

Assessing the potential distribution of insect pests under current and future climatic conditions in European forests

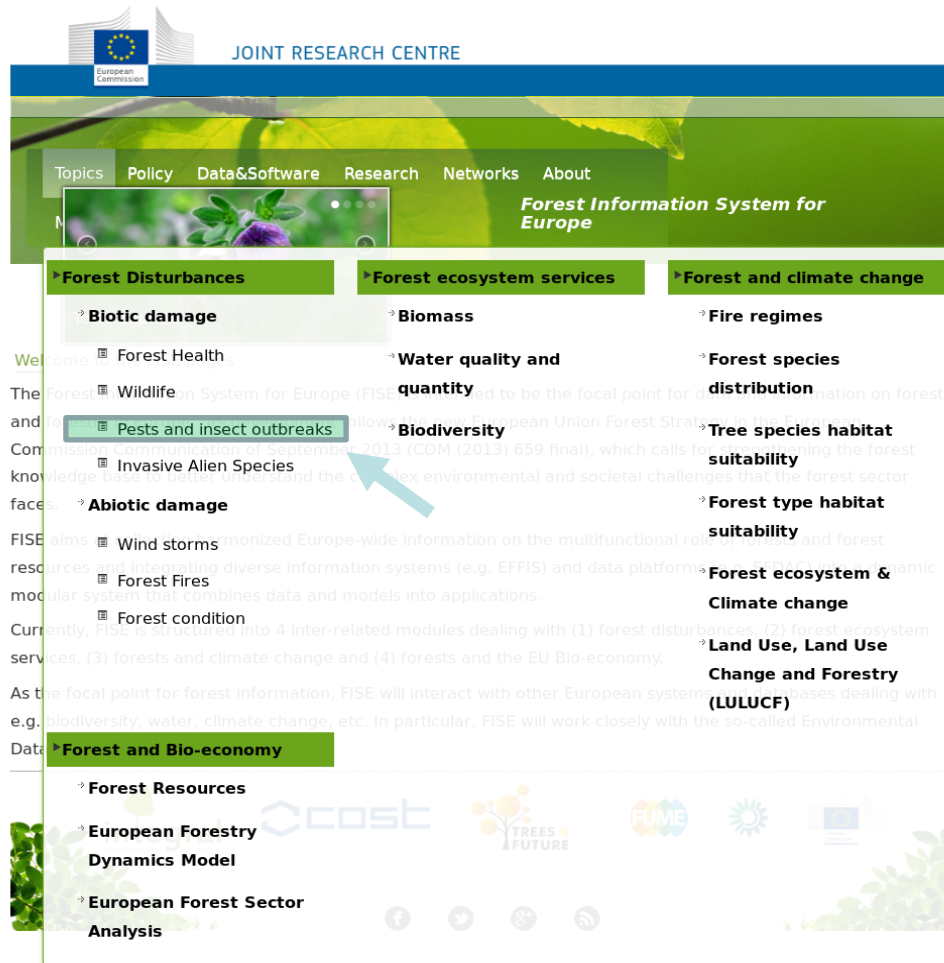
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Joint EFSA-EPPO Workshop on Data collection and information sharing in plant health
Parma, Italy – 1-3 April 2014

FISE: Forest Information System for Europe



The screenshot shows the FISE website interface. At the top, there is a header with the European Commission logo and the text "JOINT RESEARCH CENTRE". Below this is a navigation bar with links: Topics, Policy, Data&Software, Research, Networks, and About. The main content area is titled "Forest Information System for Europe" and features a large green banner with a forest image. Below the banner, there are three main sections: "Forest Disturbances", "Forest ecosystem services", and "Forest and climate change". Each section has a list of sub-topics. A blue arrow points to the "Pests and insect outbreaks" link under the "Forest Disturbances" section. At the bottom, there is a footer with logos for "cost", "TREES FOR THE FUTURE", "FLUME", and "Joint Research Centre".

JOINT RESEARCH CENTRE

Topics Policy Data&Software Research Networks About

Forest Information System for Europe

Forest Disturbances

- Biotic damage
 - Forest Health
 - Wildlife
 - Pests and insect outbreaks**
 - Invasive Alien Species
- Abiotic damage
 - Wind storms
 - Forest Fires
 - Forest condition

Forest ecosystem services

- Biomass
- Water quality and quantity
- Biodiversity

Forest and climate change

- Fire regimes
- Forest species distribution
- Tree species habitat suitability
- Forest type habitat suitability
- Forest ecosystem & Climate change
- Land Use, Land Use Change and Forestry (LULUCF)

Forest and Bio-economy

- Forest Resources
- European Forestry Dynamics Model
- European Forest Sector Analysis

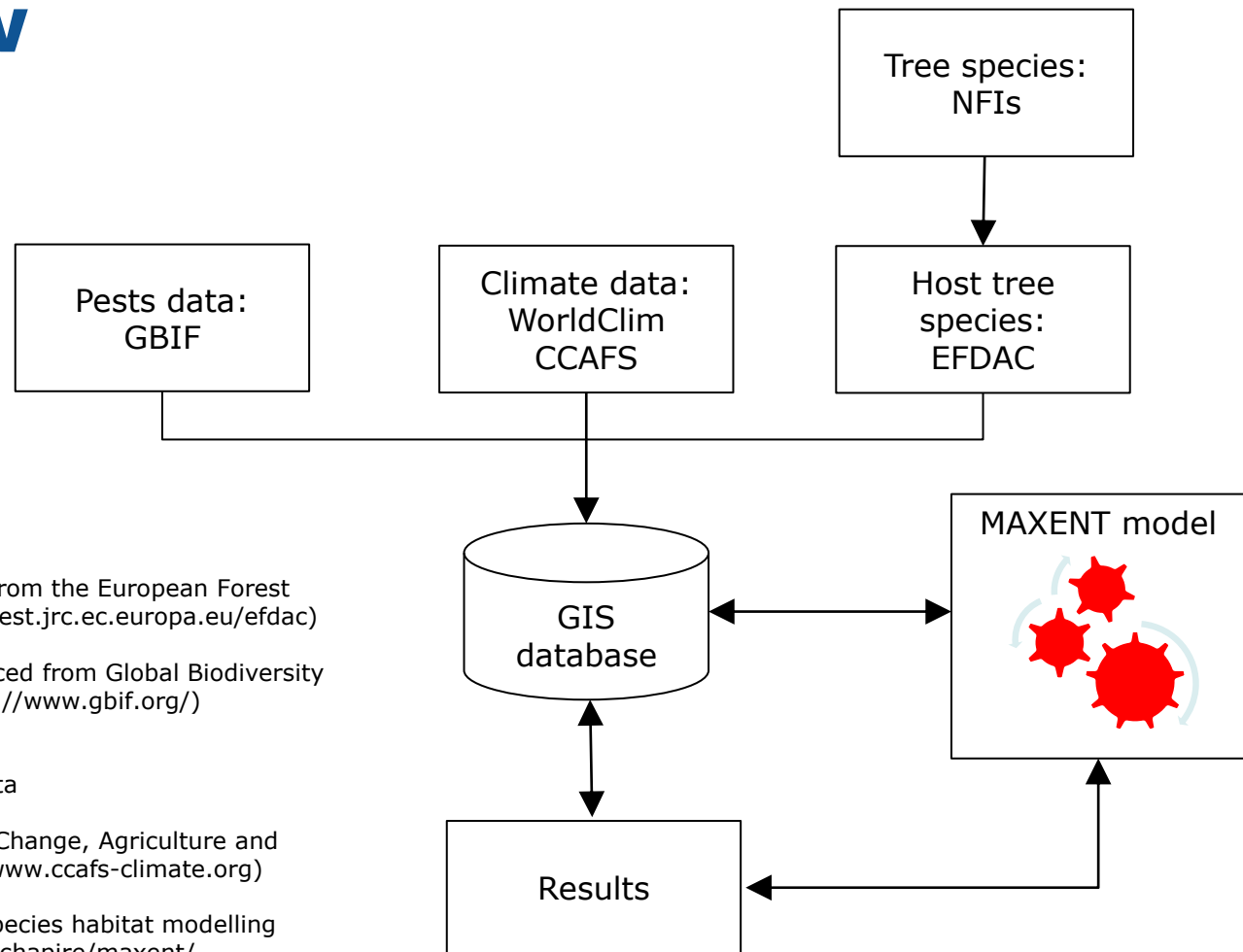
cost TREES FOR THE FUTURE FLUME

Joint Research Centre

Introduction

- We propose a methodology for assessing forest vulnerability to insect pests at pan-European level
- Pilot study using two insect pests for testing and validating a methodology that could be extended to other pests
 - Large Pine Weevil (*Hylobius abietis*)
 - Horse Chestnut Leaf Miner (*Cameraria ohridella*)
- Our results highlight the strengths of the approach, facilitate information sharing with decision makers and discuss the limitations, including data availability of forests insect pests

