

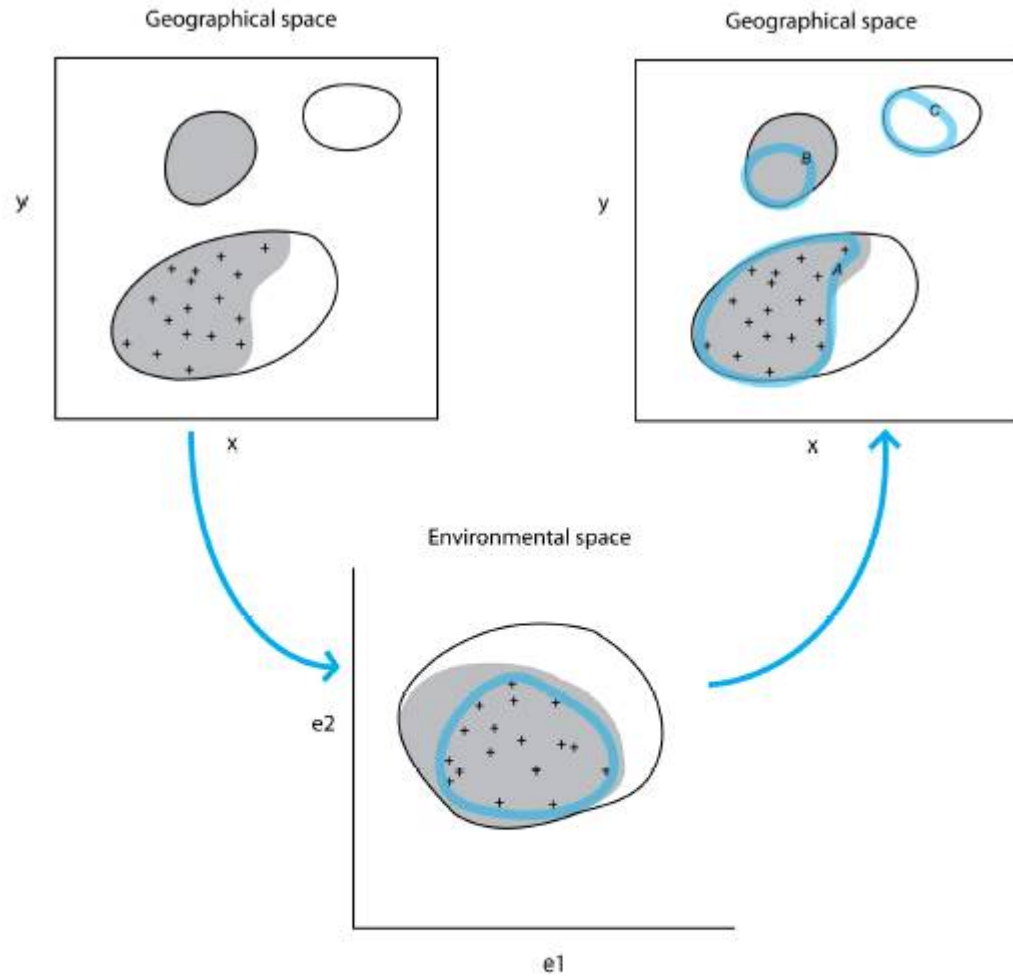
Species Distribution Modeling and GIS

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Field of Soil and Crop Sciences
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GIS Day 2009

19 November 2009



- + Observed species occurrence record
- Actual distribution (upper panels)/Occupied niche (lower panel)
- Potential distribution (upper panels)/Fundamental niche (lower panel)
- Species distribution model fitted to observed occurrence records

