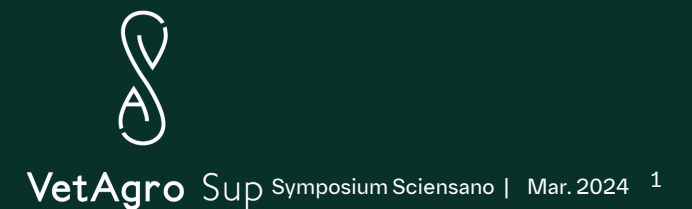




Risk of *Culicoides* dispersal by the wind : case study with EHDV in France

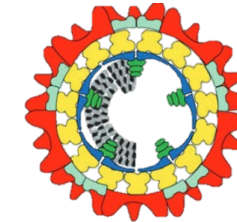
Amandine BIBARD

amandine.bibard@boehringer-Ingelheim.com

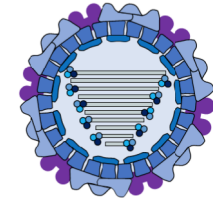


Epizootic Hemorrhagic Disease Virus (EHDV) : an emerging threat in Europe

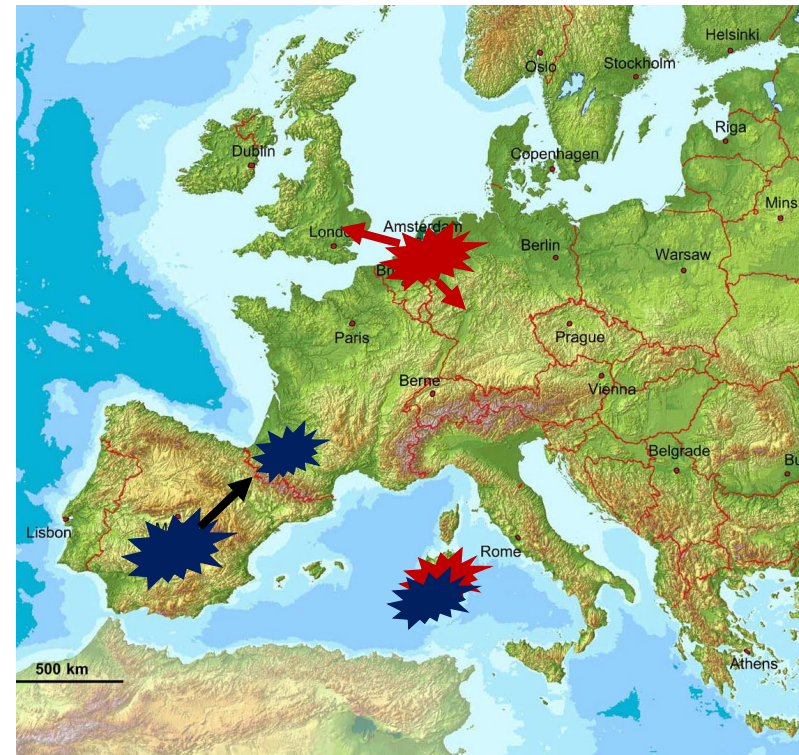
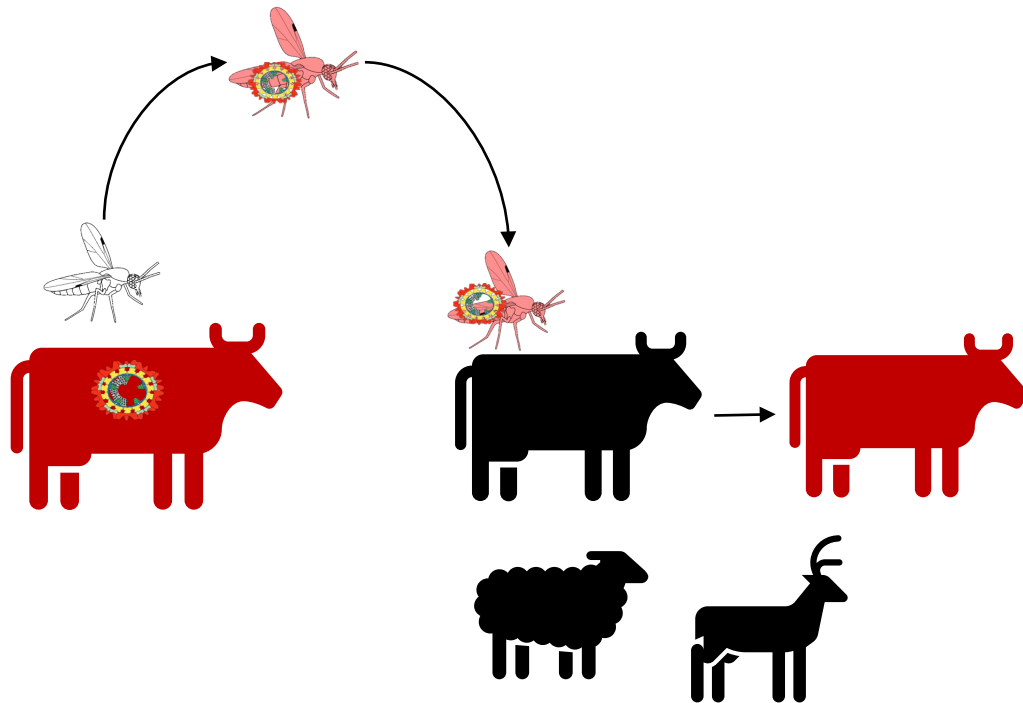
- Orbivirus like Bluetongue virus
- Morbidity/mortality variable according to serotypes
- High economical & animal health impacts mostly in the cattle farms
- Also transmitted by *Culicoides* midges



