Species Distribution Models: Models and Application for Sustainable Species Conservation

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- 1 Species Distribution Models (SDMs)
- 2 Types of SDMs and some Algorithms
- 3 Correlative SDMs
- 4 Mechanistic SDMs

Concepts

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Concept and Theory

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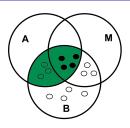
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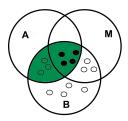
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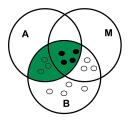


Niche Theory explained with a diagram

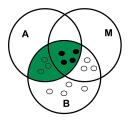


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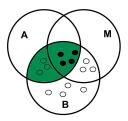


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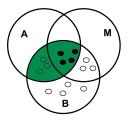
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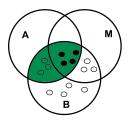
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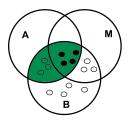


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- Fundamental niche was the set of all the states of the environment which would permit the species S to exist (Hutchinson 1957)

Type of models in SDM

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 - Generalized linear models (GLMs) = enabled pioneering regression-based SDMs that had much more sophistication and realism than was possible earlier (e.g., see Austins work in 1970s and 1980s, cited in Austin 1985). The key structural features of GLMs (non-normal error distributions, additive terms, nonlinear fitted functions) continue to be useful and are part of many current methods including RSFs (Manly et al. 2002) and maximum entropy models (MaxEnt: Phillips et al. 2006).

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Explanation: These models are used to seek insight, even if indirectly, into the causal drivers of species distributions (Mac Nally 2000). SDMs are still regularly used for such purposes, particularly in quantitative ecological studies (Leathwick & Austin 2001) and evolutionary biology (Graham et al. 2004).

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 - The model-based extrapolation to unsampled sites: The environments in these new times and places need to be carefully assessed, particularly for new combinations of predictor values or for predictor values outside their original ranges in the training data. Prediction of new geographic regions is a special case and has been termed transferability, but often without clear information on the environmental similarities and differences between the model fitting and prediction regions.

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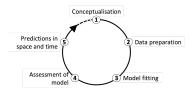
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Scale of SDMs

Scale is relevant to the distributions of both species and environments and comprises both grain and extent.

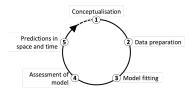
- The extent (or domain) usually reflects the purpose of the analysis. For instance, macroecological and global change studies tend to be continental to global in scope(e.g., Arau io& New 2007), whereas studies targeting detailed ecological understanding or conservation planning tend toward local to regional extents (Fleishman et al. 2001, Ferrier et al. 2002).
- The Grain usually describes properties of the data or analysis often the predictor variables and their grid cell size or polygon size, but also the spatial accuracy and precision of the species records (Dungan et al. 2002, Tobalske 2002)

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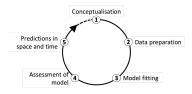
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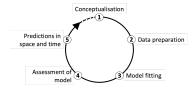


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- How to deal with multicollinearity in the environmental data?
- How many variables should be included in the model (without overfitting) and how should we select these?
- How to select the final model or average the models?
- Guisan, Thuiller, and Zimmermann (2017)
- Model assessment: Strictly, checking the plausibility of the fitted species-environment relationship by visual inspection of response curves, and by assessing model coefficients and variable importance would also be part of the model assessment.
- Predictions: Inspected model and extrapolation behavior, and assessed predictive performance, it is finally time to make predictions in space and time.

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Grouping of correlative algorithms

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 - 2 DOMAIN
 - Ecological Niche Factor Analysis (ENFA)
 - 4 Mahalanobis distance

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Family of correlative SDMs algorithms (End)

Machine learning techniques

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- Furthermore, ensemble models can be created from several model outputs to create a model that captures components of each. Often the mean or median value across several models is used as an ensemble.
- Similarly, consensus models are models that fall closest to some measure of the central tendency of all models consensus models can be individual model runs or ensembles of several models.



Description

Classification Tree Analysis (CTA), also known as Decision Tree Analysis, is a machine learning algorithm used for classification and regression tasks. In the context of SDM, CTA is applied to predict the presence or absence of a species based on environmental predictor variables. The algorithm recursively splits the dataset into subsets based on the predictor variables, creating a tree structure where each node represents a decision based on a specific variable. Here are some of the advantages of CTA:

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- Variable Importance: CTA provides a measure of variable importance, helping identify which
 environmental factors are most influential in predicting species distribution.



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(4)

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$$MaximizeH(p) = -\sum_{i} p(x_i)log(p(x_i)) + \lambda \sum_{j} w_j f_j(x)$$
 (5)

where H(p) is the entropy; $p(x_i)$ is the probability of the environmental variable x_i ; λ is a regularization parameter; w_i are weights assigned to environmental features $f_i(x)$

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Concepts

Thank You