

**Overview of a tool for assessing the risks of
introducing infectious animal diseases into
Belgium (RT 18/02 MORISKIN project).**

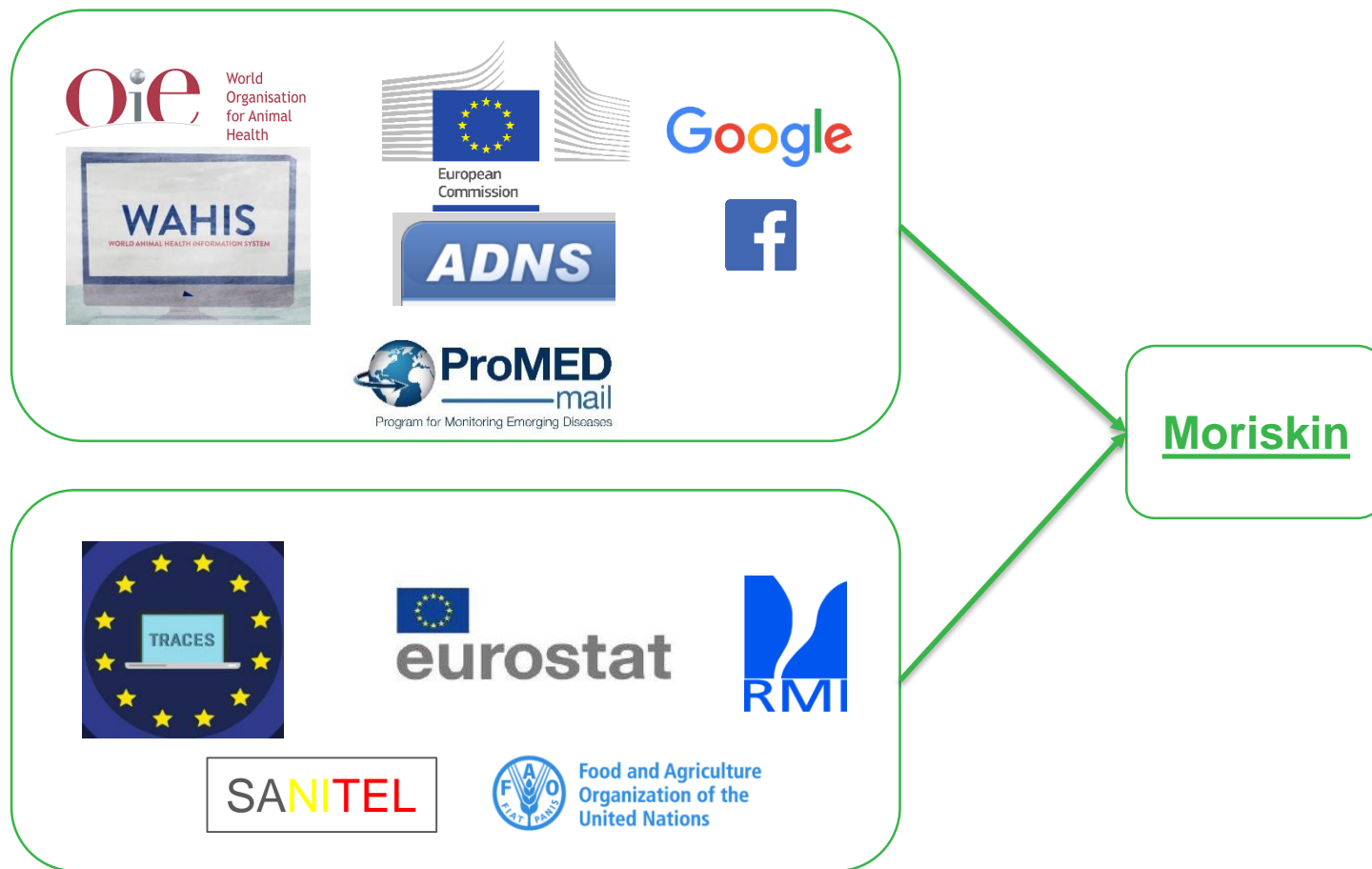
Xavier Simons

PROJECT CONTEXT

- RT project: 01/2019 – 04/2021
- Project partners:



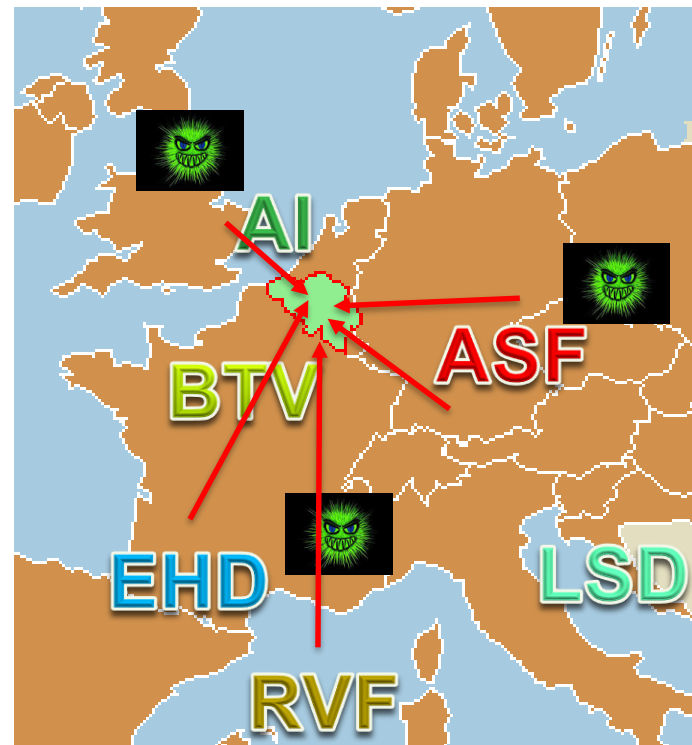
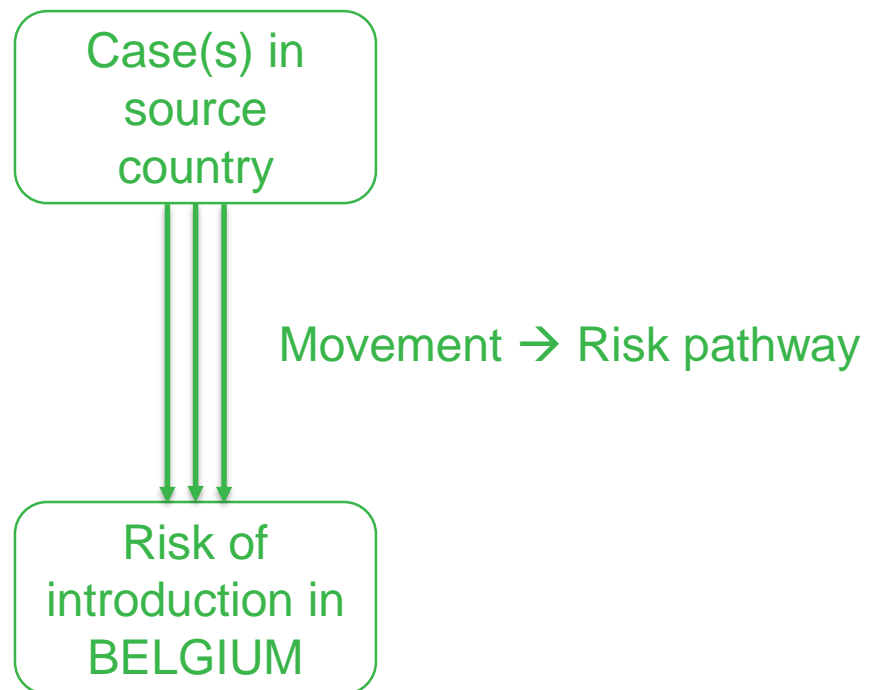
Context



GOAL: develop a method → tool (Web app) for risk assessment and threat analysis.
Focus on the **risk of entry** of infectious pathogen in BE.

→ ASF as case study

Method : simple concept



ASF – introduction pathways



AFRICAN SWINE FEVER

Trade activities: Importation (legal or illegal) of

Animal products.

- Traded freely within the EU
- Imports from third countries
- Personal imports carried into the country by travellers.

Live cultures
for example, germplasm,
vaccines, reagents, tissue
cultures, diagnostic
samples etc.

-Wildlife
spreading
the disease
to livestock
(e.g. wild
boars)

Live animals
- Live infected/Exposed
animal(s) from EU or third
countries
- Live animals carrying a
disease vector.