BTV



EHDV

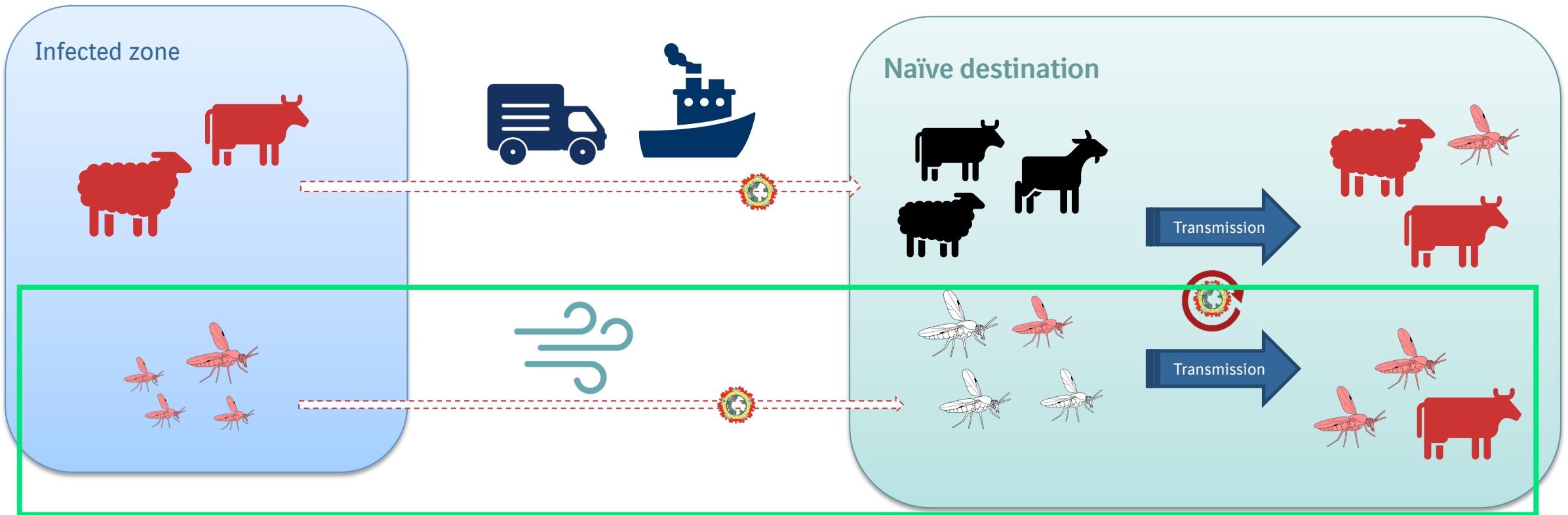


Sardinia: 2018
NL: Sept 2023



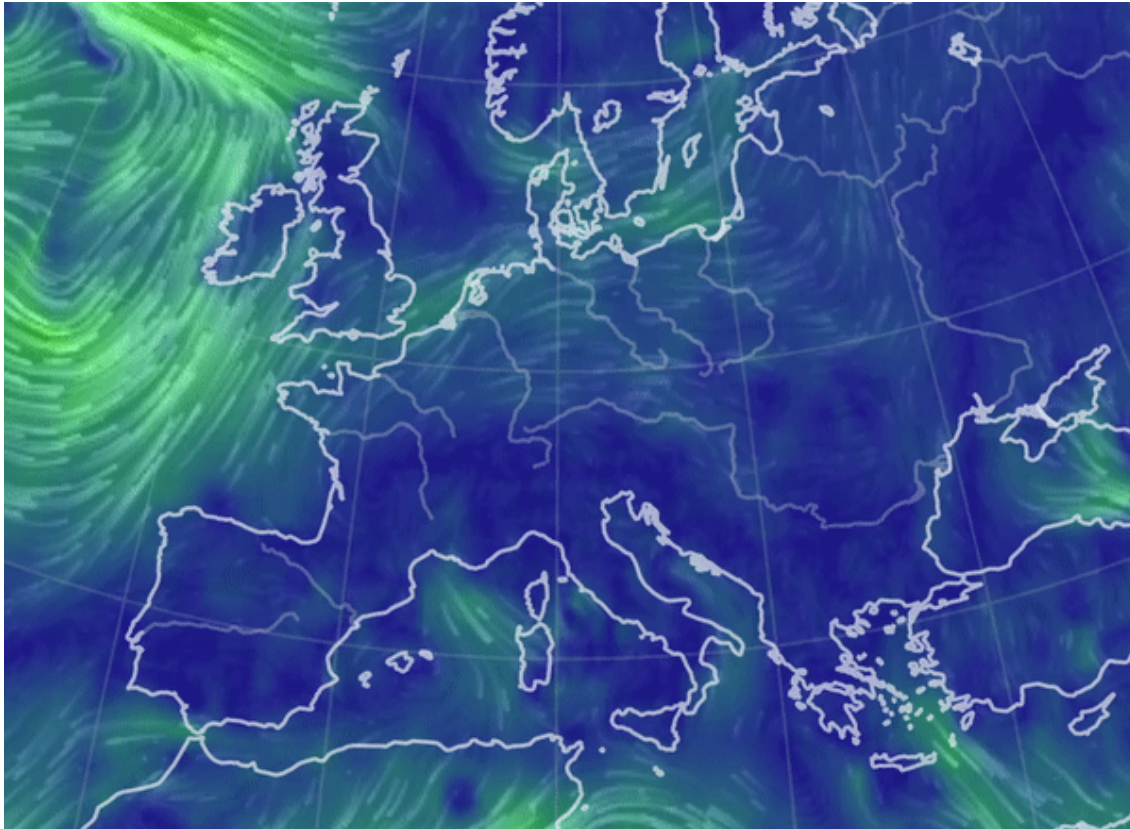
Sardinia: Oct 2022
Spain: Oct 2022
France: Sept 2023

Two main pathways of virus introduction



- Animal movements & *Culicoides* dispersal by the wind
- Dispersal by the wind difficult to control and highly variable due to the high number of parameters to control

