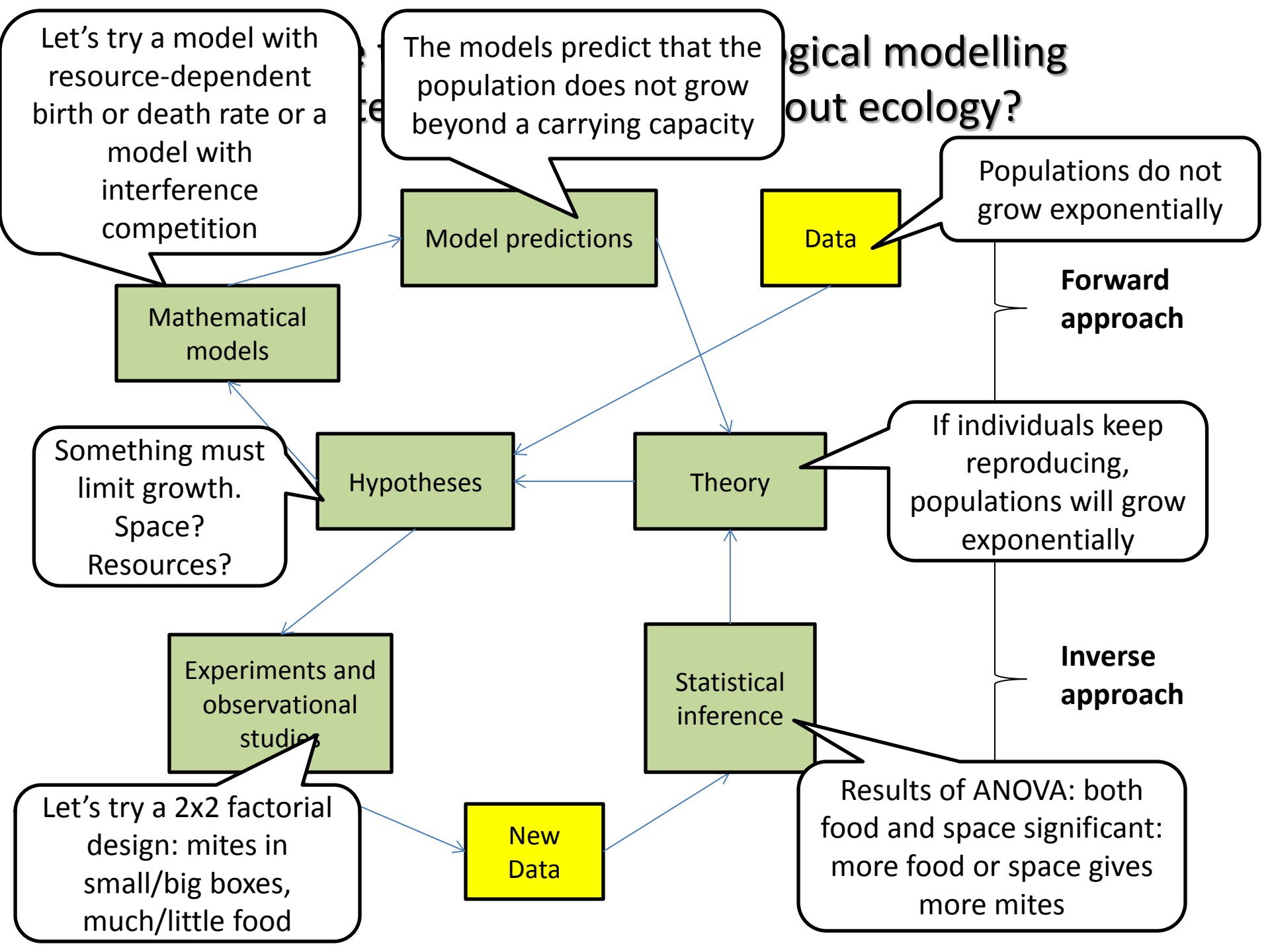


**Pattern:** Movements in a mosaic of meadows, cultivated fields, and forests

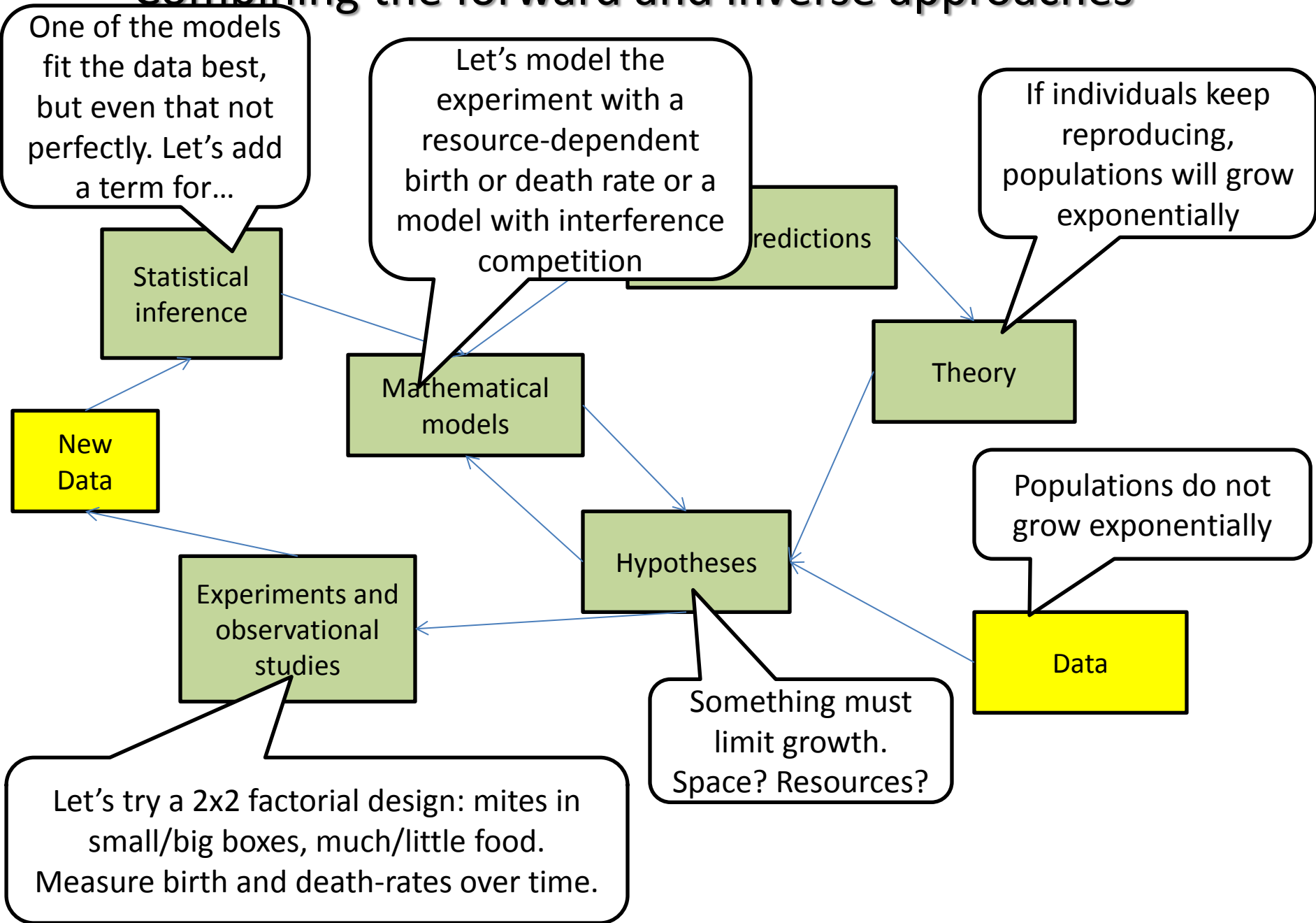
Ovaskainen, O. et al. 2008. Tracking butterfly movements with harmonic radar reveals an effect of population age on movement distance. *PNAS* **105**, 19090-19095.

Ovaskainen, O. et al. 2008. An empirical test of a diffusion model: predicting clouded apollo movements in a novel environment. *American Naturalist* **171**, 610-619.