Overview



Host tree species data sourced from the European Forest Data Centre – EFDAC (<http://forest.jrc.ec.europa.eu/efdac>)

Insect pests presence data sourced from Global Biodiversity Information facility - GBIF (<http://www.gbif.org/>)

Climate data sourced from:
- WorldClim - Global Climate Data (<http://www.worldclim.org>)
- Research Program on Climate Change, Agriculture and Food Security – CCAFS (<http://www.ccafs-climate.org>)

MAXENT: Maxent software for species habitat modelling
<http://www.cs.princeton.edu/~schapire/maxent/>

MAXENT software for species habitat modelling

- Maxent (v.3.3.3e) is a general purpose habitat modelling algorithm for estimating probability of distributions based on presence-only data (Phillips et al., 2004, 2006)
- The model is non-linear, nonparametric, and not sensitive to multicollinearity
- Freeware [!]
- Input data:
 - Environmental covariates (climate, etc.)
 - Presence data (usually point data)

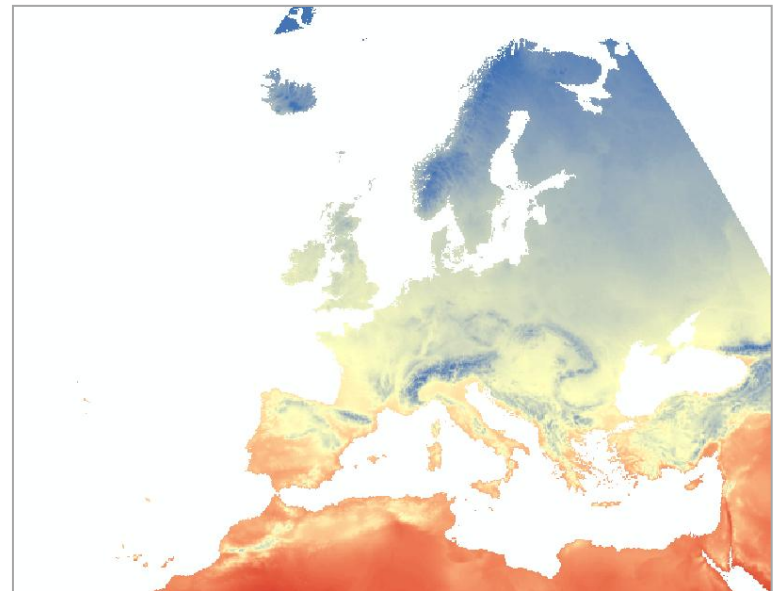
Climate datasets

- WorldClim – Observed Global Climate Data: 1960-1990
- Research Program on Climate Change, Agriculture and Food Security (CCAFS) – A1B IPCC SRES Scenario: 2100
 - Max Planck Institute (MPI): ECHAM5 (GCM)
 - Canadian Centre for Climate Modelling and Analysis (CCCMA): CGCM3 (GCM)
- Disaggregated: 10km spatial resolution

BIOCLIM variables:

- BIO1 = Annual Mean Temperature
- BIO2 = Mean Diurnal Range (Mean of monthly (max temp - min temp))
- BIO3 = Isothermality ($BIO2/BIO7$) (* 100)
- BIO4 = Temperature Seasonality (standard deviation *100)
- BIO5 = Max Temperature of Warmest Month
- BIO6 = Min Temperature of Coldest Month
- BIO7 = Temperature Annual Range ($BIO5-BIO6$)
- BIO8 = Mean Temperature of Wettest Quarter
- BIO9 = Mean Temperature of Driest Quarter
- BIO10 = Mean Temperature of Warmest Quarter
- BIO11 = Mean Temperature of Coldest Quarter
- BIO12 = Annual Precipitation
- BIO13 = Precipitation of Wettest Month
- BIO14 = Precipitation of Driest Month
- BIO15 = Precipitation Seasonality (Coefficient of Variation)
- BIO16 = Precipitation of Wettest Quarter
- BIO17 = Precipitation of Driest Quarter
- BIO18 = Precipitation of Warmest Quarter
- BIO19 = Precipitation of Coldest Quarter Annual average temperature

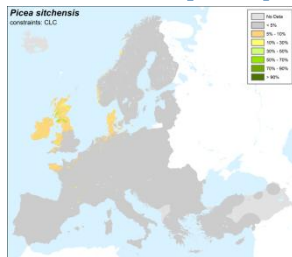
BIO1: Annual mean temperature



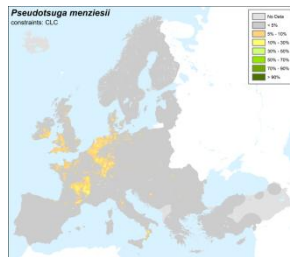
EFDAC - Tree species distribution

Host tree species of Large Pine Weevil (*Hylobius abietis*)

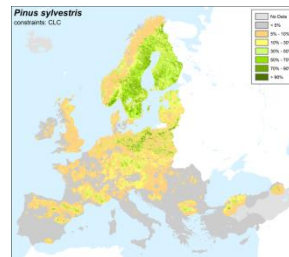
Probability of presence



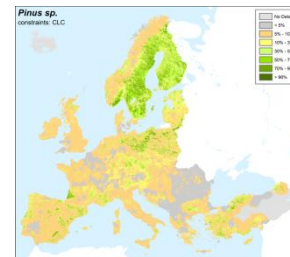
Picea sitchensis



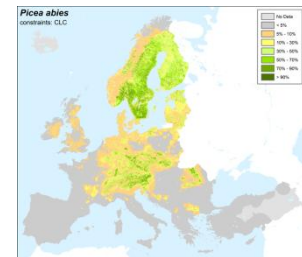
Pseudotsuga menziesii



Pinus sylvestris



Pinus sp.

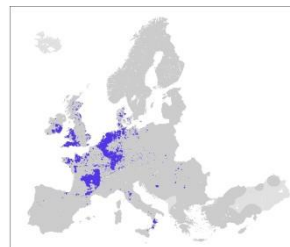


Picea abies

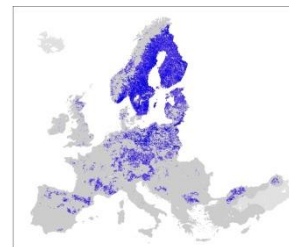
10th percentile training presence (10-TP)



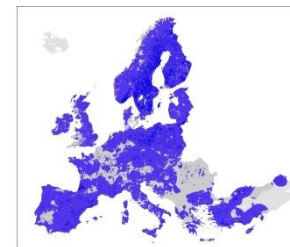
Picea sitchensis



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Pinus sylvestris



Pinus sp.