The wind: highly variable parameter



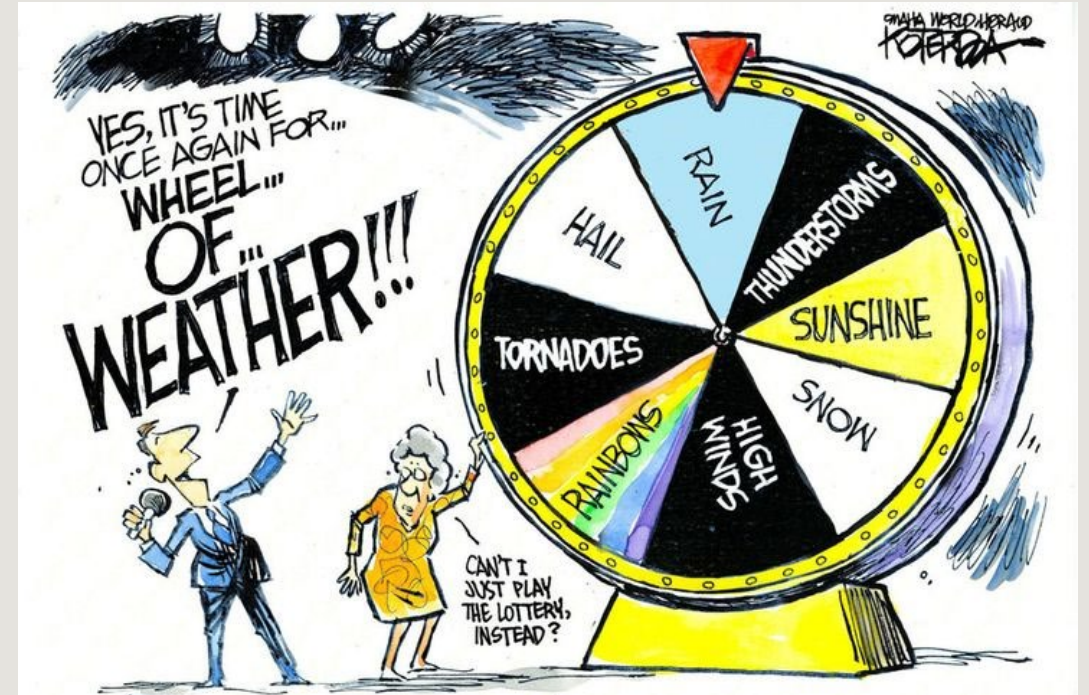
Source: iweather.net.com

- Wind direction and speed highly variable
- Influenced by the terrain
- Midge survival depends on temperature



How to predict *Culicoides* wind dispersal in Europe?

Atmospheric Dispersion Model (ADM)



HYSPLIT Model

- Originally developed for nuclear explosion and volcano eruption.
- HYSPLIT **simulates the dispersion and trajectory** of substances transported and dispersed through our atmosphere, over **local to global scales**.

1 Define the starting point(s)

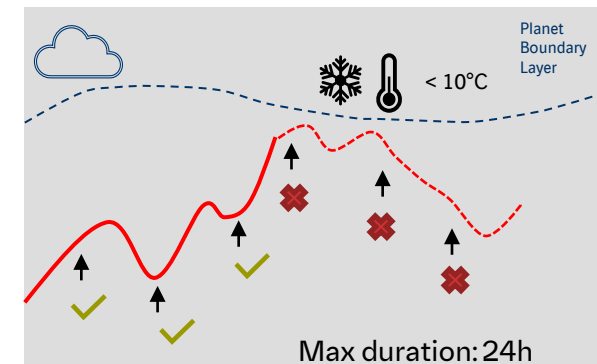


2 Initiate trajectories from it



- **2 starting time** per day (sunset & sunrise)
- **36 weeks** of simulations (mid March – Mid Nov)
- **Historical data of 2020-2021-2022-2023**

3 Filter suitable trajectories according to mites survival conditions

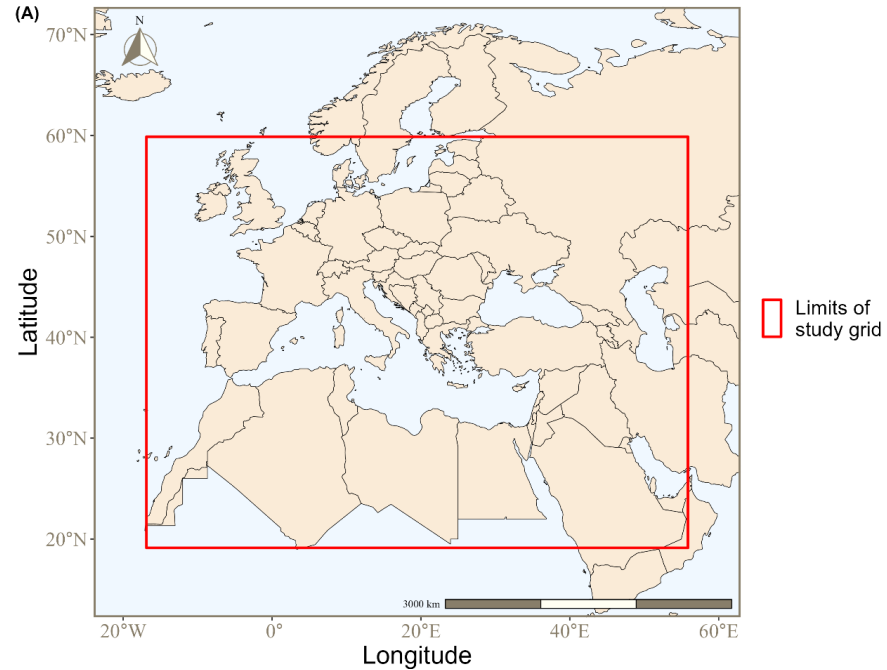
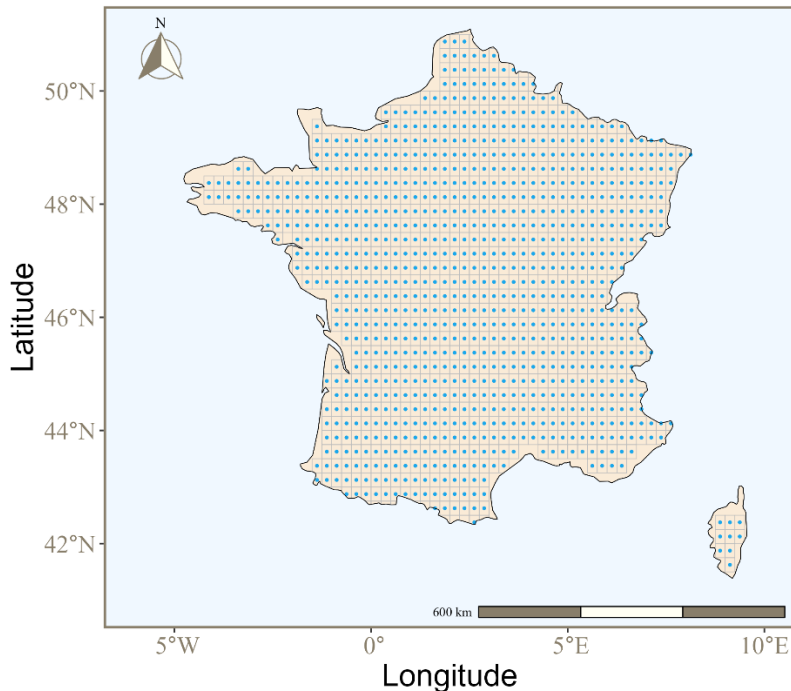


$H_{ij} \sim$ probability to reach destination from starting point

Stein A F, Draxler R R, Rolph G D, Stunder B J B, Cohen M D and Ngan F 2015 NOAA's HYSPLIT Atmospheric Transport and Dispersion Modeling System *B Am Meteorol Soc* 96 2059–77