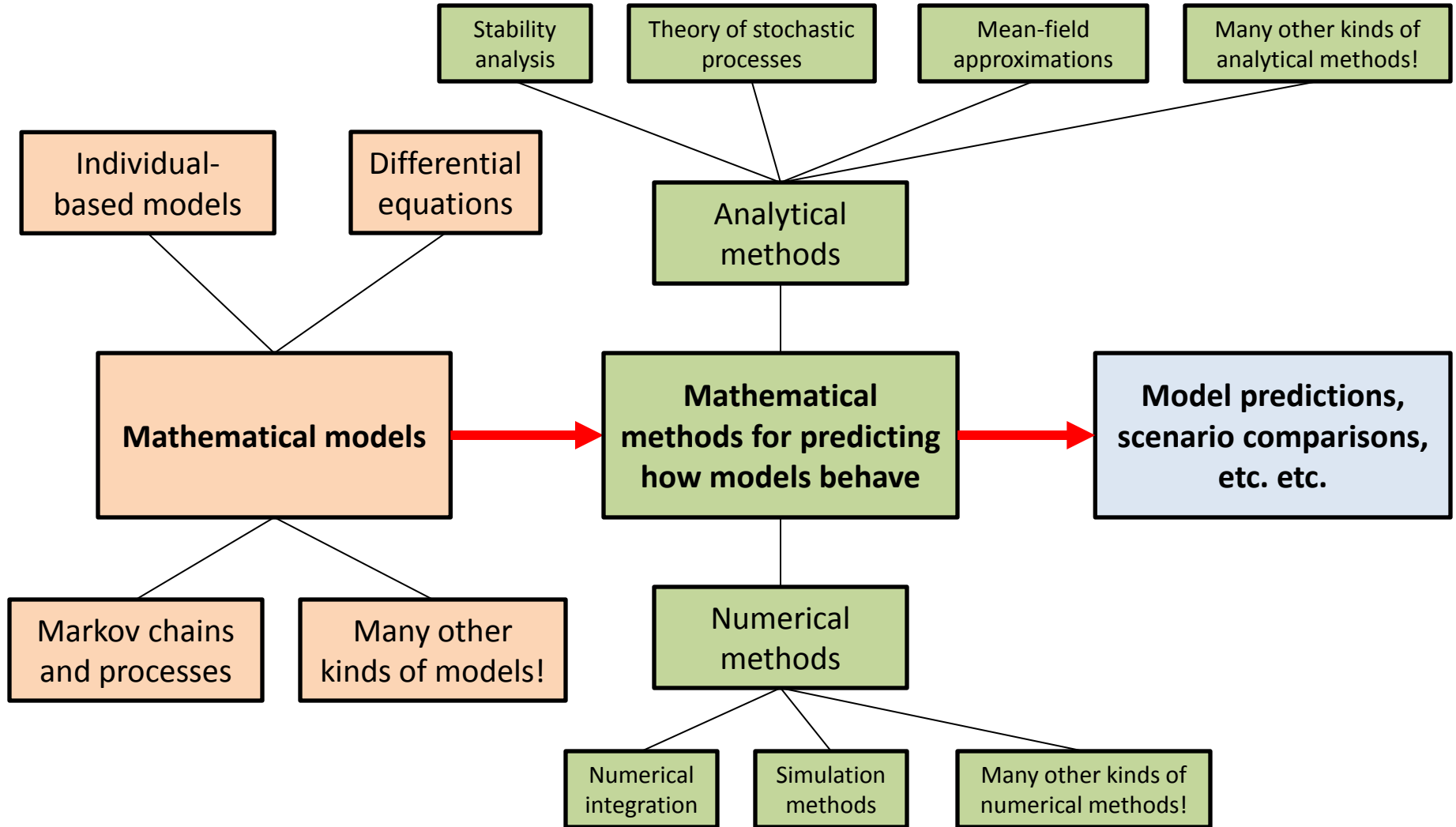
# Biological modelling out ecology?



