

# **Assessing coupled process-based models' performance in predicting tree mortality under climate change**

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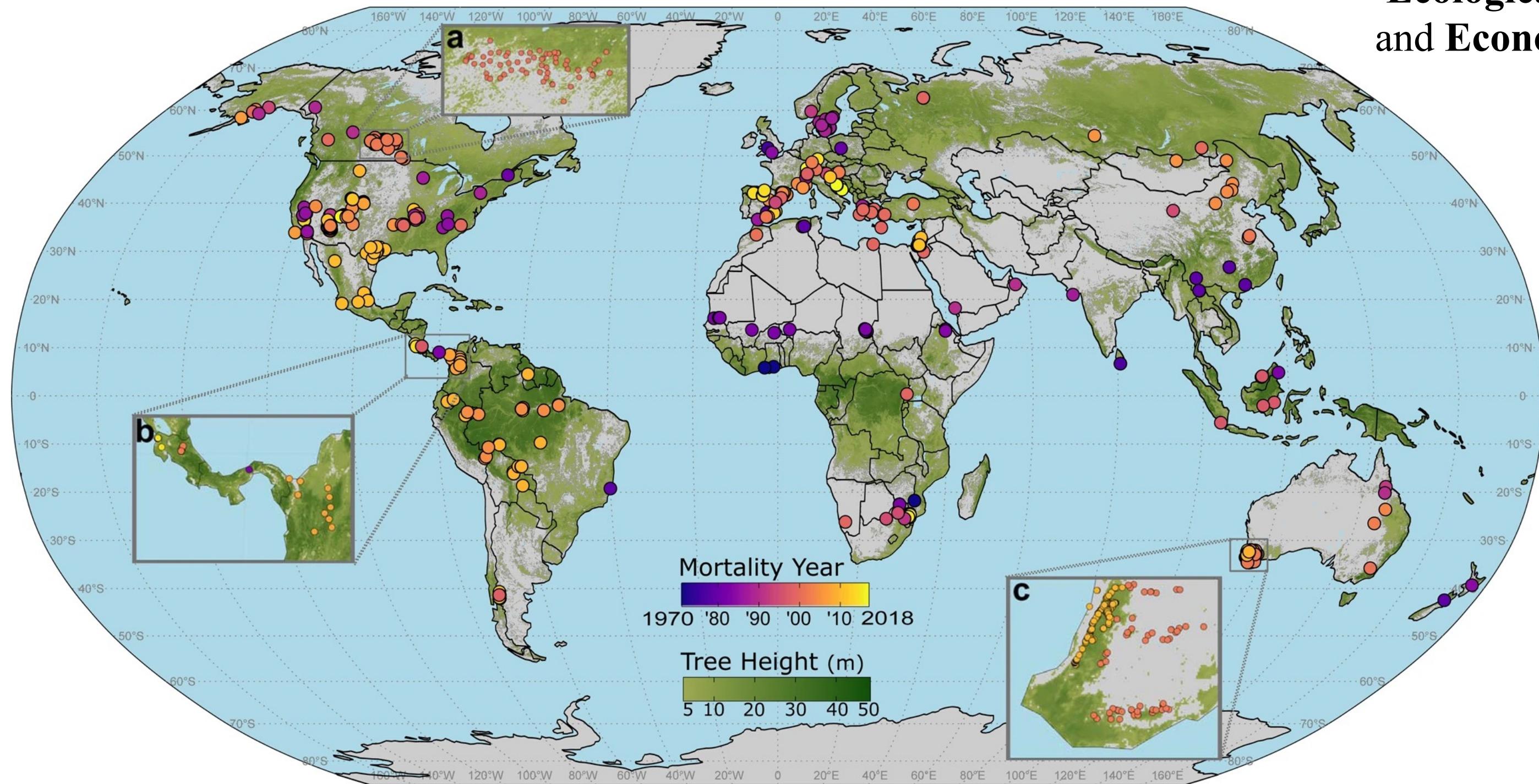
**Academic referent: Julien Sainte-Marie**

