

Neural Dynamic N-mixture Model

A deep learning framework for inferring demographic rates from count data

Speaker:

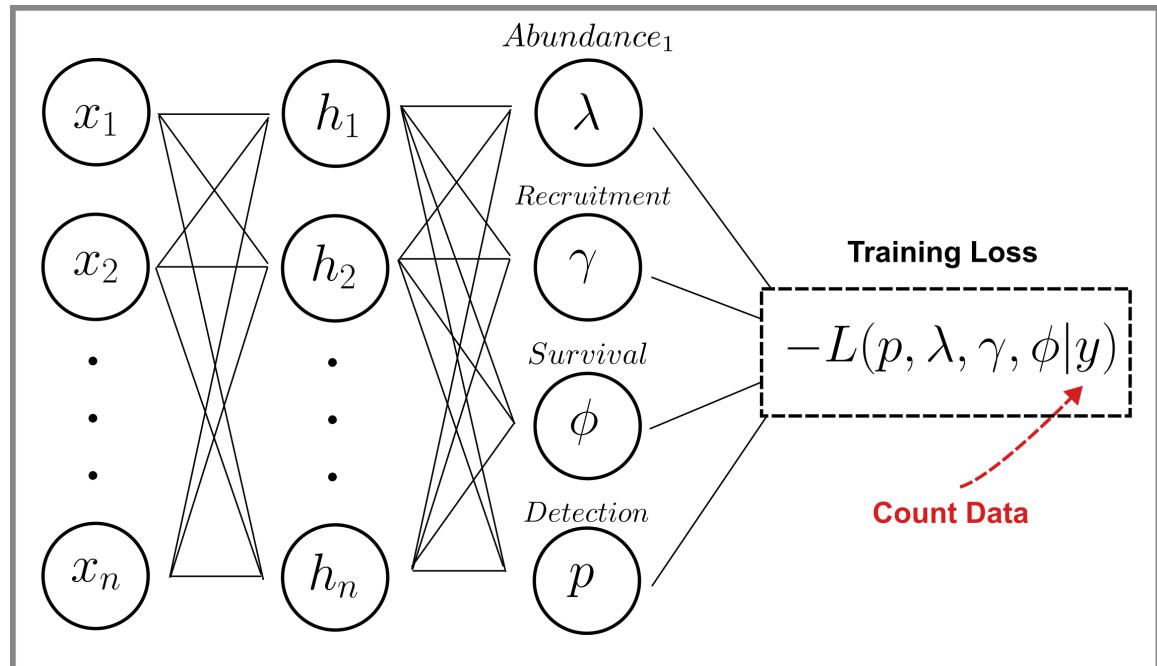
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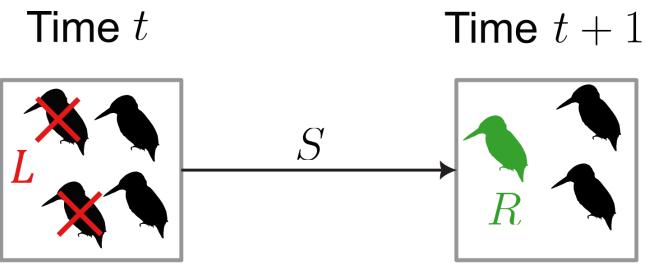
The Ohio State University
2025-09-29



Beyond abundance change

Demographic rates

- Changes in abundance are valuable, but demographic mechanisms offer deeper ecological insights
- Species' extinctions event are the result of increasing difference between birth and loss
- Demographic rates can be early warnings of species extinction
- Monitoring survival and recruitment can help anticipate species' extinction and understand better the current biodiversity crisis



$$N_t = L + S$$

$$N_{t+1} = S + R$$

$$\Delta N = N_{t+1} - N_t = R - L$$

Beyond abundance change

Problem

- Data for demographic rates models are individual based (i.e. *individual encounter history data*)
- They are costly in resources and time
- Individual identification is not equal for all taxa
- Can be invasive/traumatic
- Limited spatial and temporal extent

