



## From spatial data to mosquito models *A case study in the Benelux*

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# Introduction

Inventory endemic mosquito population  
Monitoring mosquito co-occurrence



# Modelling principle

Input: points

Result: continuous grid

Filling gaps between points



# Methodology

P/A ~ predictor variables

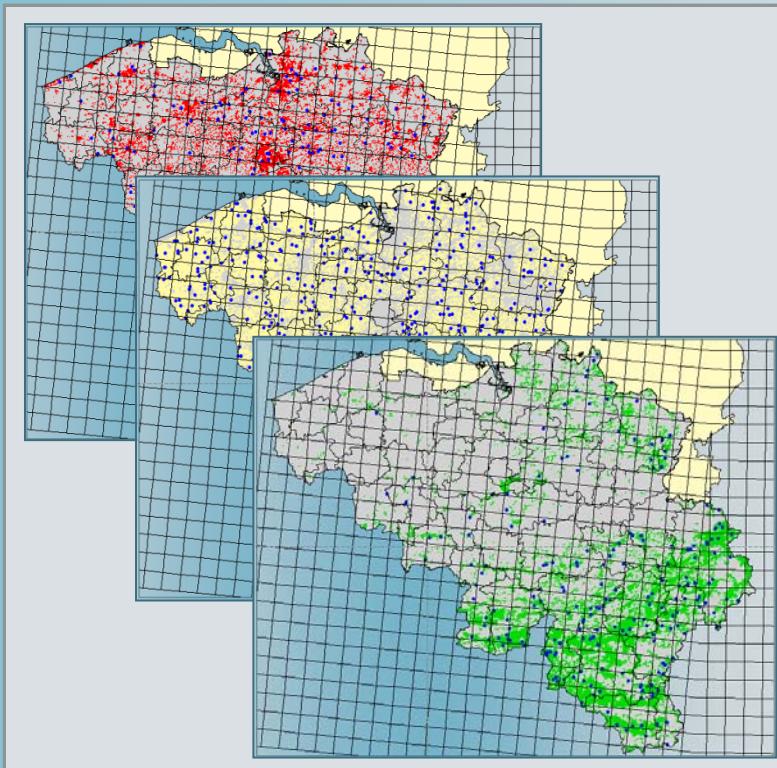
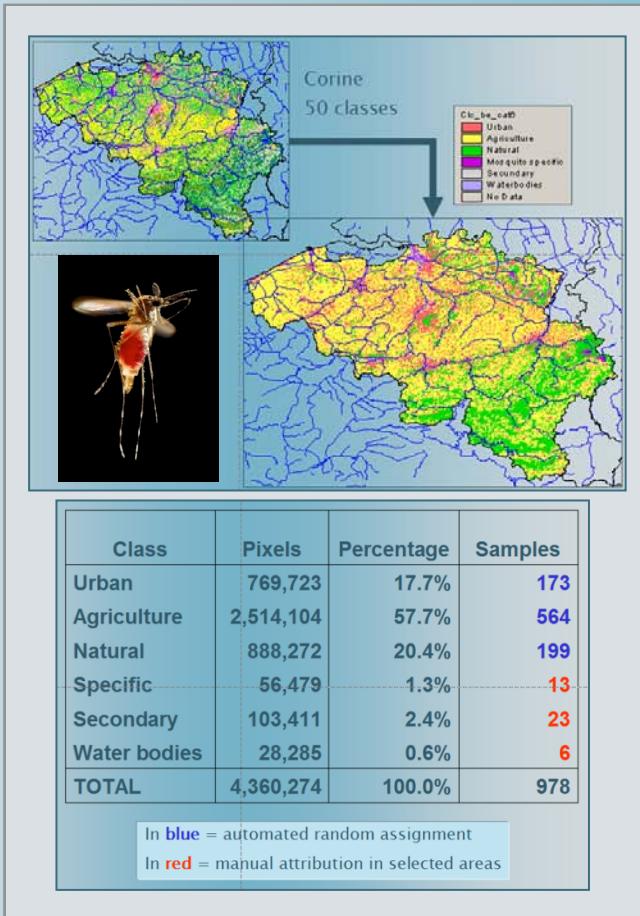
Balanced dataset:

→ nr Presences (positive traps) = nr Absences  
(negative traps)

Random Forest: collection of CART-trees



# Random stratified sampling



# Sample: institute / year / season



Avia  
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# Predictor variables

CORINE land cover : (1) urban, (2) agriculture, (3) natural, (4) risk areas referred to as *specific*, (5) secondary and (6) water bodies

Available water capacity of topsoil

Digital Elevation Model

Yearly/monthly Precipitation

Distance to particular features: (1) Waterways , (2) Protected areas

Percentage cover (% in 1 km<sup>2</sup>): (1) Broadleaved, Coniferous, Mixed forest, (2) Urban, Agriculture, Natural

Population density (inhabitants/km<sup>2</sup>)

