

*Barbara Vuillaume*

Geeklunch, Wednesday 26 June 2020



# Introduction to IPM

# First things to know about IPM

- IPM = Integrated Population Model
- Use in studies focusing on population dynamics
  - Population dynamics = changes in abundance over time and demographic causes of these changes
- Strength : combine different available data sets to get deeper insights into population dynamics and better estimates of the demographic traits

$$N_{t+1} = N_t \times g(s, f)$$

*t = year, s = survival, f = productivity*

## 3 main benefits from this method

- IPM includes information about demographic rates both from
  - Explicit data ( e.g. mark recoveries for survival)
  - Data on population size

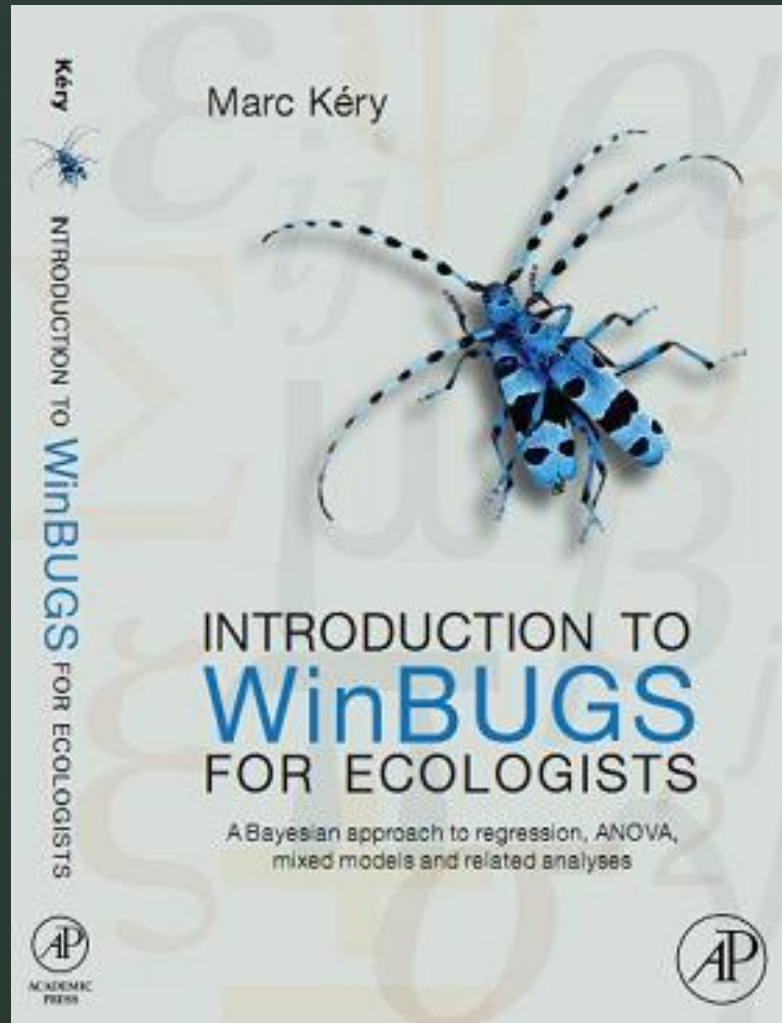
=> estimation of demographic rates with increased precision which will improve population size estimates
- Case of missing data : demographic rate can be estimated from data on population size
- Simultaneous study of demographic rates and population size allow a comprehensive assessment of the cause of population changes

# Development of an IPM : 3 theoretical steps

- 1st: develop a population model that links the demographic rates with changes in population size
- 2<sup>nd</sup>: write the likelihood of all data sets available  
Counts, Survival, Productivity, Immigration, ...
- 3<sup>rd</sup>: construct the joint likelihood of the complete model



# Want to start ?



<https://drive.google.com/file/d/14JTU0kFGgHpkMUXs6B780npSlrJ3OogG/view?usp=sharing>

Marc Kéry  
Michael Schaub



## BAYESIAN POPULATION ANALYSIS USING **WinBUGS**

A hierarchical perspective



## Step 1: Define the link between changes in population size and demographic rates



- $E$  = expected number

$$E(N_{1,t+1} \mid N_{1,t}, N_{ad,t}) = N_{1,t} * S_{juv,t} * f_t + N_{ad,t} * S_{juv,t} * f$$

$$E(N_{ad,t+1} \mid N_{ad,t}) = N_{1,t} * S_{ad,t} + N_{ad,t} * S_{ad,t}$$

## Step 2: Define the likelihoods of each Individual Data Set

### □ Population counts

Sources of uncertainty: we include demographic stochasticity using appropriate distributions to describe the number of individuals at year  $t+1$

$$N_{1,t+1} \sim \text{Poisson}( N_{1,t} * S_{\text{juv},t} * f_t + N_{\text{ad},t} * S_{\text{juv},t} * f )$$

$$N_{\text{ad},t+1} \sim \text{Binomial}( N_{1,t} + N_{\text{ad},t} , S_{\text{ad},t} )$$

