



Rating potato varieties:

30 years of experiments reanalyzed to explicit resistance and explain its variability.

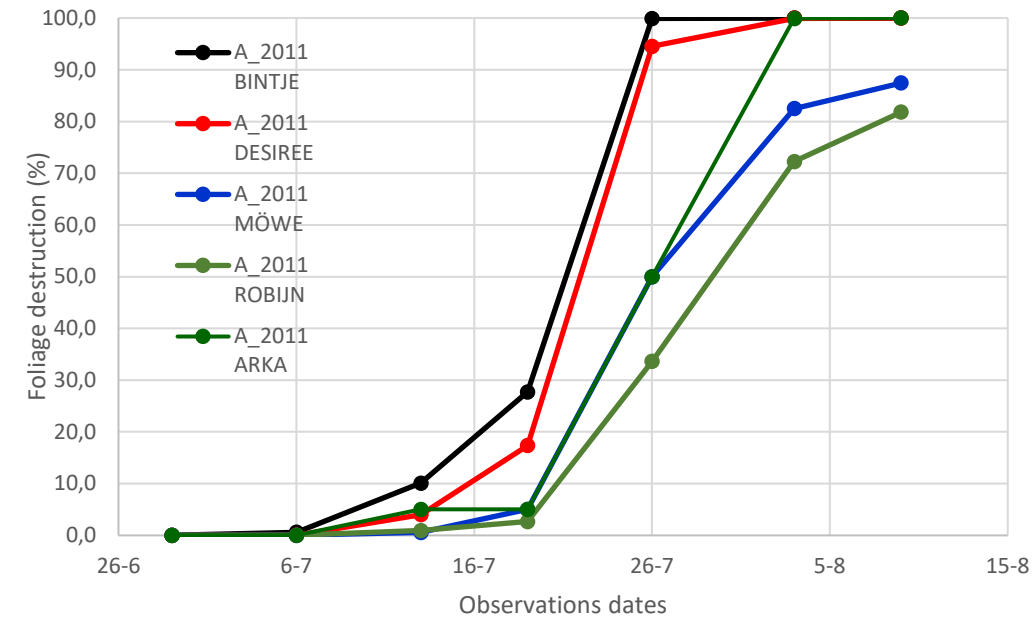
19th EuroBlight Workshop 13-16 May, 2024

# Context

- Varietal resistance : reduce disease and needs for treatments
- Variability : difficulty for prediction
- Goal :
  - explore epidemics diversity
  - create types
  - evaluate predictability
- In this presentation :
  - Epidemic diversity and discrimination types created
  - Comparison with official ratings
  - Variables importance to predict epidemic type

# Datasets

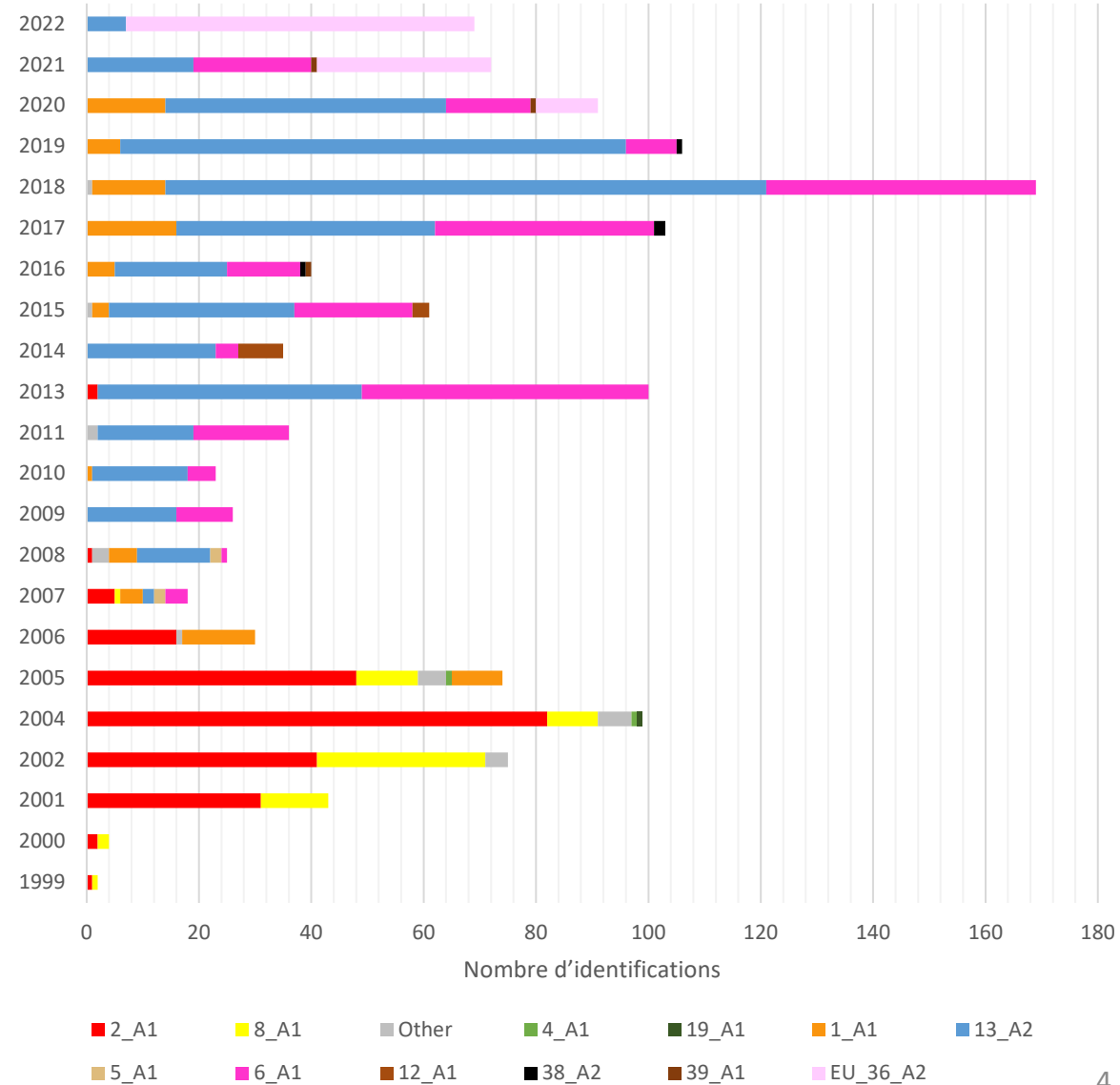
- Epidemic dataset :
  - 2526 sets of foliage destruction observations
  - 1 site : Ploudaniel, France, oceanic climate
  - 29 years, from 1994 to 2022
  - 201 potato genotypes
    - 43 varieties (1427 curves)
    - 23 differential hosts (759 curves)
    - 3+120 breeding lines (340 curves)