**Zhong Haoming    04/09/2023**

# Introduction

## Tree mortality under climate change

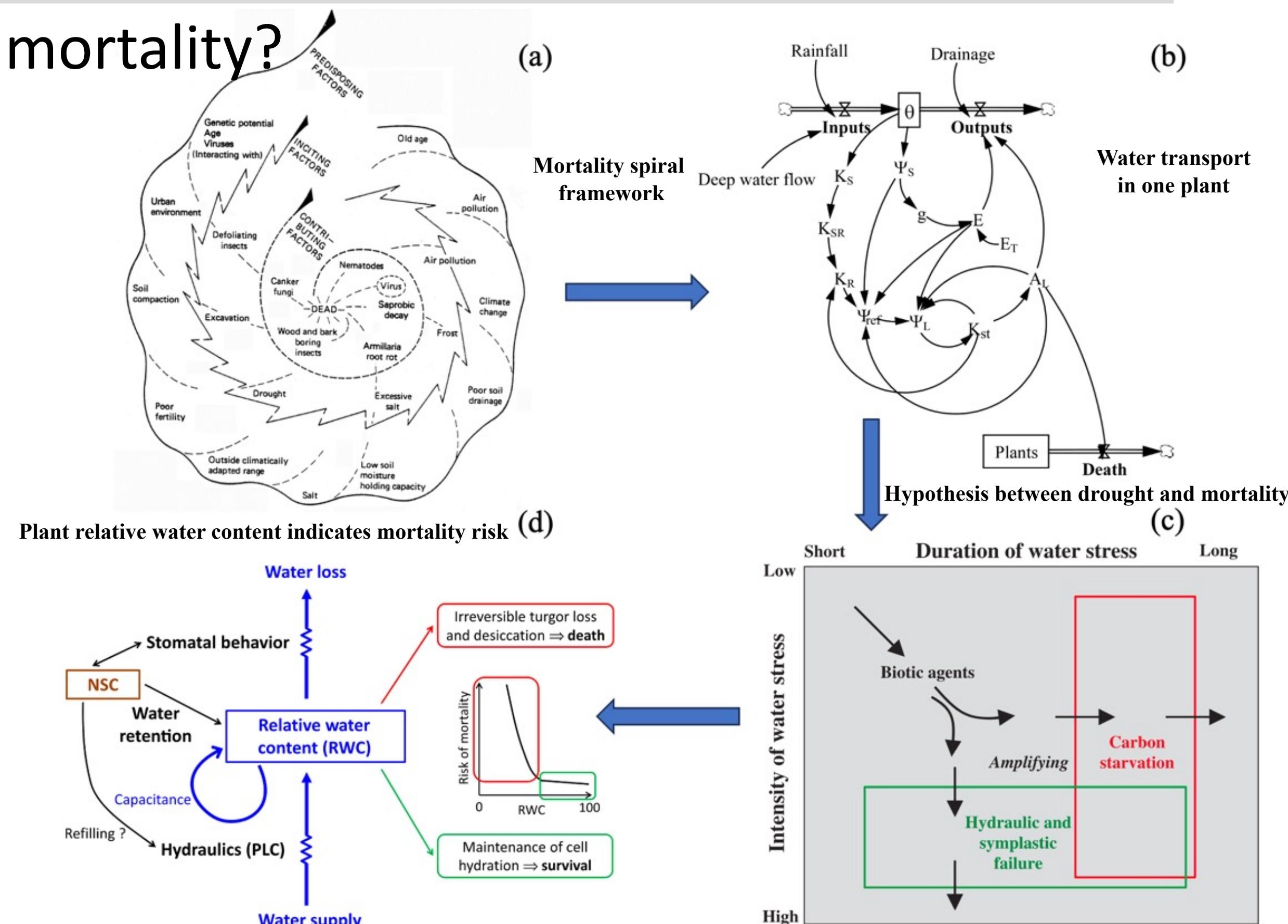
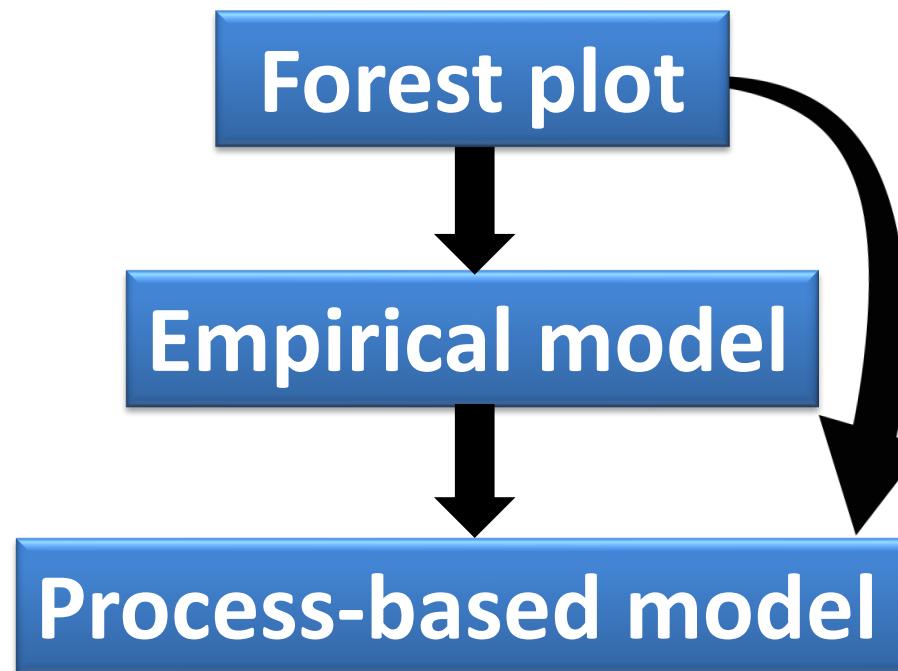
Loss of  
**Ecological benefits**  
and **Economic value**



