Modelling airborne particle movement at national scale: consequences for long-term surveillance of plant epidemics



Davide Martinetti ModStatSAP 2020

November 24, 2020

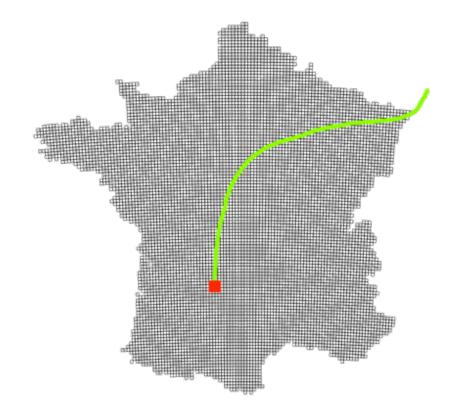


Objectives

- Model air-mass connectivity over the entire country from past observations
- Discover spatial and temporal patterns of connectivity
- How connectivity change the *"geography"* of the region
- Inform epidemic-surveillance



<u>Data</u>



Airmass trajectories

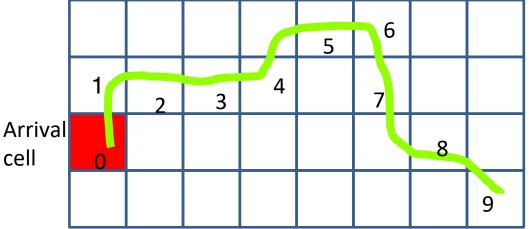
- 48-hours backward
- 500m AMSL
- Computed daily at 00-06-12-18
- From 01/01/2013 to 31/12/2018
- 8401 arriving points (8X8 km grid)
- Generated with HYSPLIT¹

TOTAL = <u>73.592.760</u> trajectories



¹ https://www.ready.noaa.gov/HYSPLIT.php 3





Connectivity matrix

$$M(i,0) = \begin{cases} 1 \forall i \in \{1, \dots, 9\} \\ 0 \text{ else.} \end{cases}$$

Averaging to grasp main signal

To reduce uncertainty, the connectivity matrices have been aggregated on yearly and hourly scales to obtain one matrix per day of the year: one day represents the average connectivity for the four moments of the days (00,06,12,18) across the 6 years of the study period (2013-2018)

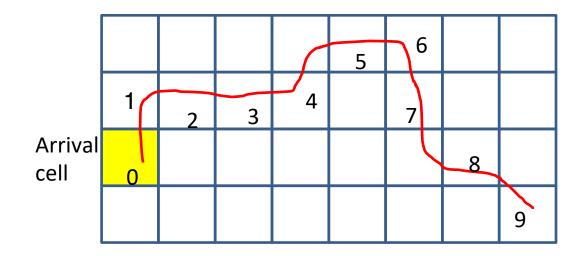




Sinks & sources

Every day, every cell is the '<u>sink</u>' of the airmass that arrives at its center. Its <u>in-</u> <u>degree</u> is the number of cells that the airmass has crossed before arriving to the destination.

In the example, the arrival cell in yellow has received an airmass that hovered 9 other cells in the previous 48 hours.

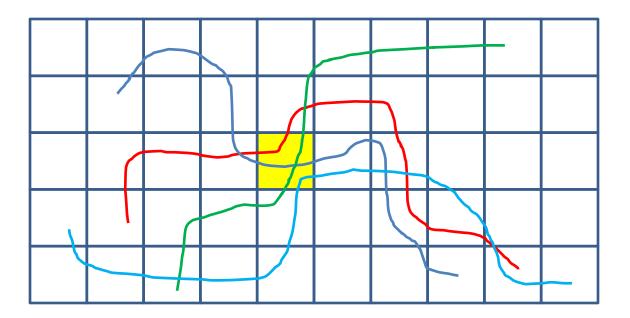






Every day, every cell is crossed by a certain number of airmasses that will end up in another cell. The number of airmasses hovering a cell is the <u>out-degree</u> and the cell is considered as a <u>source</u>.

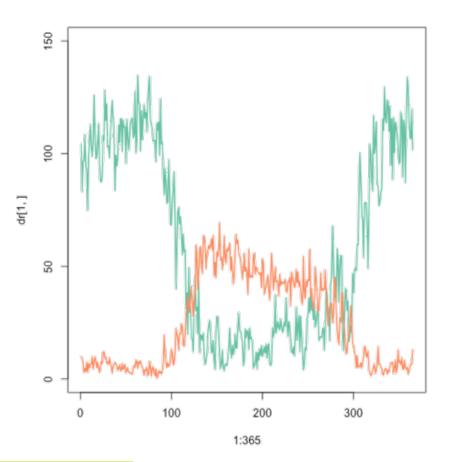
In the example, the cell in yellow has been crossed by 4 trajectories going elsewhere in the space.