# Combining the forward and inverse approaches



# Mathematical models and methods in ecological modelling

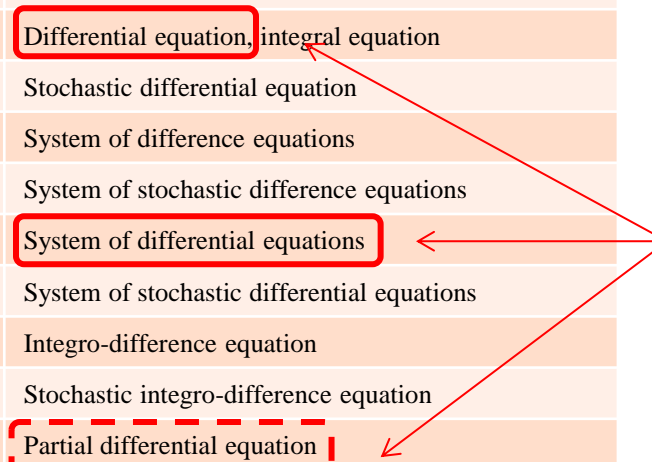




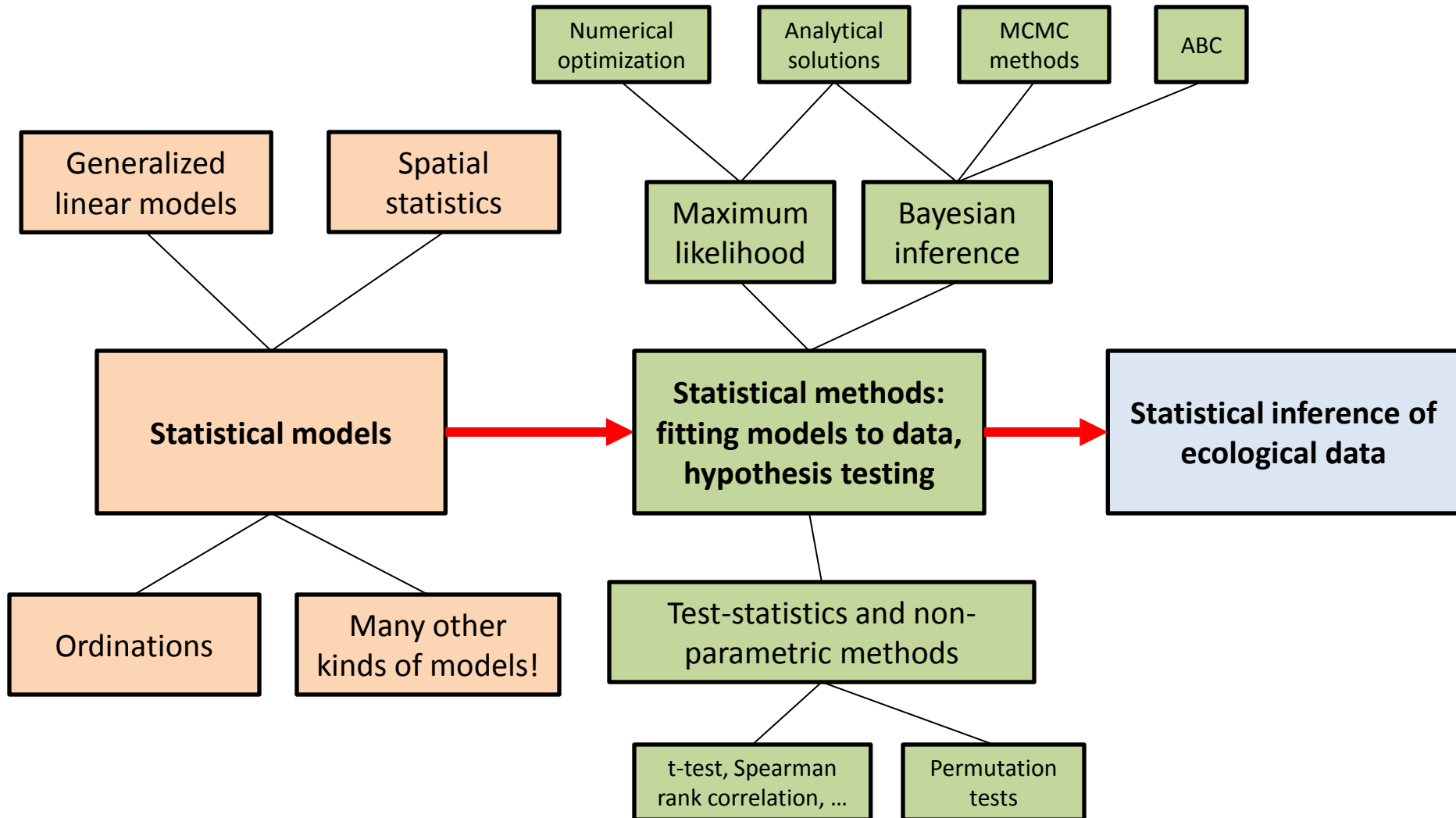
# There are many kinds of mathematical models!