# Datasets

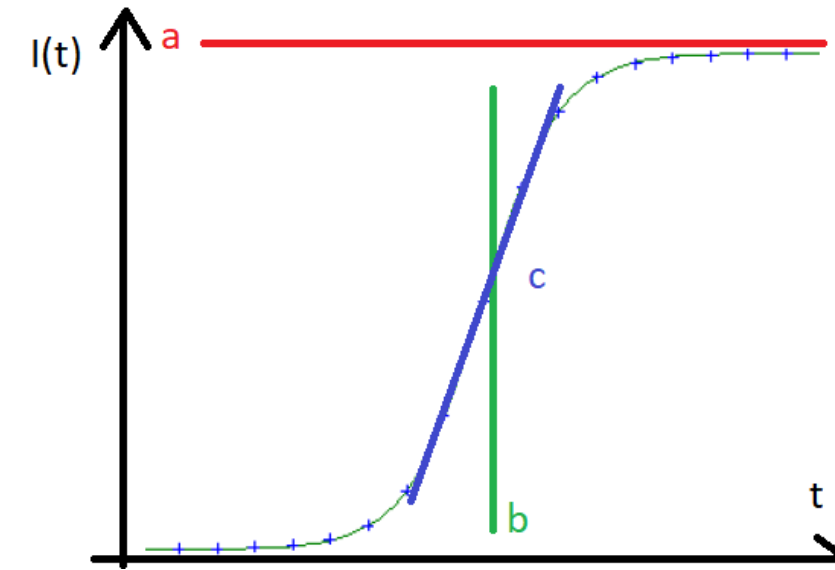
- Climate dataset :
  - Weather station in Ploudaniel : daily temperature, humidity and wind
  - Variables : means between planting and harvest
- *Phytophthora infestans* dataset :
  - From 1999 to 2022 : genotyped lineages from the fields
  - Variables : 1<sup>st</sup> and 2<sup>nd</sup> most frequent lineages, lineages diversity, % 1<sup>st</sup> most frequent lineage

Genotyped lineages in Ploudaniel



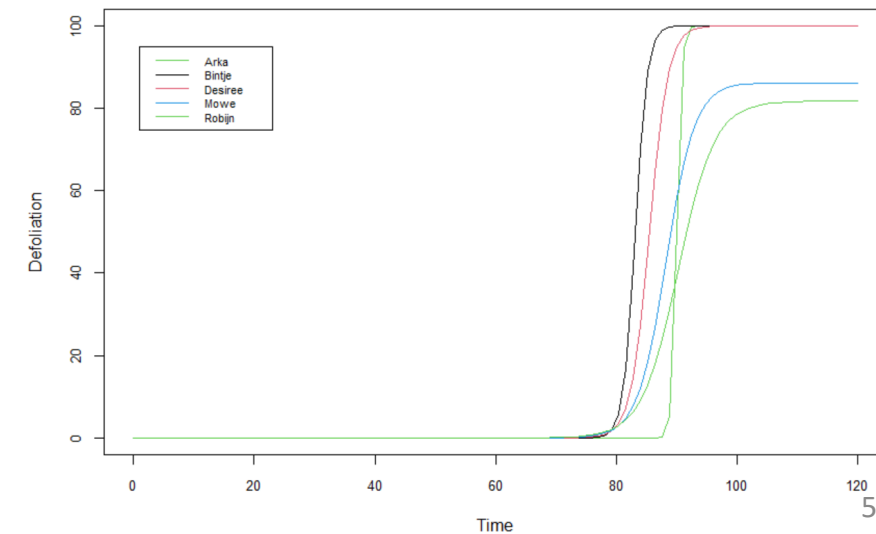
# Classification method

- Logistic regression :
  - a: asymptote, final severity
  - b: time 50% final severity (inflection point)
  - c: inverse of slope at inflection point
- PCA on a,b,c
- Hierarchical clustering on PCA results



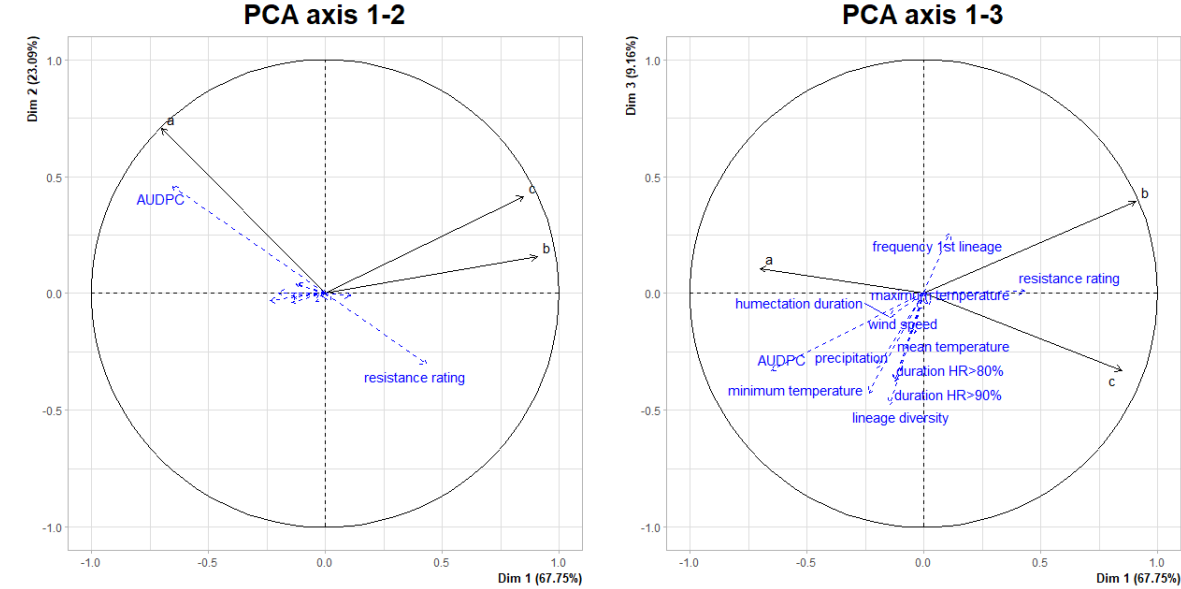
$$I(t, a, b, c) = \frac{a}{1 + \exp\left(\frac{b-t}{c}\right)}$$

Epidemic curves

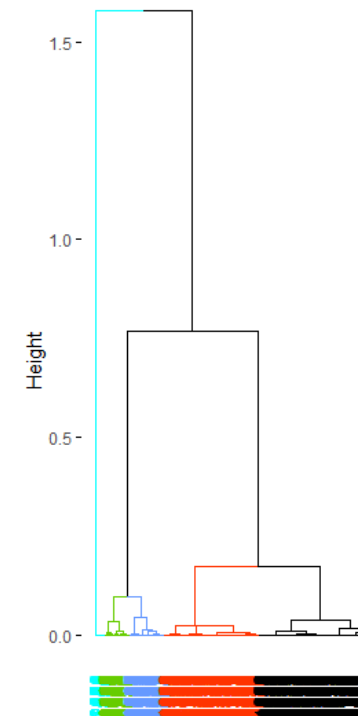


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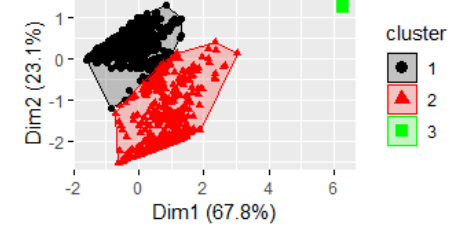
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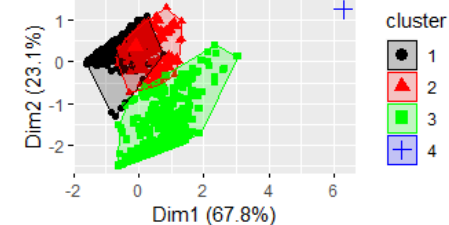
Cluster Dendrogram



