

Effects of landscape spatio-temporal heterogeneity and of food web structure on biological control for pest regulation

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Landscape complexity



Agricultural landscapes vary from **structurally simple** landscapes with **one or two cropping systems** to **complex mosaics of diverse cultivated fields** embedded in a **natural habitat** matrix.

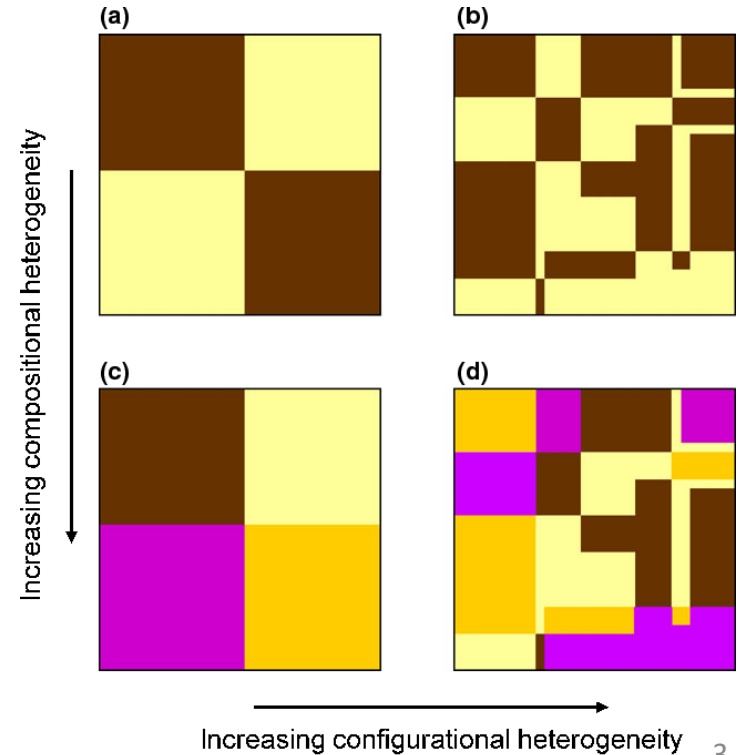
(Power, 2010)

Landscape complexity



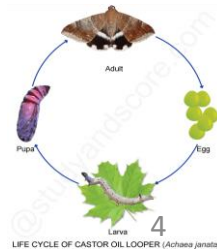
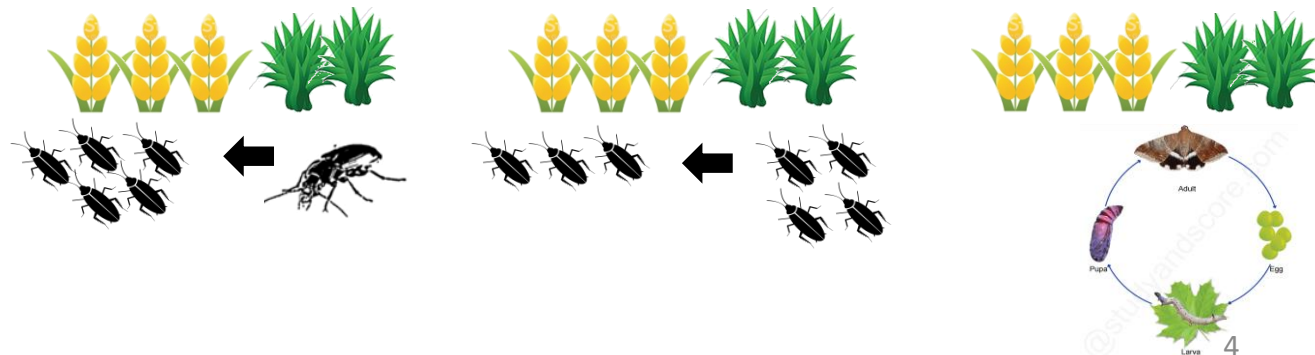
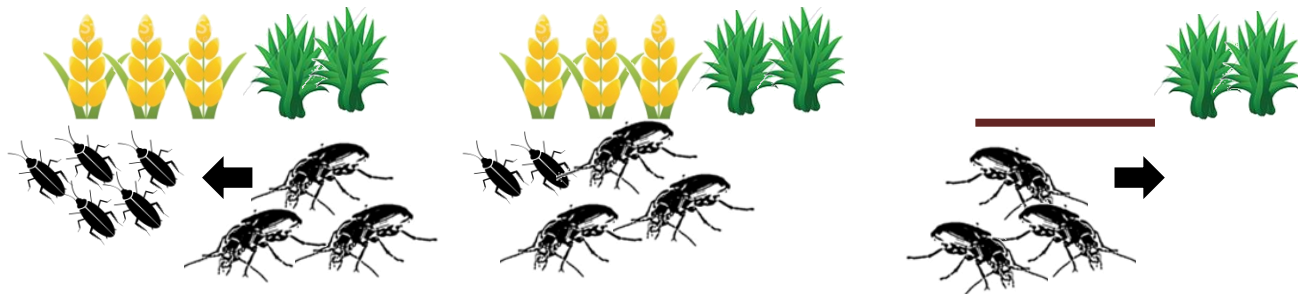
as

Landscape heterogeneity



(Fahrig et al. 2011)

However, the nexus among **agricultural landscape structure** and **the species trophic interactions** is not trivial:

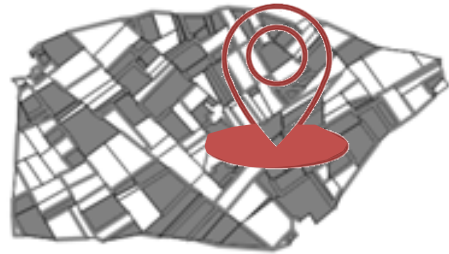


Objectives:

1) Couple **landscape features** and **species traits** to provide insights on **biological control outcomes** at landscape scale



2) To assess how the **landscape structure** is able to **locally** influence the **pest spatio-temporal** dynamic providing insight on **biological control**

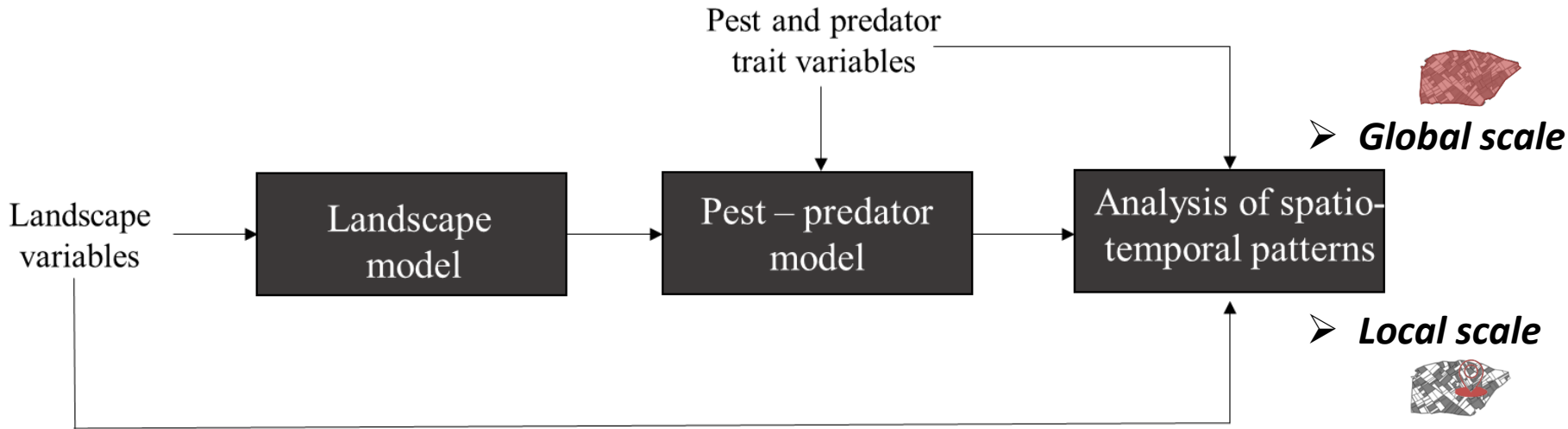


Stochastic-mechanistic prey-predator dynamic model:

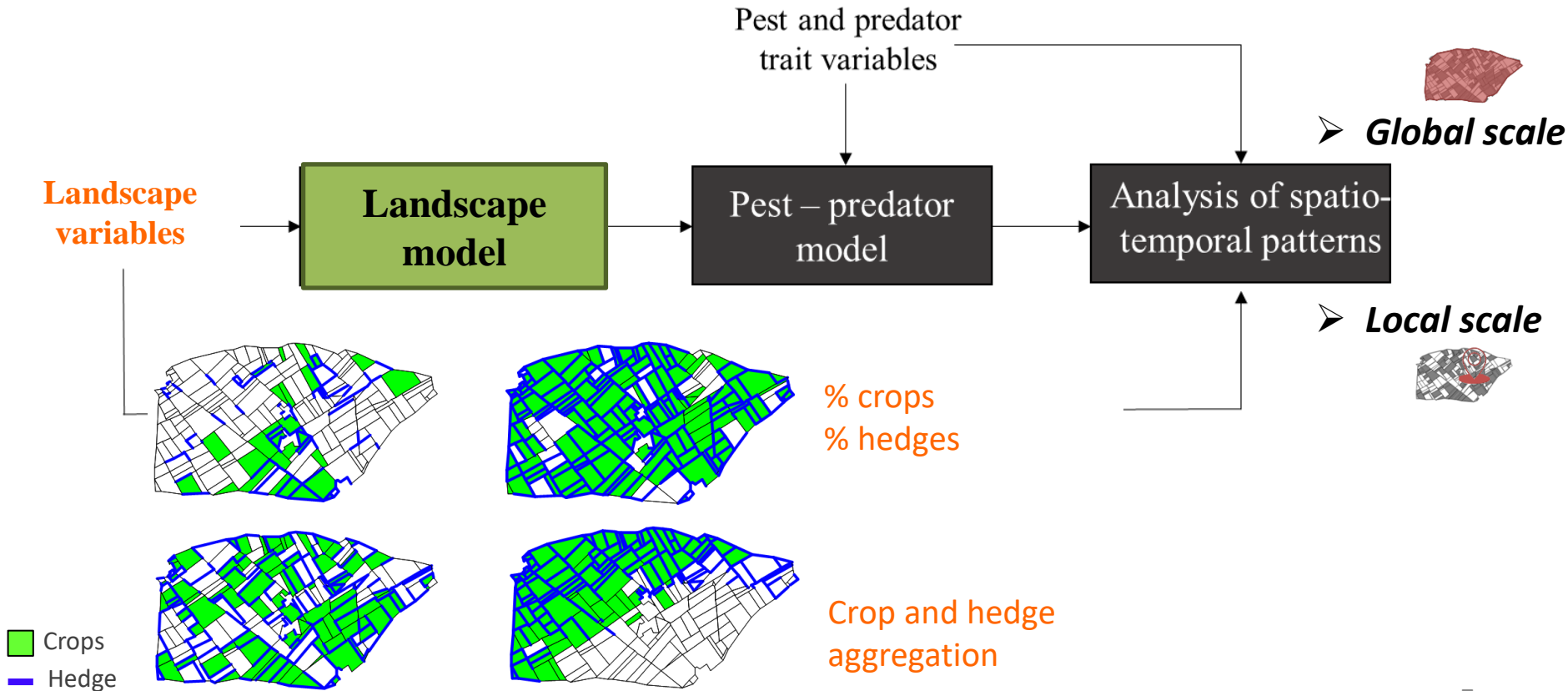
➤ *Global scale*

➤ *Local scale*

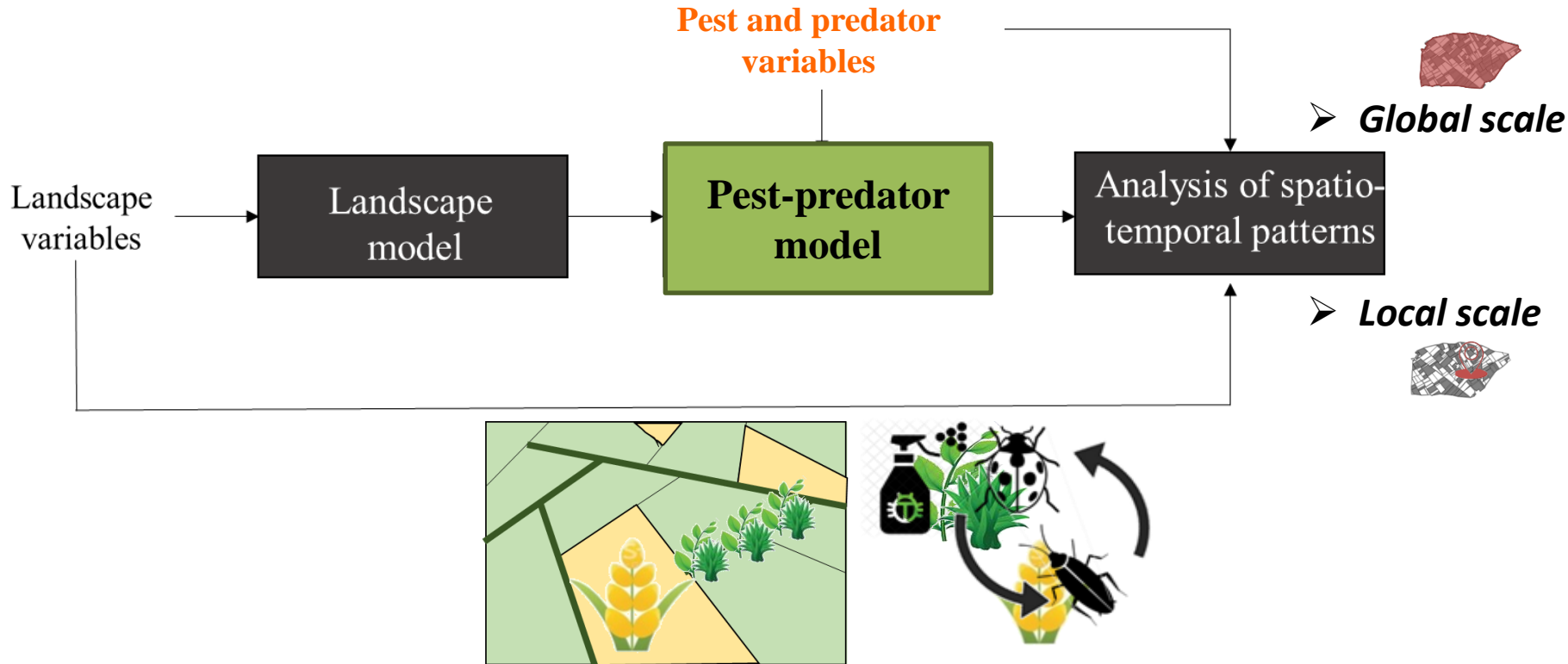
Modelling framework



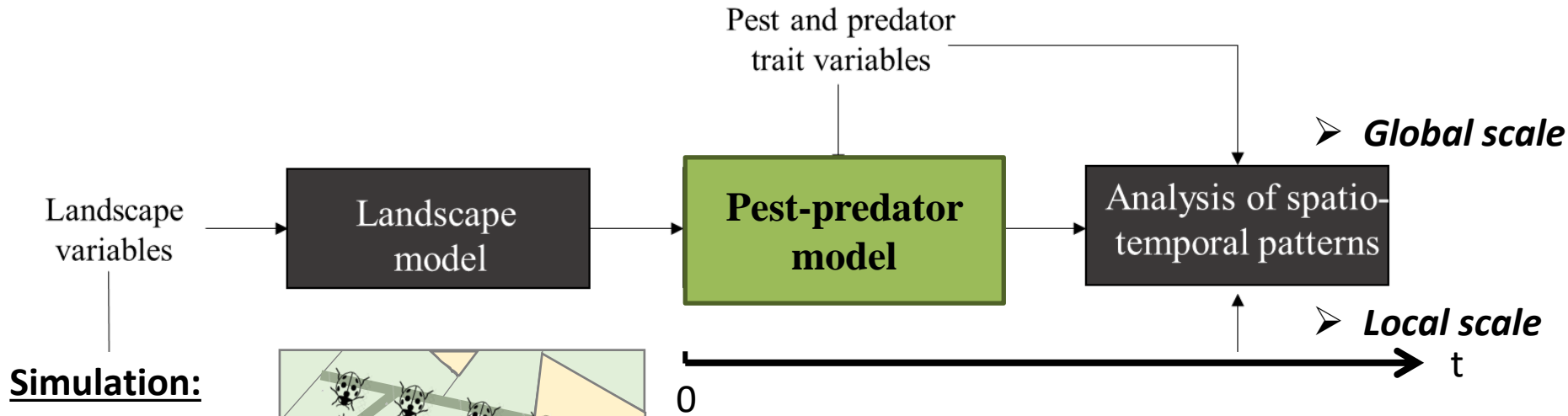
Modelling framework



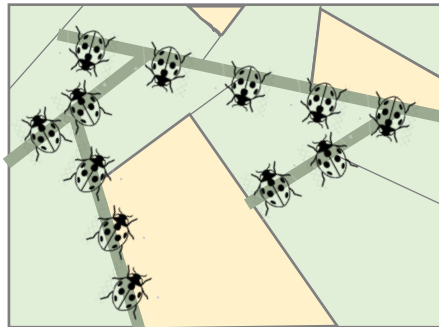
Modelling framework



Modelling framework



Simulation:



Linear elements h_i

— No hedge

— Hedge

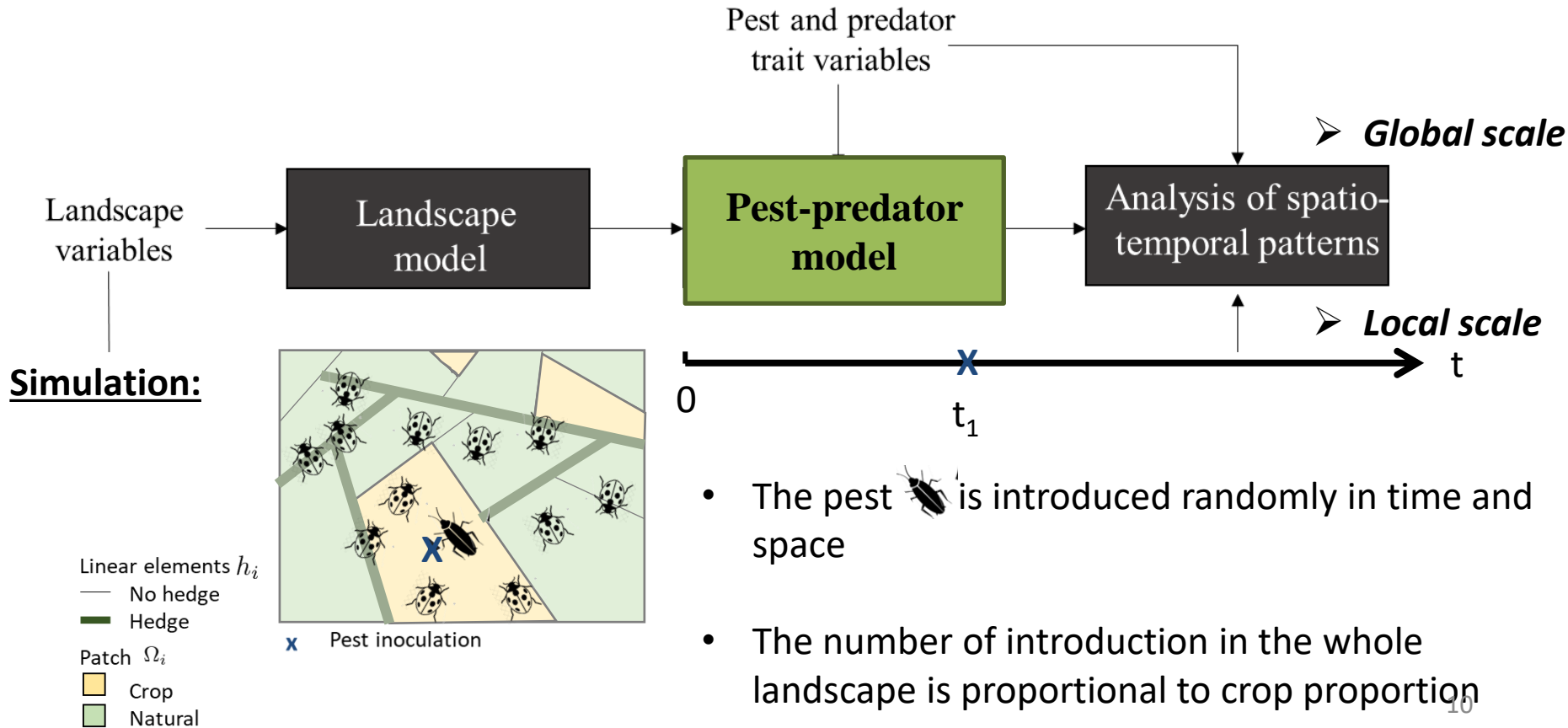
Patch Ω_i


■ Crop

■ Natural

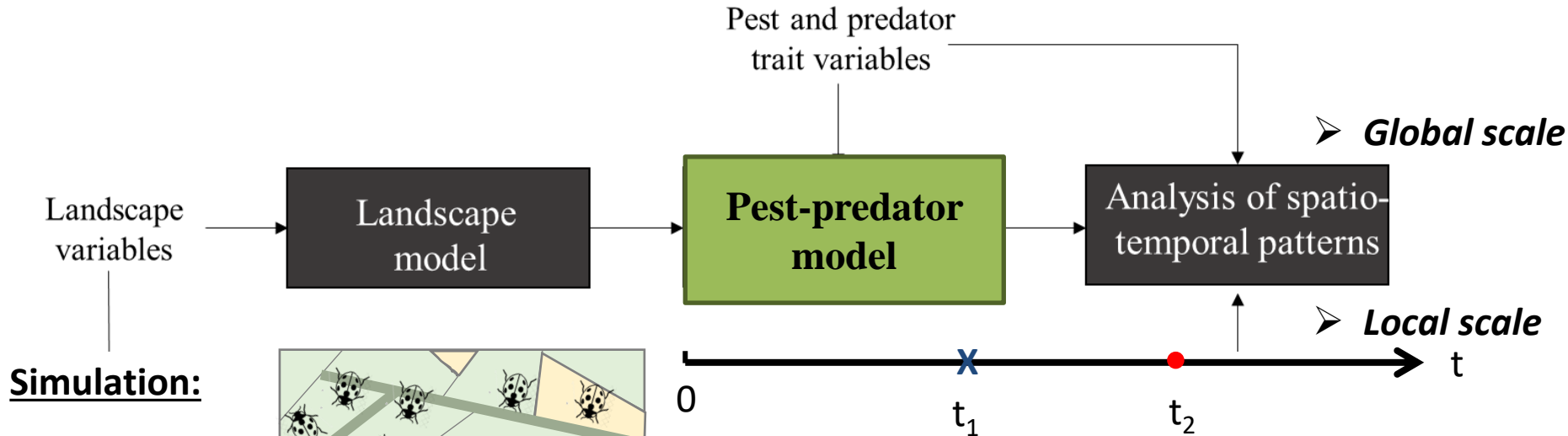
The predator  is present in all hedges at carrying capacity

Modelling framework

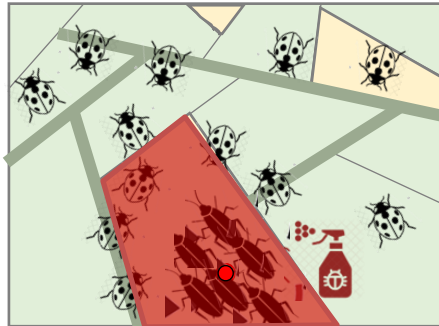


- The pest  is introduced randomly in time and space
- The number of introduction in the whole landscape is proportional to crop proportion

Modelling framework



Simulation:

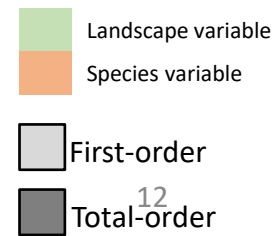
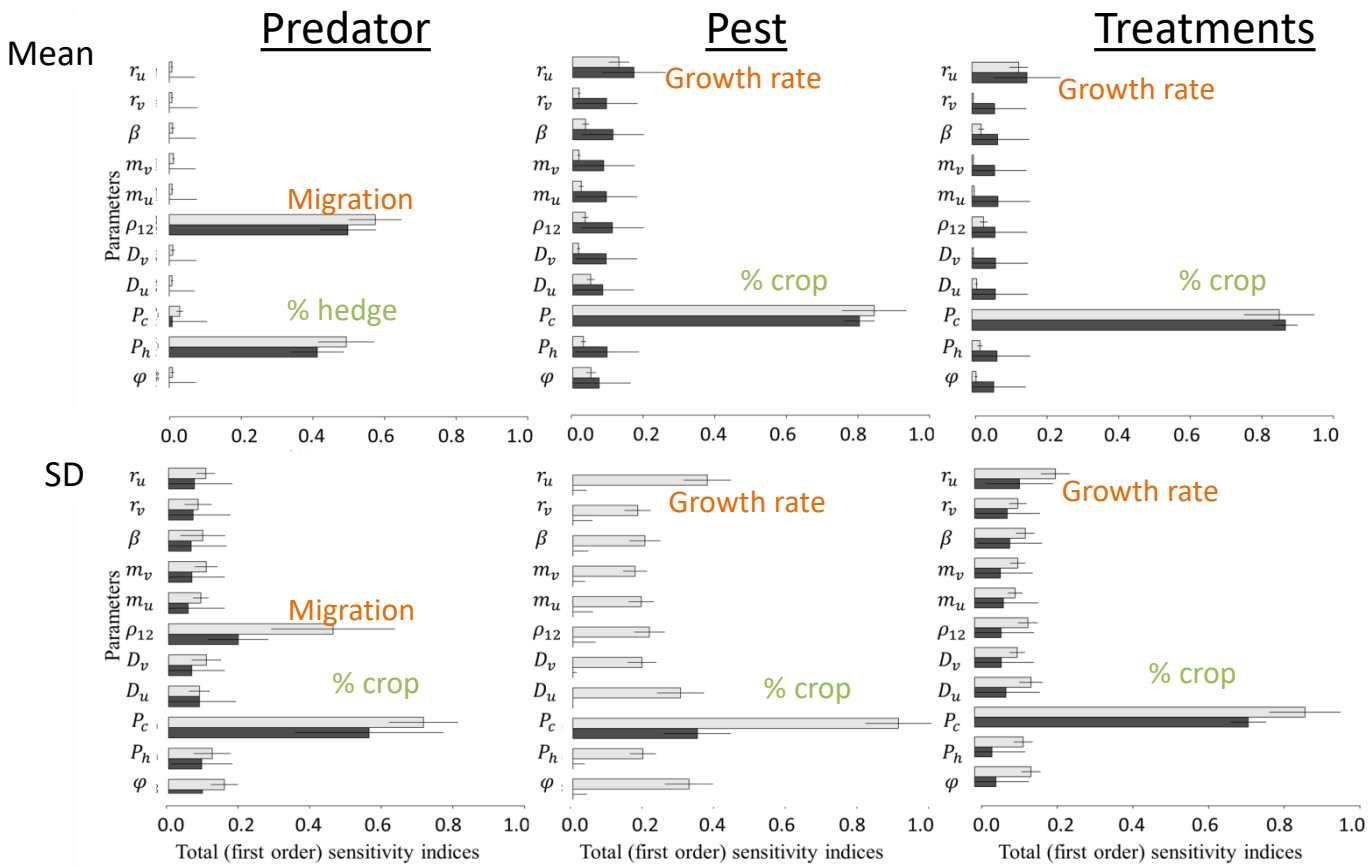


- Linear elements h_i
- No hedge
- Hedge
- Patch Ω_i
- Crop
- Natural

- x Pest inoculation
- Pest treatment

- A **pesticide treatment** is applied to a given crop field when the pest population in that field reach a give threshold
- The spatial location (x,y) and the magnitude of pest density at the moment (t) of treatment application is identified as output.