Variables	Structure (e.g. space)	Time	Stochasticity	Model type(s)
discrete	no	discrete	no	-
			yes	Markov chain
		continuous	no	-
			yes	Markov process
	discrete	discrete	no	-
			yes	IBM on grid or patch network, SPOM
		continuous	no	-
			yes	IBM on a grid or patch network, SPOM
	continuous	discrete	no	-
			yes	IBM in continuous space
		continuous	no	-
			yes	Spatio-temporal point process
continuous	no	discrete	no	Difference equation
			yes	Stochastic difference equation
		continuous	no	Differential equation, integral equation
			yes	Stochastic differential equation
	discrete	discrete	no	System of difference equations
			yes	System of stochastic difference equations
		continuous	no	System of differential equations
			yes	System of stochastic differential equations
	continuous	discrete	no	Integro-difference equation
			yes	Stochastic integro-difference equation
		continuous	no	Partial differential equation
			yes	Stochastic partial differential equation

This course



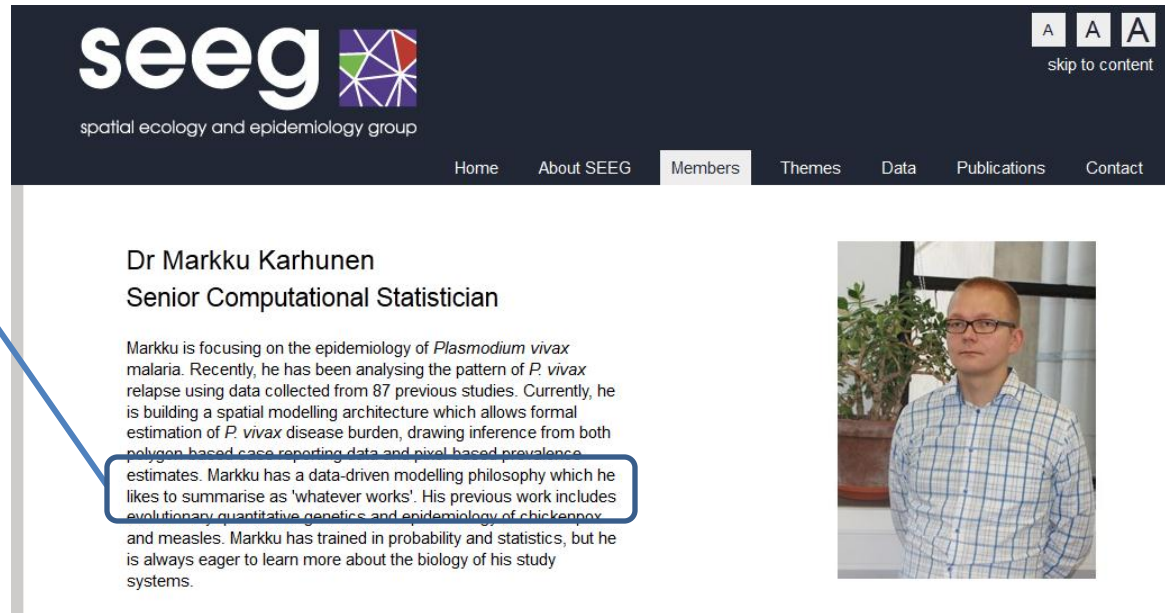
# Statistical models and methods in ecological modelling



# L1: take home messages

- There are many approaches to ecological modelling!
- Think critically why you play with mathematical models! Just because you can (and you like it), or because that helps to learn about ecology?
- Find your own modelling philosophy!

Markku has a data-driven modelling philosophy which he likes to summarise as 'whatever works'.



The screenshot shows the SEEG (spatial ecology and epidemiology group) website. The header includes the SEEG logo and navigation links: Home, About SEEG, Members, Themes, Data, Publications, and Contact. The main content area features a profile for Dr Markku Karhunen, Senior Computational Statistician. A blue callout box highlights a specific sentence in his bio: "Markku has a data-driven modelling philosophy which he likes to summarise as 'whatever works'." A photograph of Dr. Karhunen is visible on the right side of the page.

**seeg** spatial ecology and epidemiology group

Home About SEEG **Members** Themes Data Publications Contact

**Dr Markku Karhunen**  
Senior Computational Statistician

Markku is focusing on the epidemiology of *Plasmodium vivax* malaria. Recently, he has been analysing the pattern of *P. vivax* relapse using data collected from 87 previous studies. Currently, he is building a spatial modelling architecture which allows formal estimation of *P. vivax* disease burden, drawing inference from both polygon-based case reporting data and pixel-based prevalence estimates. Markku has a data-driven modelling philosophy which he likes to summarise as 'whatever works'. His previous work includes evolutionary quantitative genetics and epidemiology of chickenpox and measles. Markku has trained in probability and statistics, but he is always eager to learn more about the biology of his study systems.