Human activities

- **Movement of people.** Tourism and trade related to visit to livestock premises. Hunter
- **Air, sea or land transportation activities.** Planes, ships, train, cars and military

Natural phenomena

- Arthropod vectors (insects) blown on the wind
- Windborne spread of the infectious agent itself e.g. virus particles
- Wildlife movements/migration e.g. wild boars, birds migration



Legend

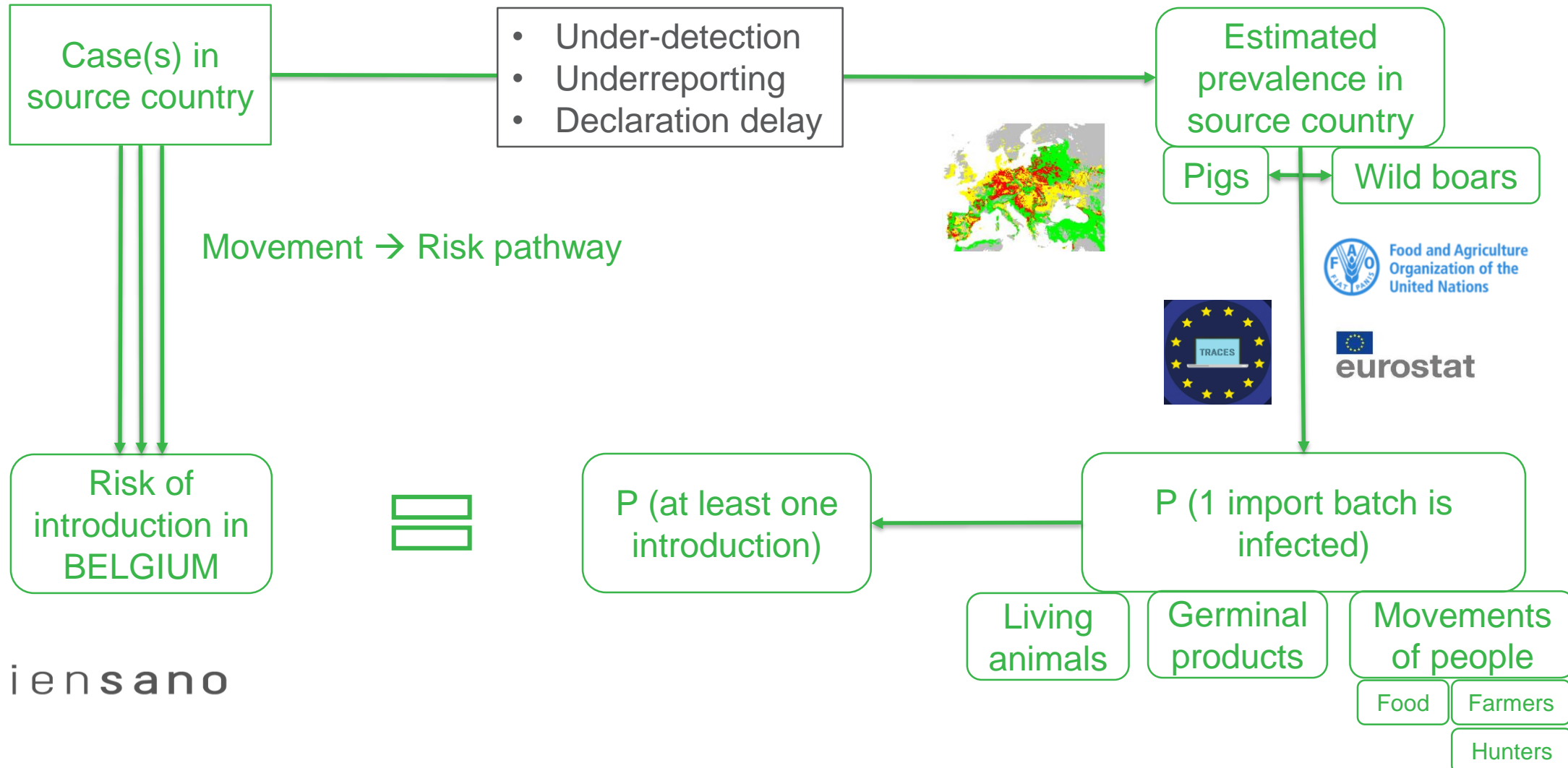
- Pathway considered in the risk calculation in the application
- - - → Pathway which is possible but not considered in the risk calculation in the application as it is not applicable for the Belgian context