Sobol sensitivity analysis



Effects:



Predator



Predator migration
% Hedge
% Crop AND Hedge
% Crop AND Predator migration

% Crop
% Hedge AND Predator migration

Pest



% Crop
% Hedge
Population growth
% Crop AND Hedge
% Crop AND Predator migration

Prey diffusion
Predator migration
Predator predation
% Hedge AND Predator migration

Traitements



% Crop

of treatments

% Hedge
Predator migration

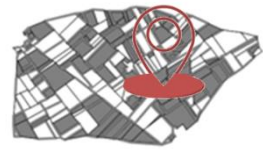
Presence of treatments

% Crop AND Hedge
% Crop AND Predator migration

Presence and number of treatments

Pest diffusion
% Hedge AND Predator migration

Similar landscapes with the same number of pest outbreaks, but with **different spatial distribution of pest density peaks**



% crop = 98%
% hedge = 0.22%
Aggregation = 4.96
N treatments = 10

% crop = 98%
% hedge = 0.35%
Aggregation = 3.22
N treatments = 10

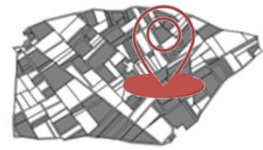


● Pest treatment

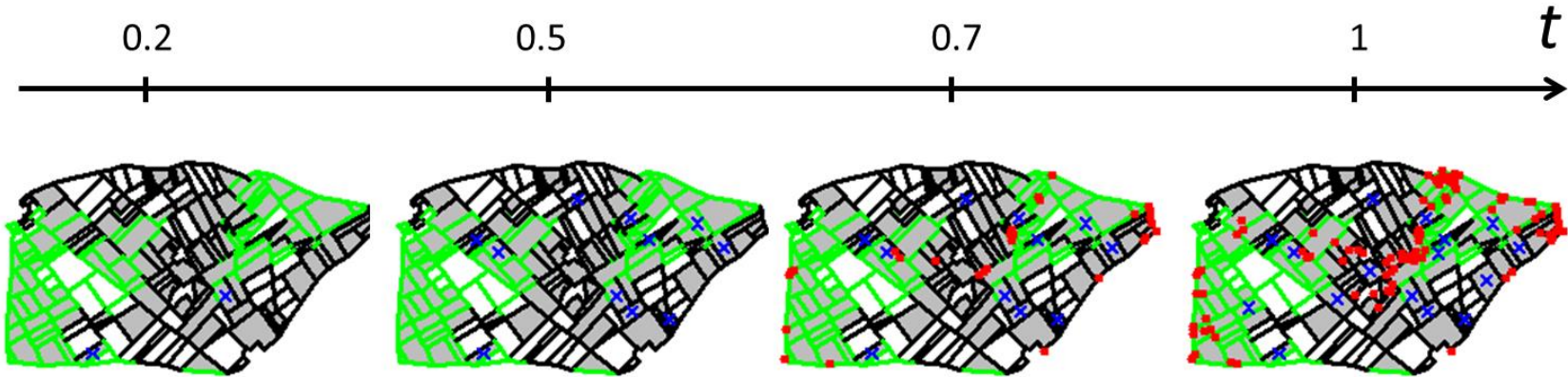
■ Culture

— Hedge

Method:



Point pattern as spatio-temporal point process



It is a collection X of pairs (s_i, t_i) , $i = 1, \dots, n$

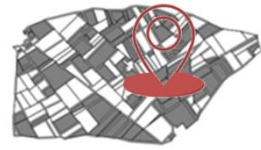
where s_i , t_i are the spatial location and time of occurrence associated with the i th event

We identify the spatio-temporal point patterns:

x pest inoculation locations

● pest peak locations

Spatio-temporal models in order to LOCALLY predict:



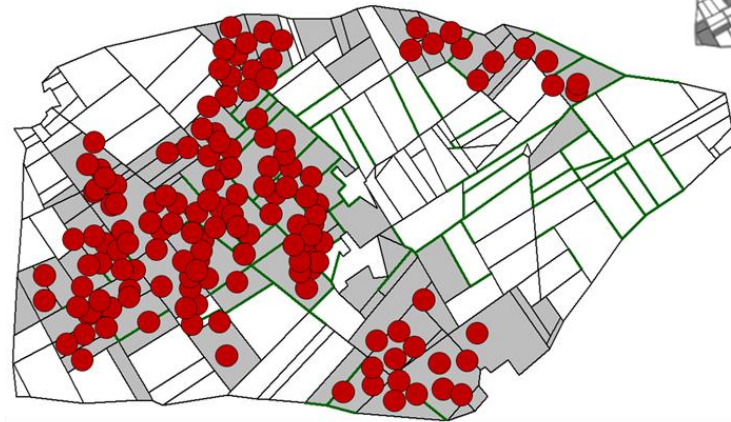
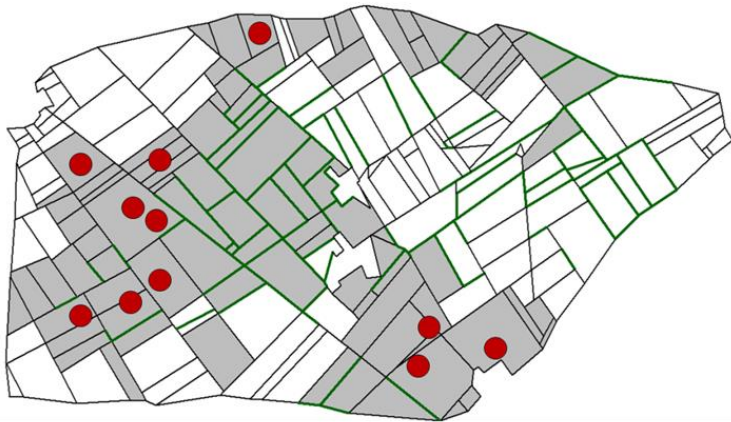
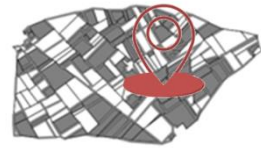
1) Number of pest density peak model $\lambda(s, t) \bullet \rightarrow \bullet \bullet \bullet \bullet$

$$\lambda(s, t) = f \left(\begin{array}{l} \text{Spatial} \\ \text{covariates} \end{array}, \begin{array}{l} \text{Spatio-temporal} \\ \text{covariates} \end{array}, \begin{array}{l} \text{Population dynamic} \\ \text{covariates} \end{array} \right)$$

2) Magnitude of pest density peak model $P_{max}(s, t) \bullet \rightarrow \bullet$

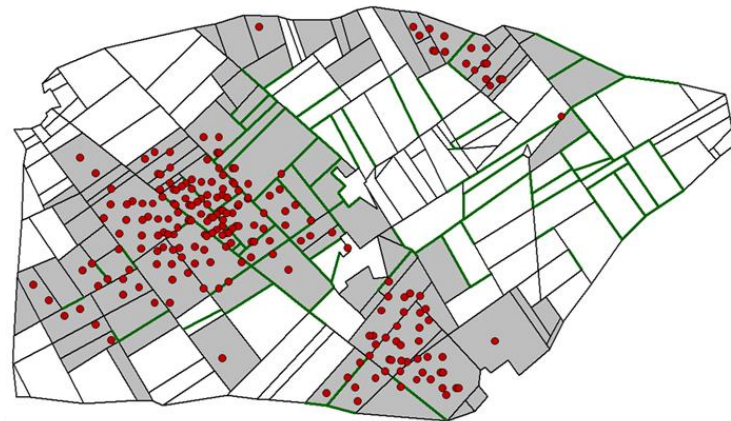
$$P_{max}(s, t) = f \left(\begin{array}{l} \text{Spatial} \\ \text{covariates} \end{array}, \begin{array}{l} \text{Spatio-temporal} \\ \text{covariates} \end{array}, \begin{array}{l} \text{Population dynamic} \\ \text{covariates} \end{array} \right)$$