Global distribution of hotter-drought tree mortality plots

# Introduction

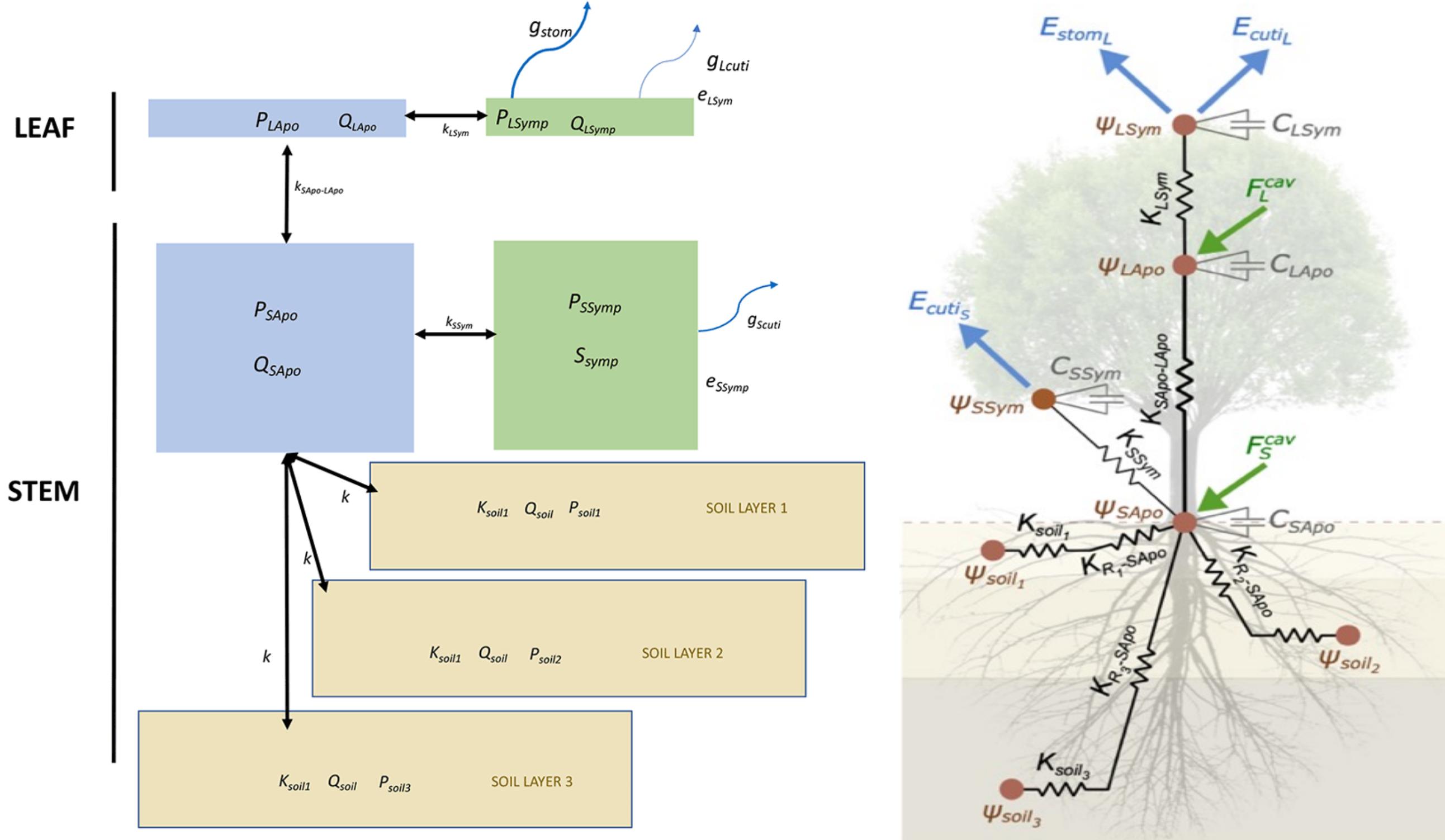
## How to predict tree mortality?