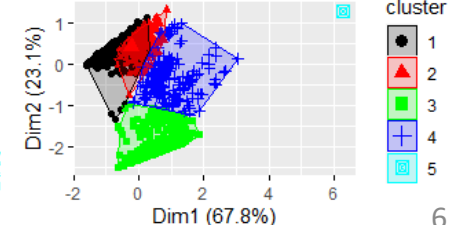
Factor map



Factor map

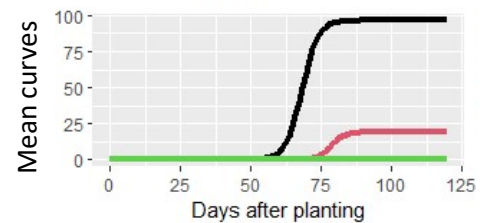
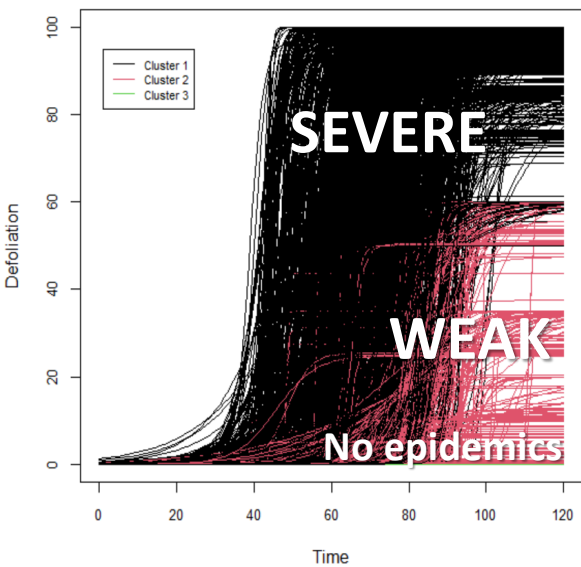


Factor map



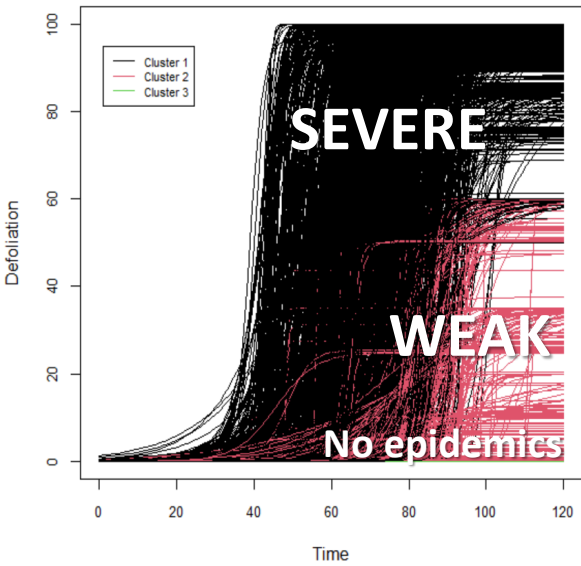
# Classification results

Epidemic curves

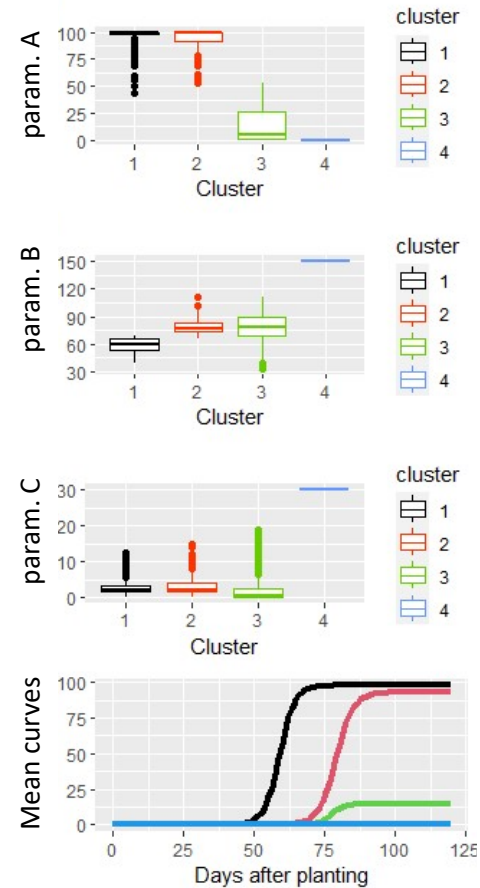
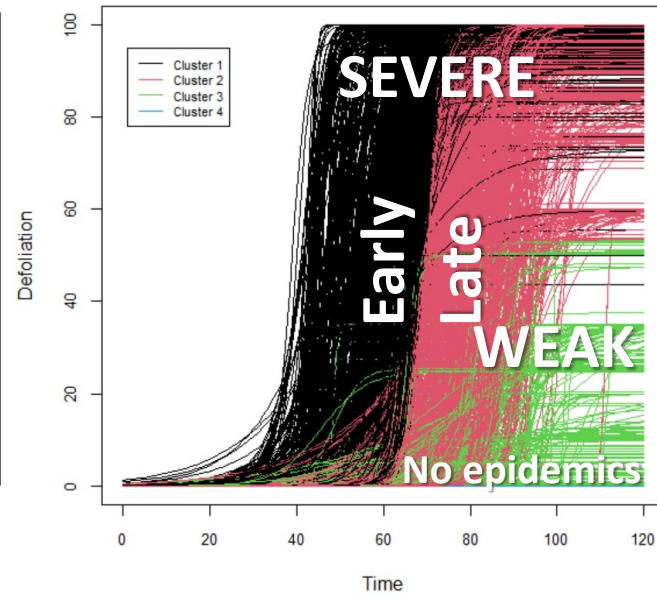