Species Distribution Models

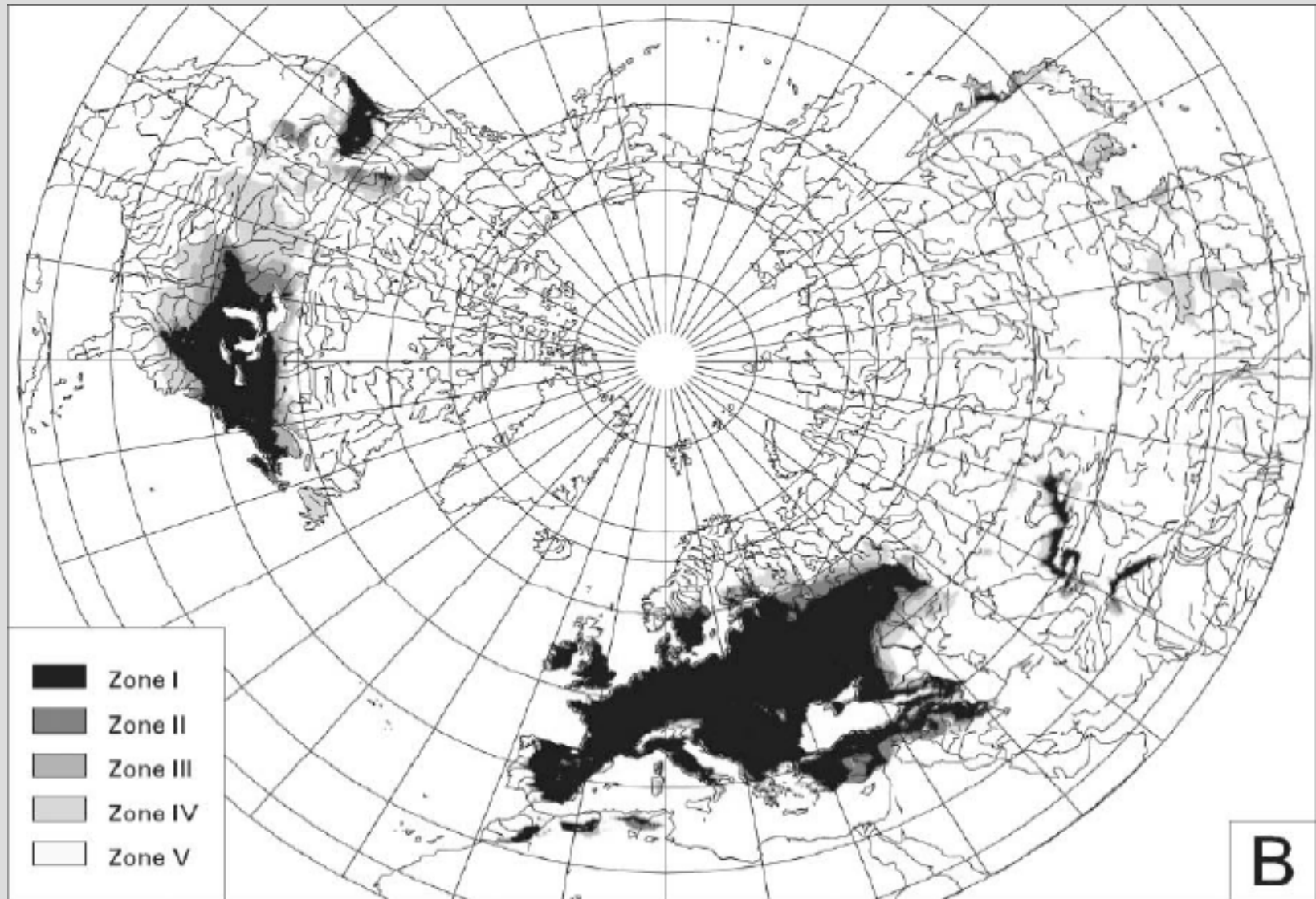
Models	Class of model	Data
• <i>BIOCCLIM</i>	envelope model	p
DOMAIN	multivariate distance	p
LIVES	multivariate distance	p
GLM	regression; generalized linear model	p/pa
GAM	regression; generalized additive model	p/pa
MARS	multivariate adaptive regression splines	p/pa
DT-GARP OM-GARP	rule sets derived with genetic algorithms	p/pa
• <i>BRT</i>	boosted regression trees	p/pa
• <i>MAXENT</i>	maximum entropy	p/b
MARS COMM	as for MARS; uses community data	p/c/b
GDM	generalized dissimilarity modeling; uses community data	p/c/b

Concerns with Species of Interest: Garlic Mustard

- Ecological threat to an endangered species
- Wide range of potential distribution
- Invades healthy, intact forest ecosystems without a disturbance