Plant hydraulic failure  
VS  
Carbon starvation

# Method

## Plant hydraulic model: SurEau-Ecos



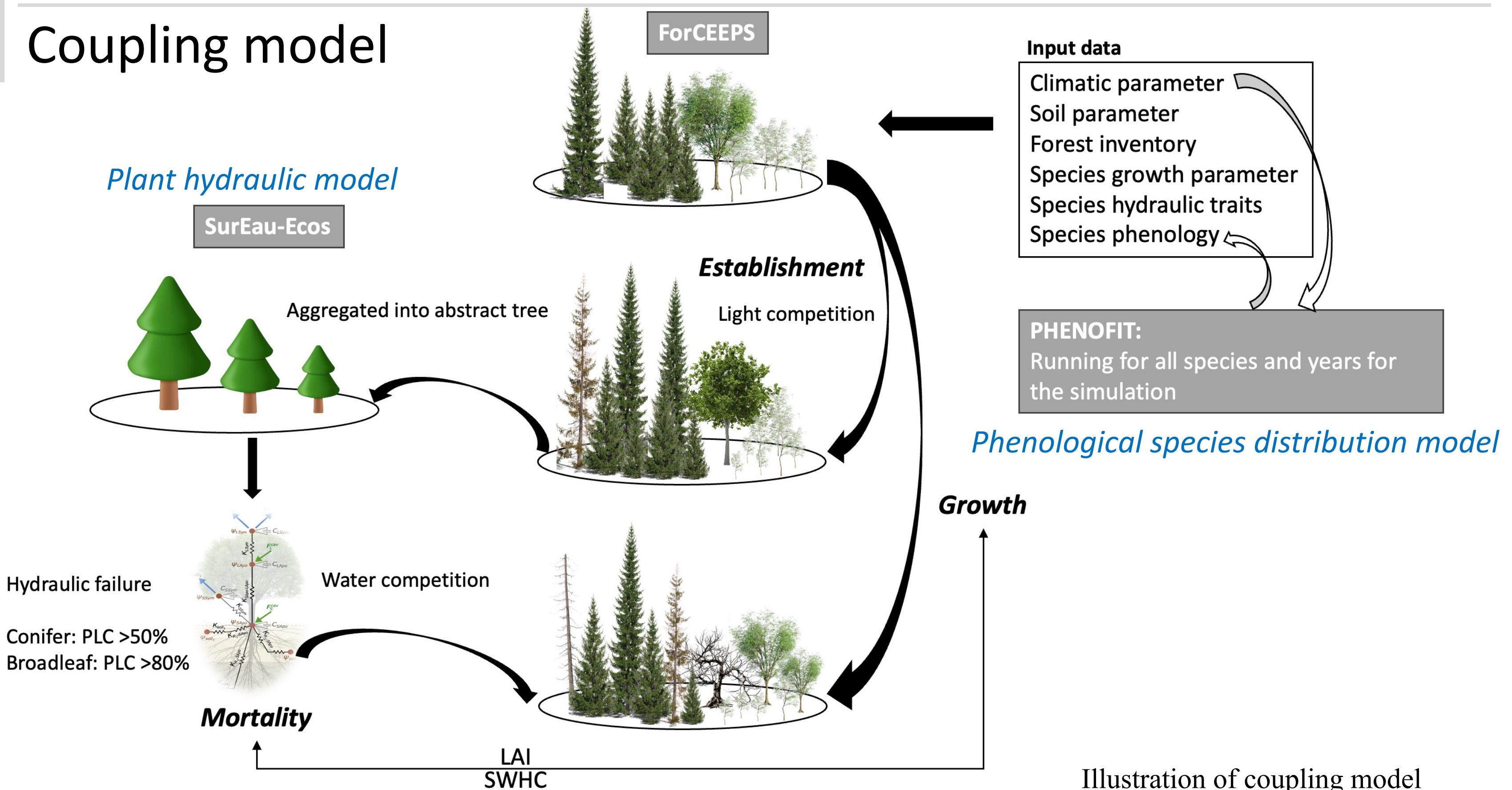
Plant architecture in SurEau-Ecos

Cochard et al., 2021; Ruffault et al., 2022

# Method

## Coupling model

### Individual-based gap model



# Introduction

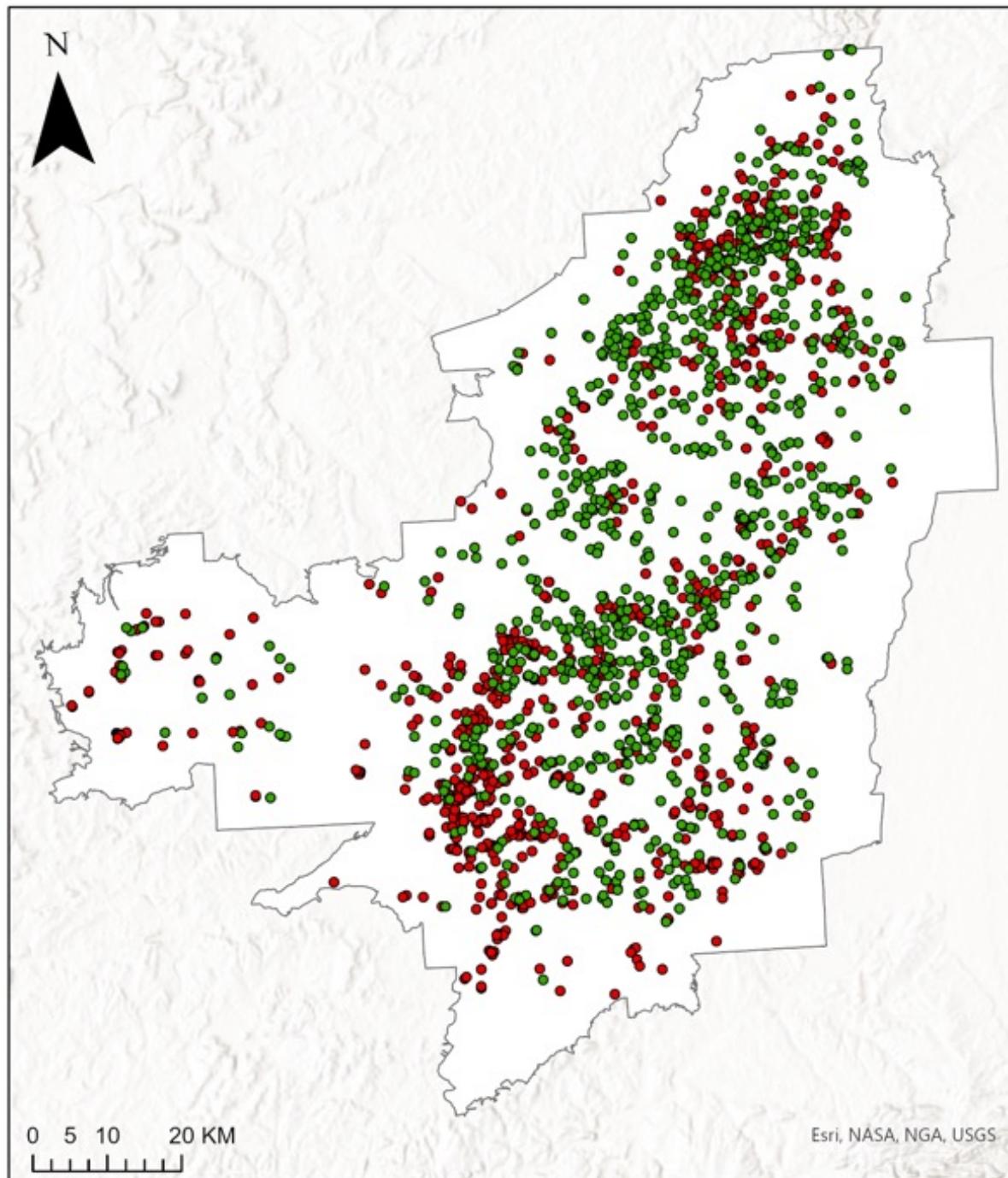
## Research objectives

- Use **Vosges and NFI validation datasets** to assess whether this coupled model could **predict mortality** at both plot and tree scale.