### □ Capture-Recapture data

Modelisation of survival, matrix of capture-recapture

### □ Reproductive success

e.g. birds survey: number of nestlings, number of surveyed broods, productivity

## Step 2: Define the likelihoods of each Individual Data Set

# 3.3. Likelihood for productivity data: Poisson regression

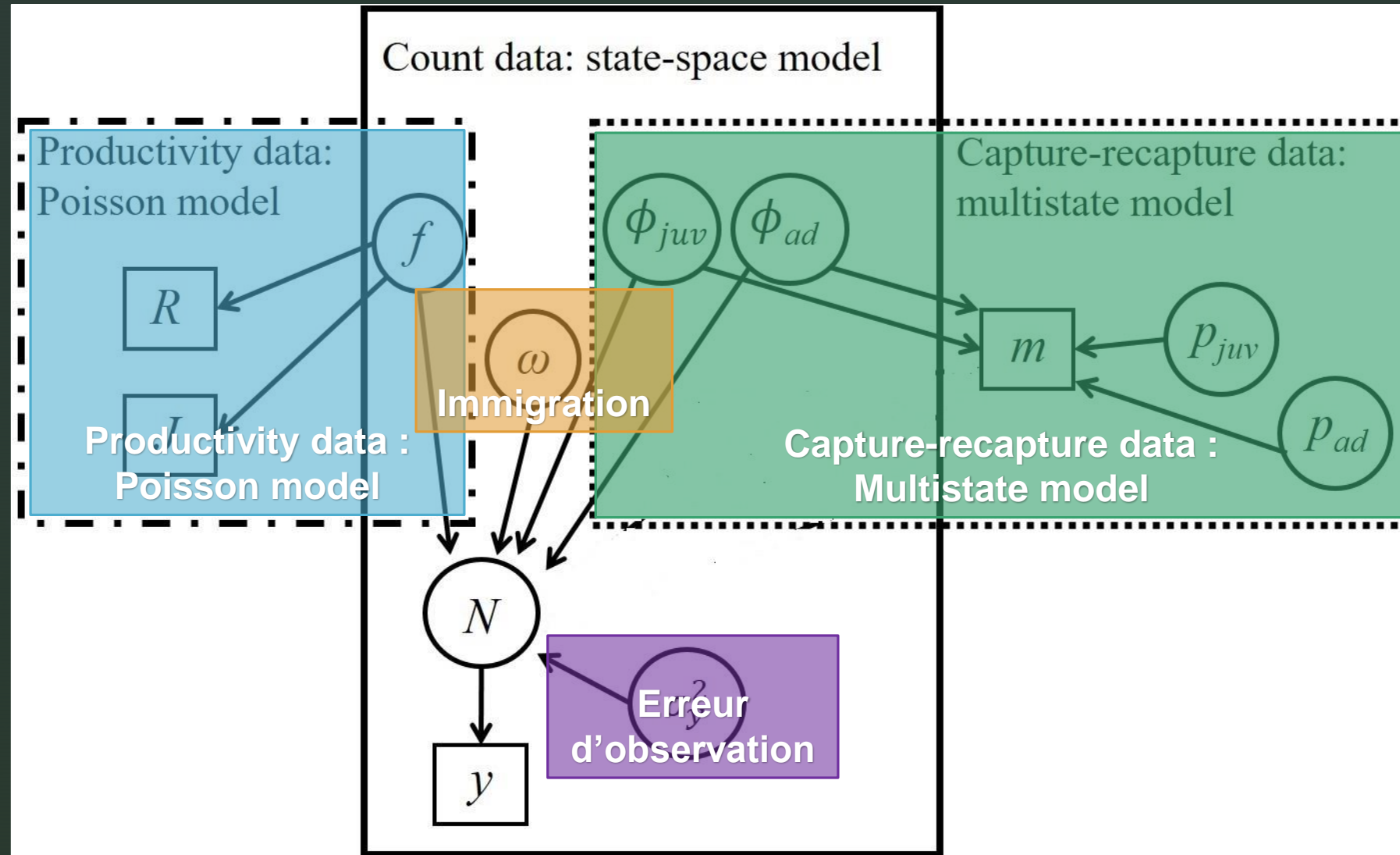
```
for (t in 1: (nyears - 1)) {  
  J[t] ~ dpois ( rho[t] )  
  rho[t] <- R[t] * f[t]  
}  
}
```