MODIS derivates: (1) Day/Night Temperature ; (2) Vegetation greenness

Fourier analysis on data of last 5 years: (1) Mean ; (2) Interannual variation

Nr of freezing days/nights



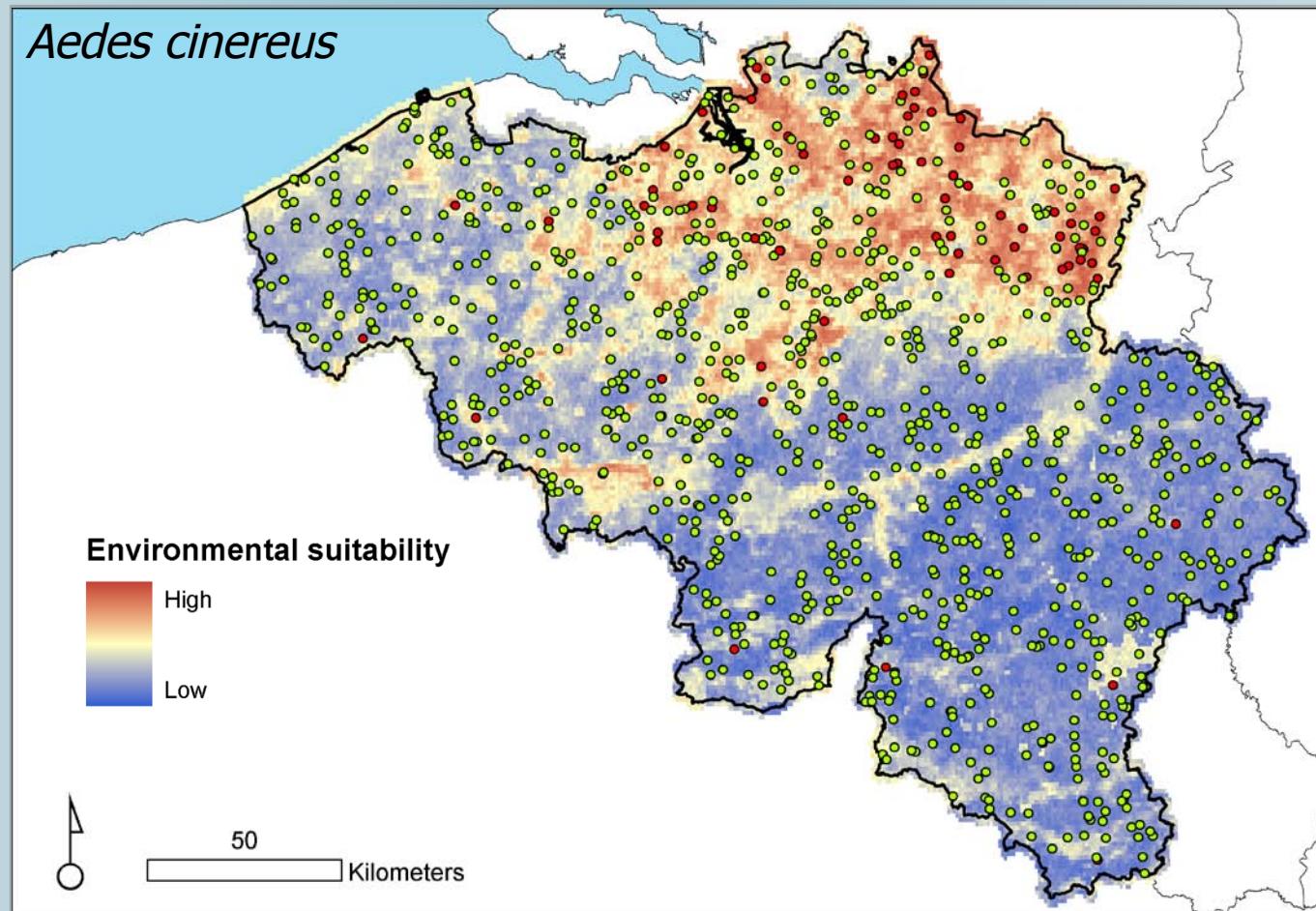
# Most frequent species

Positive traps

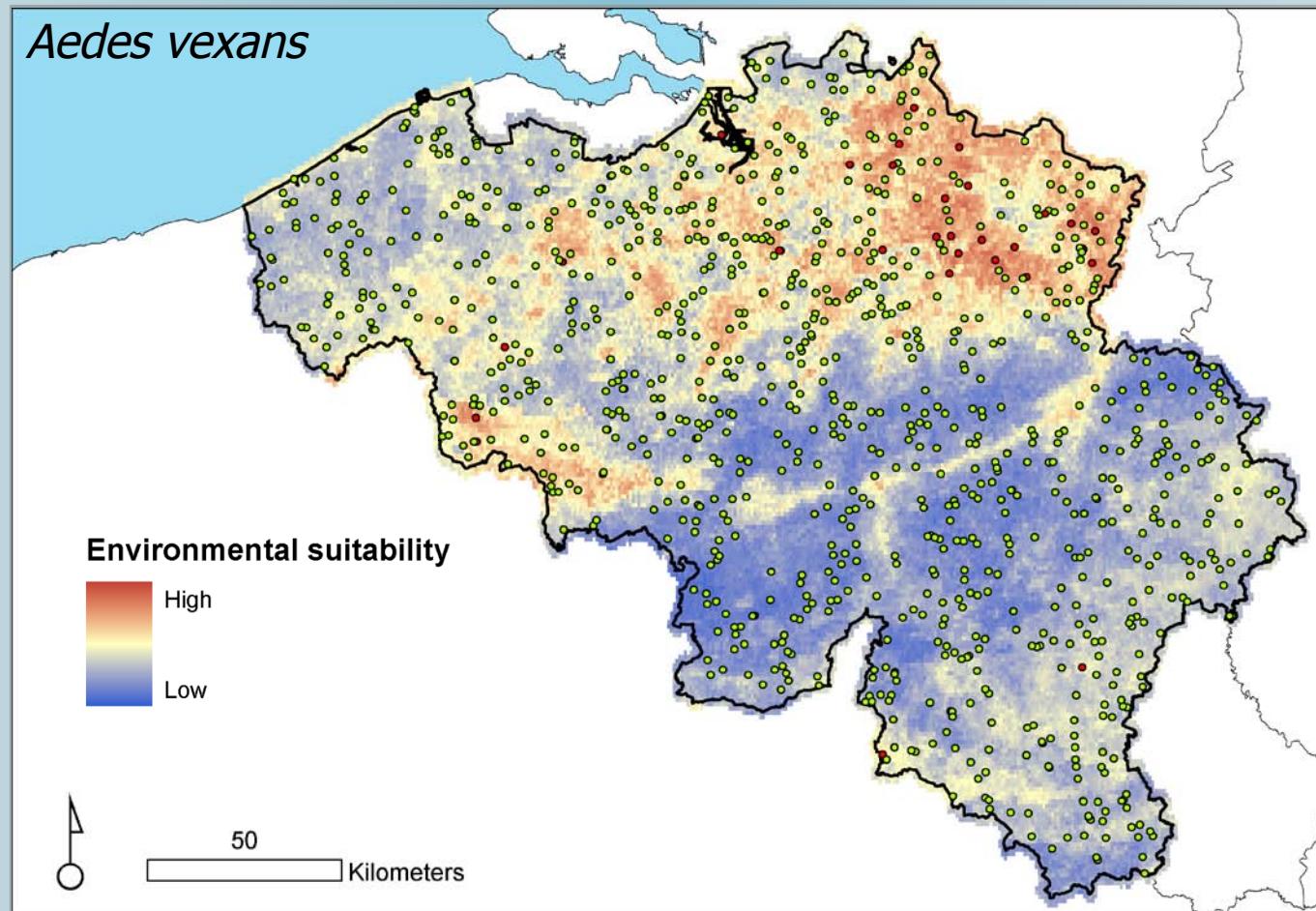
Species	Total	%	N	N/+Tr
<i>Aedes vexans</i>	<b>29</b>	2.97%	776	<b>26.76</b>
<i>Coquillettidia richiardii</i>	<b>38</b>	3.89%	4,095	<b>107.76</b>
<i>Aedes cinereus/geminus</i>	<b>74</b>	7.57%	1,328	<b>17.95</b>
<i>Culex pipiens</i>	<b>698</b>	71.37%	16,338	<b>23.41</b>