No arrow implies this is not a pathway for the disease in question

Words coloured in black are the specific category taken into account for the risk calculation

Method : simple concept

By country and time point
(week)



Moriskin: R-Shiny Web application

Risk of entry

MORISKIN project: V1.2

HOME

RISK ASSESSMENT ▾

DISEASES DISTRIBUTION ▾

TRADE AND MOBILITY ▾

POPULATION DISTRIBUTION ▾

INFOS ▾

Risk assessment for disease introduction in Belgium



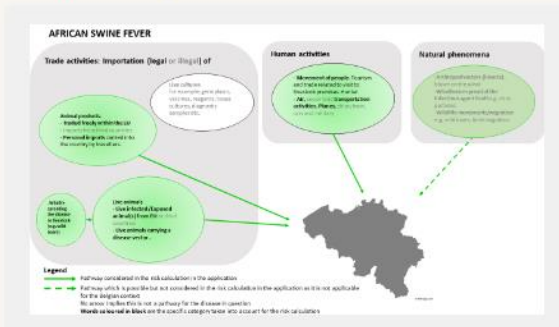
Date of last data update : 2024-01-26

Disease data source : ADIS

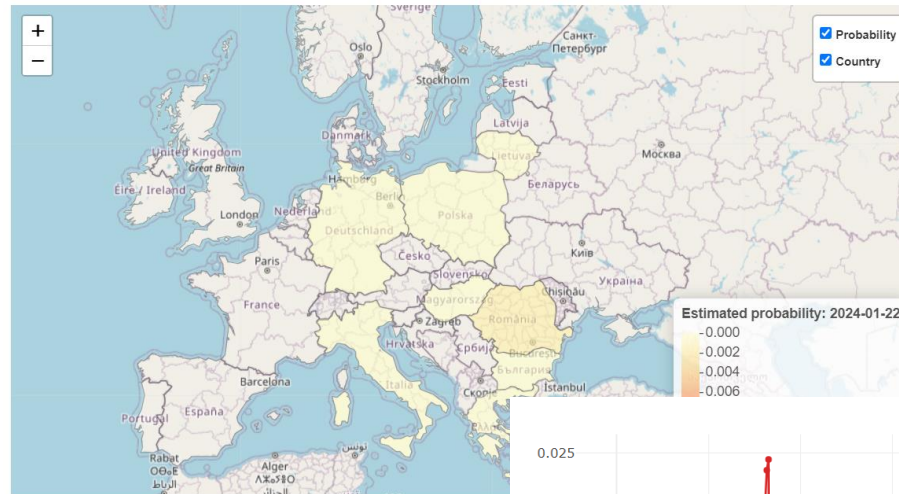
Date input for mapping

2024-01-22

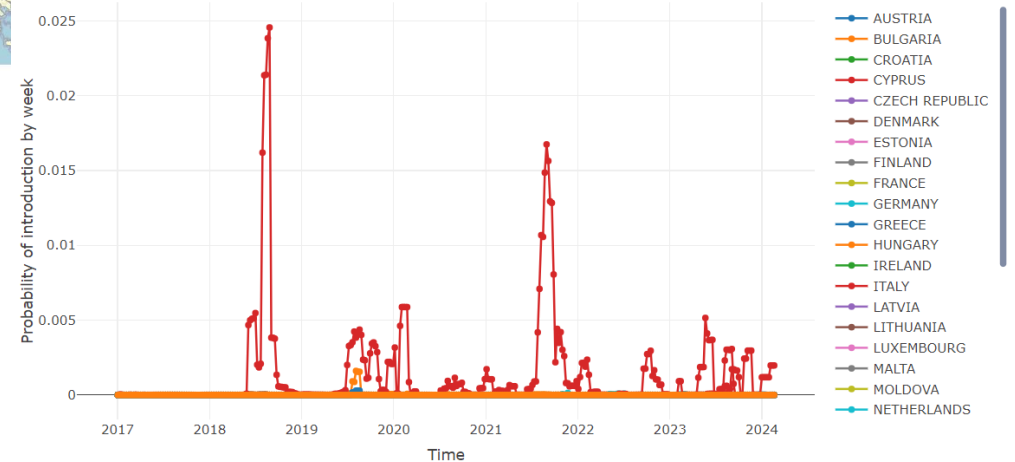
Risk pathways:



Weekly risk of introduction of ASF in BE from



All pathways combined



Moriskin: R-Shiny Web application

Disease distribution

Graph selection

Select data source (ADNS/OIE)

Select the disease

Select the species

Set time period for the map

The screenshot displays the 'MORISKIN project: V1.0' interface. The navigation menu includes: HOME, POPULATION DISTRIBUTION OVERVIEW, POPULATION DISTRIBUTION, TRADE AND MOBILITY, DISEASES DISTRIBUTION (selected), RISK ASSESSMENT, and DATA ACCESSIBILITY. The main content area is titled 'Cases report on the selected disease' and contains the following elements:

- Disease:** A dropdown menu with 'African Swine Fever' selected.
- Species:** A dropdown menu with 'Wild Boars' selected.
- Minimum date:** A date range selector from 2015-12-28 to 2020-05-04.
- DOWNLOAD:** A button to download the data.

Below the form, a note reads: 'Please consider the relationship between disease and host. An error will be sent by the system if you query any incompatible couple of host & pathogen.'

The 'ADNS database' section offers two views: 'MAP' and 'TIME SERIES'. The 'TIME SERIES' view is active, showing a line chart titled 'African Swine Fever in Wild Boars'. The y-axis is 'Amount of declared infected animals' (0-350) and the x-axis is 'Time (aggregated by week)'. The chart shows multiple colored lines representing different countries, with a legend on the right listing: ESTONIA, ITALY, LATVIA, LITHUANIA, POLAND, UKRAINE, CZECH REPUBLIC, HUNGARY, ROMANIA, BELGIUM, BULGARIA, SLOVAKIA, REPUBLIC OF SERBIA, and MOLDOVA.

The 'MAP' view shows a map of Europe with countries shaded in red and orange, indicating the number of infected animals. A legend for 'Infected Wild Boars' shows five categories: 1-10, 10-50, 50-100, 100-1,000, and 1,000-10,000. A legend for 'Number of infected animals' shows two categories: Country and Number of infected animals.

Moriskin: R-Shiny Web application

Movements of live animals

Graph selection

Set time period for the maps

Select livestock species:
Cattle/goat/sheep/pig

Select animal type:
Production/slaughter

Select type of movement:
Count of animals / movements

MORISKIN project: V1.0

Animals movements from EU countries to Belgium

To query the livestock mobility database, please define:

Select period to be considered:

Livestock species:

Type:

Time series

DOWNLOAD

TRACES database

Imports of Live swine (for Production in BE)

Count of animals imported by week

Time

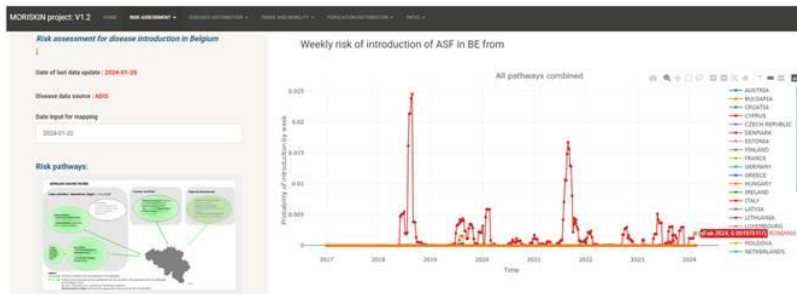
MAP

TIME SERIES

Number of traded animals

Using the tool

- Tool accessible only for FASFC and SPF
- Monthly updates
- Monthly e-mails



Résumé: Probabilité hebdomadaire moyenne d'introduction du pathogène en Belgique entre le 01/01/2024 et le 31/01/2024, par pays à risque et tenant en compte les voies d'introductions étudiées dans le projet MORISKIN.