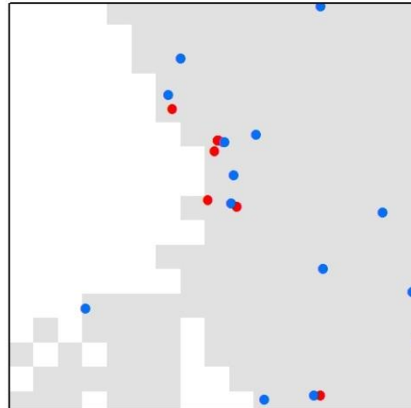
Picea abies

Pests data – Spatial filtering (sampling bias)

Cameraria ohridella - raw data



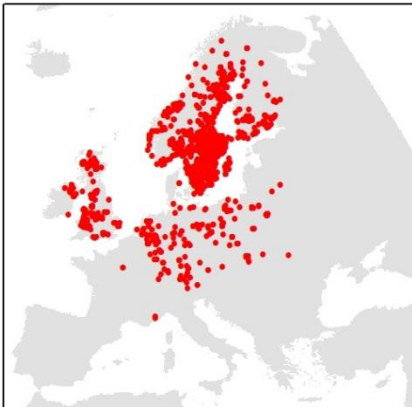
Spatial filtering - 10 Km



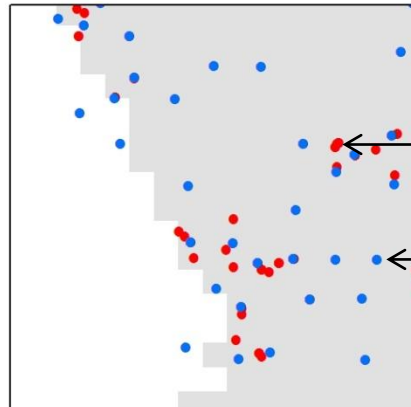
Sample data pre-processing

- 1) Acquisition: download
- 2) Spatial accuracy ≥ 0.1 DD
- 3) Spatial filtering
- 4) Avoid replication (inside 10x10km grid size)

Hylobius abietis - raw data



Spatial filtering - 10 Km



← Eliminated

← Retained

Cameraria ohridella

Raw: 1188

Final: **152**

Hylobius abietis

Raw: 2868

Final: **677**

Results

Maxent calibration (CTR): 10 replications; 20% test data

Regularisation multiplier

b=1.0 standard model

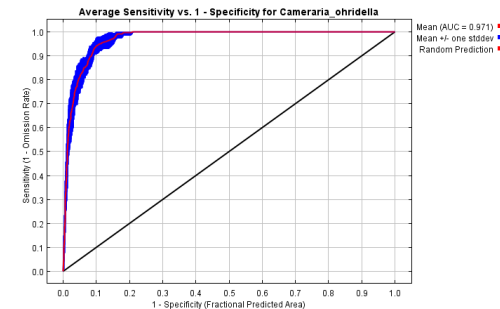
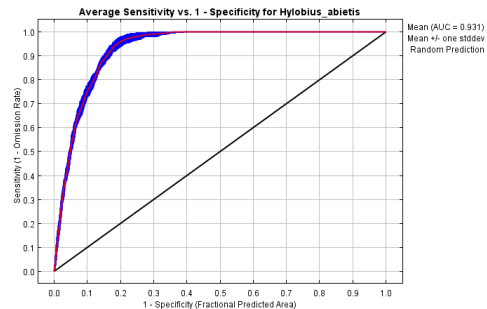
b=2.5 smooth model

	<i>Hylobius a.</i>	<i>Cameraria o.</i>
AUC:	0.931	0.971
AUC:	0.928	0.969

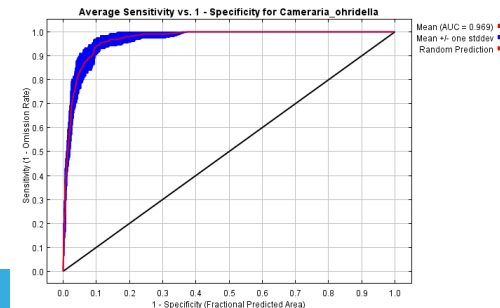
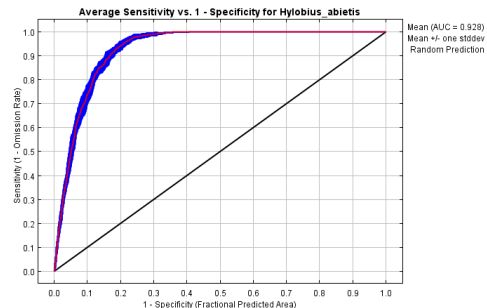
Hylobius abietis

Cameraria ohridella

b=1.0



b=2.5

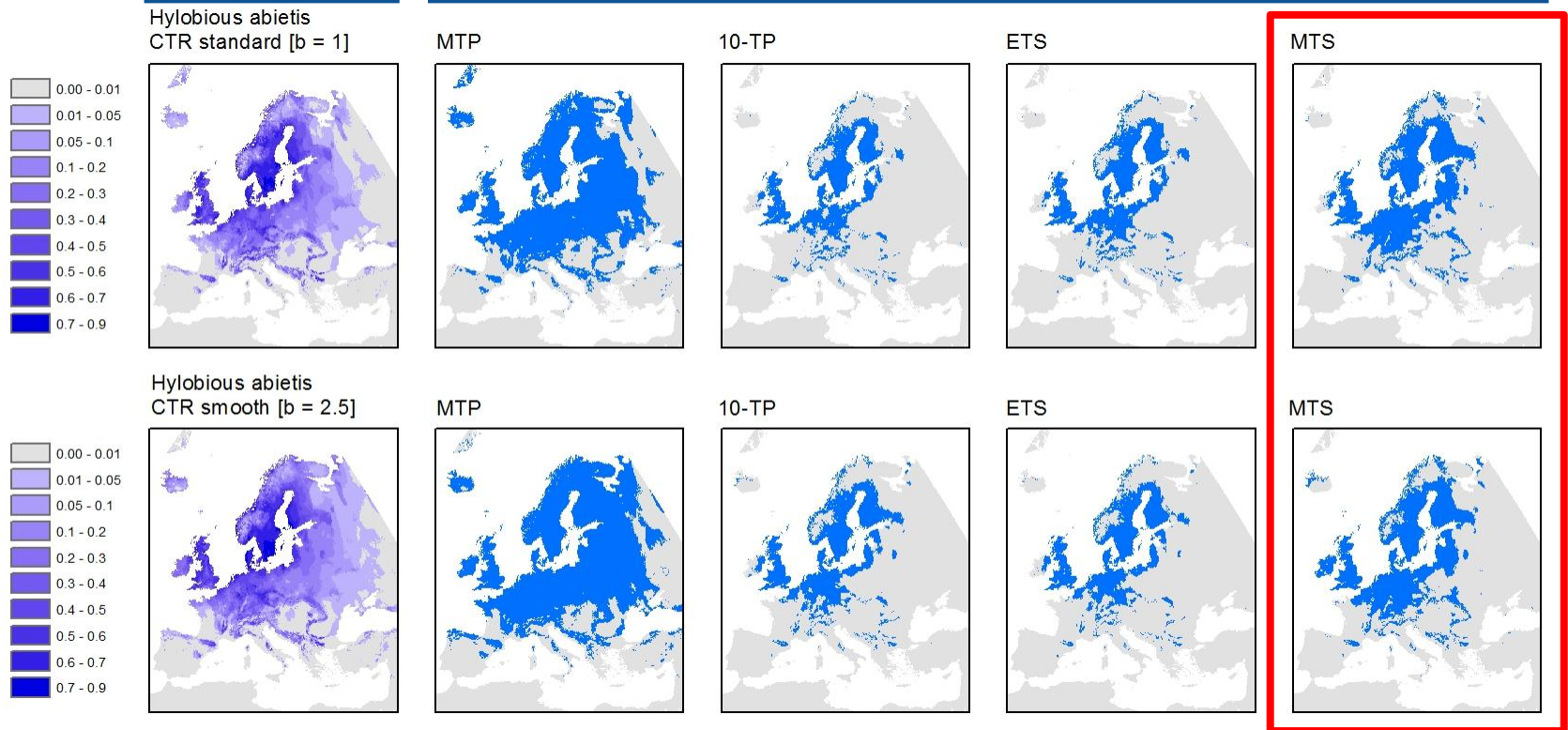


Large Pine Weevil (*Hylobius abietis*)