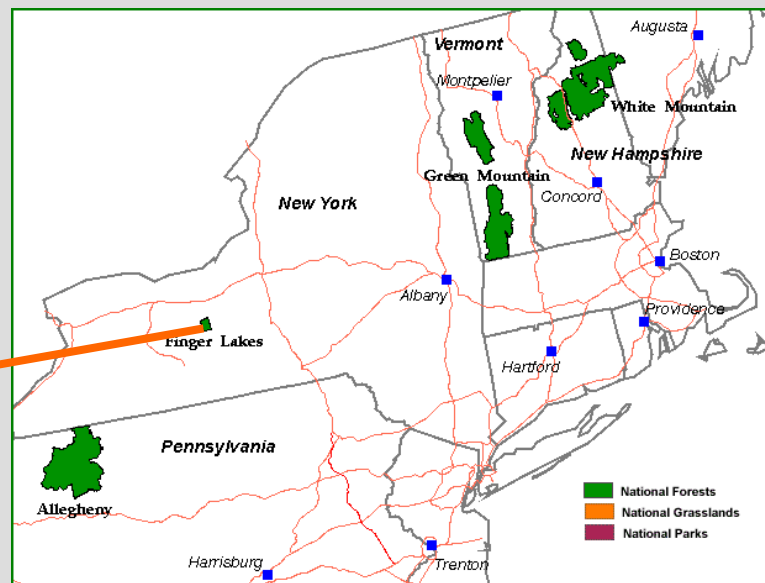


Current modeling approaches are global in nature (need local/regional approaches for effective management)



Climatically modelled range and long-term probability zones for invasion in North America.
(Welk et. al. 2002)

Study Area

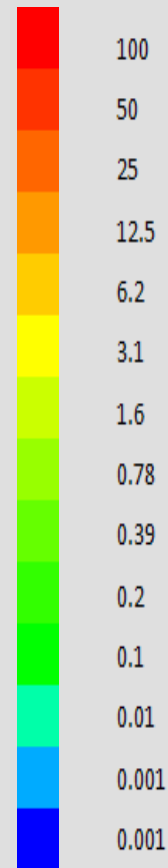
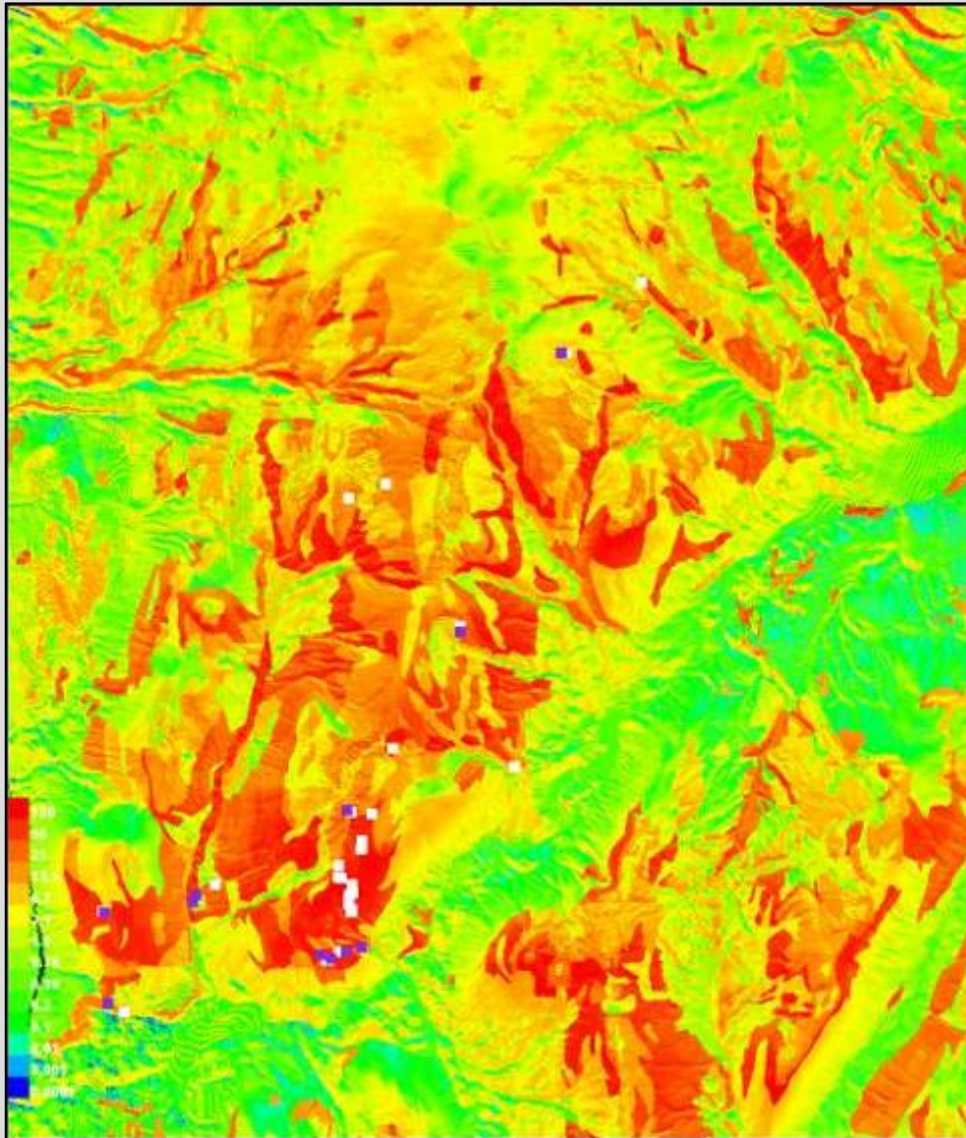