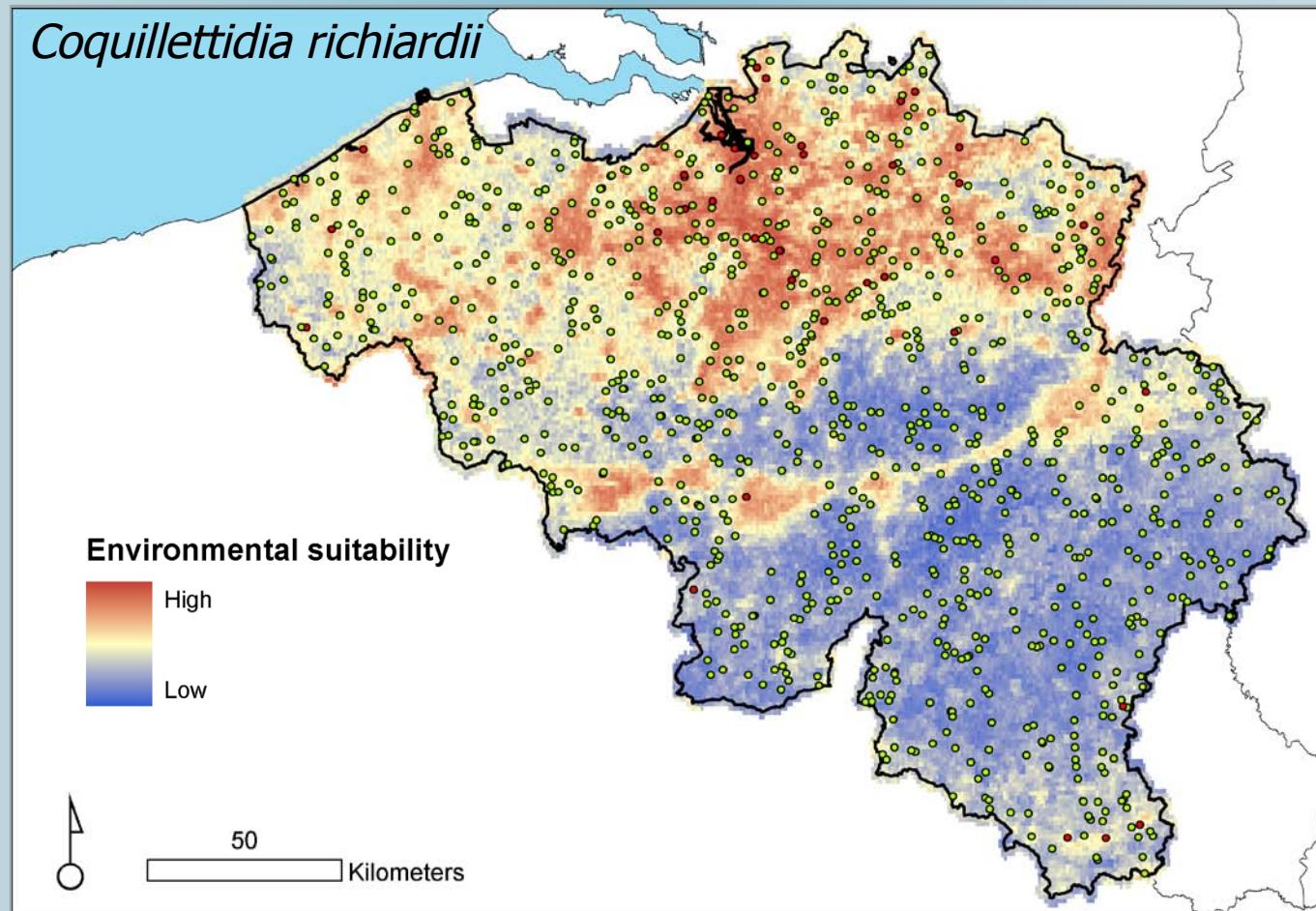
# Model results for 14 mosquito sp.