Simulated habitat – Current climate

Probabilistic map

Thresholds [0, 1]



MTP: Minimum training presence

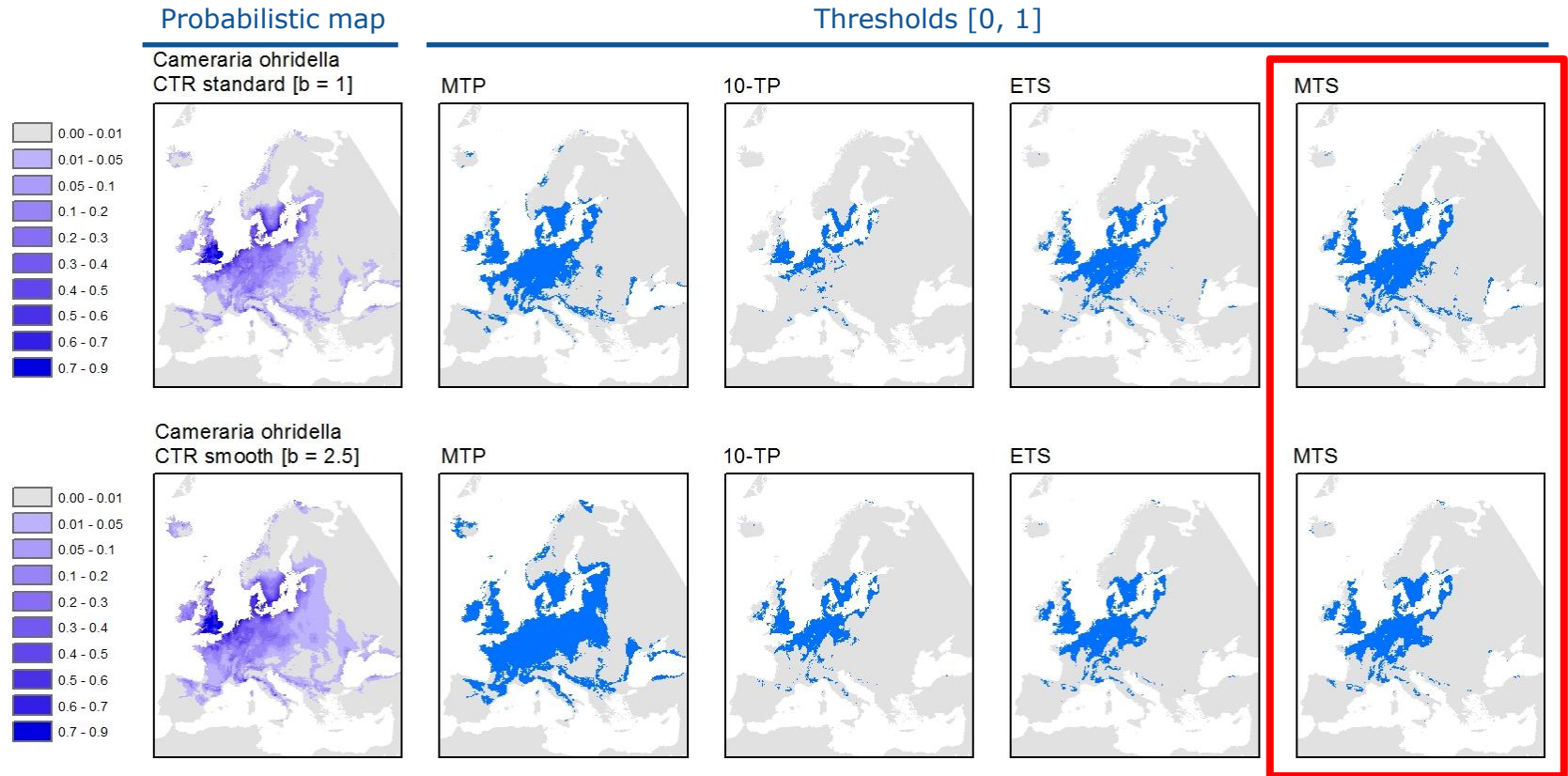
10-TP: 10th percentile training presence

ETS: Equal test sensitivity and specificity

MTS: Maximum test sensitivity plus specificity (used as reference)

Horse Chestnut Leaf Miner (*Cameraria ohridella*)

Simulated habitat – Current climate



MTP: Minimum training presence

10-TP: 10th percentile training presence

ETS: Equal test sensitivity and specificity

MTS: Maximum test sensitivity plus specificity (used as reference)

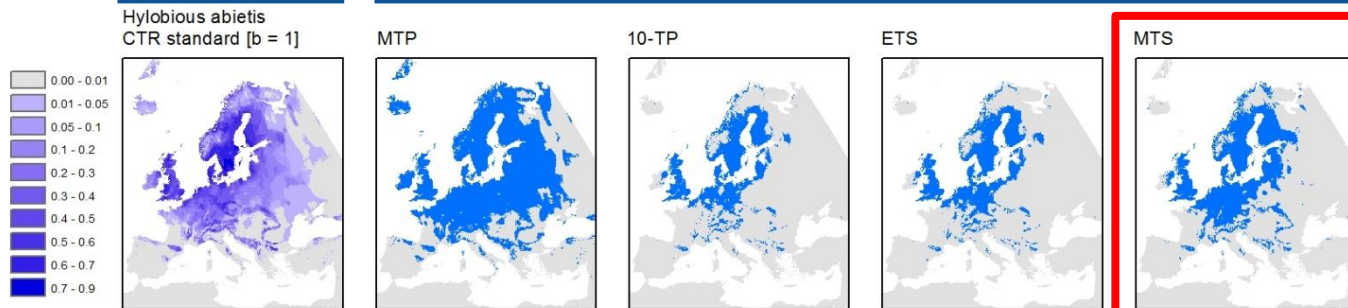
Large Pine Weevil (*Hylobius abietis*)

Simulated habitat and future scenarios

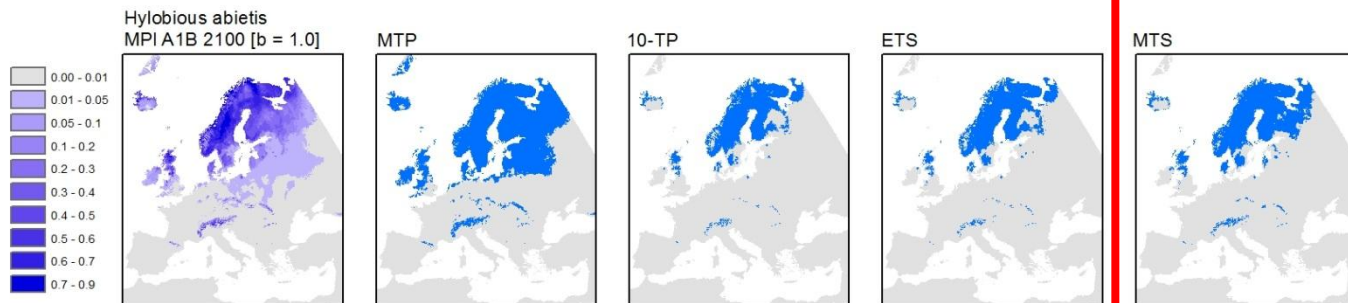
Probabilistic map

Thresholds [0, 1]