### ■ Reproductive success

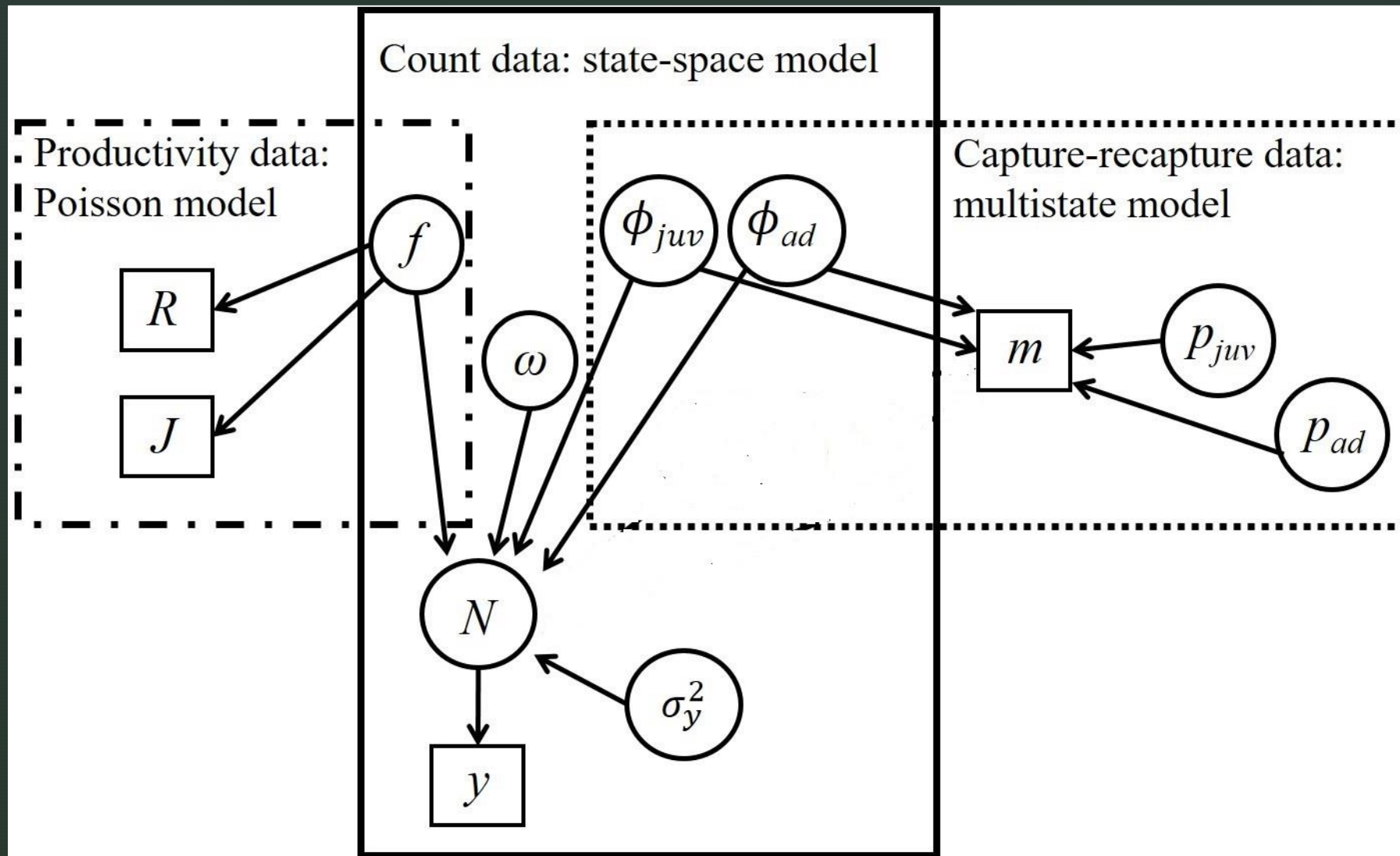
e.g. birds survey: number of nestlings, number of surveyed broods, productivity



## Step 3: Formulate the joint likelihood



## Step 3: Formulate the joint likelihood



# Example with a simple IPM including counts, capture-recapture, reproduction

- Need for WinBugs or Jags (also for Linux) !!!!
  - <https://www.mrc-bsu.cam.ac.uk/software/bugs/the-bugs-project-winbugs/>
  - <https://sourceforge.net/projects/mcmc-jags/>
- Use R instead of R Studio ! Too many errors !



# Real data example: Hoopoe population dynamics