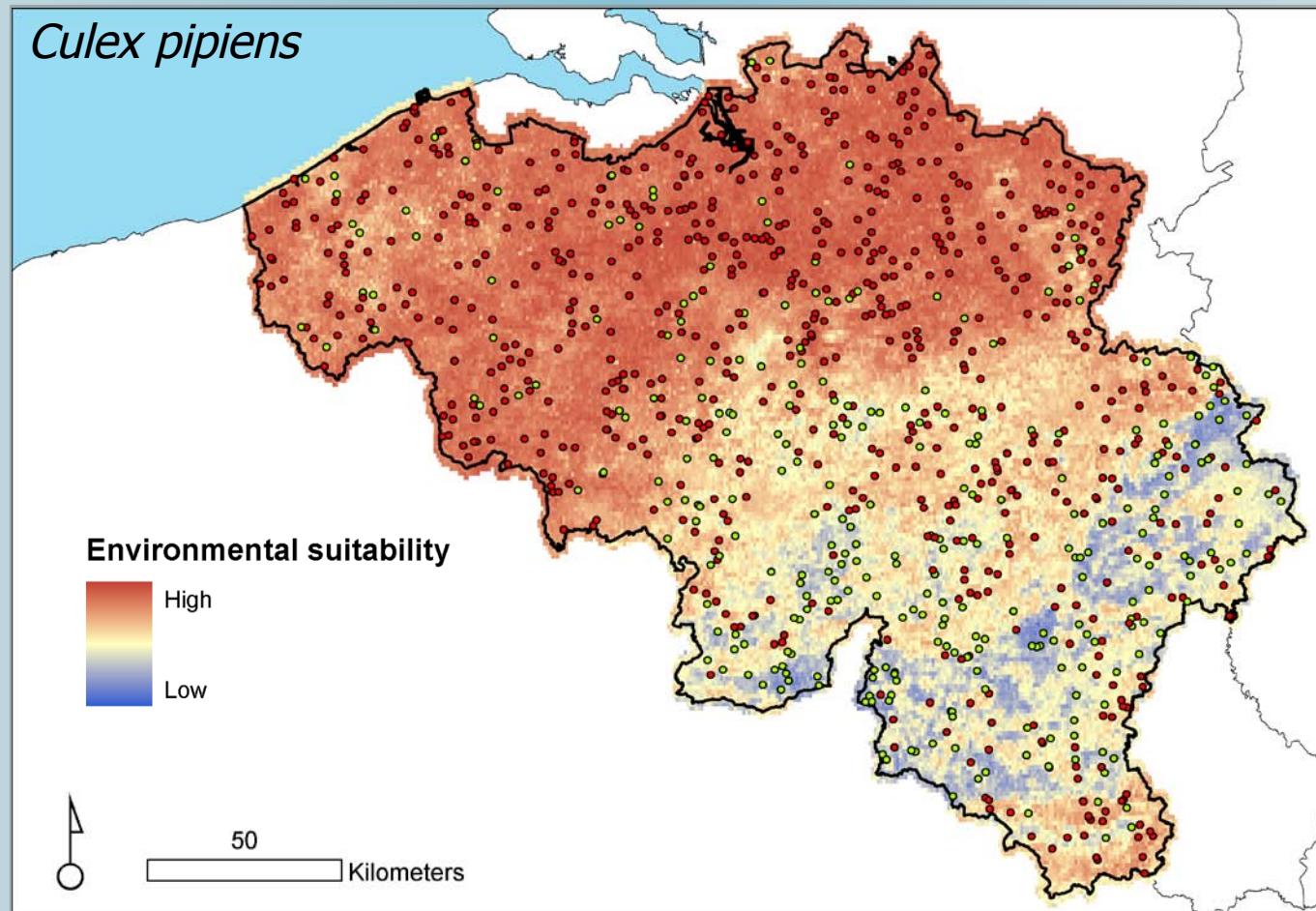
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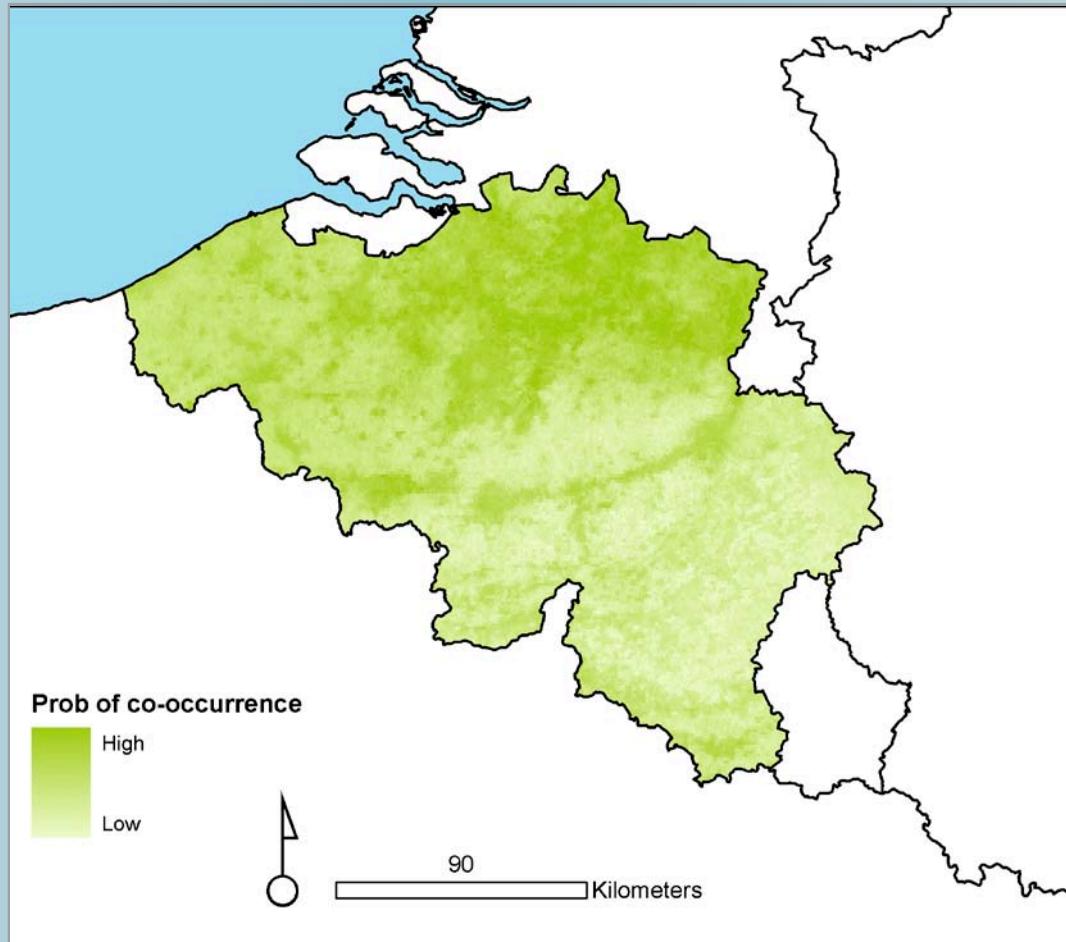
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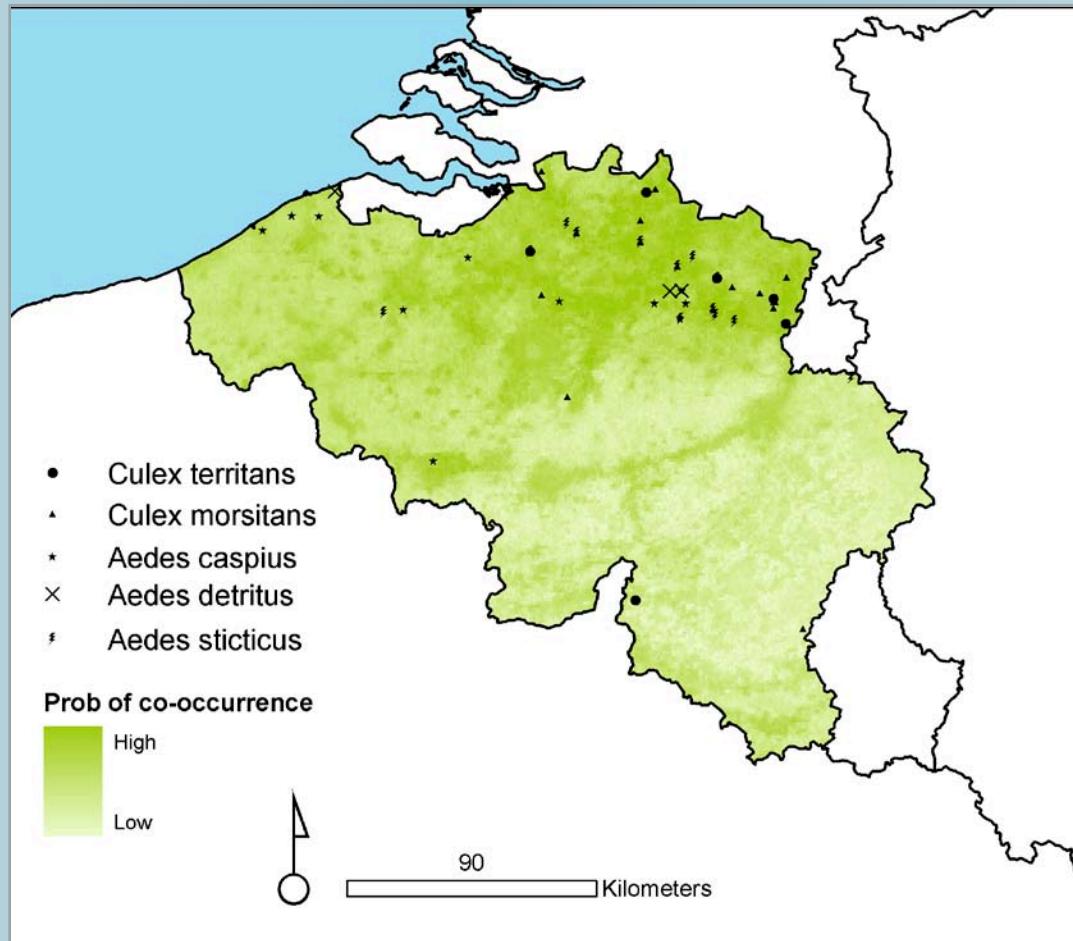
## Probability of co-occurrence for 14 species