Individual based sampling



Less invasive
Lower effort
Broadly applicable
Resource efficient

More invasive
Higher effort
Taxon specific
Resource intensive

Beyond abundance change

Using abundance to infer demographic rates

- However, abundance data are available at large spatial and temporal scale
- The idea of inferring birth/immigration (μ) and death/emigration (λ) from experimental data is not new
- While overall change in abundance gives information about $\lambda - \mu$, the volatility of the time-series provide information about $\lambda + \mu$ making λ and μ identifiable

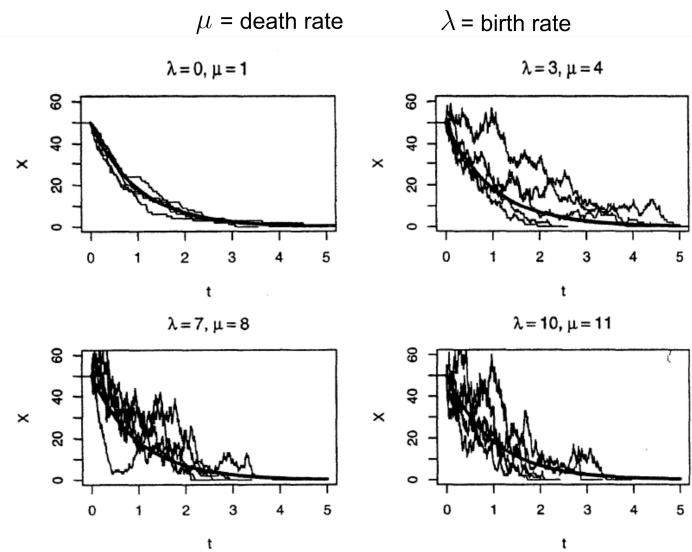


Figure 1.3 Five realisations of a stochastic linear birth-death process together with the continuous deterministic solution for four different (λ, μ) combinations, each with $\lambda - \mu = -1$ and $x_0 = 50$

Dynamic N-mixture model

Hierarchical Model

Observation process

$$y_{i,j,t} \sim \text{Binomial}(N_{i,t}, p)$$

y = observed abundance
 p = detection probability

State process

$$N_{i,1} \sim \text{Poisson}(\lambda)$$

λ = abundance at $t = 1$

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

ϕ = survival probability

$$R_{i,t+1} \sim \text{Poisson}(\gamma_{i,t})$$

γ = number of recruits

$$N_{i,t+1} = S_{i,t+1} + R_{i,t+1}$$

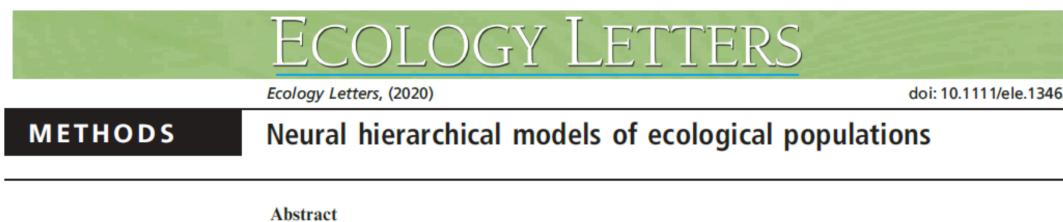
Neural hierarchical model

Limitations of hierarchical framework

- Linear or simple polynomial effects of covariates, even though true ecological responses are often unknown *a priori*
- MCMC algorithm scales poorly because inherently sequential, little possibility of parallelization
- Sensitive to priors and initial values

Limitations of Neural Networks

- Usually lack inferential power, making it less relevant for ecological insights
- Doesn't distinguish between ecological process and imperfect detection



The image shows the front cover of the journal 'Ecology Letters'. The title 'ECOLOGY LETTERS' is at the top in large white letters. Below it, the subtitle 'Ecology Letters, (2020)' and the DOI 'doi: 10.1111/ele.13462' are visible. The main article title 'METHODS Neural hierarchical models of ecological populations' is displayed prominently. The author's name 'Maxwell B. Joseph*' and an ORCID icon are at the bottom left. An abstract section is partially visible at the bottom right.