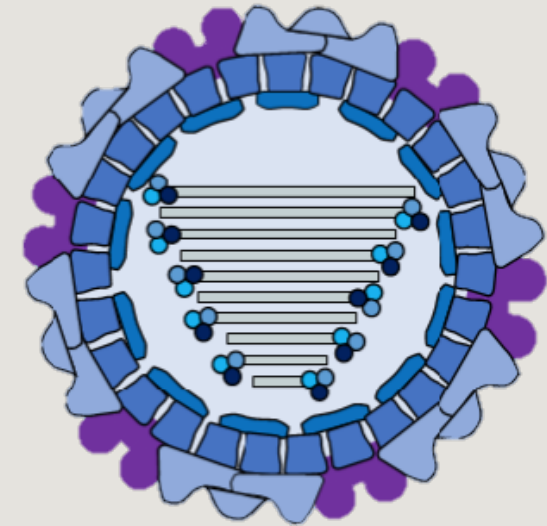
A framework developed for the whole EU

- Large study grid
- 32 292 grid cells of ~25x25 km
- 65 100 672 atmospheric simulations performed
- > 10 Go of output data

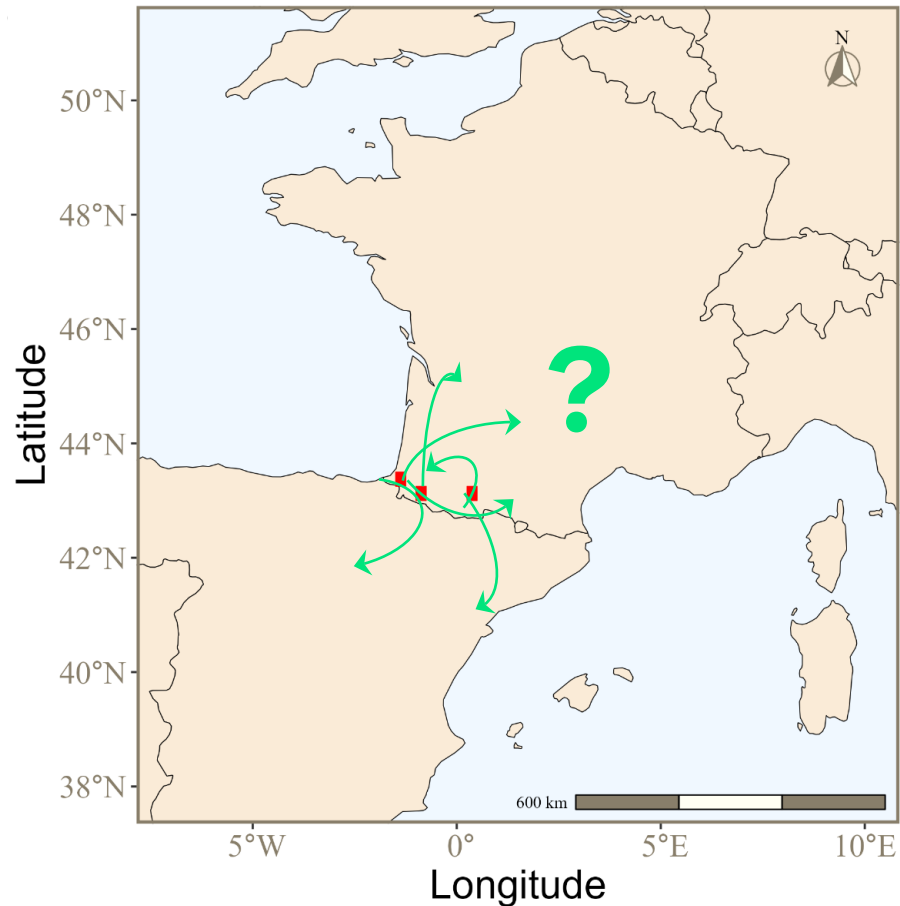


EHDV-8 in France: zone of vector wind dispersal from index cases

First introduction in EU France (Sept 2023)



First 3 outbreaks in France (early September 2023)