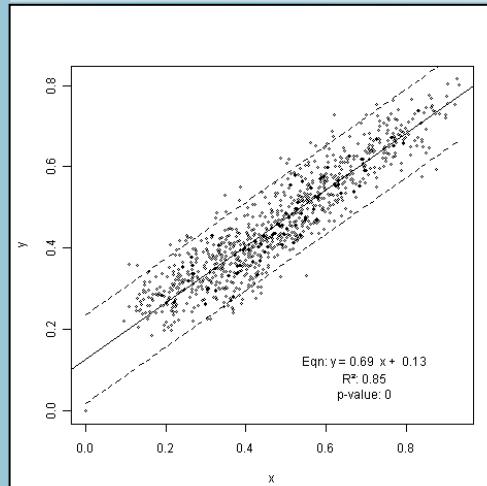
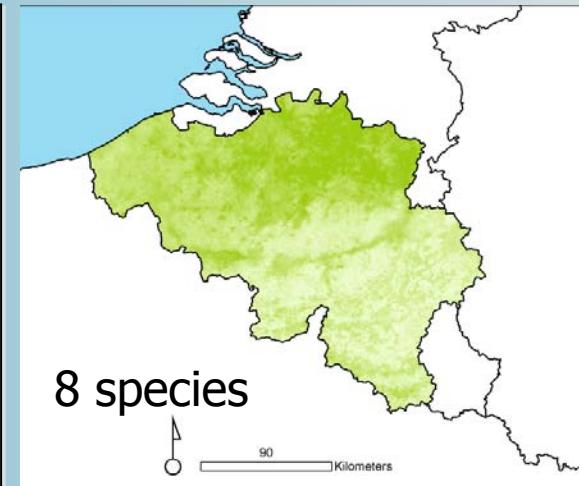
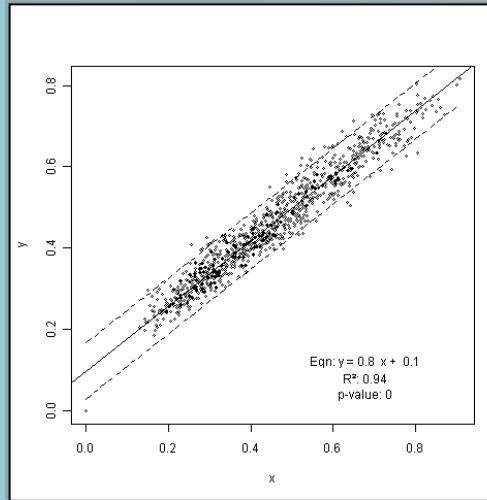
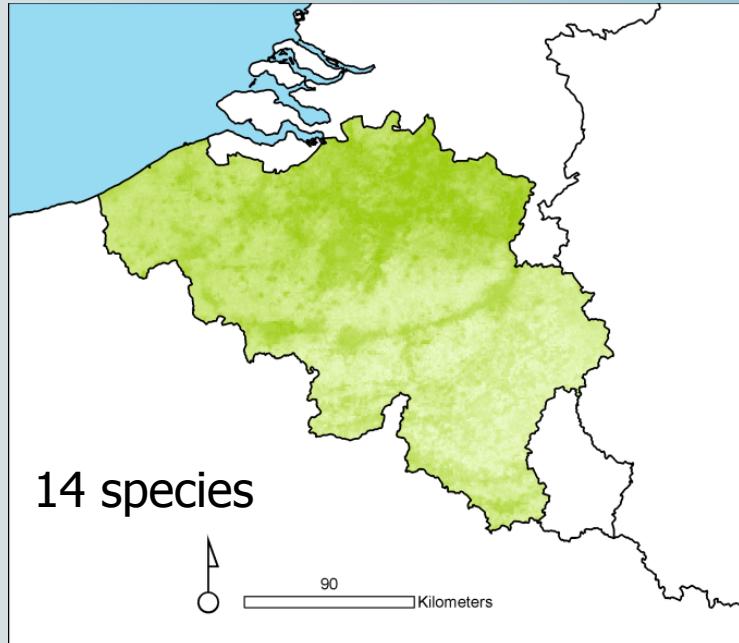
*Avia*  
GIS



## Probability of co-occurrence for 14 species

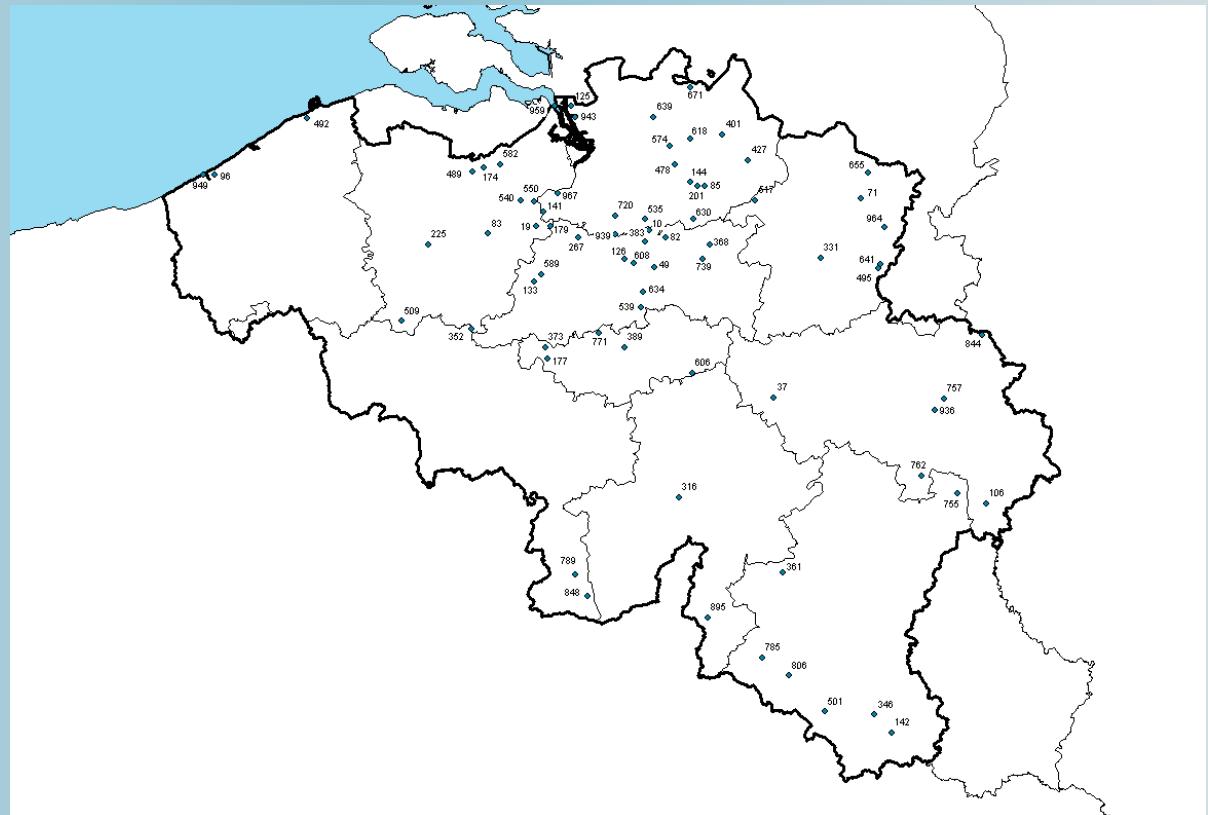


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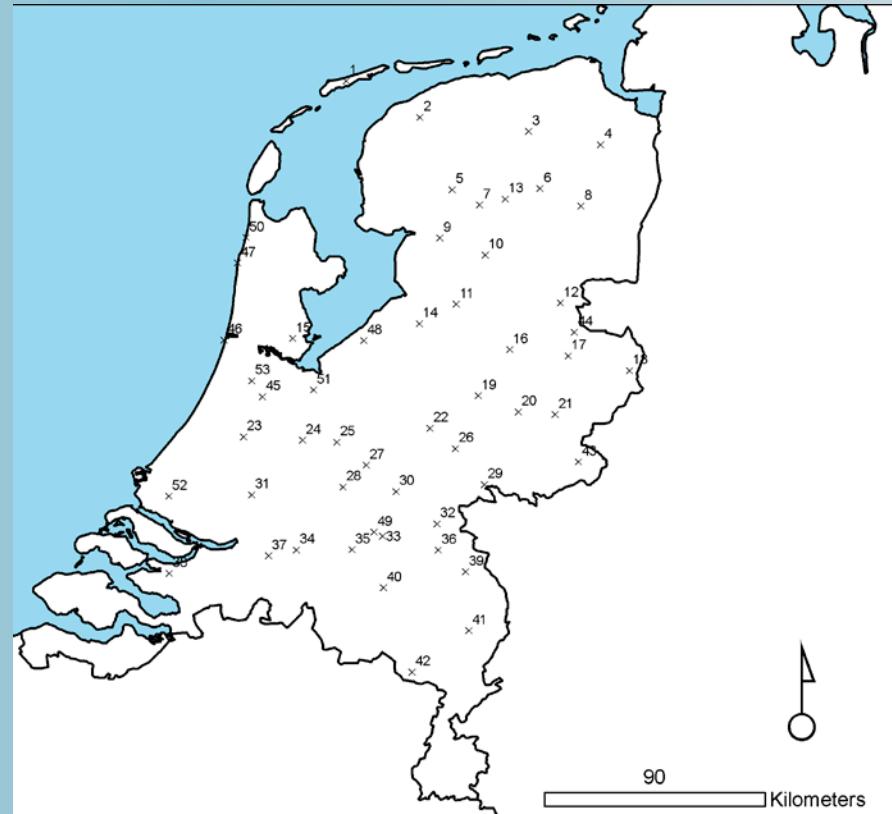
# Validation of models in Belgium