Maxwell B. Joseph* 

Abstract

Neural networks are increasingly being used in science to infer hidden dynamics of natural systems from noisy observations, a task typically handled by hierarchical models in ecology. This

Neural hierarchical model

ECOLOGY LETTERS

Ecology Letters, (2020)

doi: 10.1111/ele.13462

METHODS

Neural hierarchical models of ecological populations

Abstract

Maxwell B. Joseph* 

Neural networks are increasingly being used in science to infer hidden dynamics of natural systems from noisy observations, a task typically handled by hierarchical models in ecology. This

- Combines flexibility and scalability of neural networks with the inferential power of hierarchical models
- Output activation function according to the parameter to infer (sigmoid for probability, exponential for counts)
- Loss function: tailored from the model specific negative log-likelihood
- Combines decades of development in hierarchical modelling for ecological data with the flexibility and predictive power of Neural Networks

Neural Dynamic N-mixture model

Deep Markov Model*

State process

$$N_{i,1} \sim \text{Poisson}(\lambda)$$

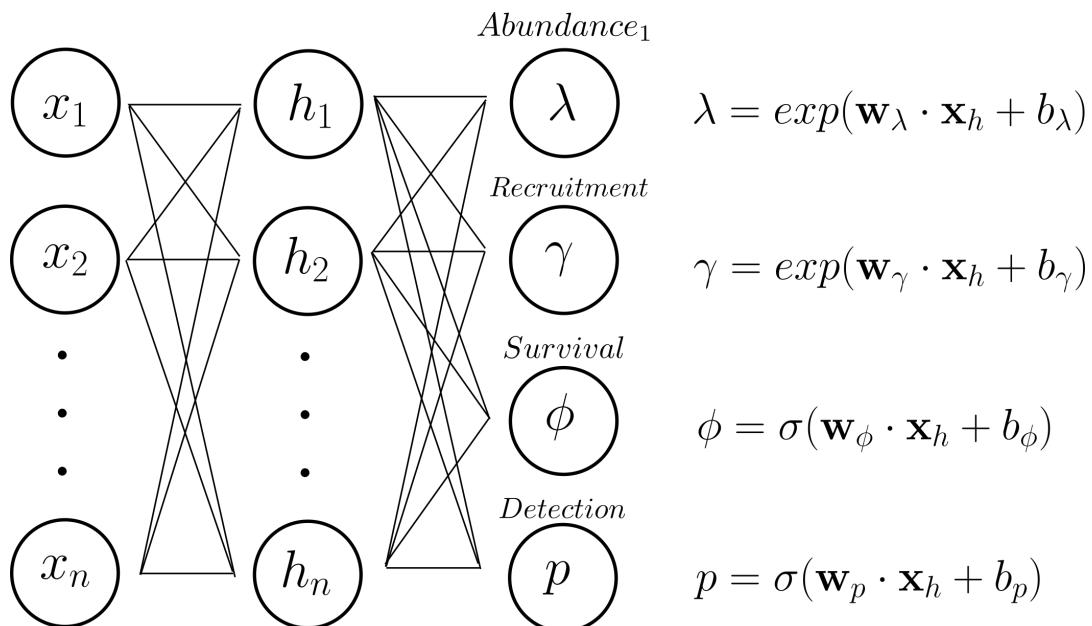
$$R_{i,t+1} \sim \text{Poisson}(\gamma_{i,t})$$