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Epidemic curves



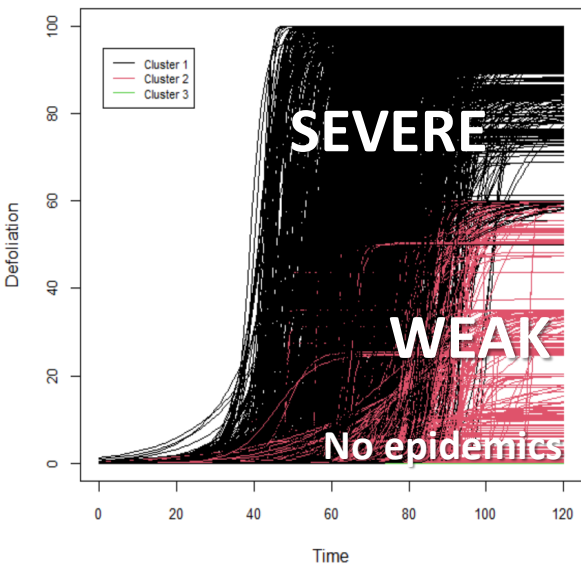
Epidemic curves



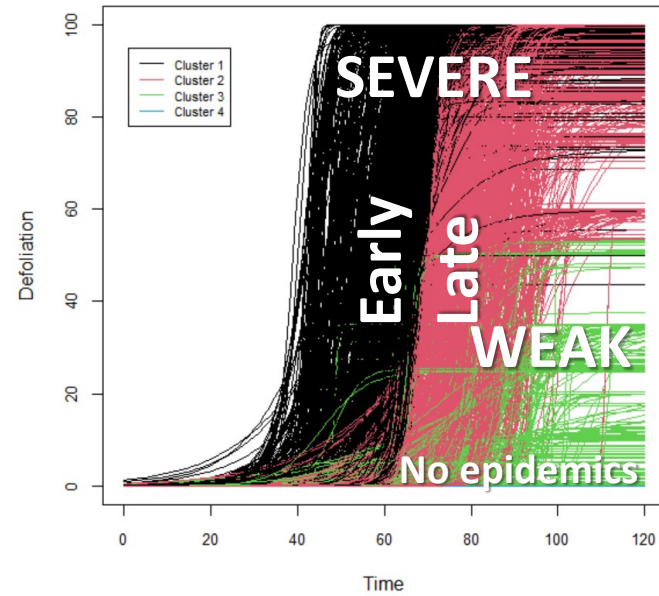


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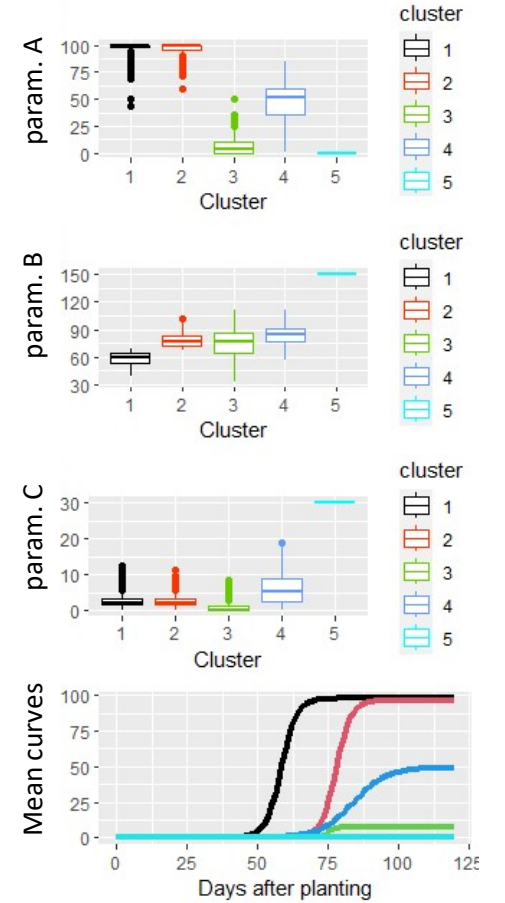
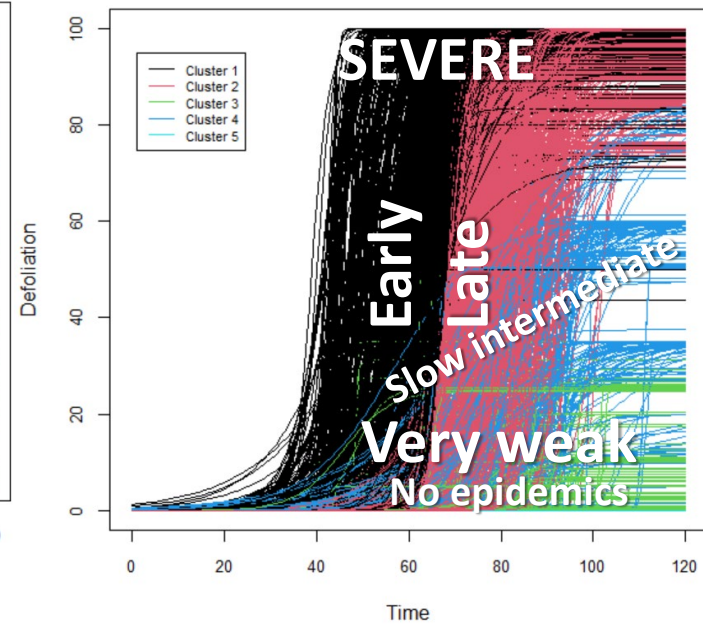
Epidemic curves



Epidemic curves



Epidemic curves

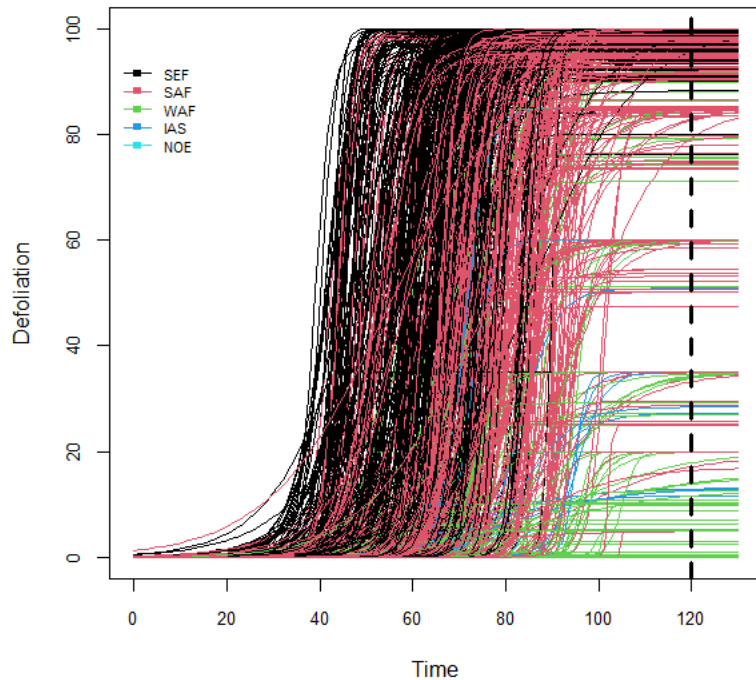


# Classification compared to official ratings

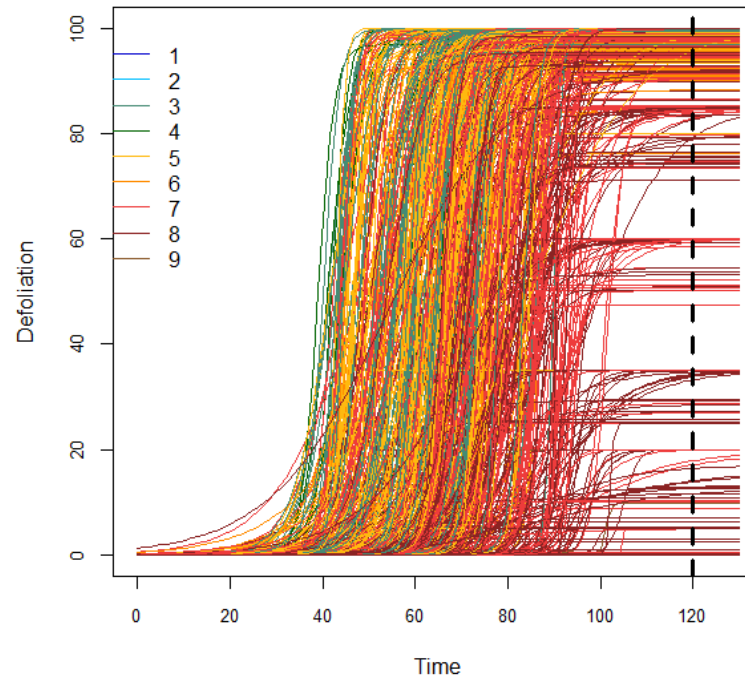
For 34 varieties with official resistance ratings