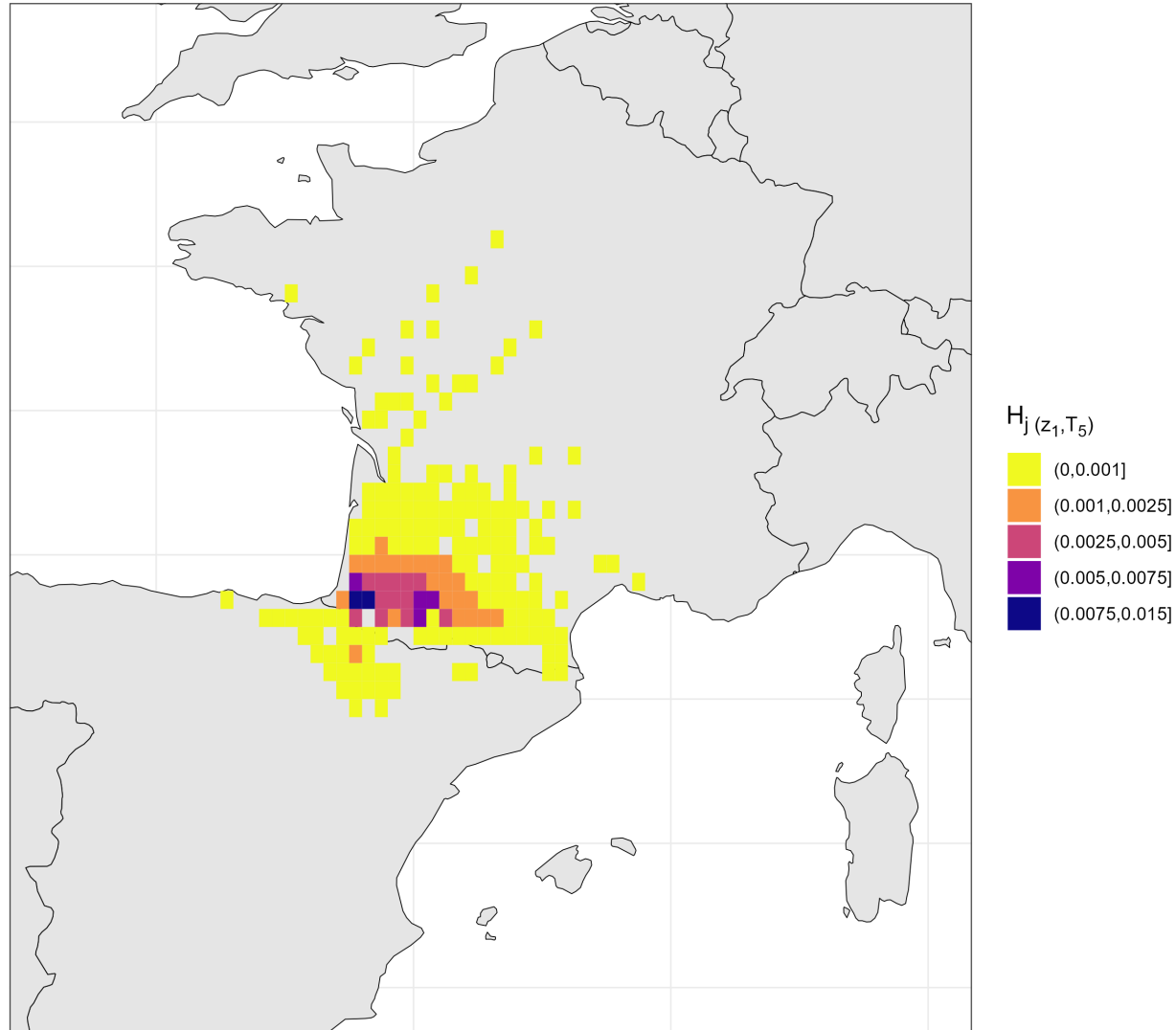
- 3 first outbreaks reported in France

What could be the magnitude of the long-distance dispersal zone from the first cases?

Is a predictive insight for emerging outbreaks?

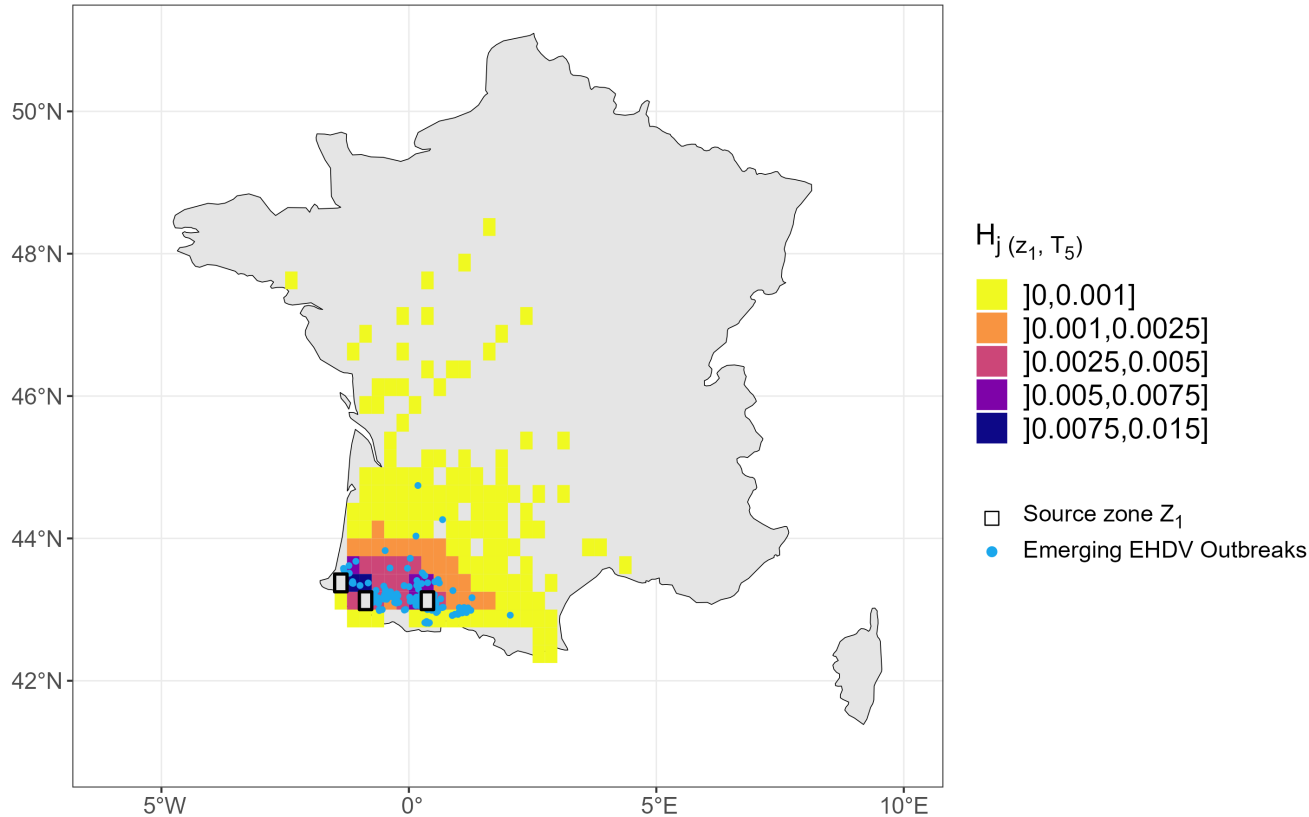
Predicted risk zone averaged over 5 week-period

Windborne dispersion from index source zone z1

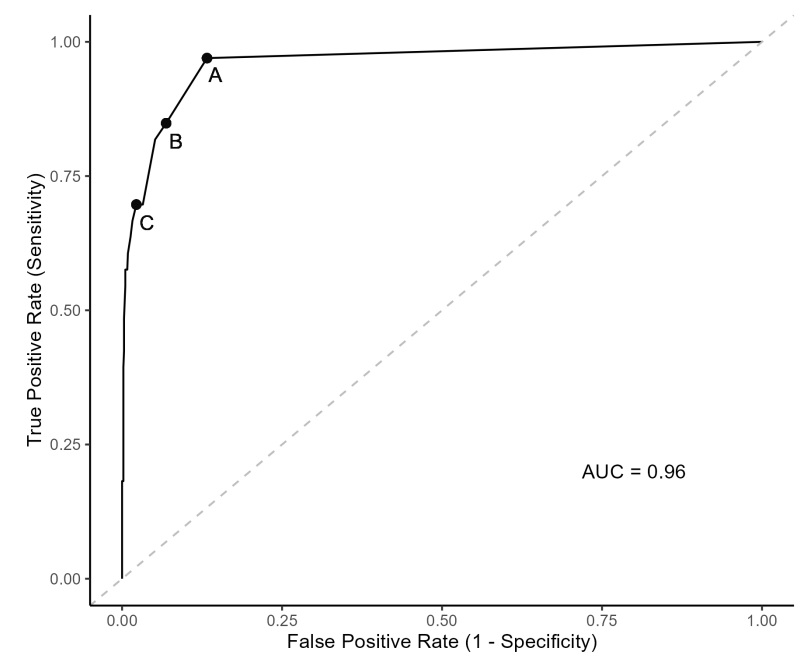


- Risk averaged from mid September to mid October
- Full risk zone (~103 000 km²)
- Remain mostly in the south western region ; (if $H_j > 10^{-3}$ ~25 000 km²)
- Limited risk of incursion in SP