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

$$N_{i,t+1} = S_{i,t+1} + R_{i,t+1}$$

Observation process

$$y_{i,j,t} \sim \text{Binomial}(N_{i,t}, p)$$



Neural Dynamic N-mixture model

Deep Markov Model*

State process

$$N_{i,1} \sim \text{Poisson}(\lambda)$$

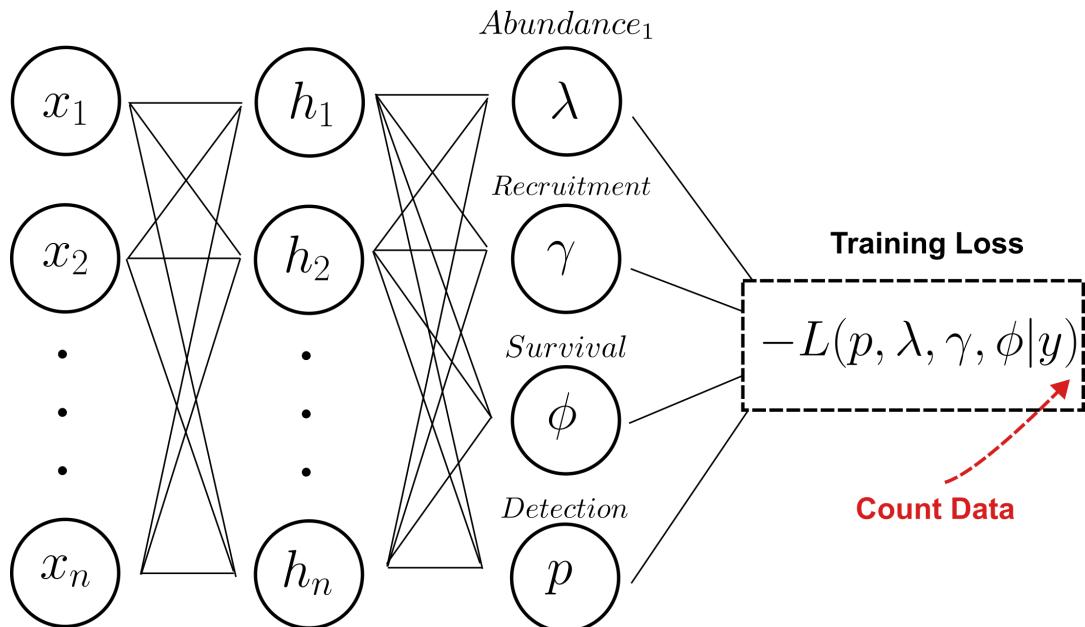
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Observation process

$$y_{i,j,t} \sim \text{Binomial}(N_{i,t}, p)$$



Example with the N-mixture model

Observation process

$$y_{i,j,t} \sim \text{Binomial}(N_{i,t}, p)$$

State process

$$N_i \sim \text{Poisson}(\lambda)$$

Integrated likelihood:

$$L(p, \lambda \mid y_{it}) = \prod_{i=1}^R \left(\sum_{N_i=0}^{\infty} \left(\prod_{t=1}^T \text{Bin}(y_{it}; N_i, p) \right) \text{Pois}(N_i; \lambda) \right)$$

Example with the N-mixture model

Observation process

$$y_{i,j,t} \sim \text{Binomial}(N_{i,t}, p)$$

State process

$$N_i \sim \text{Poisson}(\lambda)$$

Integrated likelihood:

$$L(p, \lambda \mid y_{it}) = \prod_{i=1}^R \left(\sum_{N_i=0}^K \left(\prod_{t=1}^T \text{Bin}(y_{it}; N_i, p) \right) \text{Pois}(N_i; \lambda) \right)$$

- Marginalize over $K >> y_{it}$

Dynamic N-mixture model

State process

$$\begin{aligned}N_{i,1} &\sim \text{Poisson}(\lambda) \\R_{i,t+1} &\sim \text{Poisson}(\gamma_{i,t}) \\S_{i,t+1} &\sim \text{Binomial}(N_{i,t}, \phi_{i,t}) \\N_{i,t+1} &= S_{i,t+1} + R_{i,t+1}\end{aligned}$$

Observation process

$$y_{i,j,t} \sim \text{Binomial}(N_{i,t}, p)$$

Integrated likelihood

$$\begin{aligned}\mathcal{L}(p, \lambda, \gamma, \phi \mid y_{it}) &= \prod_{i=1}^R \left[\sum_{N_{i1}=0}^{\infty} \cdots \sum_{N_{iT}=0}^{\infty} \left\{ \left(\prod_{t=1}^T \text{Bin}(y_{it}; N_{it}, p) \right) \right. \right. \\&\quad \times \text{Pois}(N_{i1}; \lambda) \cdot \prod_{t=2}^T P_{N_{it-1}, N_{it}} \Big\} \Big] \end{aligned}$$

The transition matrix

- As it is a Hidden Markov Model, the likelihood involves a transition between time t and $t + 1$
- This is the bottleneck

Optimization

- Hidden Markov Model \Rightarrow Deep Markov Model (Krishnan et al., 2017)
- Trade-off between memory use and speed
- I choose fast implementation with heavy memory use

Simulated data

Results of the simulation

Fitting on real data

Next step

- So far it is a simple MLP
- Promising to use a CNN

Limitations