2) Magnitude of pest peaks at t



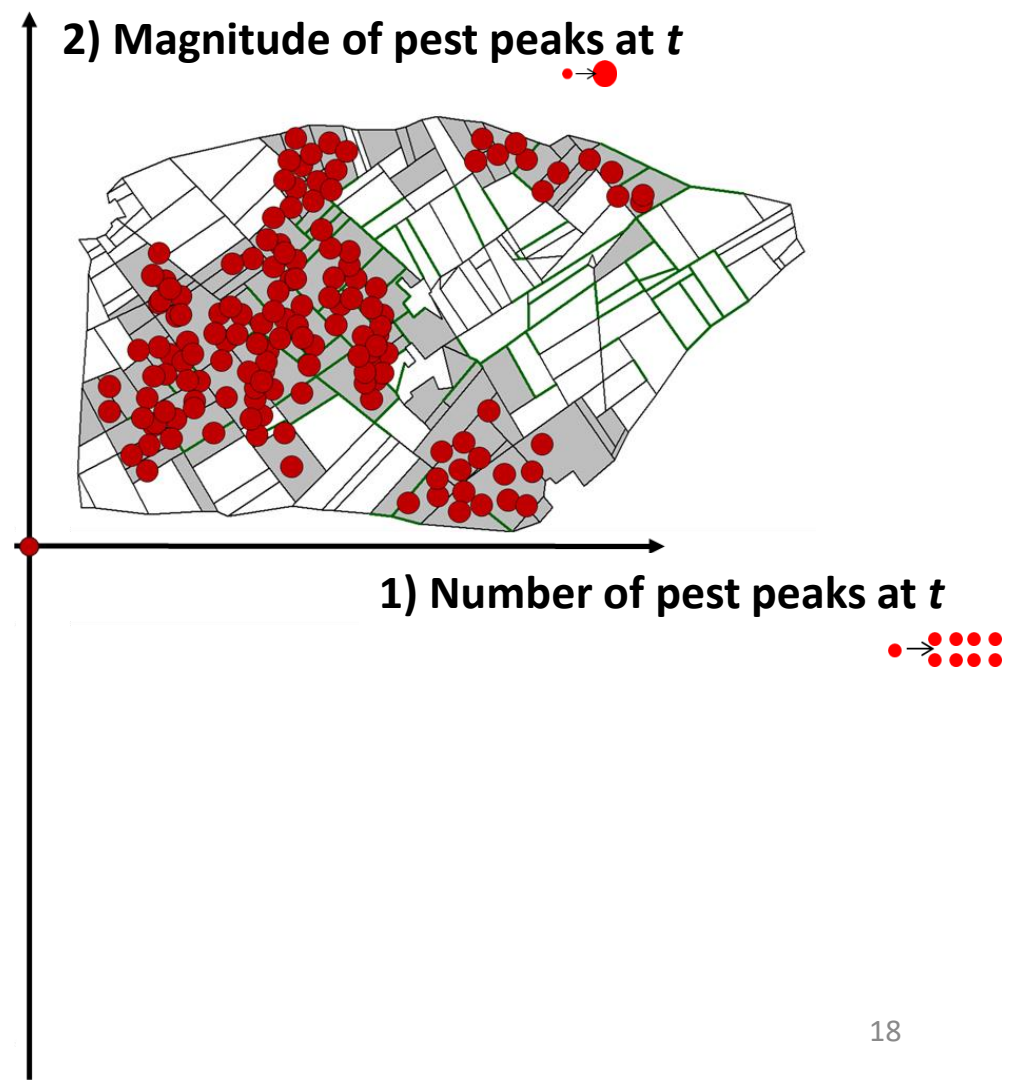
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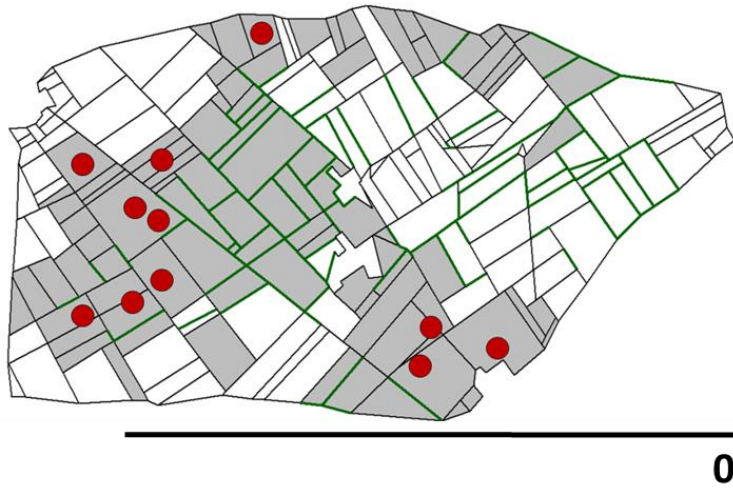
1) Number of pest peaks at t



These variables favour high number of pest peaks with elevated pest concentration.

- **Presence of peaks in previous time steps** in the same position or in the surrounding ones.
- An **elevated number of pest introductions** in the neighborings
 - **Among two patches**





These variables favour high pest concentration value, but in a low number. This happens when pests are clustered and limited in space, so they cannot diffuse, or they have a fast dynamic.

- An elevated **number of peaks in neighborings** only at one previous time step is not enough to lead to a high peak number at T.
- In the **middle of the patch** where the inoculation takes place.
 - Low **% culture**
- **Hedge proportion at landscape scale** can favour the max peak value since predator presence helps to keep pests just under the threshold, but if predators are missing in one spot pests increase rapidly and quickly reach a high density value.
- The **speed of predator spillover from hedge** mostly impacts the area close by the hedge.



These variables decrease the number of peaks and also the pest maximum value.

- Application of **pesticide treatments in same loactions or in close ones** at T-1
- The **local** presence of a **high % of hedge** in the buffer highligh a high predator presence in the local area that decrease pest peaks and their density value
- The **diffusion of predator** in culture negatively impact the pest dynamic
- A **high pest diffusion** favors diluition effect decreasing both peak value and peak number

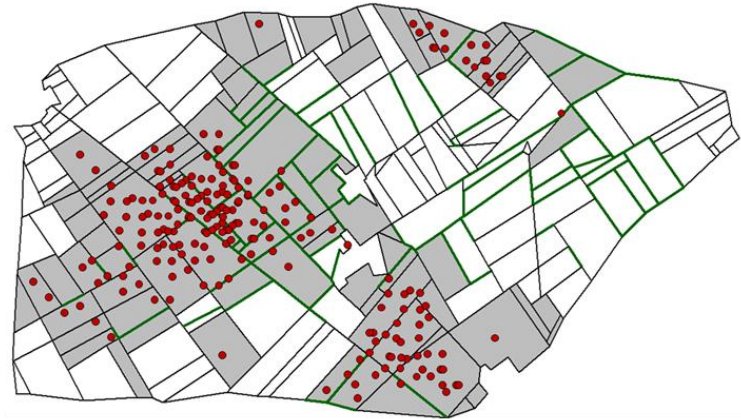
These variables leads to a high number of pest peaks, but not with an elevated pest value

- **% of crop coverage** that is natural pest habitat, so naturally favors pest peaks. However, those peaks are not characterised by a so high density due to the dilution effect.
- The **crop aggregation** contribute to cause pest spread phenomenon as said before.
- **Cells among more than 3 patches** usually host a high number of pest peaks due to high spillover. The maximum value of pest is negatively influenced since those cell are in the **periphery of the patch**.

2) Magnitude of pest peaks at t



1) Number of pest peaks at t



Take home messages:

GLOBALLY:



- **Landscape simplification** is the main driver of **pest population abundance** causing the highest number of pesticide treatments;
- The presence of **semi-natural stripes** enhances **predator presence**. However, the main positive effect on predator density results from its **ability of spilling-over from hedge to crop field** leading to an efficient biological control.
- **Different biological control outcomes** are obtained through the **combination** of both **landscape** and **species trait variables**.

***Stochastic-mechanistic
prey-predator
dynamic model***

Take home messages:

LOCALLY:



- A well **connected hedge network** could locally reduce pest density and treatment, due to the presence of pest predators;
- Local high **number** of pest peaks (1) is favoured by high pest introduction at previous time steps and situation that are the most suitable for pest dynamics, while they are decreased by dilution effect and predator presence or when treatment are previously applied;
- Local high **magnitude** of pest (2) is favoured by pest introduction and situations that constrain pests in a limited area, pest fast dynamics, low predator presence.