Very good predictability of the model



- 99,9% locations with at least 1 outbreak were predicted at risk
- ROC curve of 0,96 (Se:97%, Sp:86,7%)
- Reducing the risk threshold leads to a significant drop in sensibility

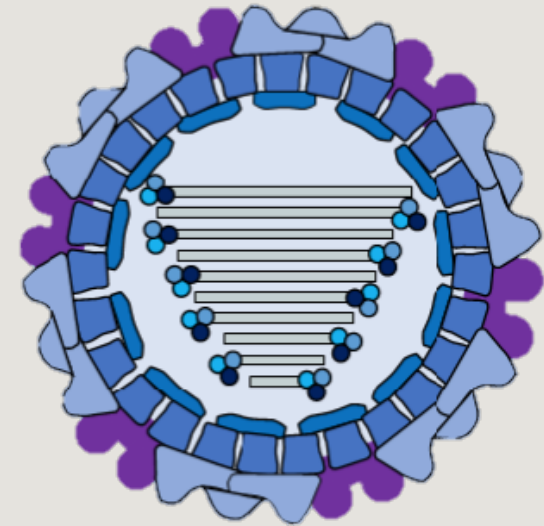


But under estimation of the short range dispersion
(26.3% Ob remain within the source zone)

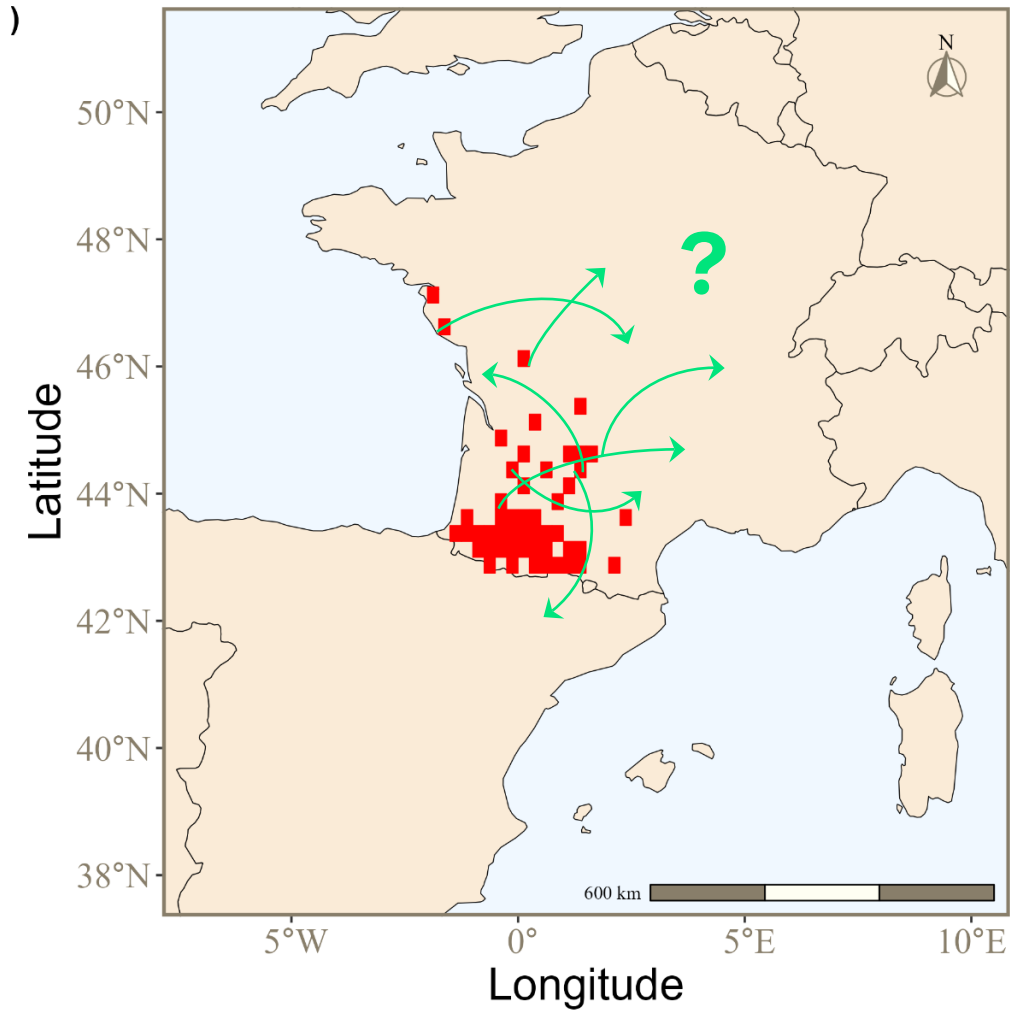
Situation	θ	TPR	FPR	Area (km ²)	Proportion
A	1.10^{-4}	97%	13.3%	103,125	100%
B	3.10^{-4}	84.8%	6.9%	60,625	58.8%
C	9.10^{-4}	69.7%	2.2%	28,750	27.9%

And what about next year ?

What could be the magnitude of the long-distance dispersal zone in early 2024?



EHDV-infected area (early December 2023)