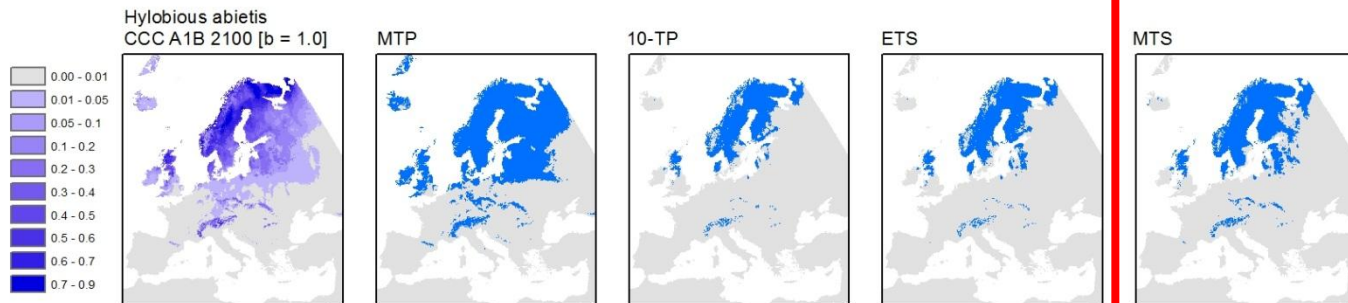
Current
climate



A1B 2100
MPI



A1B 2100
CCC



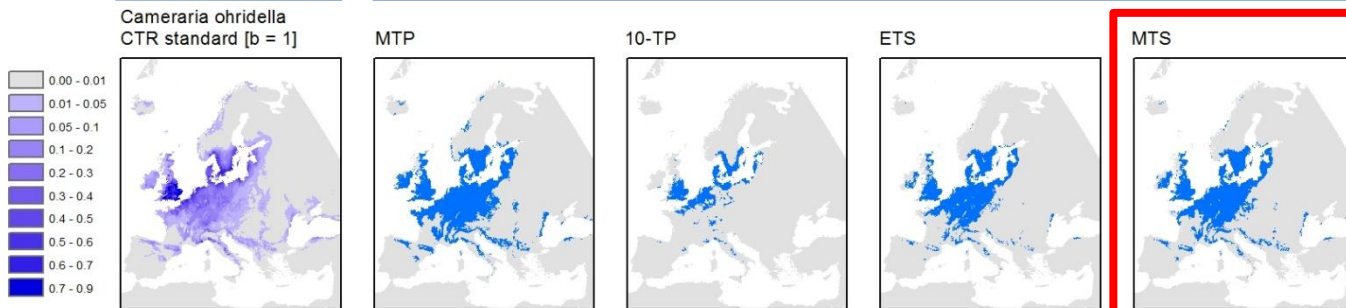
Horse Chestnut Leaf Miner (*Cameraria ohridella*)

Simulated habitat and future scenarios

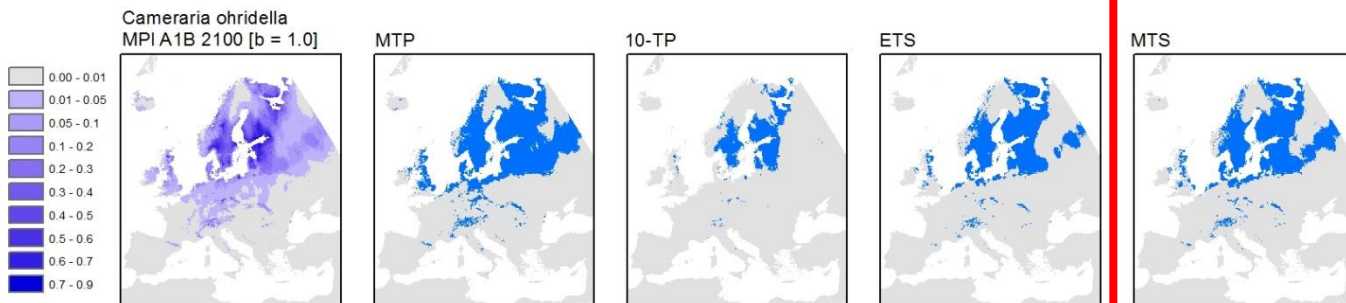
Probabilistic map

Thresholds [0, 1]