Predictor Variables (MaxEnt)

Infestation Susceptibility:

f (soil type, elevation, slope aspect, slope gradient)

MaxEnt Model Results



Probability surface for
Alliaria petiolata
infestation

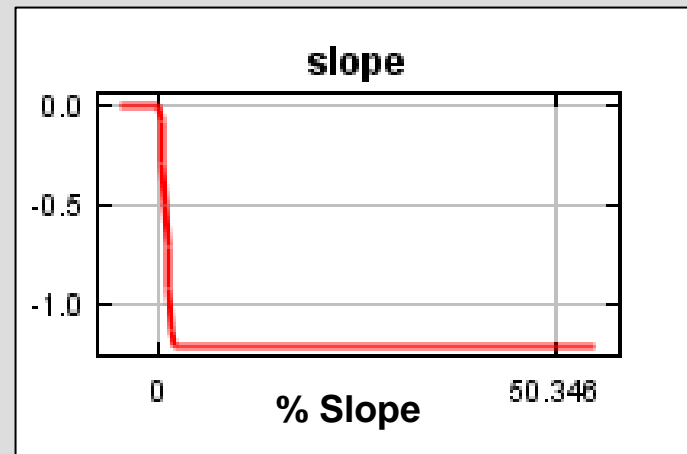
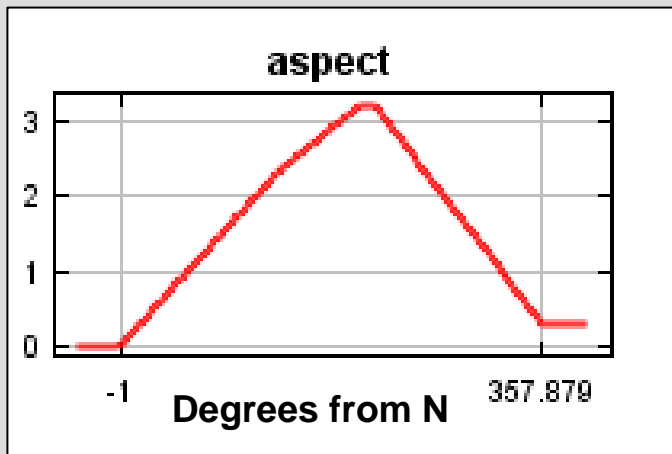
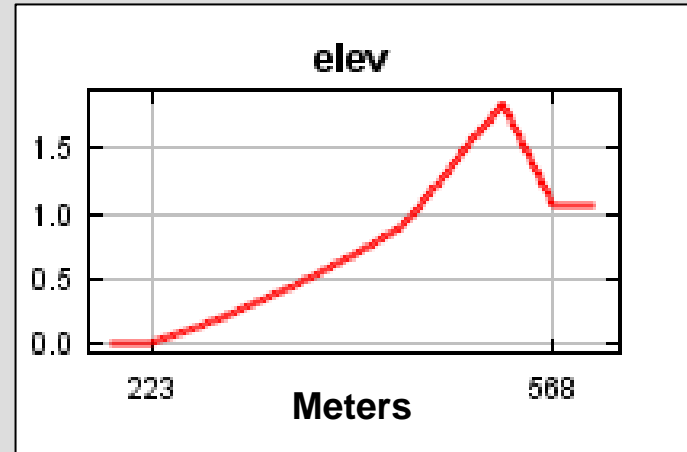
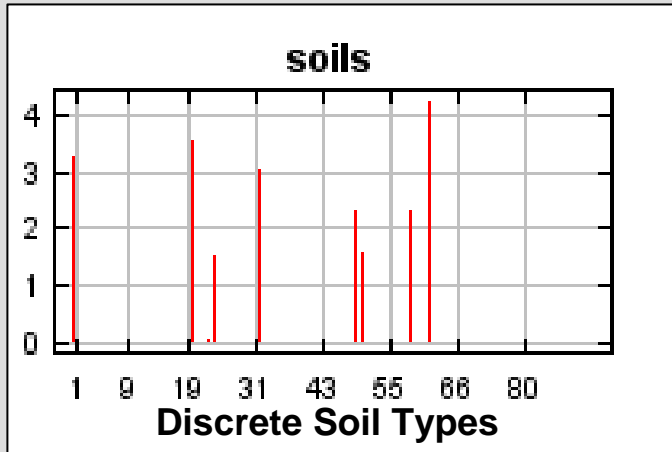
Warmer colors show
areas with higher
probability of
infestation.