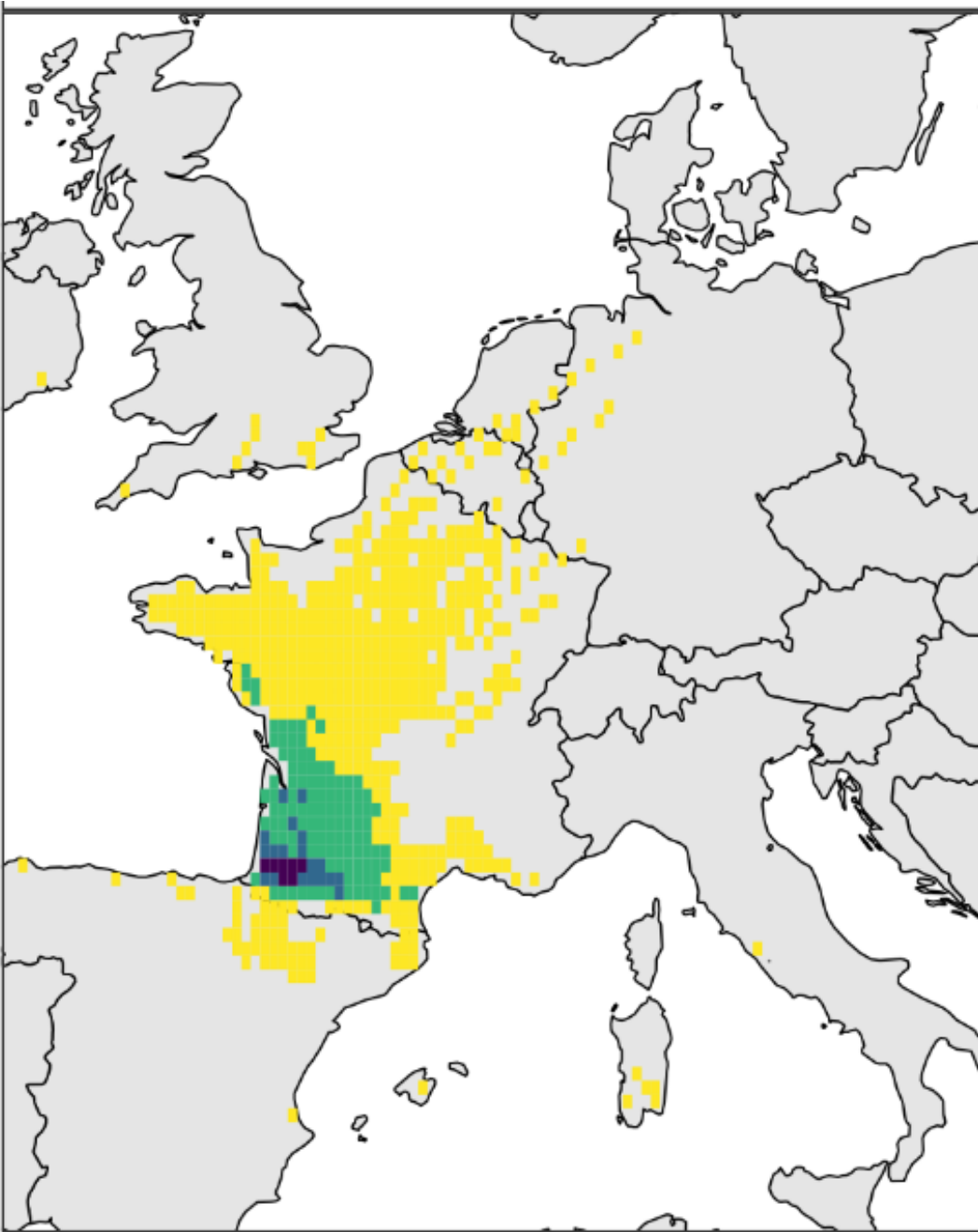


- Areas where at least 1 EHDV outbreak occurred in France
- WOAHA data at December 1st
- Main assumption: all red locations start to be infectious again all at the same time, mid of March (week 11)

What are the locations at risk of long-distance dispersal by the wind?

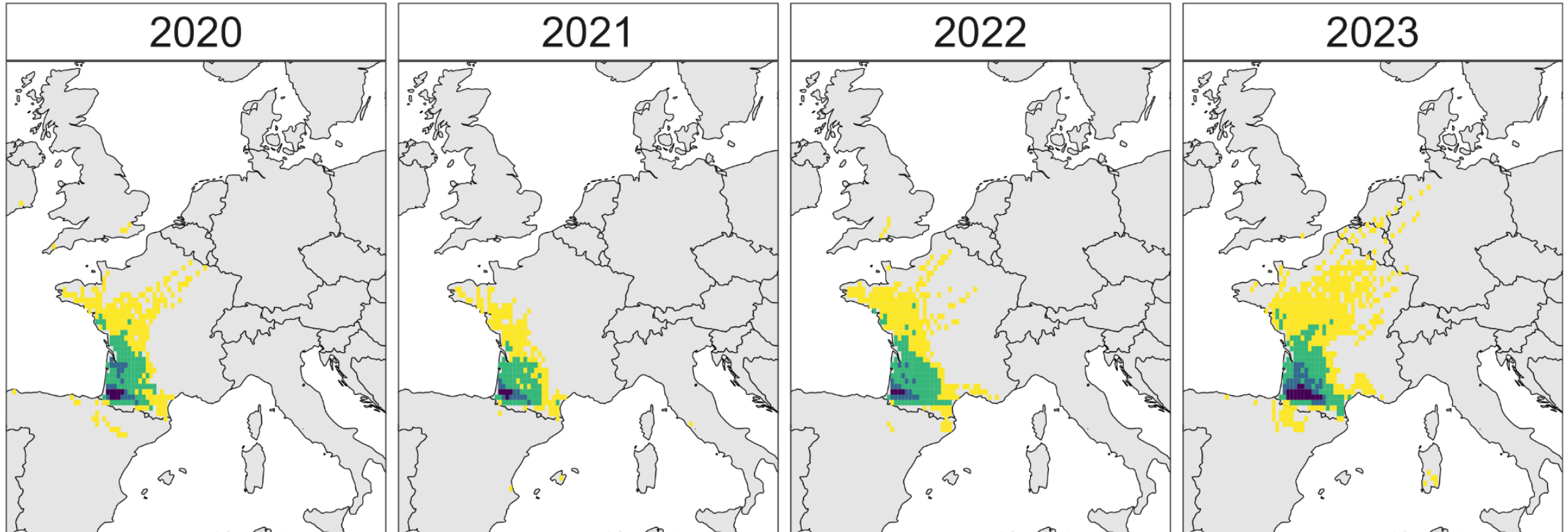
Risk zone averaged over the 2020-2023 period

- Mostly the western half of France
- Extreme destinations in UK, Ireland, Belgium, the Netherlands, Germany, the Balearic Islands, southern Sardinia, and the western border of Italy
- Pyrénées & Massif Central mountains act as orographic barriers



$H_j(z_2, T_5)$ ■]0,0.0001] ■]0.0001,0.0005] ■]0.0005,0.001] ■]0.001,0.005]