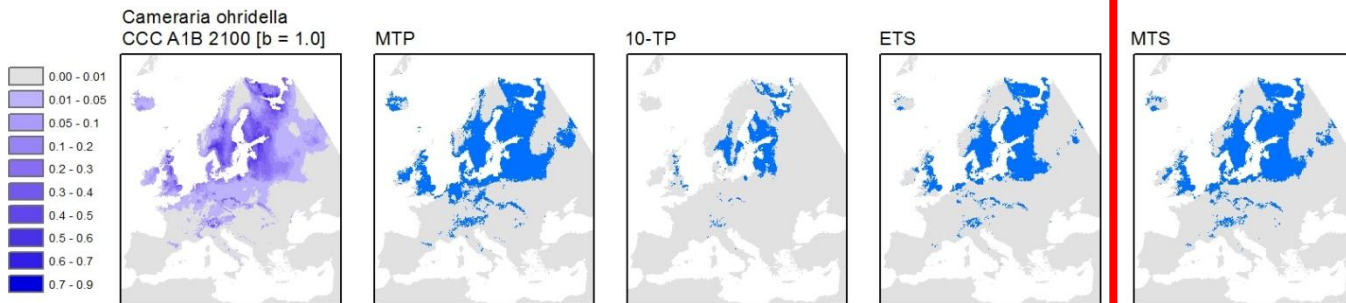
Current
climate



A1B 2100
MPI

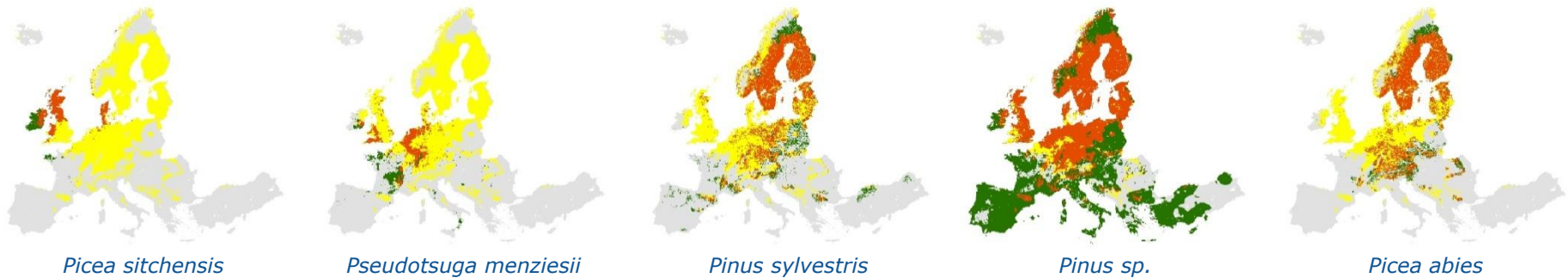


A1B 2100
CCC

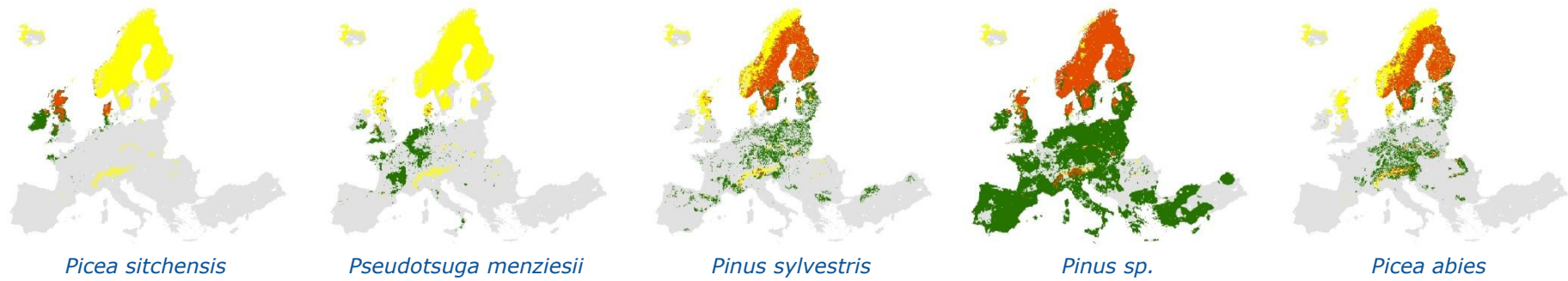


***Hylobius abietis*: current vs A1B-MPI 2100 habitat and current host tree species**

***Hylobius abietis* – Current habitat (MTS) on host tree species domain (10-TP)**



***Hylobious abietis* – A1B-MPI 2100 habitat (MTS) on current host tree species domain (10-TP)**



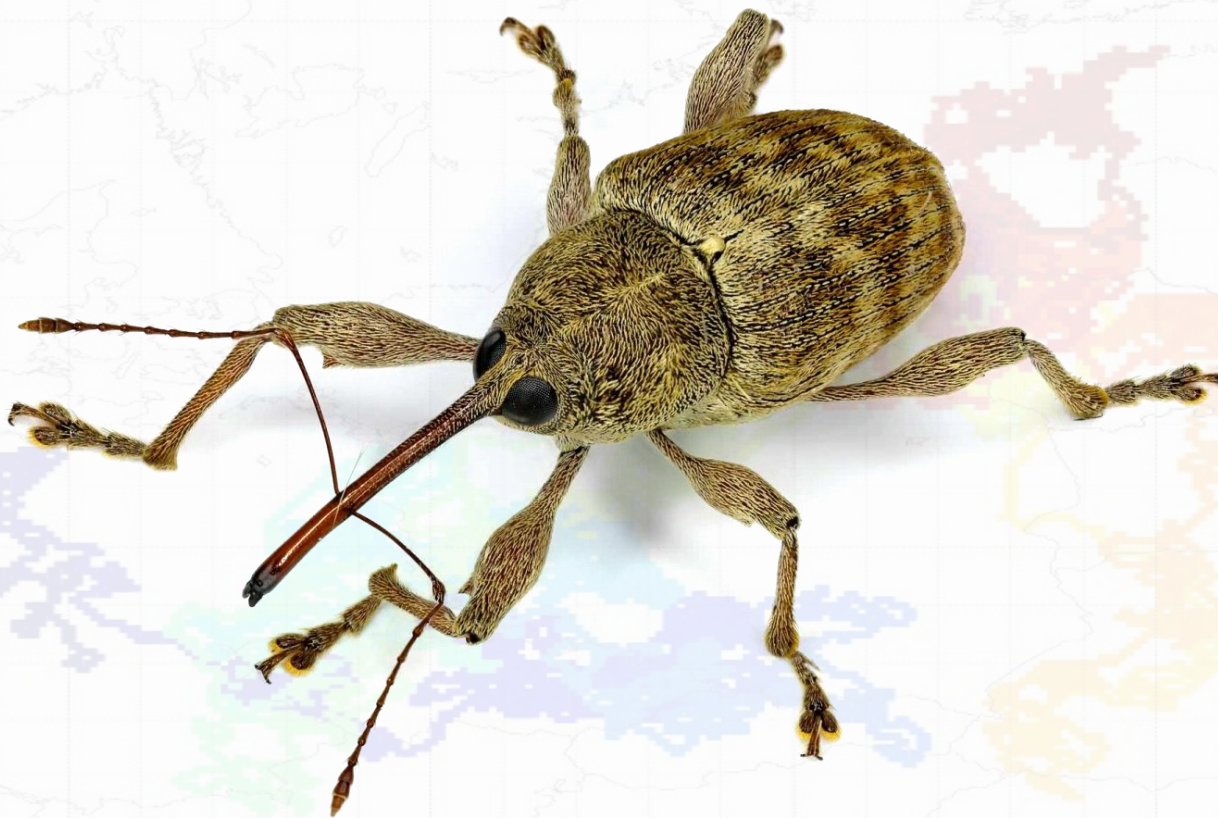
- Hylobius abietis*: habitat yes / host no
- Hylobius abietis*: habitat yes / host yes
- Hylobius abietis*: habitat no / host yes

Conclusions

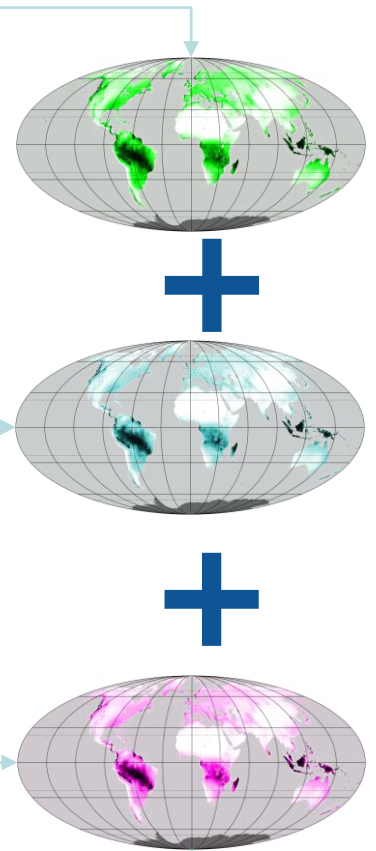
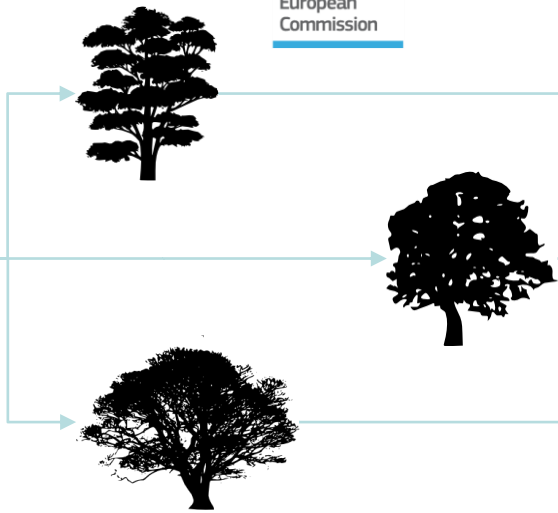
- GBIF data limitations: coordinate precision, time of observation, amount of observations, sampling bias, lack of data for most insect pests
- Uncertainty (many sources): model, climate data, pests data
- More climate runs/models, bias correction, ensemble approach
- Suitable habitats for *Hylobius abietis* will decrease under future climate conditions ...
- **Need of a public database on forest pests:** georeferenced data of observed presences, X and Y coordinates, standard format, open dissemination, metadata, Country/Region input data, other providers...



