Context and objective	Labelled DBN	SBM for prior on communities	Structure learning algorithm	Experiments

Labeled dynamic Bayesian network for learning ecological network

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Context and objective ${\scriptstyle { \bullet } \odot }$

Labelled DBN 0000 SBM for prior on communities

Structure learning algorithm

Experiments 000

Ecological context and objective

Context

- Management of biodiversity within an ecological network
- Interactions are poorly known
- Few data, but expert knowledge



Objective

Developing a method for learning the structure of an ecological network using presence/absence temporal data

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Structure learning algorithm

Experiments

Ecological network modeling

Ecological network

- Directed graph. Nodes represents species.
- Edges labeled according to the type of interaction :
 - + : Positive influence for survival
 - - : Negative influence for survival



Associated labelled Dynamic Bayesian Network model

- Binary variables (presence/absence)
- Survival and recolonization depend on previous year



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Context and objective

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Labeled dynamic Bayesian network model

Definition

- Each edge is labelled
- The transition probability distribution of a variable X^t_i only depends on its number of parents of each state and label
- Two variables with the same numbers of parents of each state and each label have the same transition probability distributions
- Transition probability distribution : function of a small vector of parameters θ. The size of θ independent from the graph structure.



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Model of sp	ecies dyn	amics		

Data

- $X_t^i \in \{1,0\}$ presence or absence of the species i $(i \in \{1,...,n\})$ at year t $(t \in \{1,...,T\})$.
- $A^t \in \{1,0\}$ protection or absence of protection at year t.

Parameters

- Recolonization probability ε .
- Probability of success of each positive influence ρ .
- Probability of success of each negative influence τ .

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Transition F	Probabiliti	es		

Recolonization

Species absent at year t - 1: probability of recolonization at year t: $P(X_i^t = 1 | X_i^{t-1} = 0) = \varepsilon$

Survival

Species present at year t - 1: probability of survival at year t: $P(X_i^t = 1 | X_i^{t-1} = 1) = (1 - (1 - \rho)^{N_{i,+}^t}) (1 - \tau)^{N_{i,-}^t}$

 $N_{i,l}^t$: number of "l" labeled parents of the species *i* present at year *t*.

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SBM model for prior on communities

Definition

- Known blocks (communities) of variables in the network
- The probability of presence of a labelled edge i → j is a function of its label and the blocks of i and j parameterized by ψ.



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SBM model for prior on trophic levels

Hypothesis knowing trophic levels

- No top down positive edge
- Positive edges more likely on the closest superior trophic level
- Top down negative edges more likely than bottom-down or intra-level negative edges



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Probabilities of edges presence

Positive edges

Trophic levels TL(i) and TL(j) determine the probability of presence of the labelled edge G'_{ij} .

• Top-down and intra-level : $P(G_{ij}^+|TL(i) \ge TL(j)) = 0.$

• Bottom-up :
$$P(G_{ij}^+|TL(i) \ge TL(j)) = \frac{e^{\alpha \Delta_{ij}}}{1+e^{\alpha \Delta_{ij}}}$$

with $\Delta_{ij} = TL(i) - TL(j)$ and $\alpha > 0$

Negative edges

•
$$P(G_{ij}^-|TL(i) \leq TL(j)) = \beta_2$$

•
$$P(G_{ij}^-|TL(i) > TL(j)) = \beta_1$$

with $\beta_1 > \beta_2$

 $\psi = (\alpha, \beta_1, \beta_2)$

Score-based method

- Number of parameters independent from structure : likelihood as score
- Greedy algorithm
 - Step 1 (Estimation) : Parameters estimation by likelihood maximization, graph structure known
 - Step 2 (Restoration) : Learning network structure maximizing likelihood, parameters known
 - Back to step 1 until convergence

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Ecological network learning algorithm

Decomposability of the likelihood

$$\log P(x^1, \dots, x^T \mid x^0, a, \theta, \mathcal{LG}_{\rightarrow}) = \sum_{i=1}^n score(i)$$

$$score(i) = \sum_{\substack{t=0\\t=0}}^{T-1} (1-x_i^t) \log \varepsilon \\ + \sum_{t=0}^{T-1} x_i^t \sum_{\substack{0 \le d^+ + d^- \le k}} \log \left(\left(1 - (1-\rho)^{d^+} \right) (1-\tau)^{d^-} \right) R_i^{t,d^+,d^-}$$

with $R_i^{t,d^+,d^-} = 1$ iff the species *i* has d^+ positive labelled parents and d^- negative labelled parents present at year t

Likelihood term for SBM

SBM term :
$$logP(\mathcal{LG}_{\rightarrow}|\psi) = \sum_{j} score^{SBM}(j)$$

$$\begin{split} & \textit{score}^{\textit{SBM}}(j) = \sum_{i, \Delta_{ij}=0} g_{ij}^{-} \log \beta_2 + (1 - g_{ij}^{-}) \log (1 - \beta_2) \\ &+ \sum_{i, \Delta_{ij}<0} \alpha \Delta_{ij} g_{ij}^{+} - \log (1 + \exp^{\alpha \Delta_{ij}}) + (1 - g_{ij}^{+}) (g_{ij}^{-} \log \beta_2 + (1 - g_{ij}^{-}) \log (1 - \beta_2)) \\ &+ \sum_{i, \Delta_{ij}>0} g_{ij}^{-} \log \beta_1 + (1 - g_{ij}^{-}) \log (1 - \beta_1) \end{split}$$

Experiments

Ecological network learning algorithm

Integer linear programming (ILP) 0-1

- Linearisation of the problem : addition of binary variables defined by linear constraints
- Optimization of the score using ILP
- One independent ILP per variable

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Tested methods

- L-DBN, no additionnal knowledge
- L-DBN, SBM prior
- L-DBN, 20% of known edges, no SBM prior
- MIT^a score method and qualitative network^b

^aMutual Information Test. Vinh, 2011 ^bWellman, 1990

Simulations

• Synthetic networks built from SBM model $(\alpha = 1/\sqrt{20}, \beta_1 = \alpha/2, \beta_2 = \beta_1/2)$

•
$$\varepsilon = \rho^+ = \rho^- = \mu = 0.8$$

 10 networks simulated, 10 datasets per network, 20 species, 30 years, last 18 years protected

Synthetic n	etwork re	sults		
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Real networ	·k			

Arthropods dataset (Bohan et al, 2013)

Arthropods trapped in experimental fields

- 66 Beetroot (41 species)
- 59 Maize (29 species)

- 67 Summer rape (40 species)
- 65 Winter rape (29 species)



Conclusion				
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Labeled dynamic Bayesian network

- Few data available
- DBN with few parameters
- Structure learning using ILP
- Inclusion of expert knowledge
- SBM prior improves learning quality

Application of LDBN and perspectives

- Adaptable for "propagation per contact" models (rumor propagation, network security, disease propagation, fire propagation...)
- Managing while learning : Factored reinforcment learning with MDP and LDBN transition structure