The behaviour in terms of sink and source of a watershed can vary across the year.



Green : in-degree of a cell Orange: out-degree of the <u>SAME</u> cell.

The horizontal axis represents the days of the year.

Notice how this cell behaves as:

- a strong <u>sink in winter</u> months
- as a moderate <u>source during summer</u>

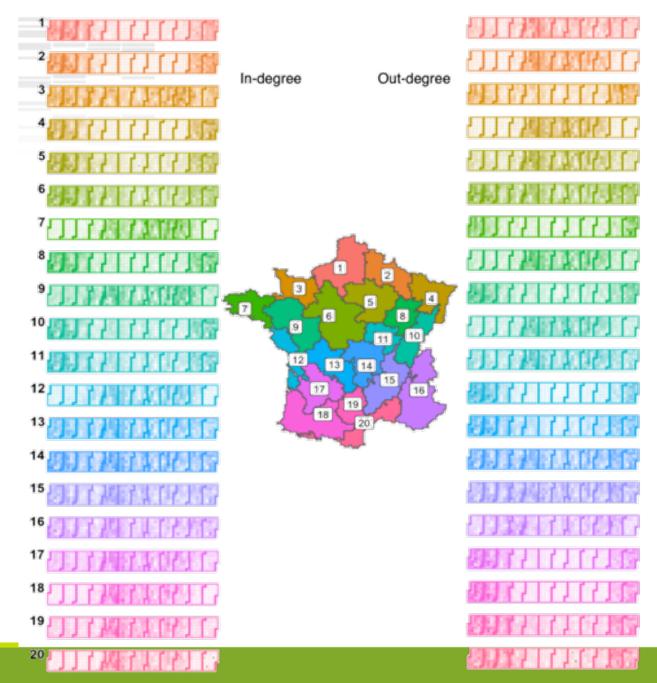


From observation to result

- Consider the daily in- and out-degree of each cell as a time series
- Compute Dynamic Time Warping (DTW) distance between in- and out-degree time series
- Sum the obtained distance matrices and obtain dendrogram to perform hierarchical clustering