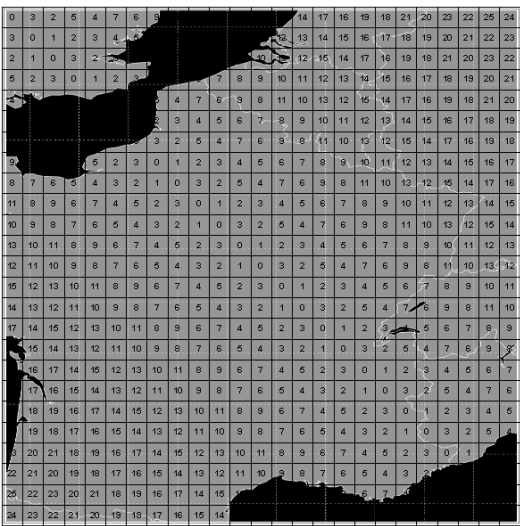
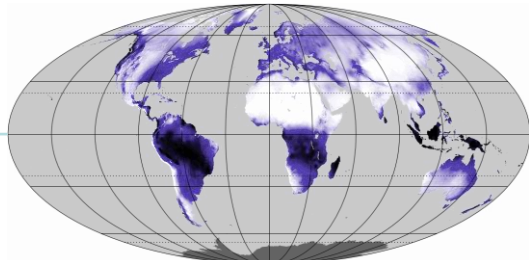
Pest Spread Dynamic Modelling



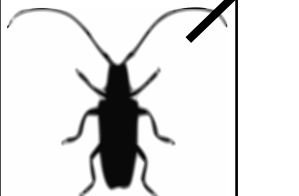
- 1) The user selects a pest
- 2) Niches of known hosts for the selected pest are combined in order to model the pest niche
- 3) The pest niche is then used to identify local environmental suitability
- 4) The user selects a starting point for the spread and setups a parameters about pest dispersal ability
- 5) A dynamic model based on pest movements through suitable area is then built



Modelling Pest Niche

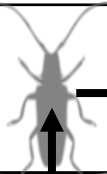
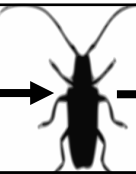
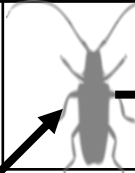
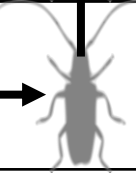



Modelling Pest Spread

0	2	5
1		2
0	3	1

1) User selects a starting point

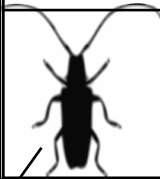

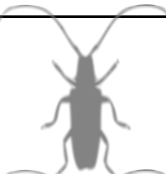
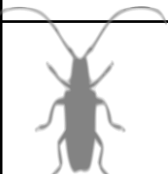

2) The pest moves towards the most suitable grid cells

1	0	0	1	1	0
1	2	2	1	5	0
0	2	1			5
0	2			2	5
1		2	1	3	2
0	3	1	0	3	1

The pest does not move here, Because the other cell with value 5 is surrounded by cells with higher Suitability.

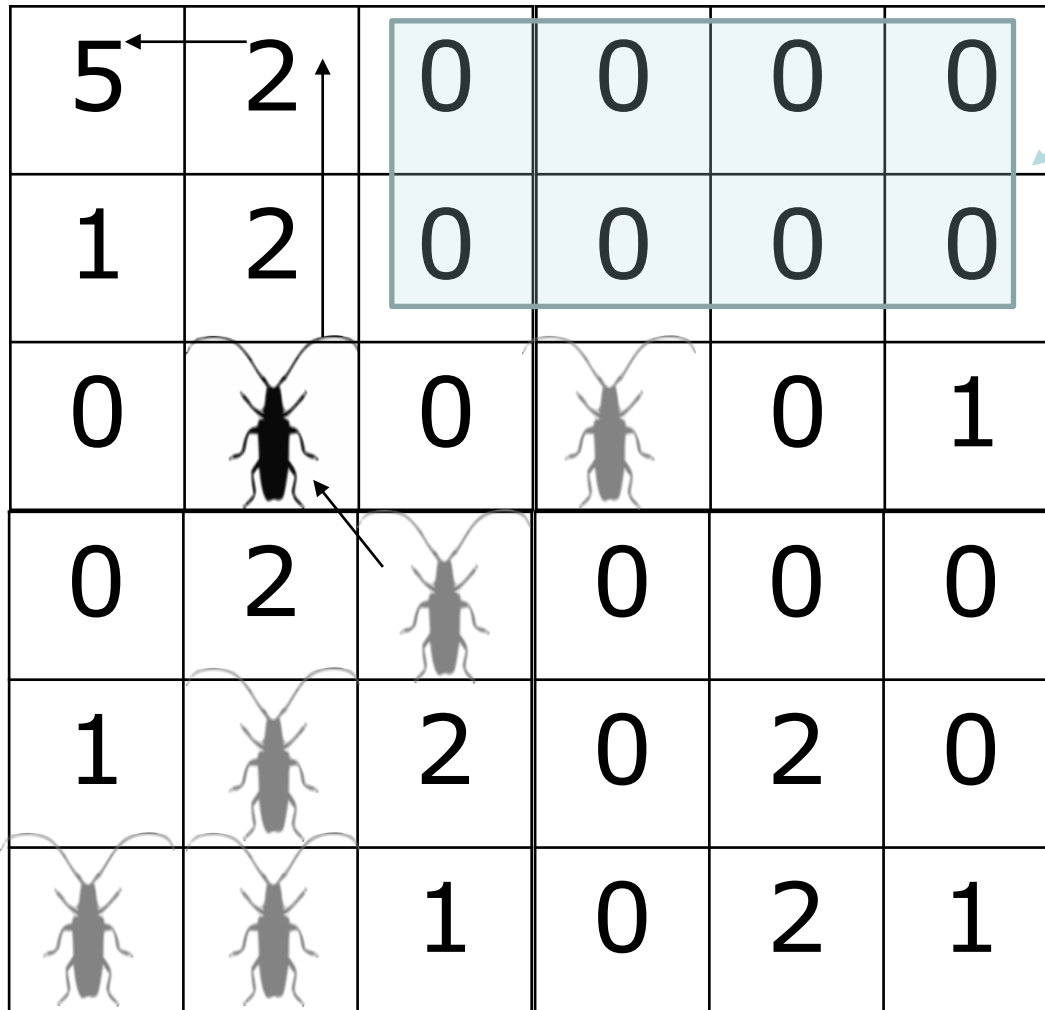
If there are several cells with equal values:

- 1) Overall suitability is evaluated in the area closer to these cells
- 2) The pest moves towards the most suitable areas

5	2	0	0	0	0
1	2	0	0	0	0
0	3	0		0	1
0	2		0	0	0
1		2	0	2	0
		1	0	2	1

This area is not suitable for pest dispersal; thus, it steps back and tries to spread through a different pathway

If the pest gets stuck in an unsuitable area, it steps back and searches for different spread pathways.

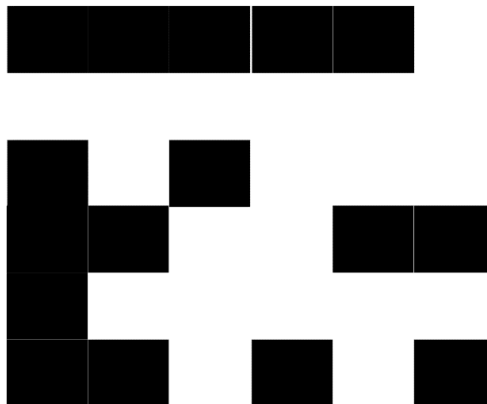
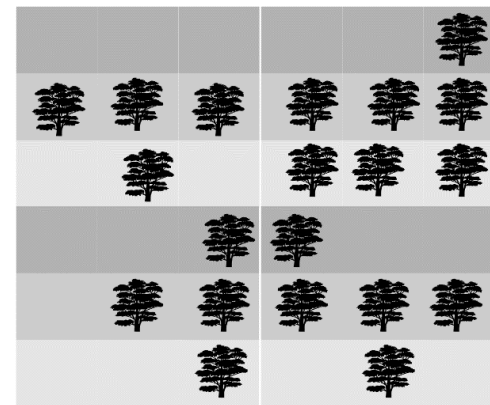


This area is not suitable for pest dispersal; thus, it steps back and tries to spread through a different pathway