1 color = 1 variety : most frequent epidemic type or official rating

**Epidemic curves for epidemic types**



**Epidemic curves for ratings**

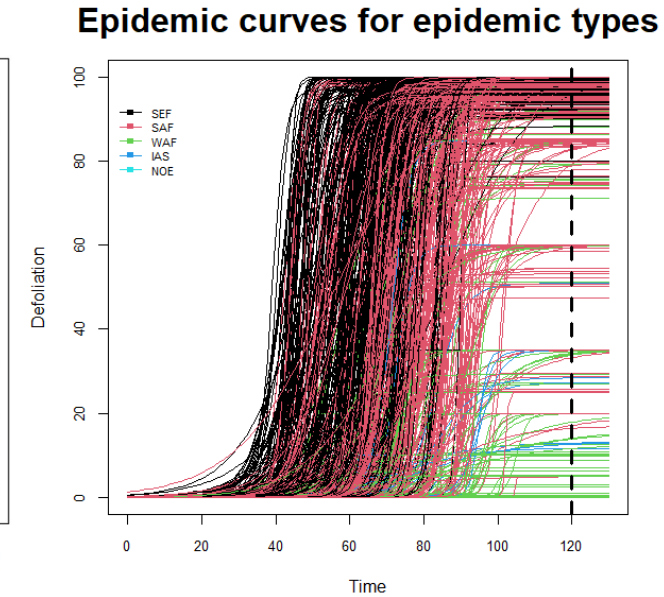
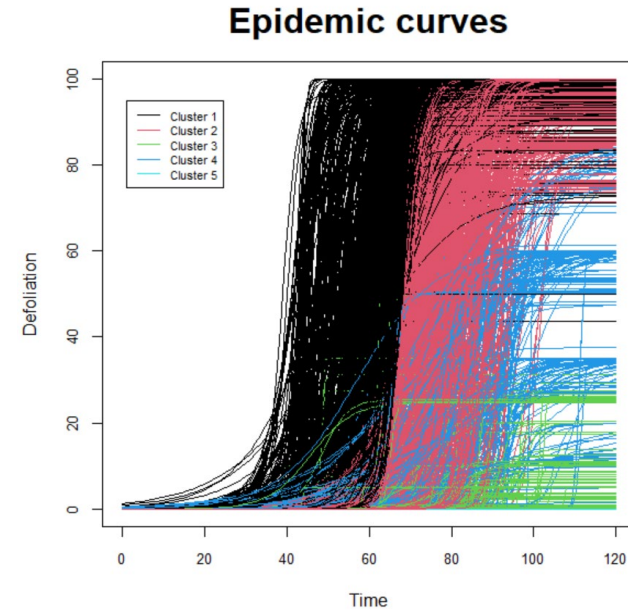


Discrimination between types clearer :

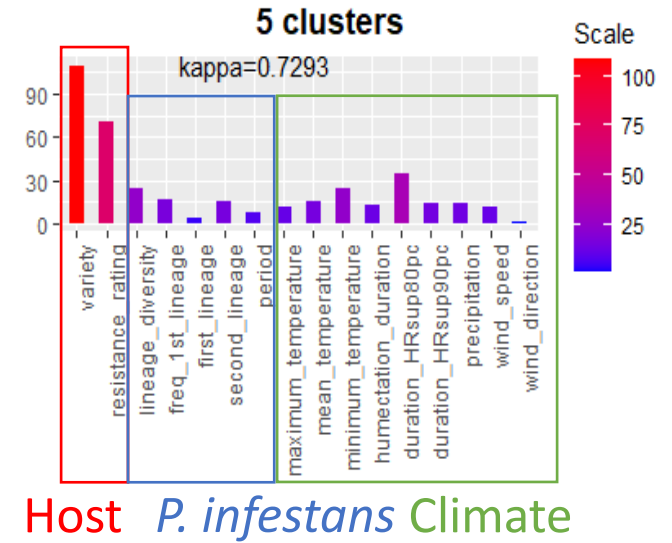
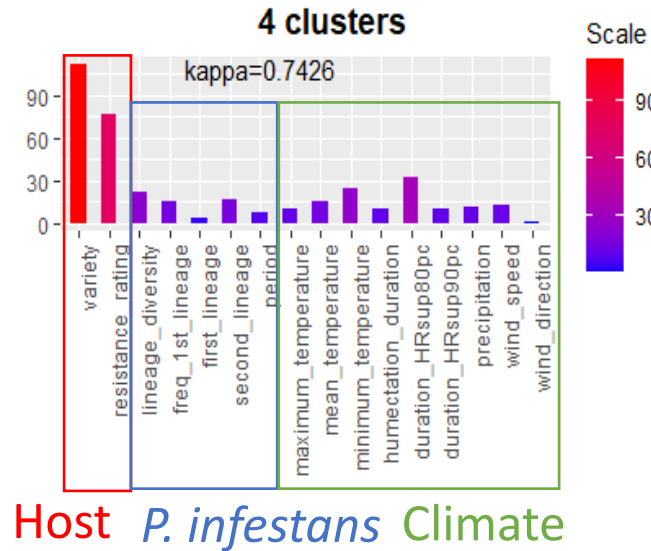
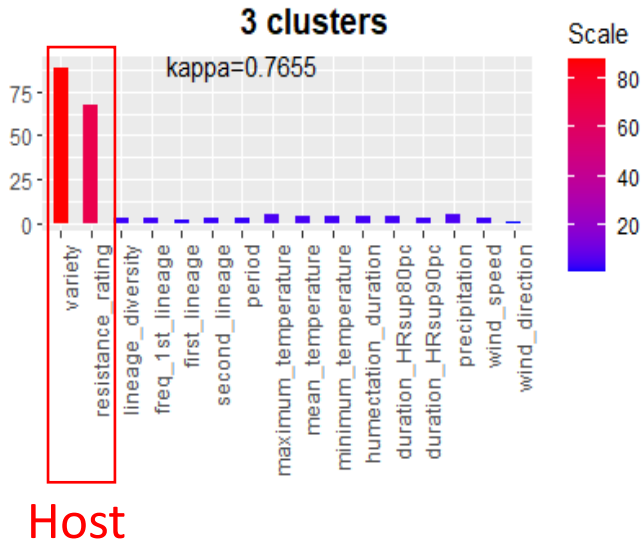
- Less groups
- Type assignment co-occurs observations and larger time period VS evaluations different ages and over 2 years

# Predicting epidemic types

- 5 epidemic types :
  - Linked to epidemic parameters
  - Apparently discriminative enough
  - Variability of varieties
- Predictability of types ?
- Method : random forest (machine learning) and importance of variables

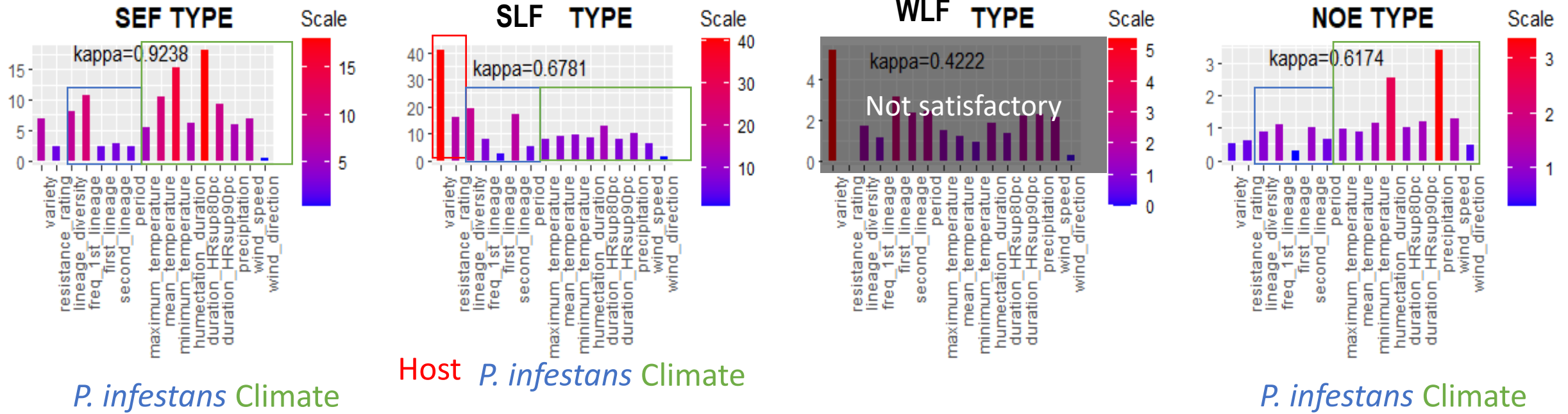


# Predicting epidemic types



- 3 cluster : host sufficient to predict
- 4 cluster and more : importance of pathogen and climate