- To **distinguish** the mortality extent among **different species**, especially Spruce and Fir in Vosges dataset, and to reproduce **mortality spatial pattern** in this dataset.
- Study the **similarity** between LAI generated by model and LAI from two external sources, and evaluate the influence of **Soil Water Holding Capacity (SWHC)** on predictions.

# Method

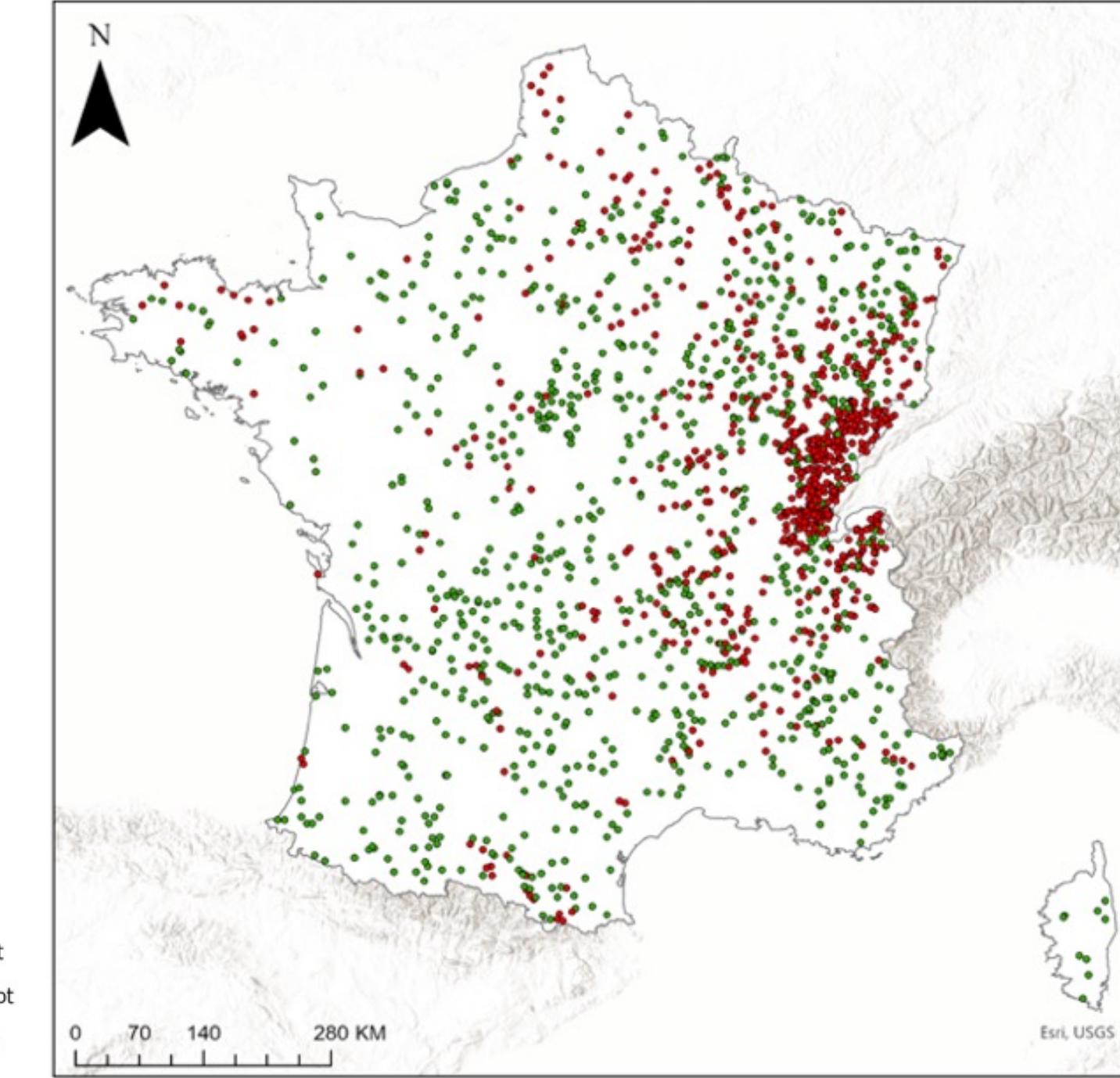
## Validation dataset: Vosges and NFI with observed mortality