***Stochastic-mechanistic
prey-predator
dynamic model***



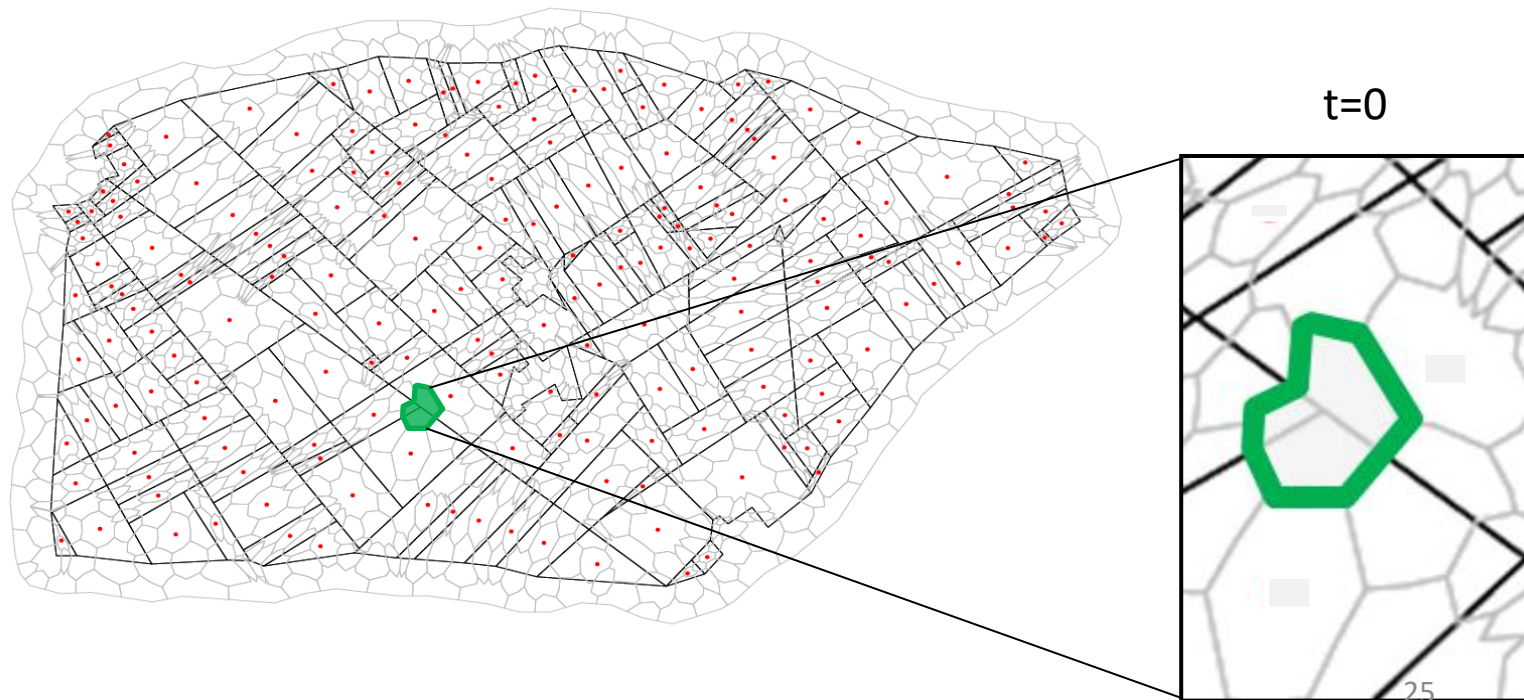
Thank you for your attention!

Spatio-temporal point process

The study region is partitioned into cells to capture the **local temporal dynamic** by models that:


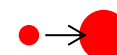
1) Count the **number** of pest peak 

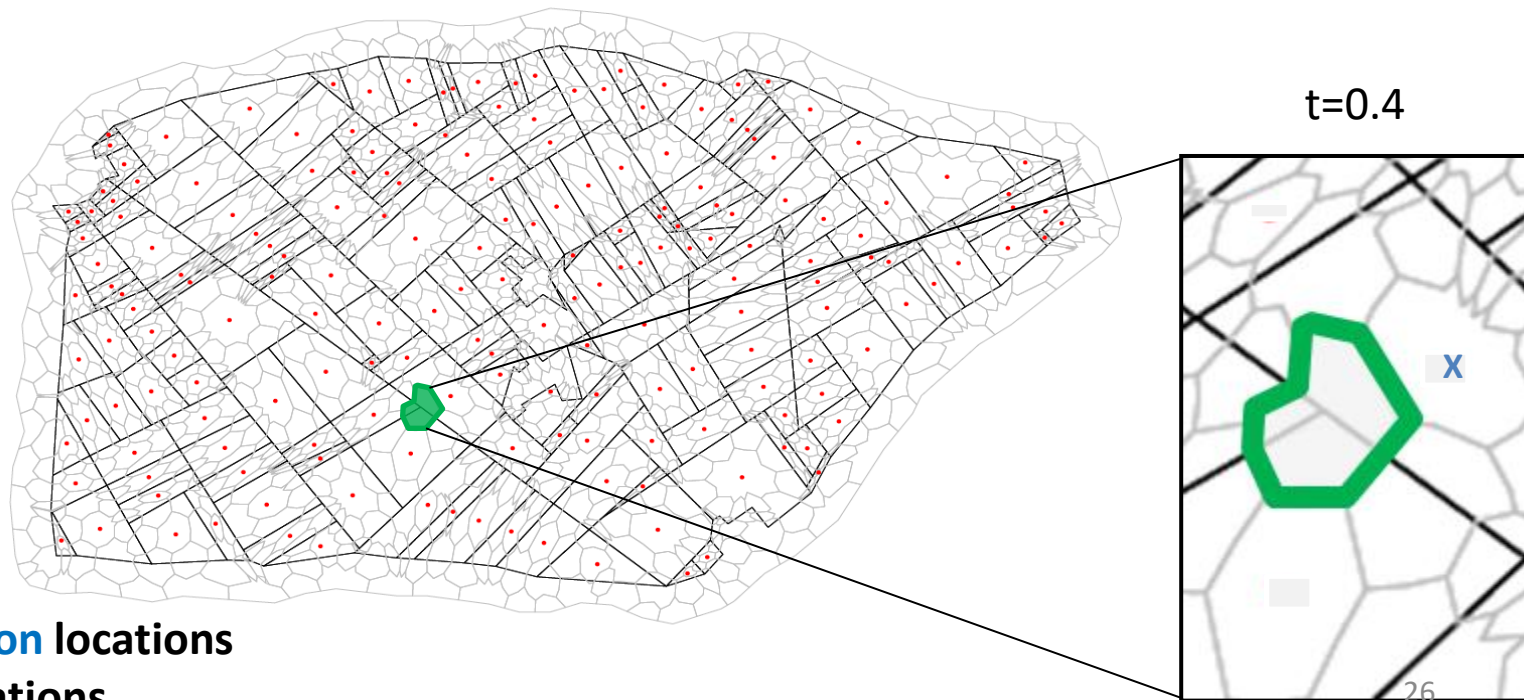
2) Consider the **magnitude** of pest peak 



Spatio-temporal point process


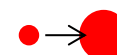
The study region is partitioned into cells to capture the **local temporal dynamic** by models that:

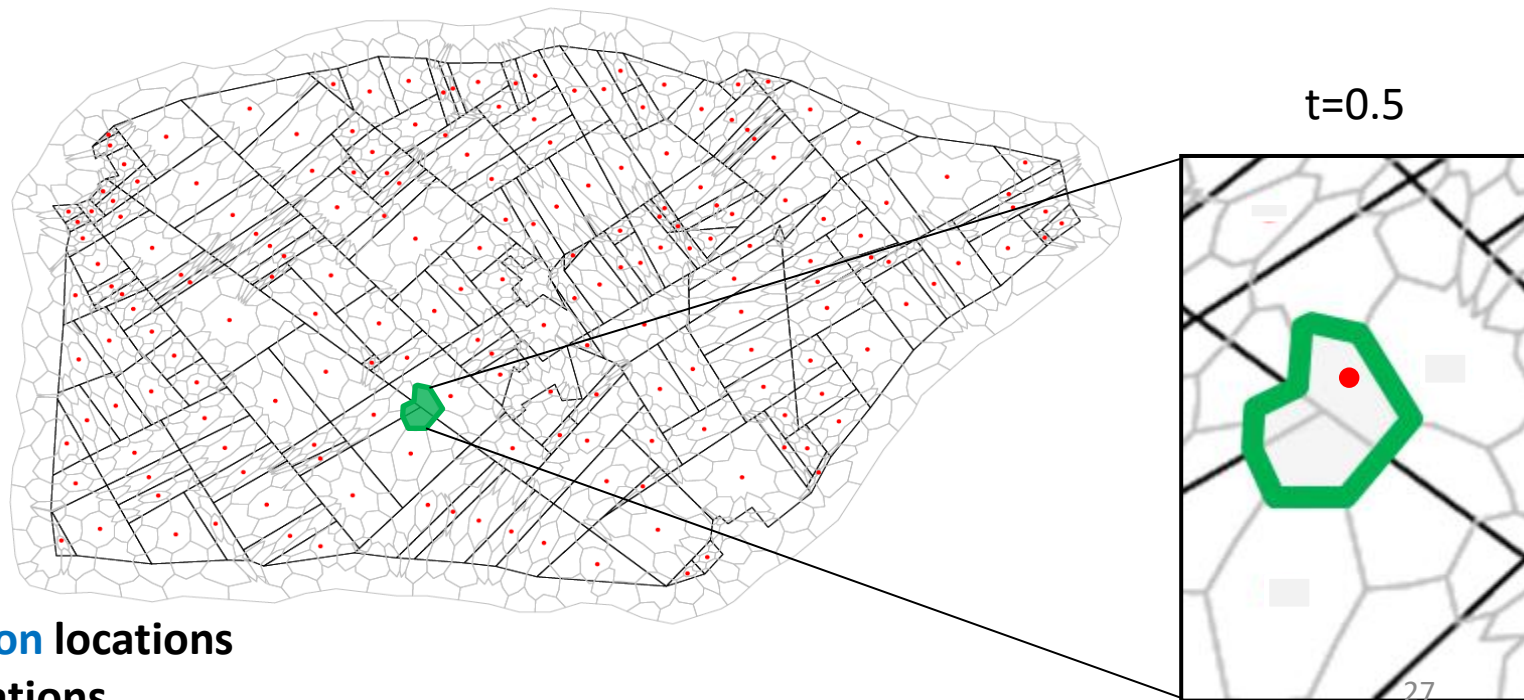
- 1) Count the **number** of pest peak 
- 2) Consider the **magnitude** of pest peak 