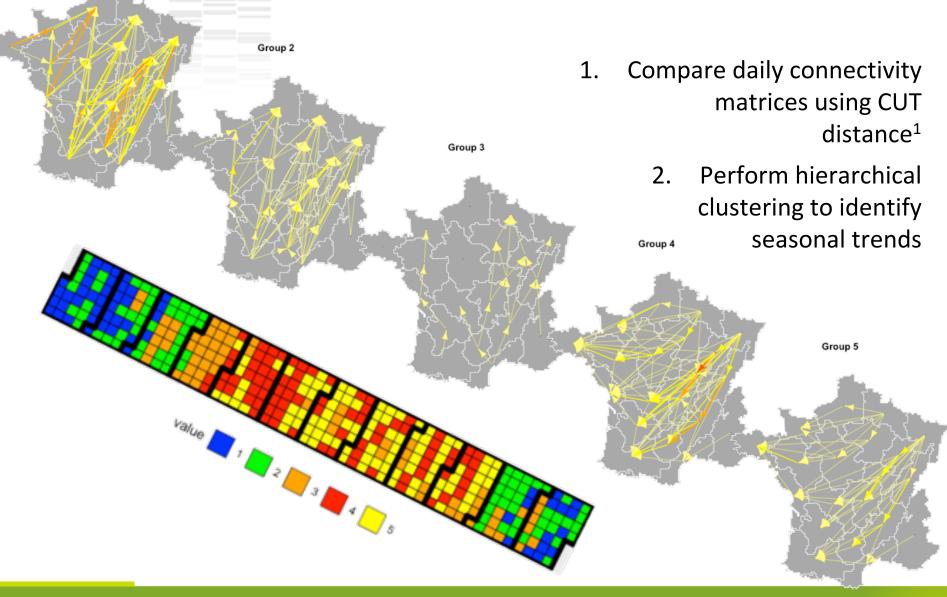








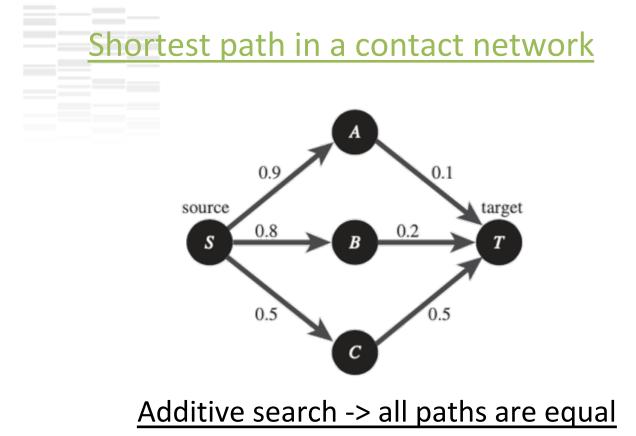
From observation to result





Group 1

¹Liu et al. (2018). Cut based method for comparing complex networks. Scientific Report 8 10



P(S,A)+P(A,T) = P(S,B)+P(B,T) = P(S,C)+P(C,T)=1

 $\frac{\text{Multiplicative search -> best path through node C}}{P(S,A)*P(A,T)=0.09 < P(S,B)*P(B,T)=0.16 < P(S,C)*P(C,T)=0.25}$





Effective distance

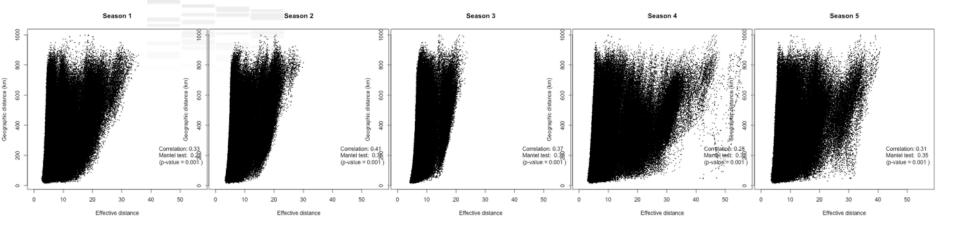
Simple transformation to transform contact probabilities into effective distances¹:

$$d_{ij} = 1 - \log(p_{ij})$$

Properties:

- $d_{ij} \ge 0$
- $d_{ij} = 1$ iff $p_{ij} = 1$
- $d_{ij} = \infty$ iff $p_{ij} = 0$
- $d_{ij} + d_{jk} \ge d_{ik}$
- Shortest-path network search: Dijkstra algorithm²

Geographic vs. effective distance

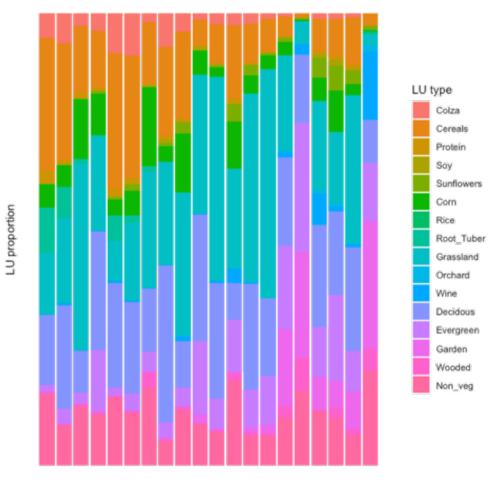


Season	Correlation	Mantel test
1	0.33	0.22***
2	0.41	0.36***
3	0.37	0.38***
4	0.28	0.32***
5	0.31	0.35***



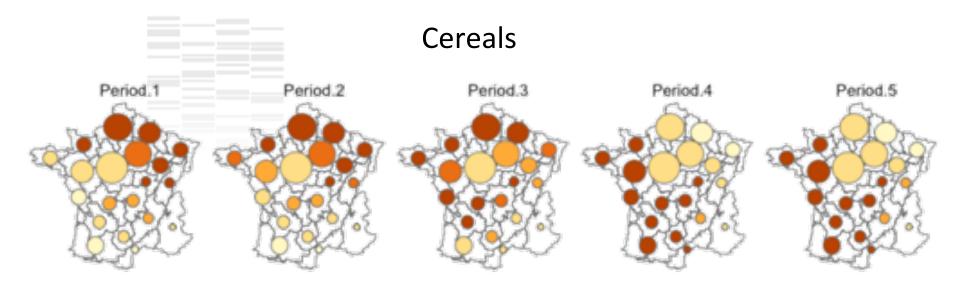
How airmass connectivity could inform epidemic surveillance