1866 plots      Norway spruce; Silver fir



Simulation period: 2008-2019

1694 plots      65 common species (Simulated 35)



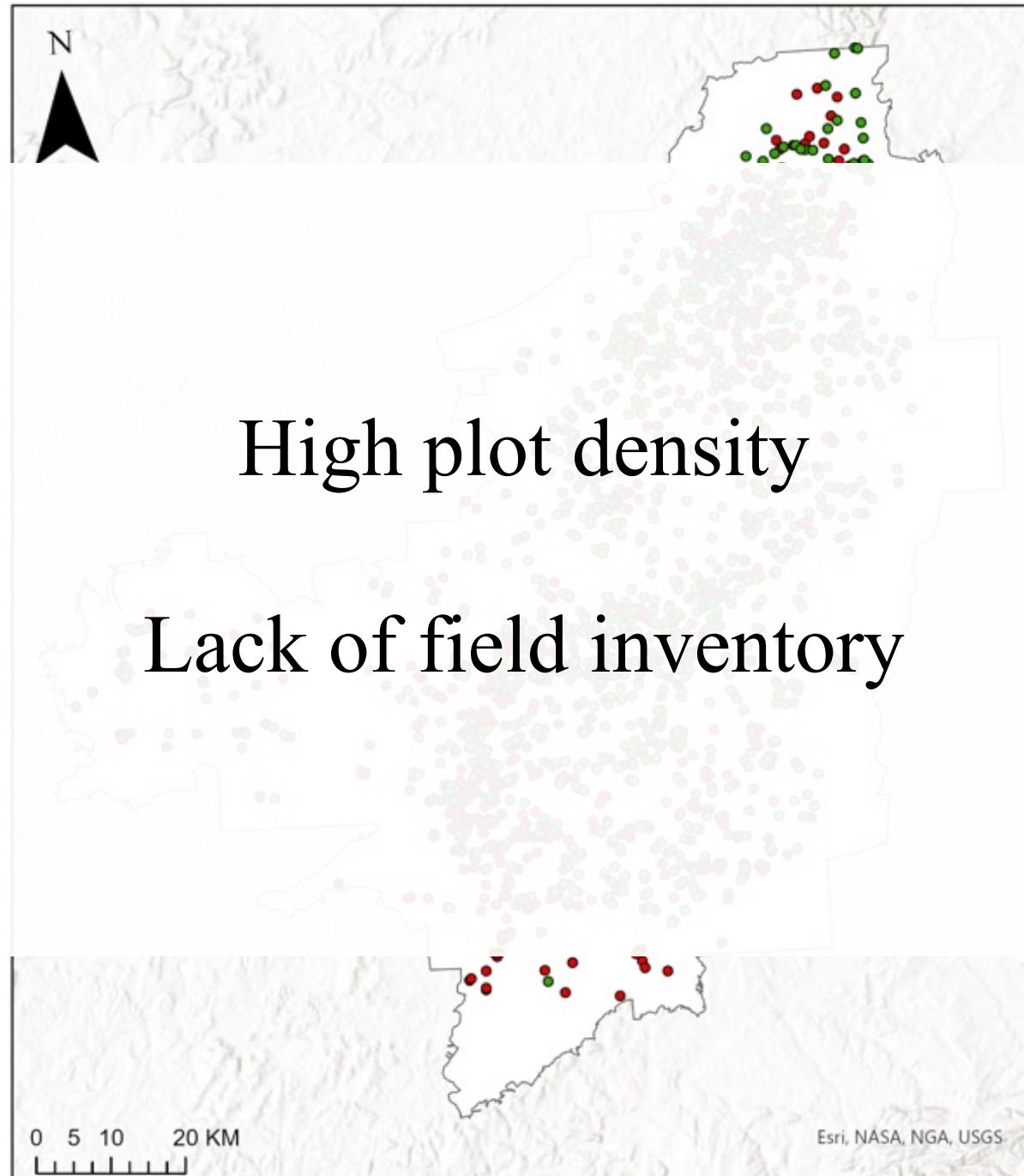
Simulation period: Rolling 5y period from 2010-2020

Distribution of Vosges (left) and NFI (right) plots

# Method

## Validation dataset: Vosges and NFI with observed mortality

1866 plots      Norway spruce; Silver fir



Pros:

High plot density

Cons:

Lack of field inventory

Simulation period: 2008-2019

1694 plots      65 common species (Simulated 35)



Simulation period: Rolling 5y period from 2010-2020

Distribution of Vosges (left) and NFI (right) plots

# Method

## Model execution



Each plot has two output files

#date	nTrees(/ha)	totalBiomass(t/ha)	totalBasalArea(m <sup>2</sup> /ha)	totalHeight(m/ha)	totalVolume(m <sup>3</sup> /ha)	LAI	droughtIndexAnnual	droughtIndexSeasonal	winterTMIN(deg)	GDDan
2008	810	180.56381	30.0175	14116.7793	212.32758	11.33557	0	0	0	0
2009	810	180.23171	30.0175	14116.7793	212.32758	10.95691	0.4733	0.54092	-0.9	
2010	810	180.27413	30.0175	14116.7793	212.32758	11.00527	0.30568	0.30568	-2.58998	
2011	800	179.37987	29.81855	13988.04334	211.29217	11.03405	0.34428	0.44265	-1.1	
2012	800	179.43624	29.81855	13988.04334	211.29217	11.09833	0.32285	0.36897	-3.6	
2013	790	178.53846	29.61961	13859.30737	210.25676	11.12308	0.27505	0.27505	-2.01643	
2014	790	178.58476	29.61961	13859.30737	210.25676	11.17588	0.3153	0.40539	2.9	
2015	790	178.62443	29.61961	13859.30737	210.25676	11.22111	0.37072	0.47664	0.1	
2016	790	178.65797	29.61961	13859.30737	210.25676	11.25935	0.33781	0.33781	2.01314	
2017	790	178.68638	29.61961	13859.30737	210.25676	11.29174	0.38348	0.43826	-2.44752	
2018	780	176.84993	29.27297	13700.12893	208.10091	11.23912	0.48859	0.55838	-2	
2019	780	176.881	29.27297	13700.12893	208.10091	11.27456	0.48375	0.52574	-0.74708	