White dots show the
presence locations
used for model
development

Violet dots show model
test data point
locations.

MaxEnt Response Curves

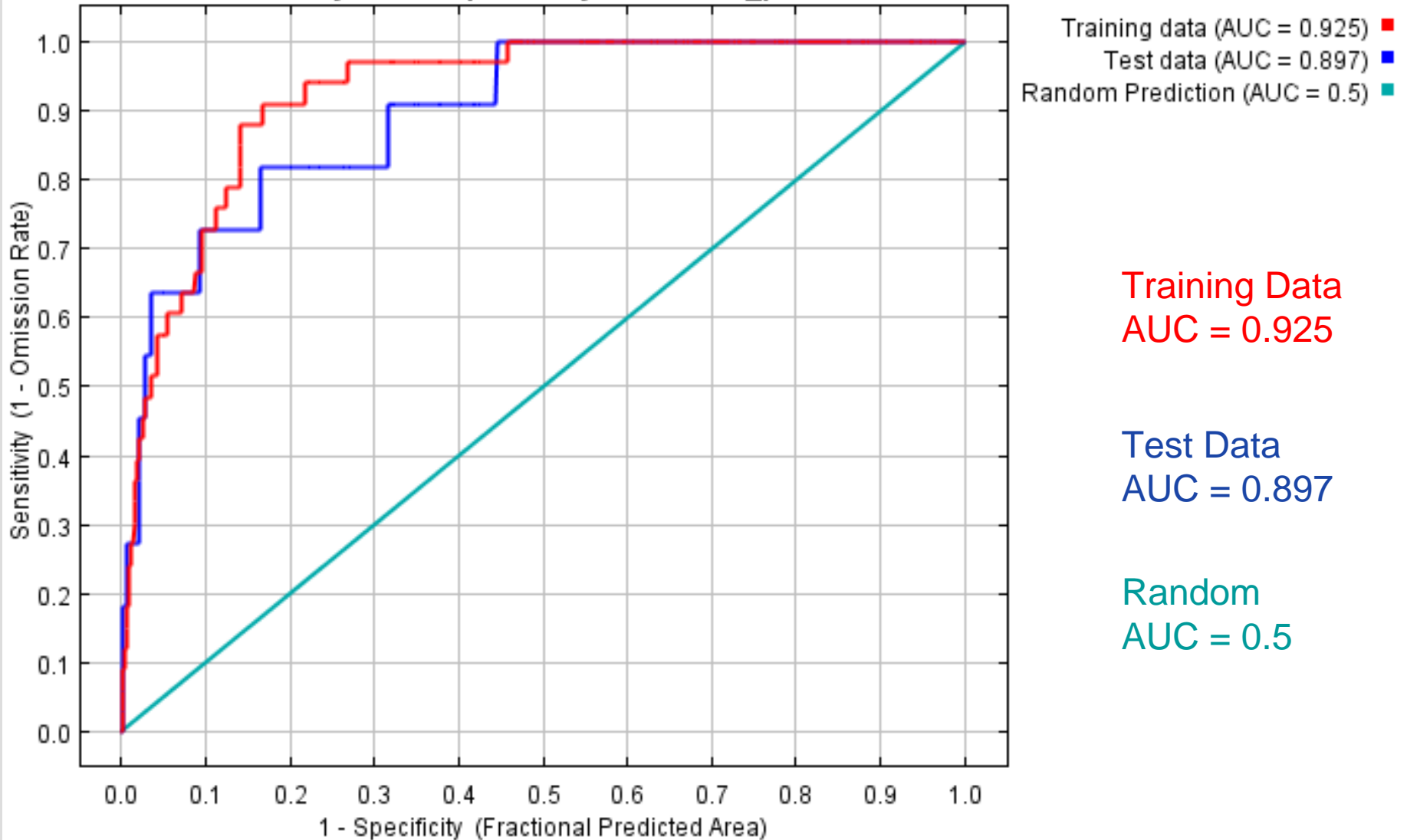
Additive contribution to exponent