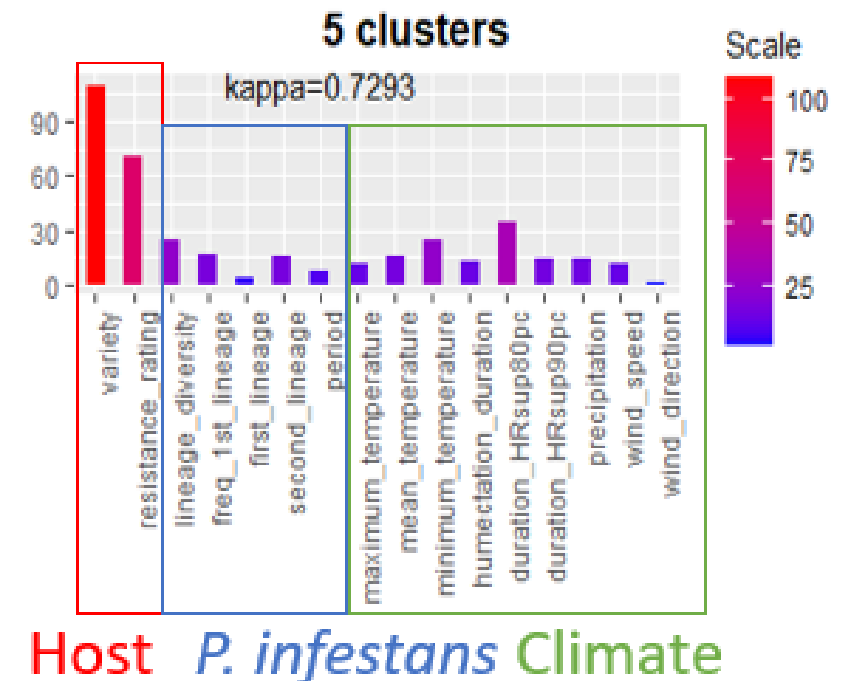
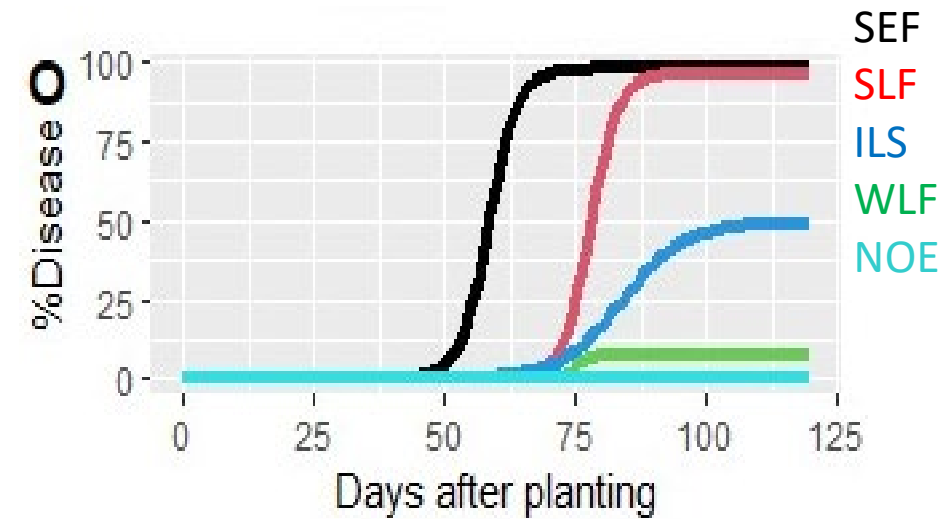
# Predicting epidemic types



- Severe early type : climate > *P. infestans* > host
- Severe late type : host > *P. infestans* > climate
- No epidemic type : climate > *P. infestans* > host

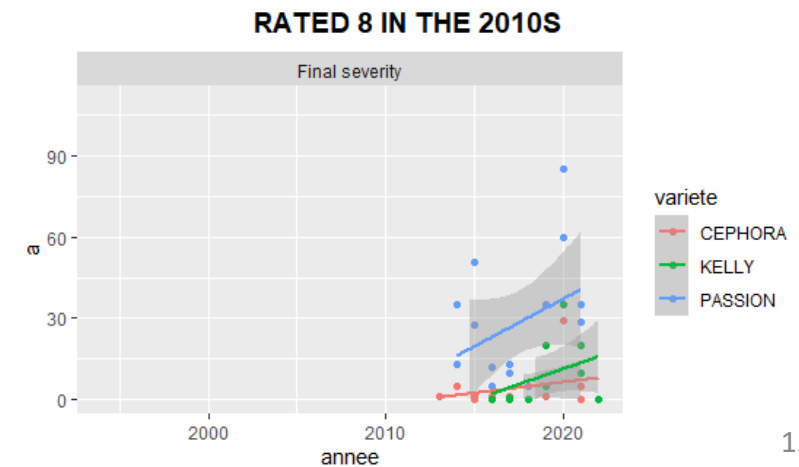
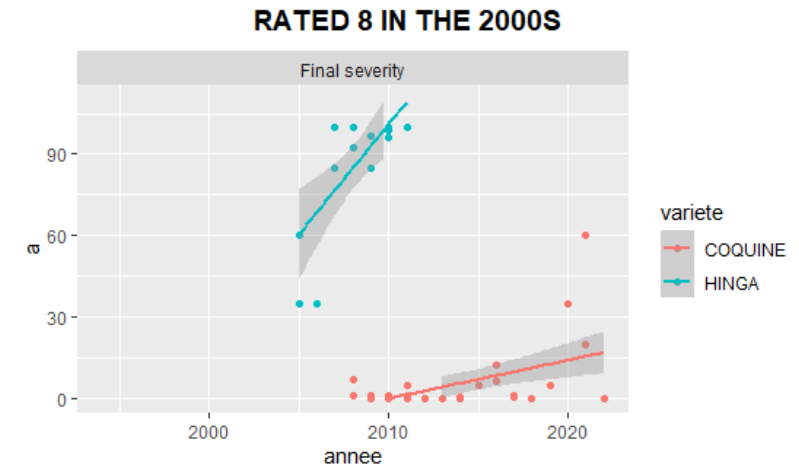
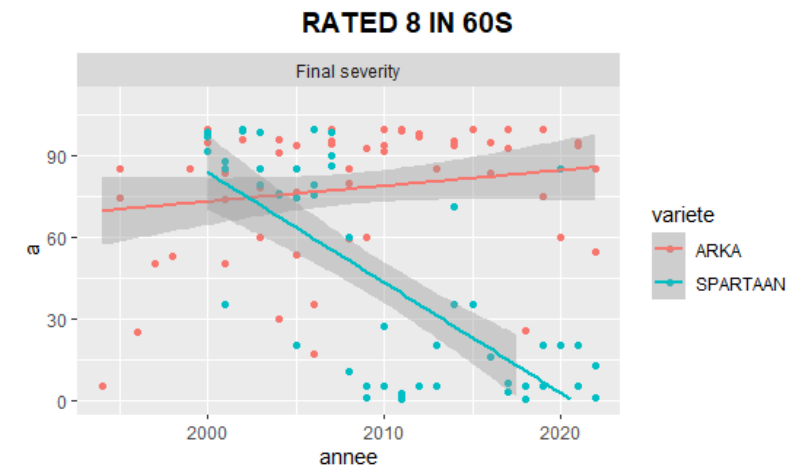
# Conclusion