Spatio-temporal point process

The study region is partitioned into cells to capture the **local temporal dynamic** by models that:


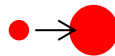
- 1) Count the **number** of pest peak 
- 2) Consider the **magnitude** of pest peak 

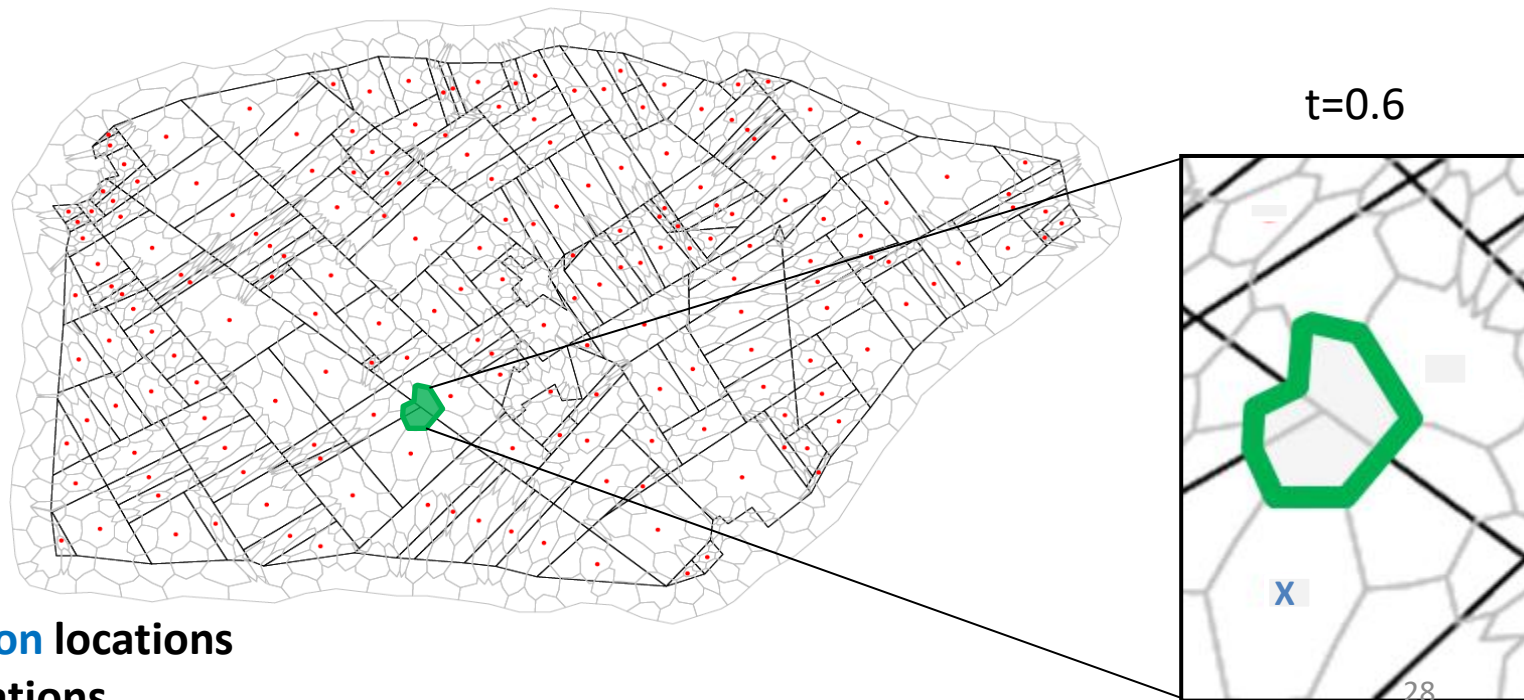


- x pest inoculation locations
- pest peak locations

Spatio-temporal point process


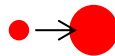
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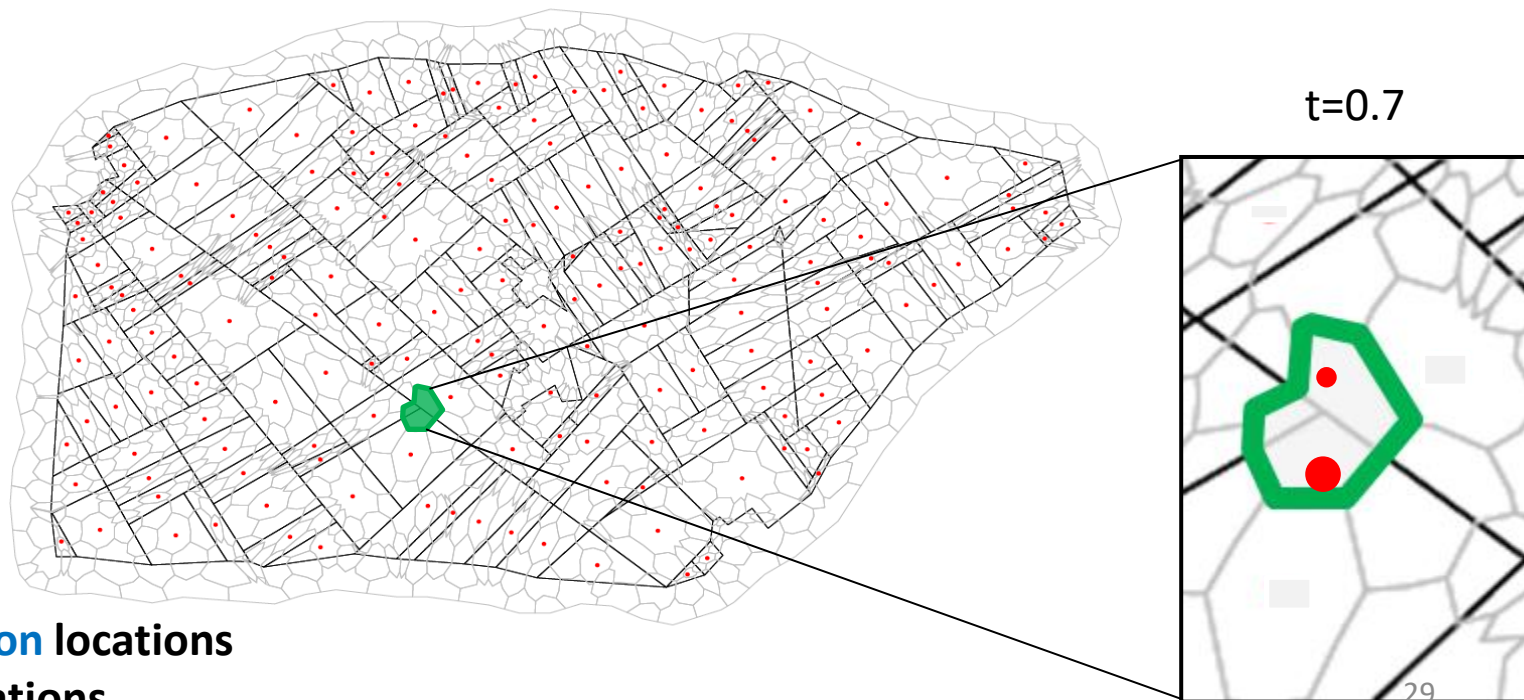
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- 2) Consider the **magnitude** of pest peak 



Spatio-temporal point process

The study region is partitioned into cells to capture the **local temporal dynamic** by models that:

- 1) Count the **number** of pest peak 
- 2) Consider the **magnitude** of pest peak 



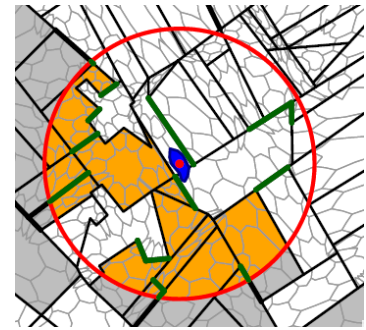
culture/hedge
proportion within
the landscape