Pays	ASF	LSD
Allemagne	$1.2 * 10^{-7}$	0
Bulgarie	$4.0 * 10^{-9}$	0
Grèce	$4.5 * 10^{-6}$	0
Hongrie	$1.2 * 10^{-9}$	0
Italie	$8.5 * 10^{-8}$	0
Lithuanie	$2.0 * 10^{-9}$	0
Pologne	$4.0 * 10^{-9}$	0
République Tchèque	0	0
Roumanie	0.0012	0
Suède	0	0
Tous les pays	0.0012	0

Code couleur:

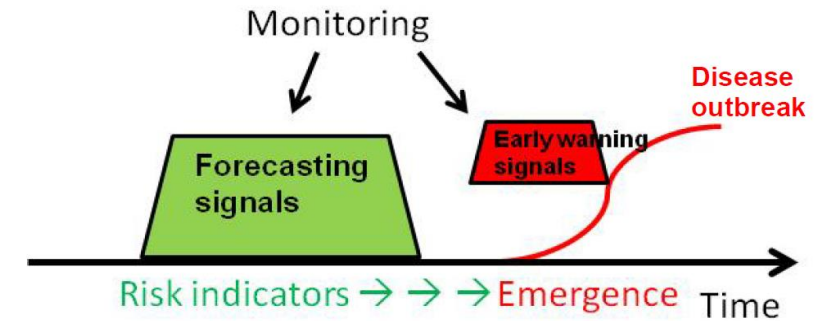
- Augmentation par rapport au mois précédent

- Alarme 1: Augmentation pendant plus de deux mois consécutifs

- Alarme T1: Au-dessus du seuil de 0.01

Appendix 1: Diagram describing the position over time of the different types of forecasting or early warning system of emergence of animal diseases

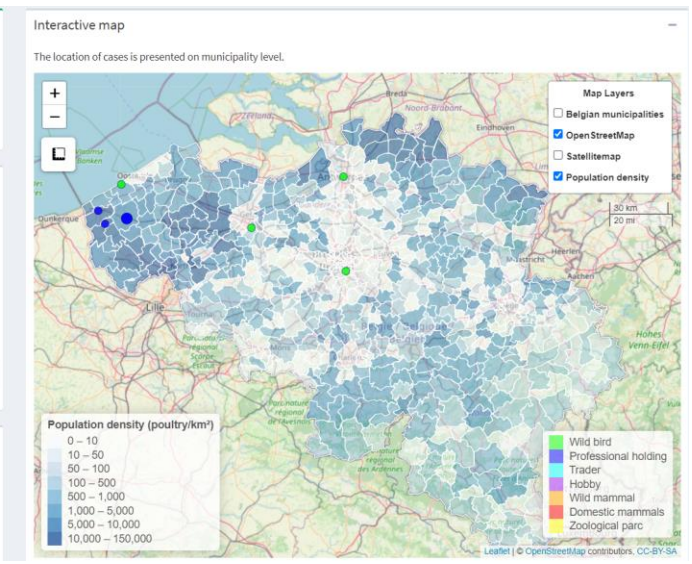
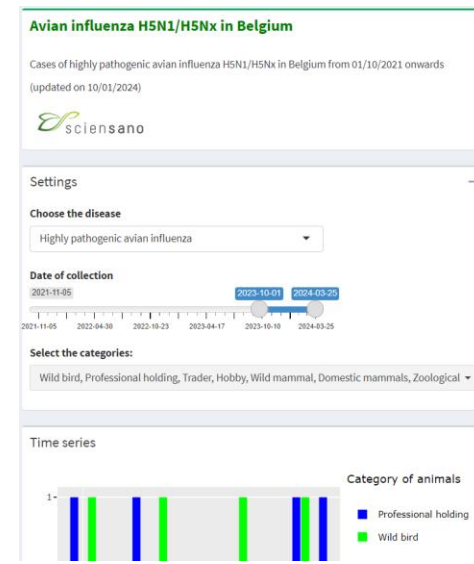
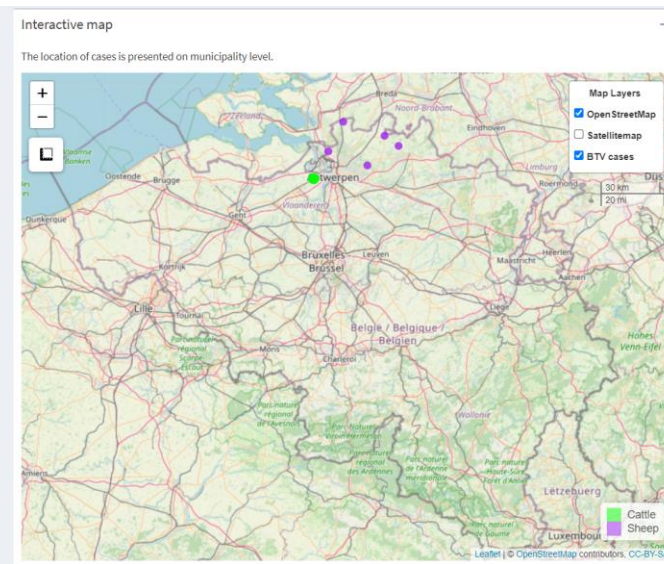
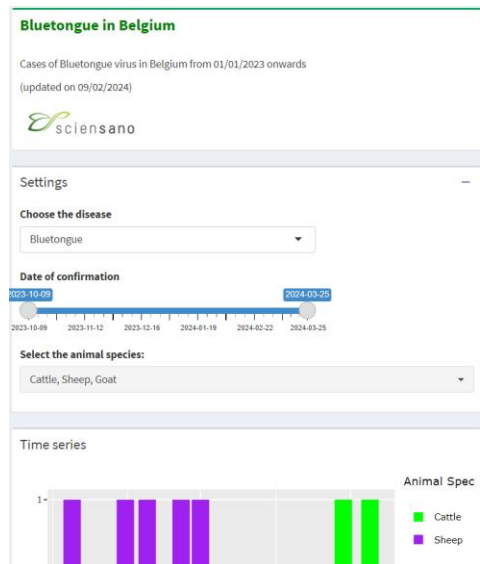
Emergence is represented by the red line (increase in incidence). Early warning is the detection of new cases of disease in the early stages of emergence. The generic and specific forecasting is a monitoring of risk indicators before actual emergence. The specific forecasting already target diseases, unlike the generic forecasting that target a priori any disease.