- Typology based on severity, timing and slope of epidemics
- Combining varietal resistance with predictive tools faces :
  - Creating discriminative and predictable types
  - Knowing varietal specific features
  - Taking into account pathogen population



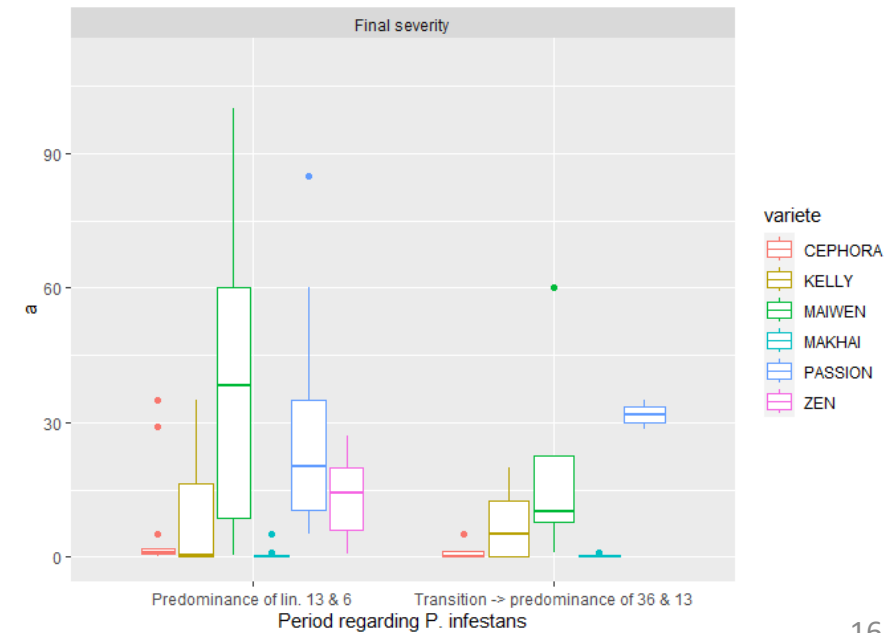
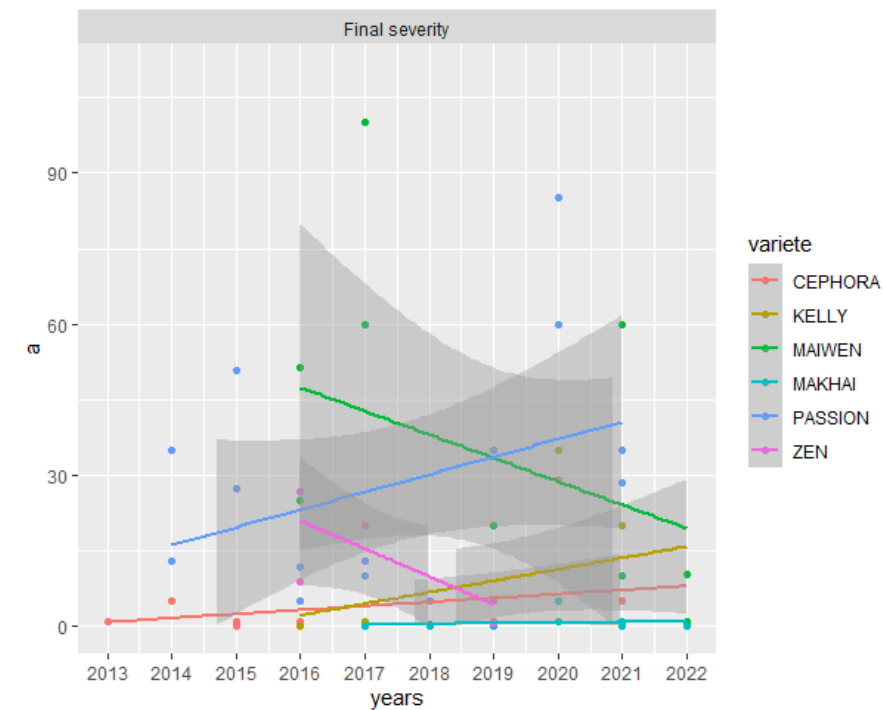
# What's next ?

- For individual registered varieties :
  - Changes of epidemic parameters over time
  - Steps or gradual loss of resistance ?
  - A common rule ?



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- For individual registered varieties :
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  - Steps or gradual loss of resistance ?
  - A common rule ?
  
- Challenge to untangle resistance erosion, specific year effects, lineages effects etc...





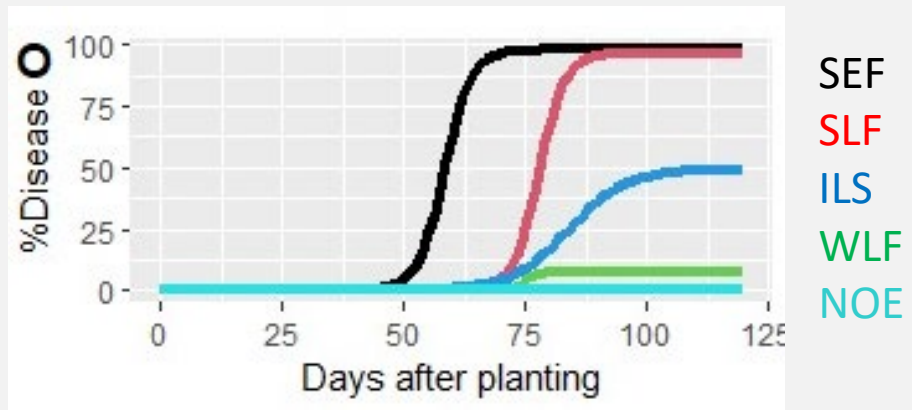


# Thanks you for listening and thanks to every person that made this work possible :

- Thanks to GEVES and Roland Pellé for providing the main dataset and to RGCO for the agronomical management of the experiments
- Thanks to Roselyne Corbière and Romain Mabon for providing the *Phytophthora infestans* dataset
- Thanks to Melen Leclerc, Didier Andrivon and Denis Gaucher for their supervision on this work