75 sites in 2009  
75 sites in 2010

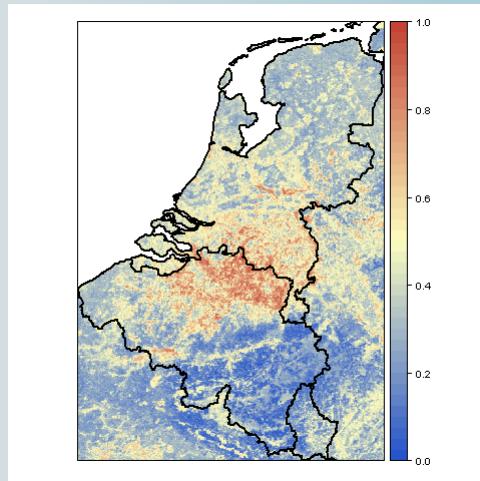


# Validation of models in the Netherlands

53 sites in 2009  
53 sites in 2010

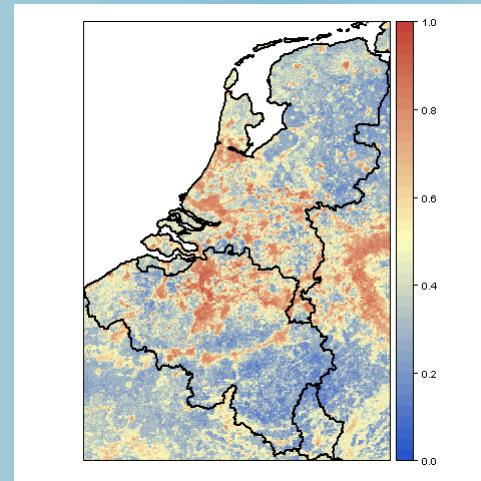


# Validation of models in Netherlands (53 sites)



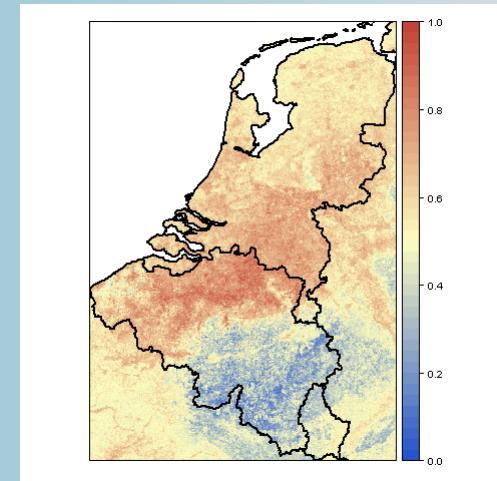
*Ae. cinereus* (58% PCC)

	prediction	
trap	PRES	ABS
PRES	3	3
ABS	19	28



*Cq. richardii* (57% PCC)

	prediction	
	PRES	ABS
	PRES	ABS
6	10	
13	24	



*Cx. pipiens* (77% PCC)

	prediction	
	PRES	ABS
	PRES	ABS
41	10	
2	0	



# Cost-efficient sampling strategy

- Stratified random sampling:
  - 1000 points => 100 points
  - 100% => 10%
  - 1 trap/30 km<sup>2</sup> => 1 trap/300 km<sup>2</sup>



# Evaluation

Is modelling output satisfactory

- AUC
- PCC, Sensitivity, Specificity
- Spatial pattern



# Accuracy measures

PCC

species	1/30km <sup>2</sup>	1/40km <sup>2</sup>	1/50km <sup>2</sup>	1/75km <sup>2</sup>	1/150km <sup>2</sup>
<i>Ae cinereus</i>	0.75	0.79	0.77	0.75	0.71
<i>Ae vexans</i>	0.75	0.71	0.65	0.60	0.54
<i>Cq richardii</i>	0.73	0.70	0.69	0.67	0.65
<i>Cx pipiens</i>	0.87	0.83	0.80	0.75	0.71

Sensitivity

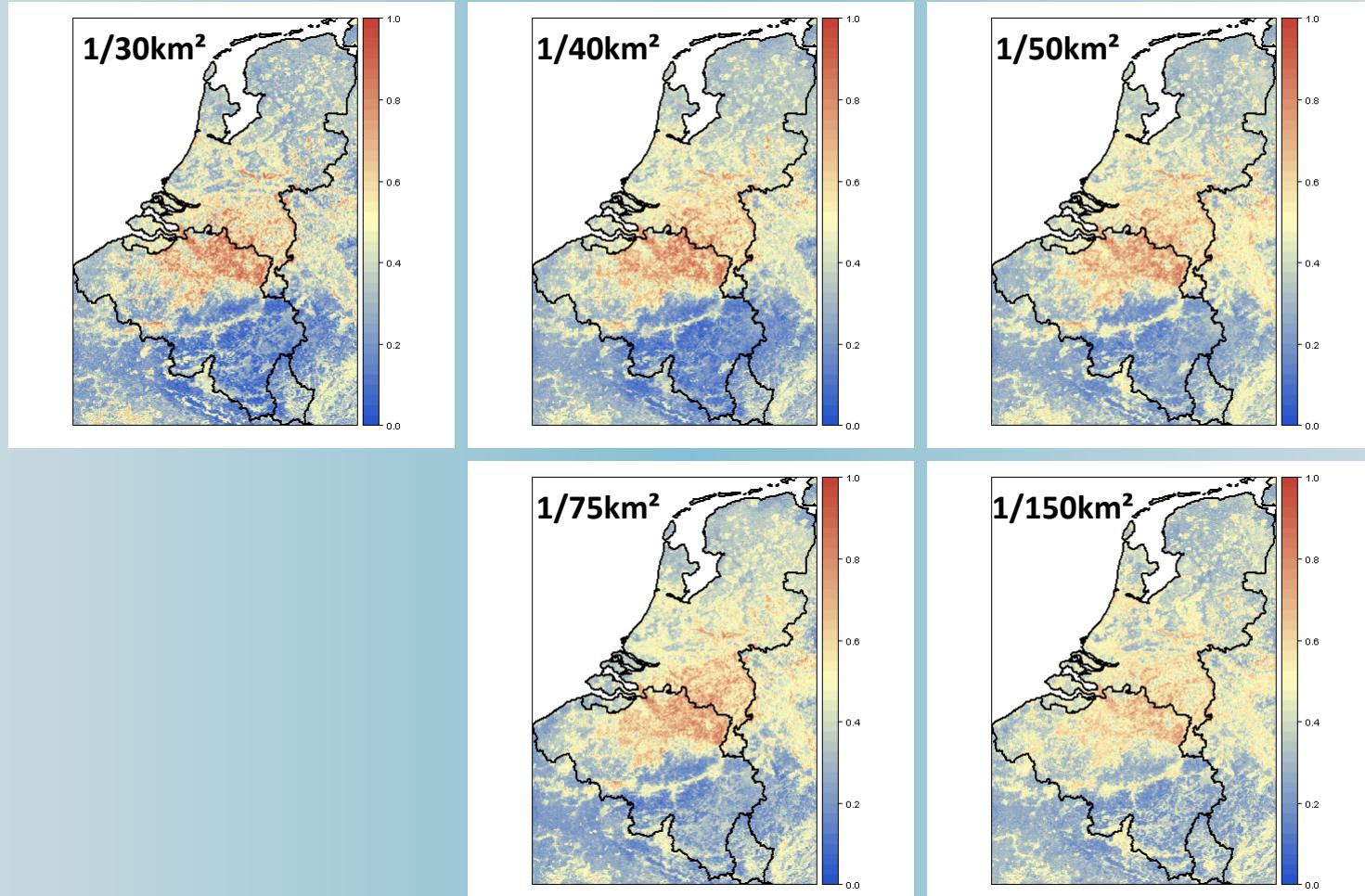
species	1/30km <sup>2</sup>	1/40km <sup>2</sup>	1/50km <sup>2</sup>	1/75km <sup>2</sup>	1/150km <sup>2</sup>
<i>Ae cinereus</i>	0.99	0.96	0.91	0.85	0.89
<i>Ae vexans</i>	1.00	0.97	0.95	0.92	0.90
<i>Cq richardii</i>	1.00	0.94	0.88	0.82	0.76
<i>Cx pipiens</i>	0.82	0.79	0.76	0.72	0.67