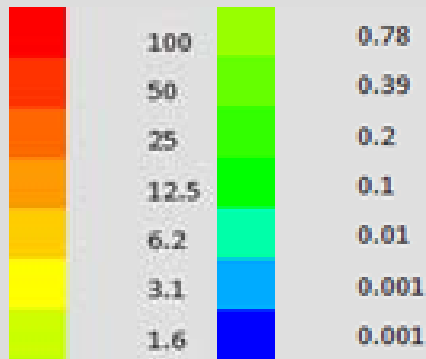
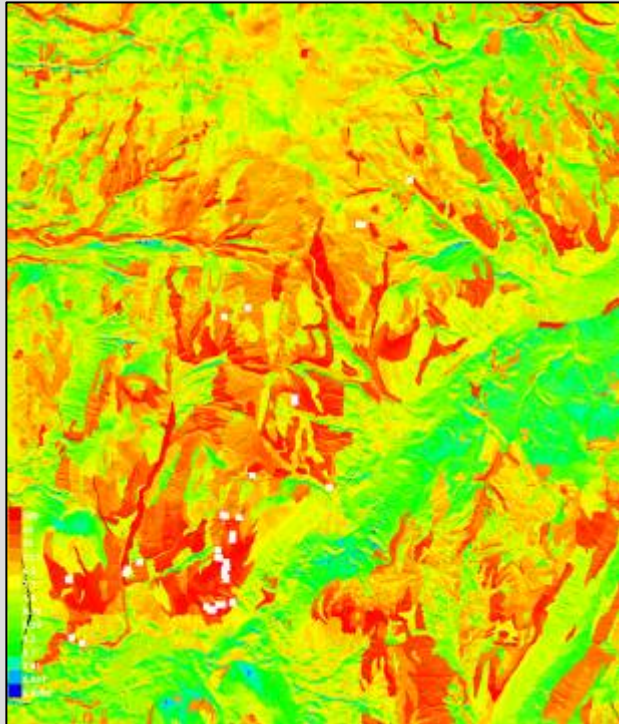
Value of environmental variable

ROC Curve

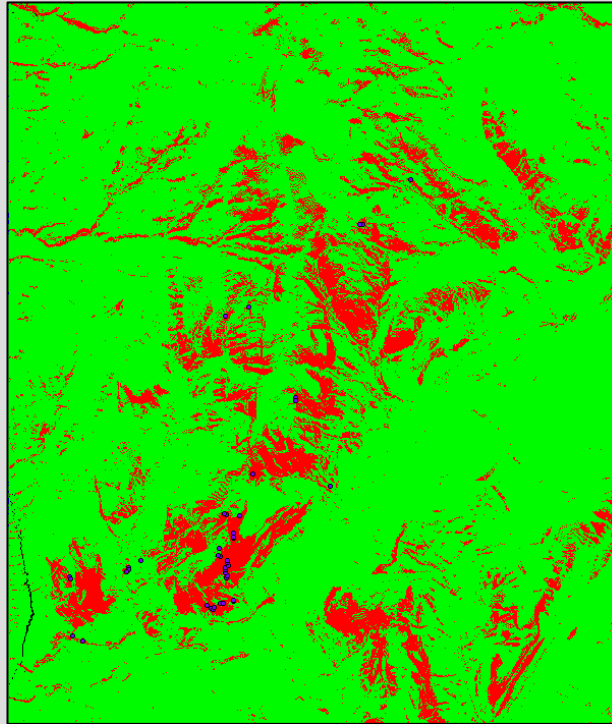
Sensitivity vs. 1 - Specificity for *Alliaria_petiolata*