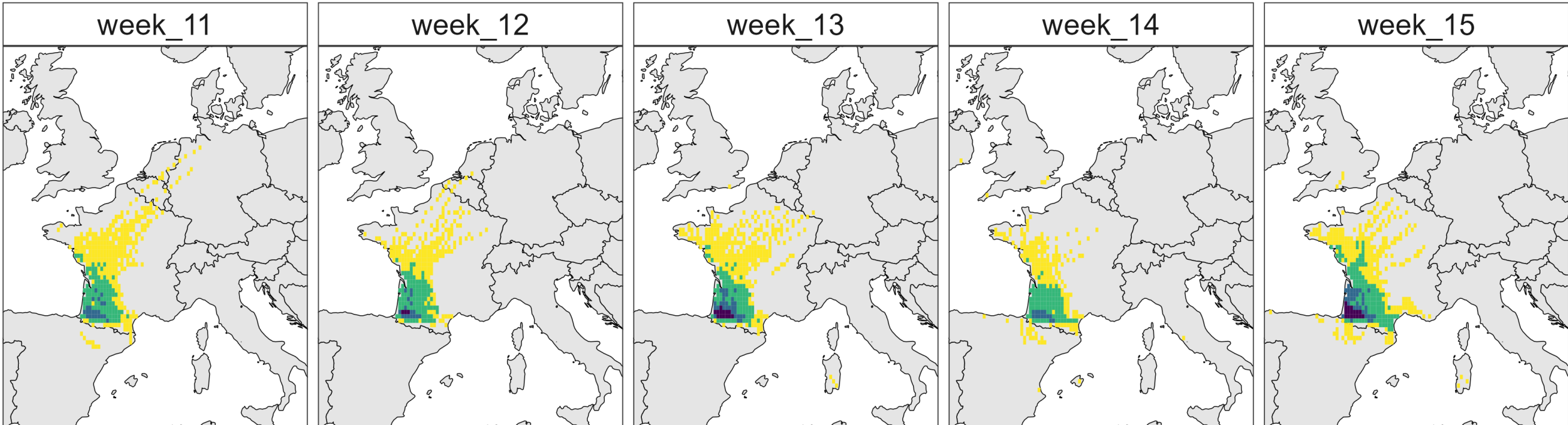
Variability over years



$H_j(z_2, T_5)$ ■]0,0.0001] ■]0.0001,0.0005] ■]0.0005,0.001] ■]0.001,0.005]

H_j : probability of long-distance dispersal for each destination j over a 5 week period (from W11 to W15; from mid-March to mid-April)

Variability over weeks



$H_j(z_2, T_1)$ ■ (0,0.0001] ■ (0.0001,0.0005] ■ (0.0005,0.001] ■ (0.001,0.005]

H_j : probability of long-distance dispersal for each destination j over a 5 week period (from W11 to W15; from mid-March to mid-April) – averaged over 2020-2023



Conclusions & perspectives

- Our model is able to **rapidly assess the risk of *Culicoides* long-range dispersal** by the wind
- It consider **historical variations** of wind speed, directions at **large scale** of Europe
- Provides insights to **identify at-risk locations** at risk from a known area with a very **good predictability** (over 5 weeks)
- **Flexible** tool (source locations / *Culicoides* borne disease /time period..)

However...,

- Risk of *Culicoides* wind dispersal \neq disease introduction
- Under estimate short-range dispersal
- Source area to be updated according to the disease epidemiology

Windi App

A web application for data visualisation and analysis



WINDi

Home

Initiation Area

Specific Week

Map per year

Map per week

Initiation Area

Select a country

Netherlands

Select an option

Forward

Select a color parameter

Change

Run

Run For Weeks Run For Years

Specific

Break 1

0

Break 3

0,005

WINDi



Home

Initiation Area

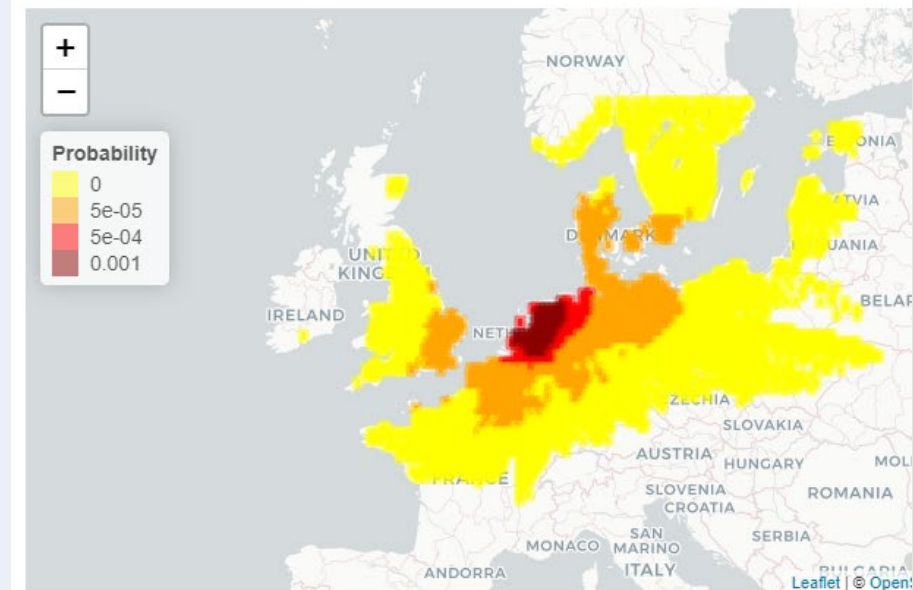
Specific Week

Map per year

Map per week

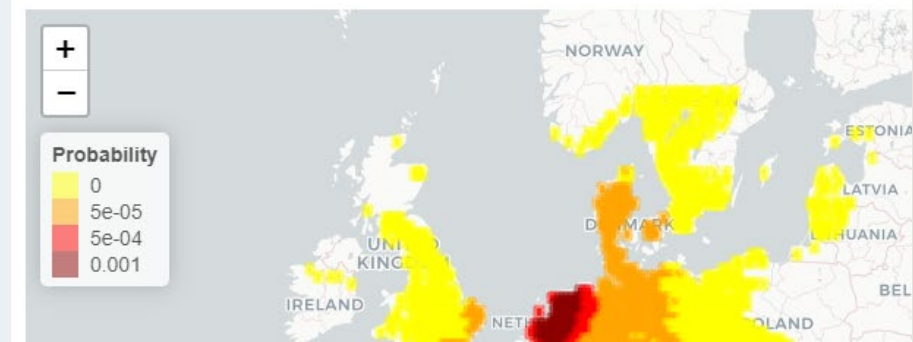
Plot per year

2020



Download

2022



Thank you for your attention

Thanks also to all the contributors



Thibaud Porphyre



Karine Chalvet-Monfray



Albert Picado



Davide Martinetti



Dorian Rollin



Melody Temperville



Anna Artiges