Specificity

species	1/30km <sup>2</sup>	1/40km <sup>2</sup>	1/50km <sup>2</sup>	1/75km <sup>2</sup>	1/150km <sup>2</sup>
<i>Ae cinereus</i>	0.73	0.77	0.76	0.74	0.70
<i>Ae vexans</i>	0.74	0.70	0.64	0.59	0.53
<i>Cq richardii</i>	0.72	0.69	0.69	0.67	0.65
<i>Cx pipiens</i>	1.00	0.93	0.88	0.82	0.79



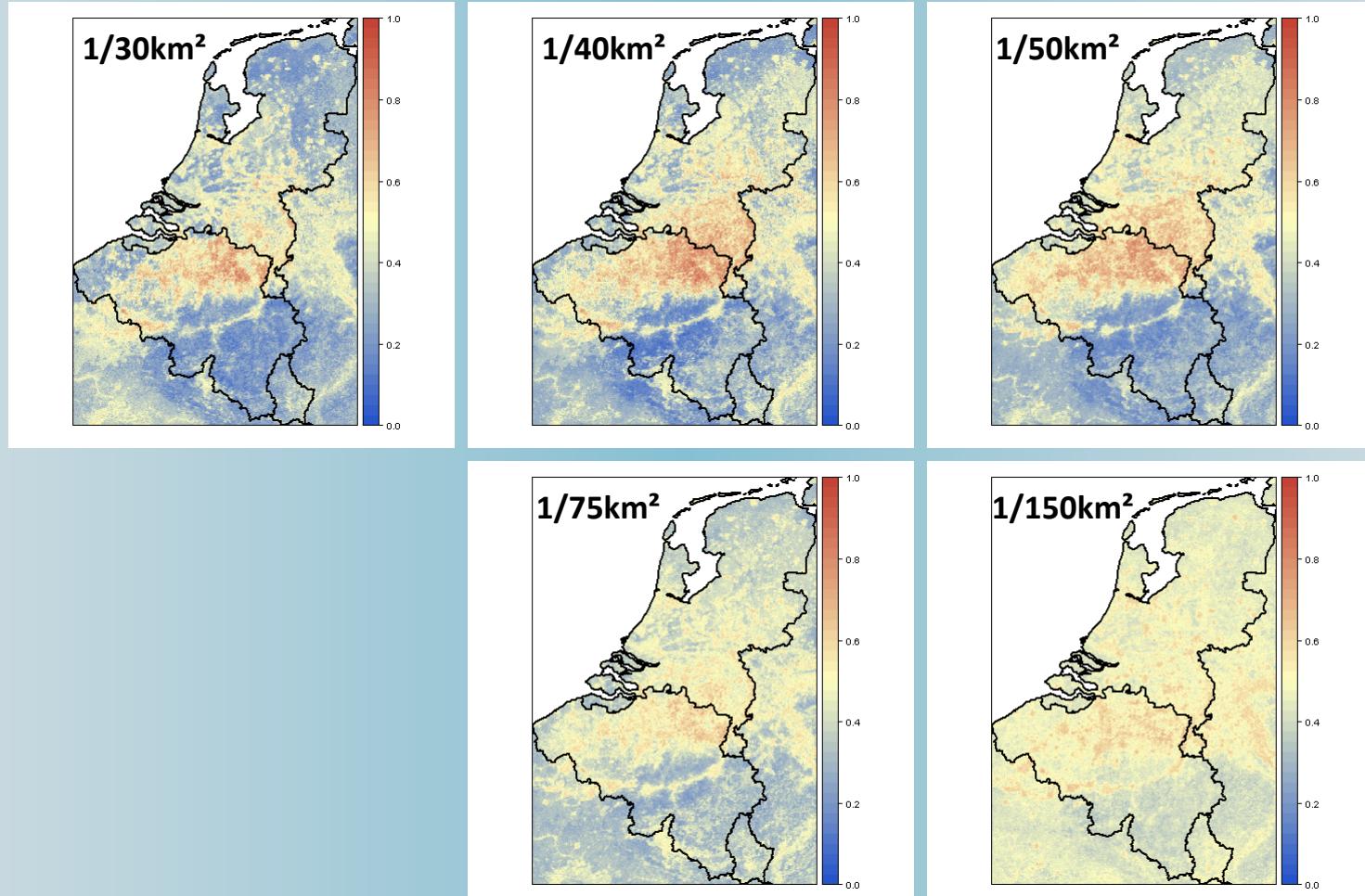
# *Ae. cinereus*



*Avia*  
GIS



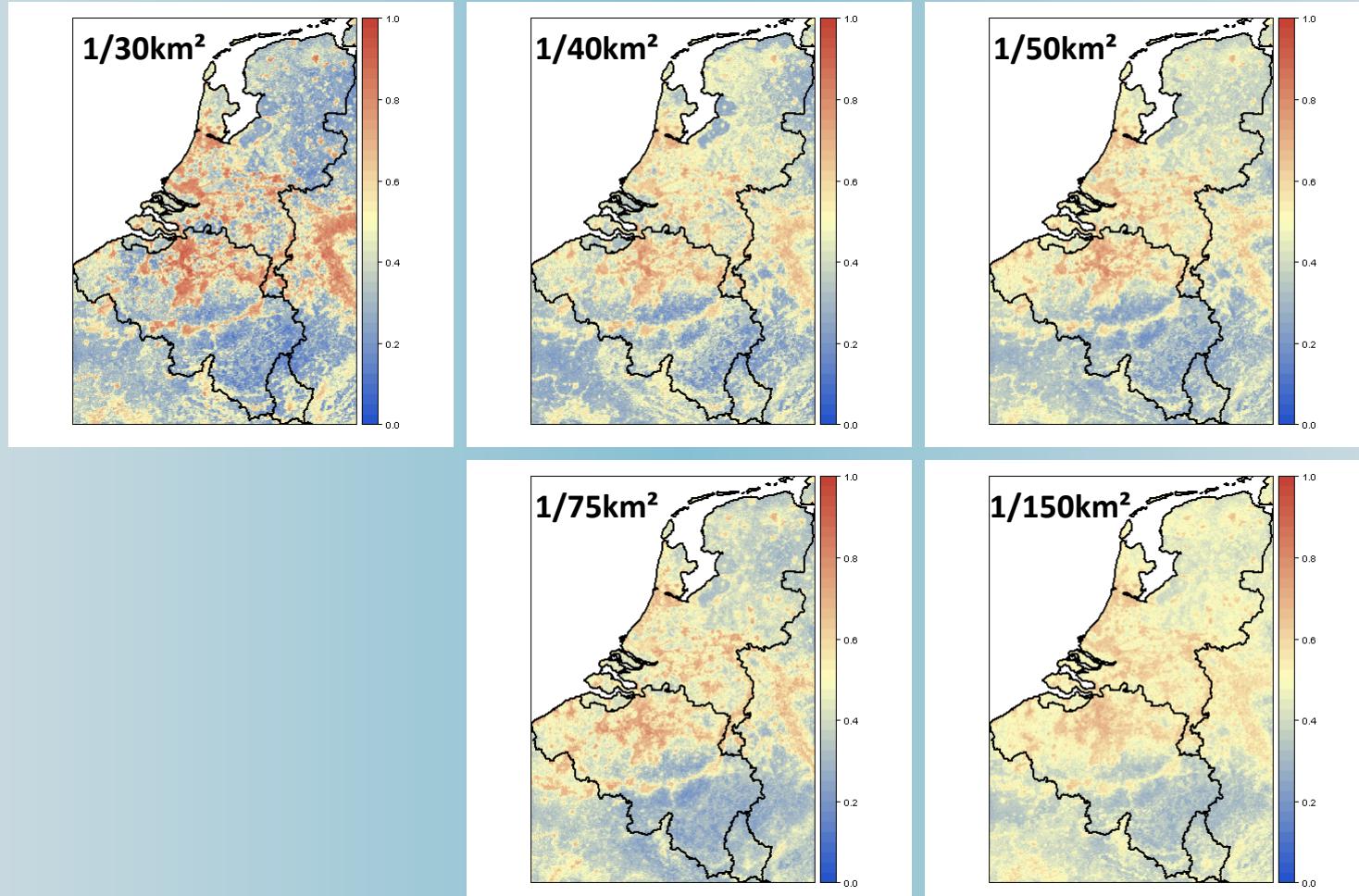
# *Ae. vexans*



*Avia*  
GIS



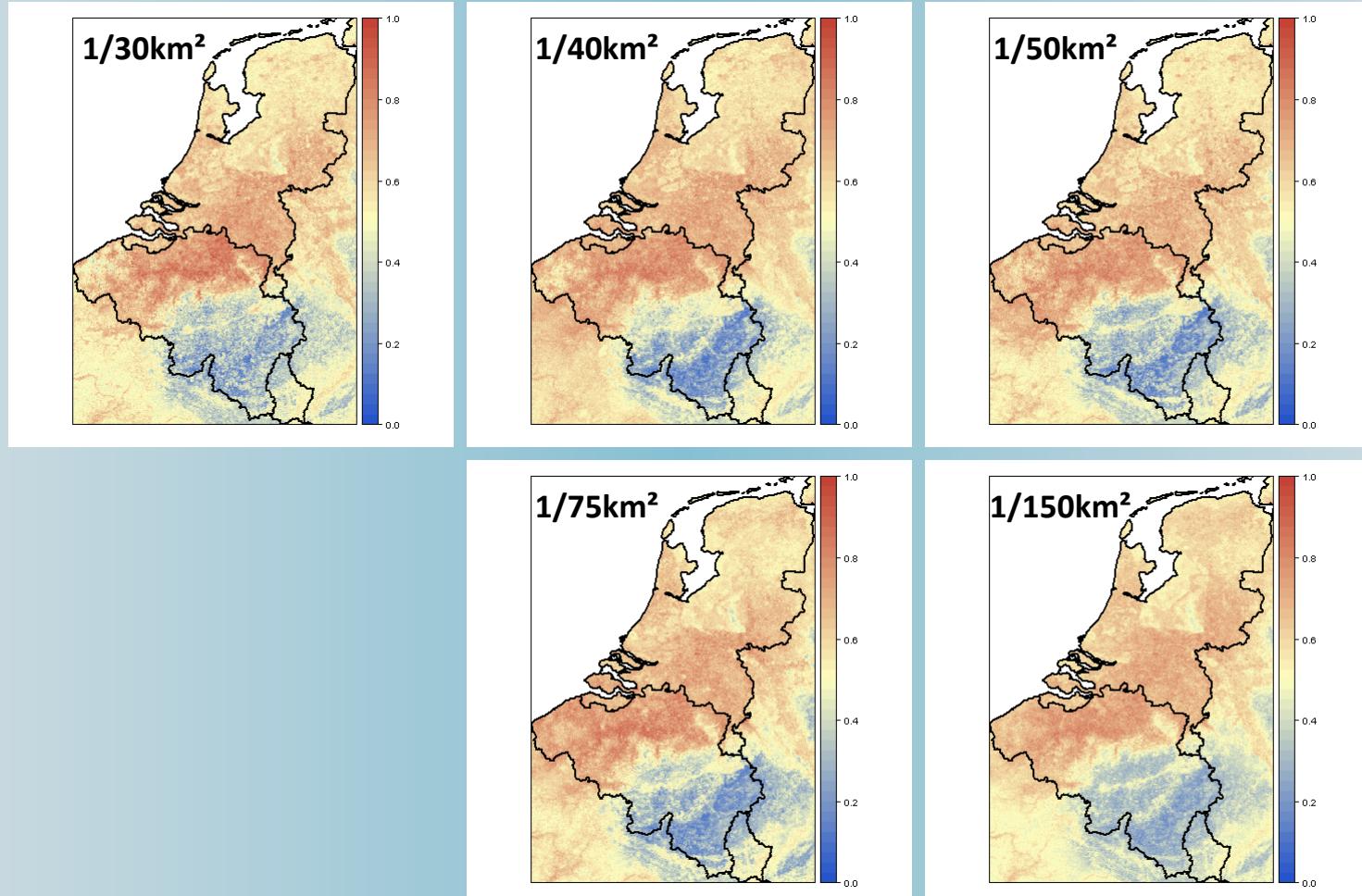
# *Cq. richardii*



*Avia*  
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# *Cx. pipiens*



*Avia*  
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# Conclusions

Hotspot for species co-occurrence

Four indicator species:

Wide-spread

Easy to catch

Stratified random sampling:

Good results up to 1 trap/150 km<sup>2</sup> at a 1km resolution



# Ongoing

- False absences
- Uncertainty
- Resolution



# Thank you for your questions

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