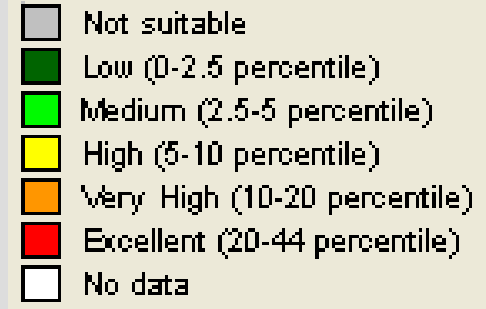
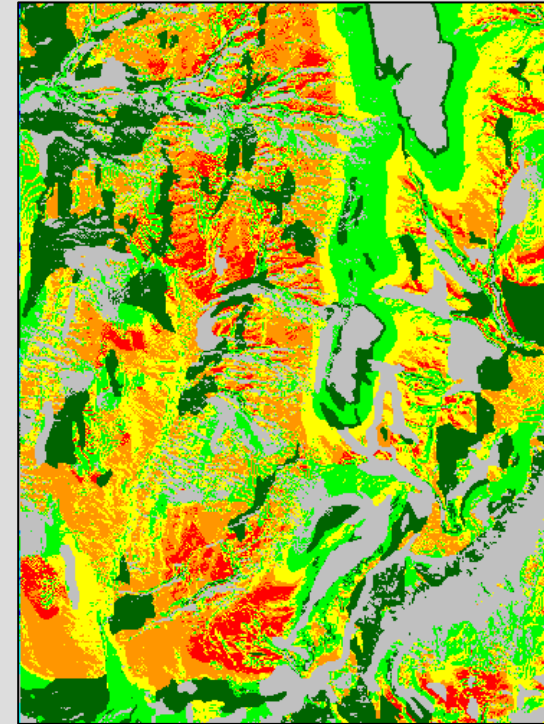
Maxent



BRT in R



Bioclim Diva GIS



Software

- MaxEnt –
 - <http://www.cs.princeton.edu/~schapire/maxent>
 - <http://groups.google.com/group/Maxent>
- Boosted Regression Trees in R <http://www.r-project.org>
- Diva-GIS – Incorporates distribution modeling tools – <http://www.diva-gis.org>
- OpenEV – Open Source Image Processing <http://openev.sourceforge.net>
- openModeller – <http://openmodeller.sourceforge.net>
- SAGA-GIS – System for automated geoscientific analyses – <http://www.saga-gis.org>

WorldClim - Global Climate Data

- <http://www.worldclim.org/> – Interpolated global climate surfaces
- **Time Periods**
 - **Current** conditions (interpolations of observed data, representative of 1950-2000)
 - **Future** conditions (downscaled from global climate model (GCM) output, IPCC 3rd assessment; 4th assessment coming soon)
 - **Past** conditions (downscaled global climate model output)
- **Grids - Generic or ESRI**
- **Resolution**
 - **30 arc-seconds (~1 km)**
 - **2.5 arc-minutes (~20 sq km @ equator)**
 - **5 arc-minutes**
 - **10 arc-minutes**

Species Distribution Modeling



Guide ○ Online Resources ○ Training ○ Search

Training

Species Distribution Modeling > Training

AMNH Training Workshop: Species Distribution Modeling

COURSE DATES: March 29 - April 2, 2010



McPhearson © AMNH CBC
The Southwestern Research Station, Arizona.

The American Museum of Natural History's Center for Biodiversity and Conservation runs a week-long workshop on **Species Distribution Modeling**.

Applications are now being accepted for the training workshop **Species Distribution Modeling**, to be held at the American Museum of Natural History's **Southwestern Research Station, Arizona**.

Models that predict species' potential distributions by combining observed occurrence records with digital data layers of environmental variables have great potential for application across a range of biogeographical analyses. Applications include

guiding field surveys to detect unknown populations, projecting potential impacts of climate change, predicting species' invasions, supporting reserve planning, and investigating niche evolution. The workshop focuses on the theoretical and practical aspects of this approach (sometimes termed 'ecological niche' or 'bioclimate envelope' modeling) and is designed for students, researchers and practitioners of conservation biology. Using a mixture of lectures, hands-on computer lab applications, discussions, and case studies, participants will learn to:

- Obtain and process data necessary for species distribution modeling;
- Run distribution models using a variety of approaches;
- Validate and interpret model results;
- Apply these techniques to a range of applications.