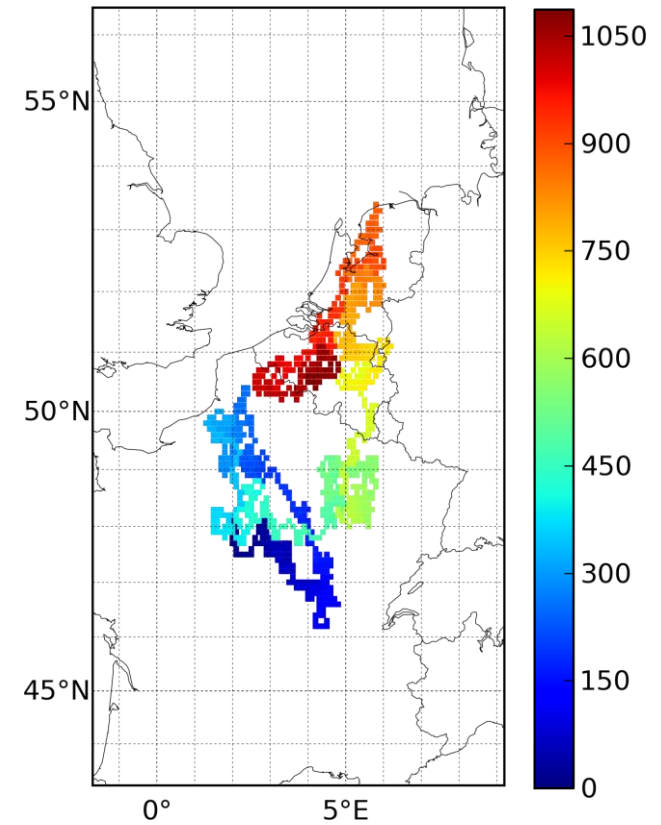
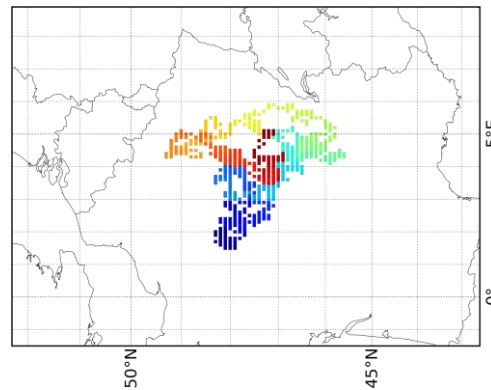
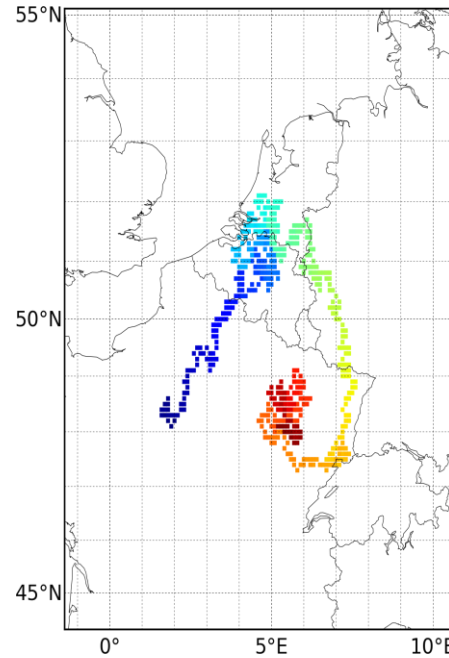
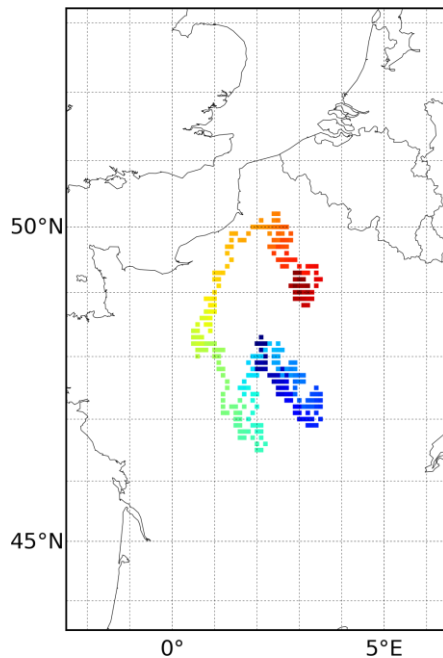
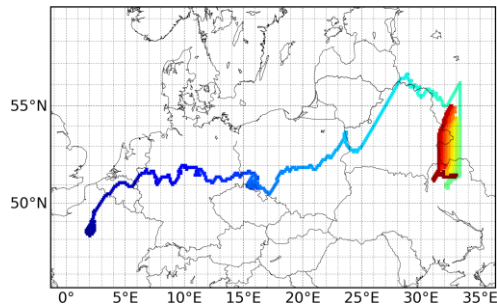
If the pest gets stuck in an unsuitable area, it steps back and searches for different spread pathways.

Constrain to Host Distribution

1	0	0	1	1	0
1	2	2	1	5	0
0	2	1	5	3	5
0	2	1	0	2	5
1	1	2	1	3	2
0	3	1	0	3	1



					0
1	2	2	1	5	0
	2		5	3	5
		1	0		
	1	2	1	3	2
		1		3	



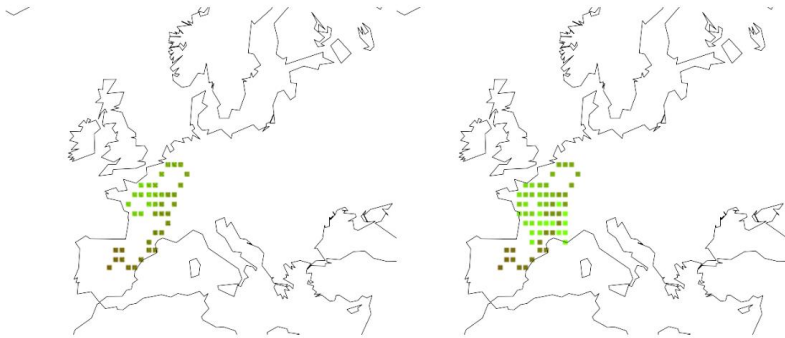
Color map indicates
model step sequence
(i.e. time scale)



Alternaria mali

Tree Pest Dispersion Model

Estimated Area of Occupancy (number of 1 lat x 1 lon grid cells): 59.



Known Hosts:

Malus domestica



Two software versions
will be available from
FISE:

- 1) Low spatial resolution
online demo version
- 2) High spatial resolution
static version (both as a
Python library and as a
standalone executable)

Thank you

Insect pests sample data (GBIF)

- Large Pine Weevil (*Hylobius abietis*)
 - The Large pine weevil is economically the most important pest of coniferous forest regeneration in Europe. Planted seedlings are frequently damaged or killed by adult weevils feeding on stem bark (SLU)
 - Heavy damage can completely girdle stems and cause plant death (IMPACT Project)
 - Adult weevils have a broad host range, feeding on a wide variety of conifer and broadleaved trees, but pine is preferred as a food source
 - Moderate concern (not EPPO)
- Horse Chestnut Leaf Miner (*Cameraria ohridella*)
 - It was concluded that due to rapid and natural spread of the pest, no practical phytosanitary measures can be taken. In 2001 it was therefore removed from the EPPO Alert List (less concern)
 - Damage: Mines in the leaves. Heavy infestations lead to brown discoloration and death of the leaves, and finally defoliation of the tree



Climate scenarios

- IPCC SRES A1B scenario
- Future world of very rapid economic growth and a balanced share between fossil and non-fossil energy sources
- Often considered BAU scenario

A1B: Temperature change in the 2071-2100 period, compared to the 1961-1990 period (°C)

	Reference	Reference variant 1	Reference variant 2
Northern Europe	3.8	4.8	3.4
UK & Ireland	2.1	2.9	1.7
Central Europe north	2.8	3.7	2.0
Central Europe south	3.0	3.8	2.0
Southern Europe	3.2	3.7	2.4
EU	3.1	3.9	2.4

PESETA II Project: http://peseta.jrc.ec.europa.eu/climate_scenarios.html

Aesculus hippocastanum