# Classification compared to official ratings

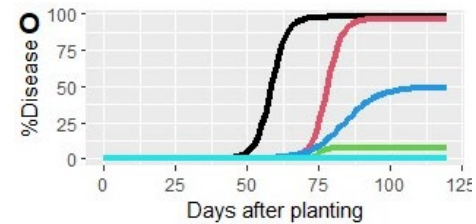
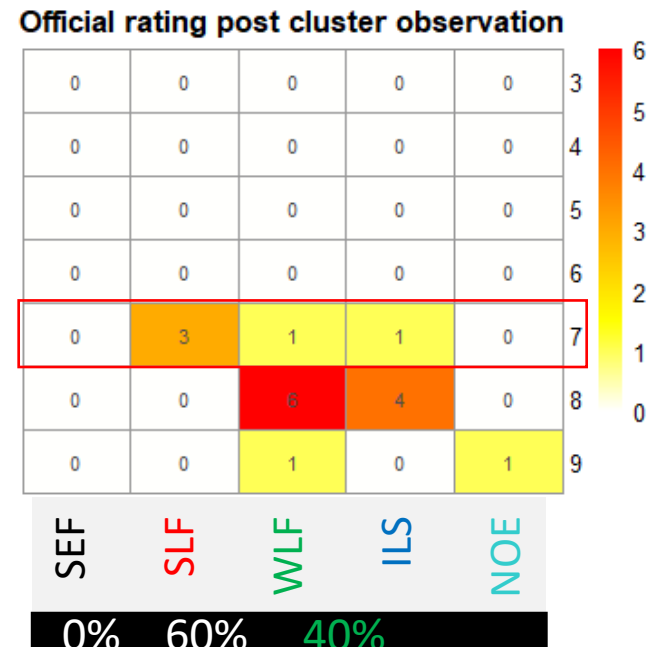
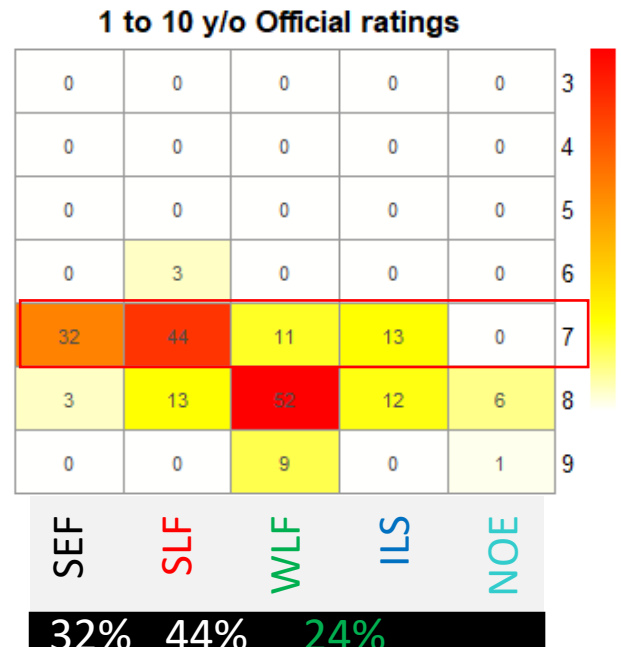
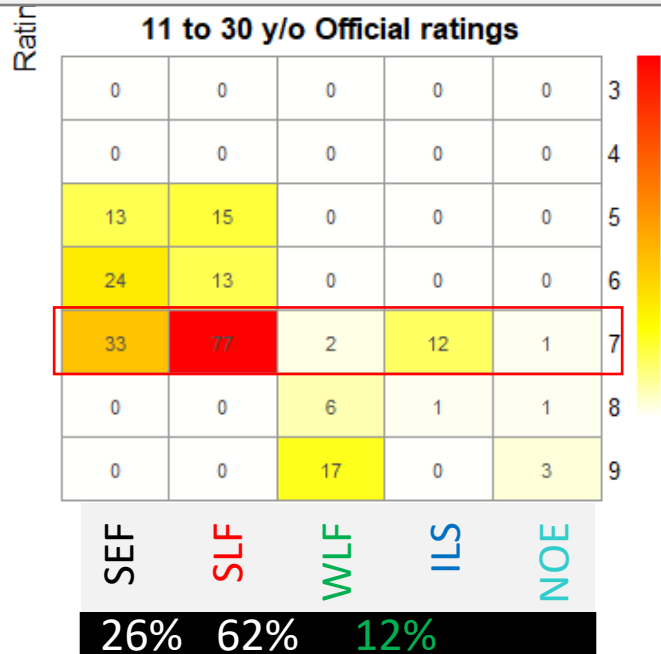
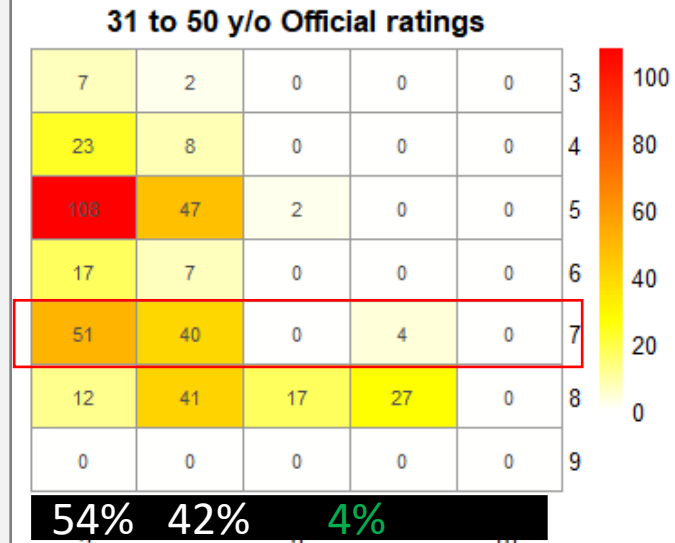
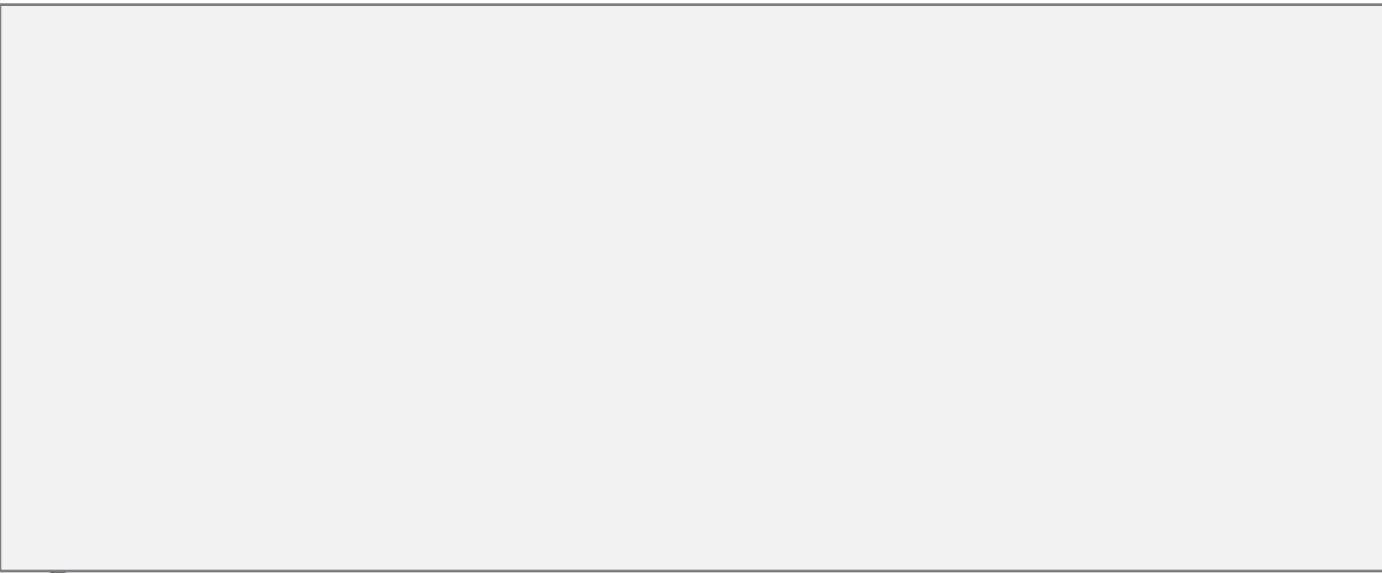


Official rating post cluster observation

0	0	0	0	0	3
0	0	0	0	0	4
0	0	0	0	0	5
0	0	0	0	0	6
0	3	1	1	0	7
0	0	6	4	0	8
0	0	1	0	1	9
SEF	SLF	WLF	ILS	NOE	

Clusters

# Classification compared to official ratings



# Classification compared to official ratings