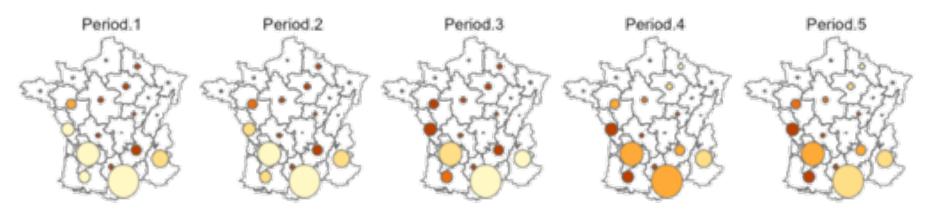


1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 Region





Wine



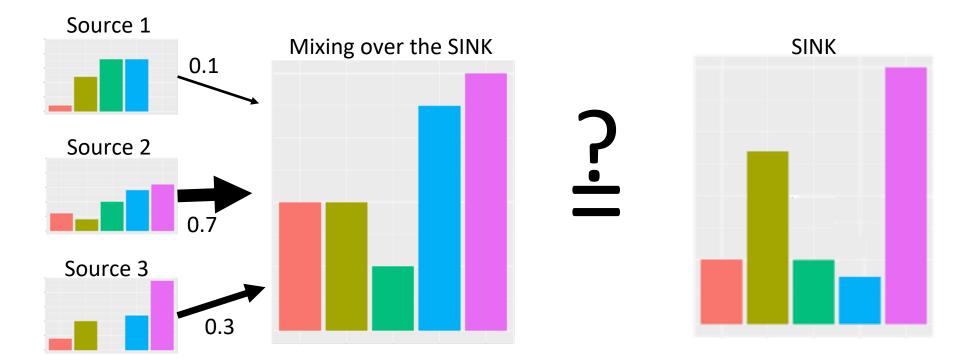


Circle size proportional to surface of LU type Colour according to sink receptivity

15

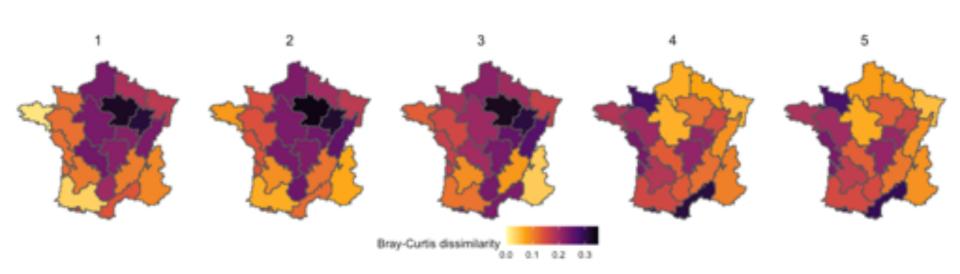


Bray-Curtis dissimilarity quantifies the compositional dissimilarity between two different populations

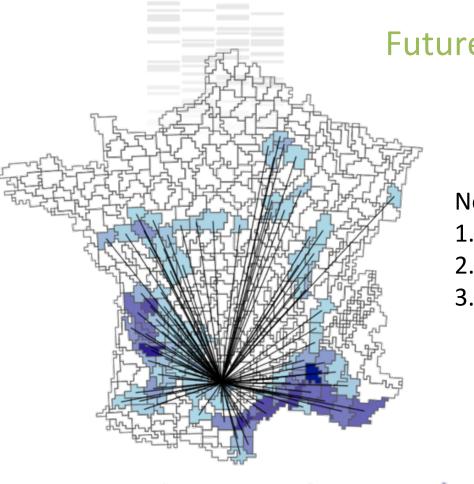




An ecological approach



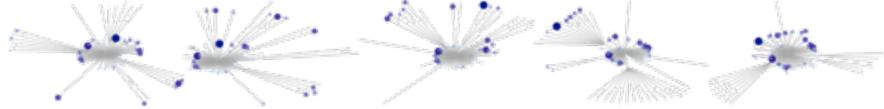




Future works

Network topology favours epidemic spread:

- 1. Surveillance: where & when to monitor?
- 2. Dangerous sites & seasons
- 3. Most probable paths of dissemination after outbreak







Thanks for your attention!