Cell-patch
intersections



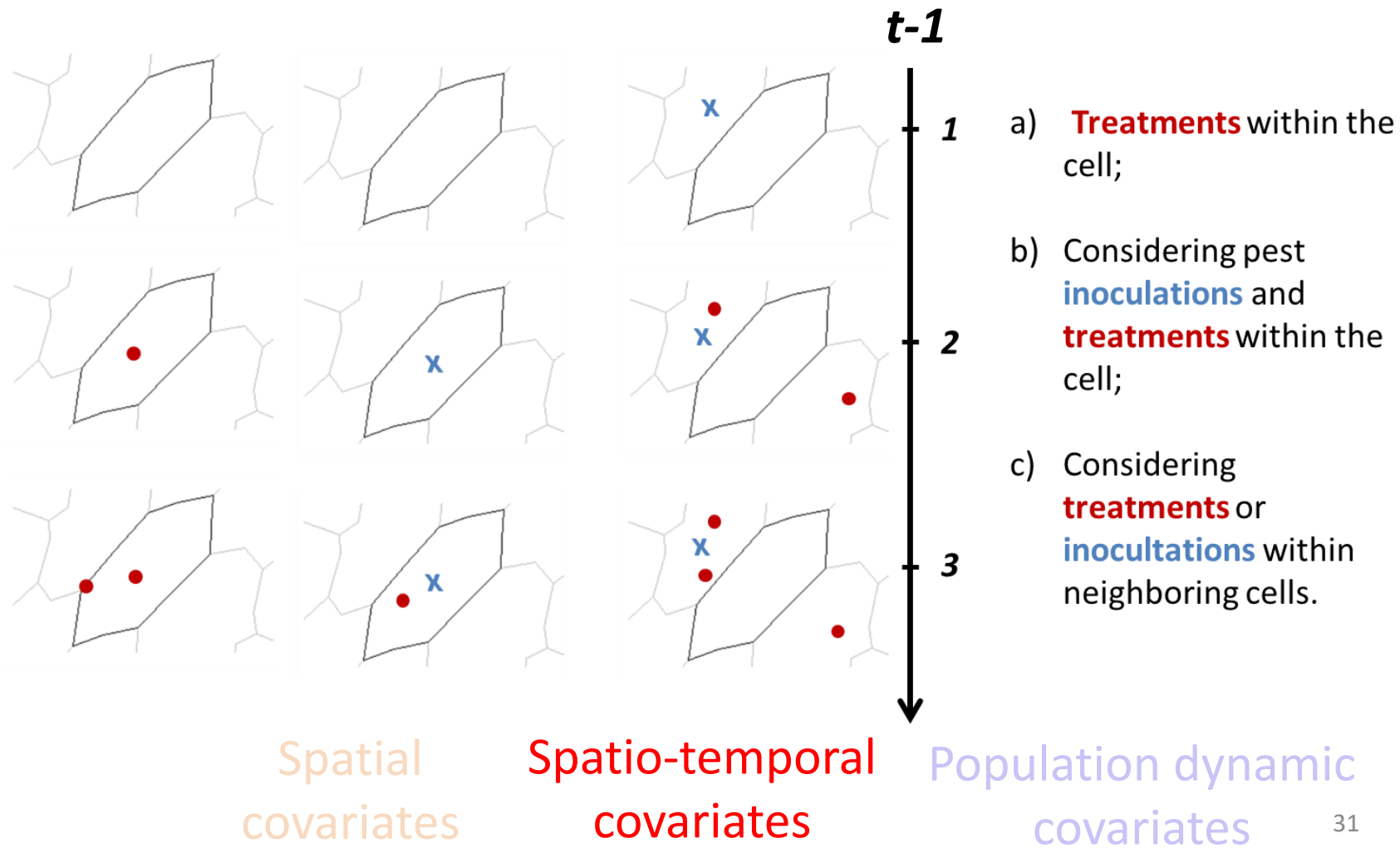
Patch/culture
proportion within
buffers



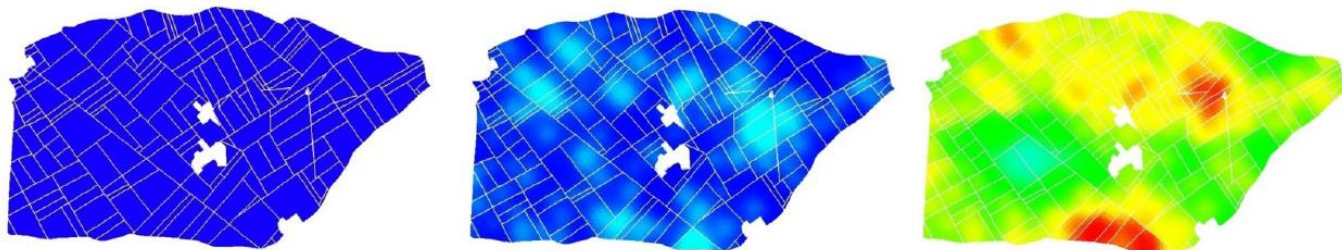
Spatial
covariates

Spatio-temporal
covariates

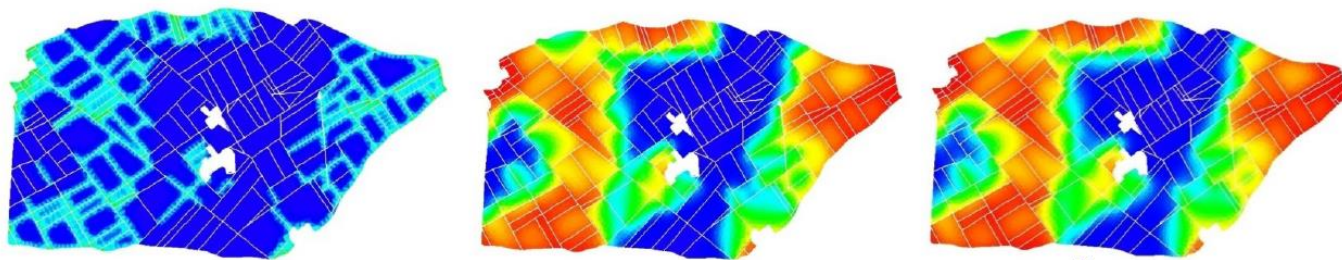
Population dynamic
covariates



Pest diffusion



Predator diffusion
and spillover from
hedge



T_1

T_2

T_3

Spatial
covariates

Spatio-temporal
covariates

Population dynamic
covariates

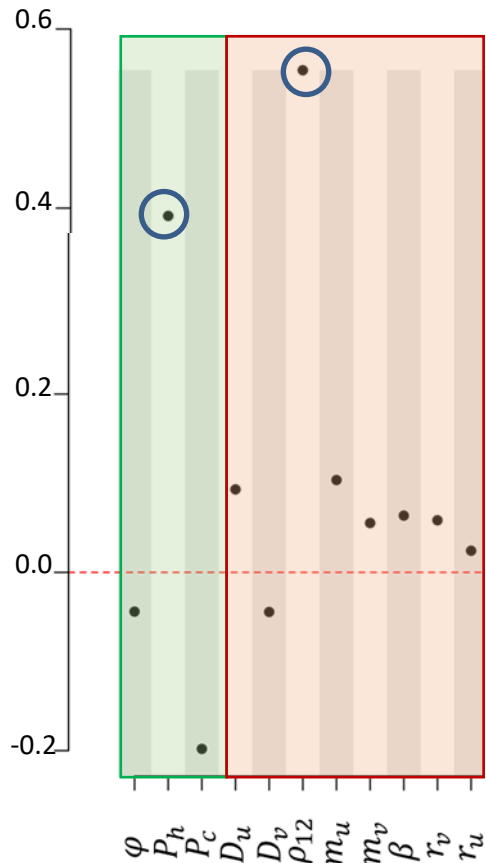
Predator population density



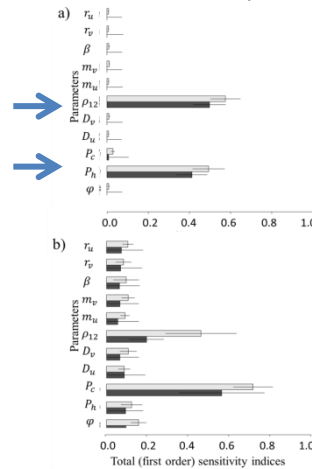
Predator spillover

Hedge proportion

Estimated effects

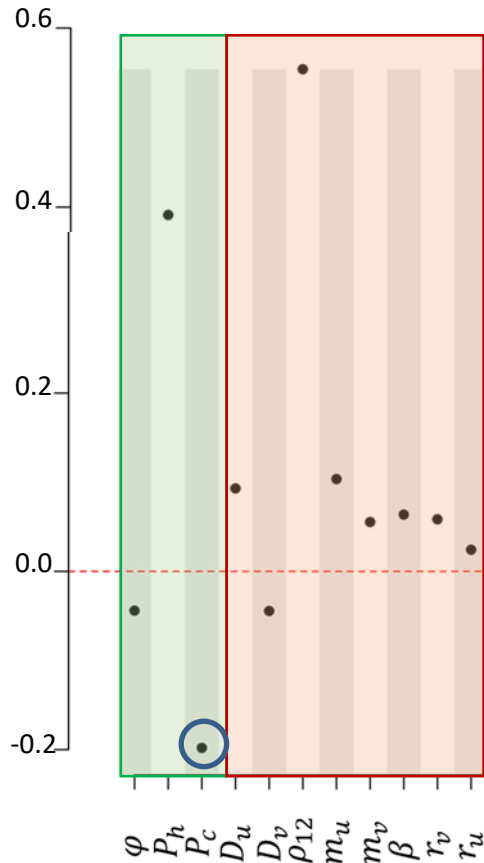


Sobolj



Landscape variable
Species variable

Estimated effects



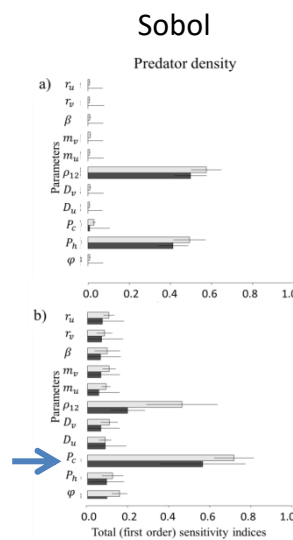
Predator population density



Predator spillover
Hedge proportion



Crop proportion



Landscape variable
Species variable

Predator population density

Sobol

Predator density

Estimated effects