Avis 2016-10 SciCom AFSCA

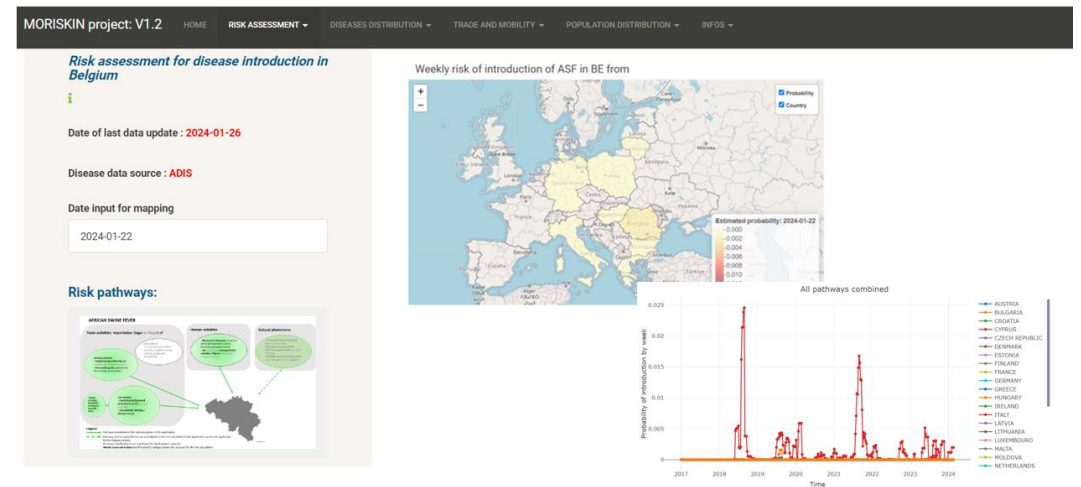
Other uses

- Moriskin R-Shiny server used to communicate on other diseases:
- See Sciensano Website



Conclusion

- Advantages:
 - Data from **various sources** in a unique tool → time saving!
 - **Frequent updates**
 - Dynamic and **user friendly** visualization
 - Risk maps + time series
 - Downloadable outputs
- Limitations:
 - Quality of the input data
 - Introduction ≠ infection
- Future perspectives
 - To monitor **more diseases**
 - To include more pathways (people by road, wind for vectors)
 - International collaborations





THANK YOU !



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