Mean output file for Vosges dataset: **Plot average information for each year**

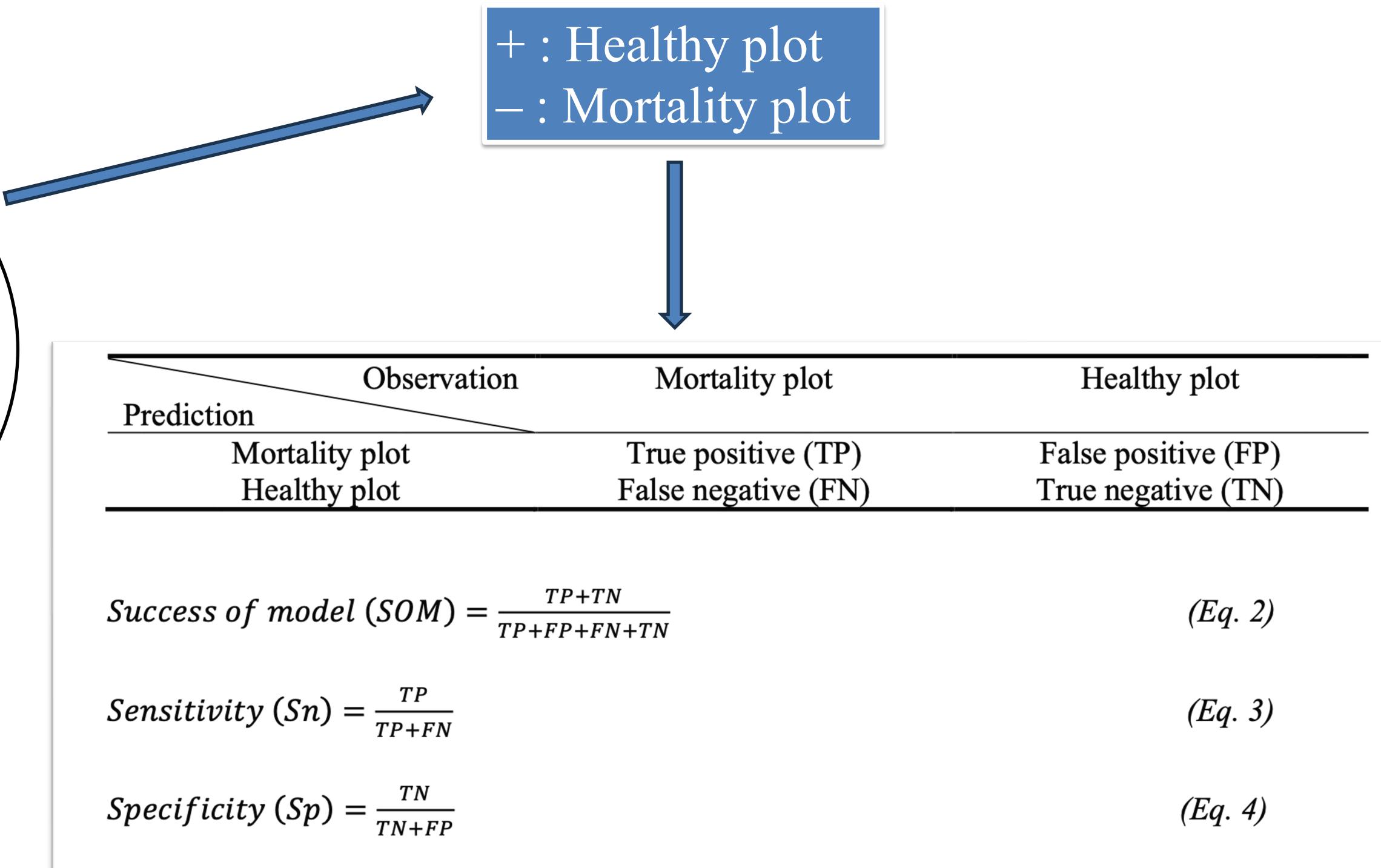
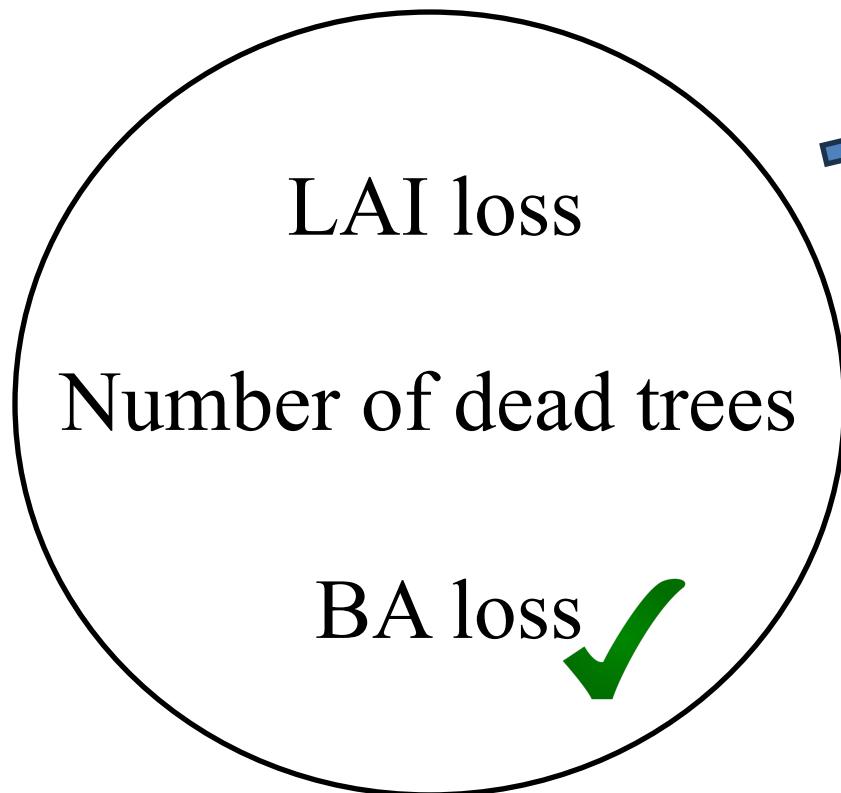
#Trees	#date	patchId	id	speciesId	speciesShortName	genotypeName	age	dbh(cm)	basalArea(m <sup>2</sup> )	height(m)	volume(m <sup>3</sup> )	crownA1	crownLength dbhl
	2008	1	10149821	2	PAbi	2-Genotype-1	30	33.4225	0.087734	22.2784	0.7317	0.4116	9.1692
	2009	1	10149821	2	PAbi	2-Genotype-1	31	33.4225	0.087734	22.2784	0.7317	0.4116	9.1687
	2010	1	10149821	2	PAbi	2-Genotype-1	32	33.4225	0.087734	22.2784	0.7317	0.4085	9.1004
	2011	1	10149821	2	PAbi	2-Genotype-1	33	33.4225	0.087734	22.2784	0.7317	0.4077	9.0834
	2012	1	10149821	2	PAbi	2-Genotype-1	34	33.4225	0.087734	22.2784	0.7317	0.4079	9.0869
	2013	1	10149821	2	PAbi	2-Genotype-1	35	33.4225	0.087734	22.2784	0.7317	0.408	9.0902
	2014	1	10149821	2	PAbi	2-Genotype-1	36	33.4225	0.087734	22.2784	0.7317	0.4083	9.0973
	2015	1	10149821	2	PAbi	2-Genotype-1	37	33.4225	0.087734	22.2784	0.7317	0.4087	9.1047
	2016	1	10149821	2	PAbi	2-Genotype-1	38	33.4225	0.087734	22.2784	0.7317	0.409	9.1118
	2017	1	10149821	2	PAbi	2-Genotype-1	39	33.4225	0.087734	22.2784	0.7317	0.4093	9.1186
	2018	1	10149821	2	PAbi	2-Genotype-1	40	33.4225	0.087734	22.2784	0.7317	0.4096	9.1248
	2019	1	10149821	2	PAbi	2-Genotype-1	41	33.4225	0.087734	22.2784	0.7317	0.41	9.1352

Complete output file for Vosges dataset: **Detail information for each tree**

# Method

## Mortality definition

At plot scale



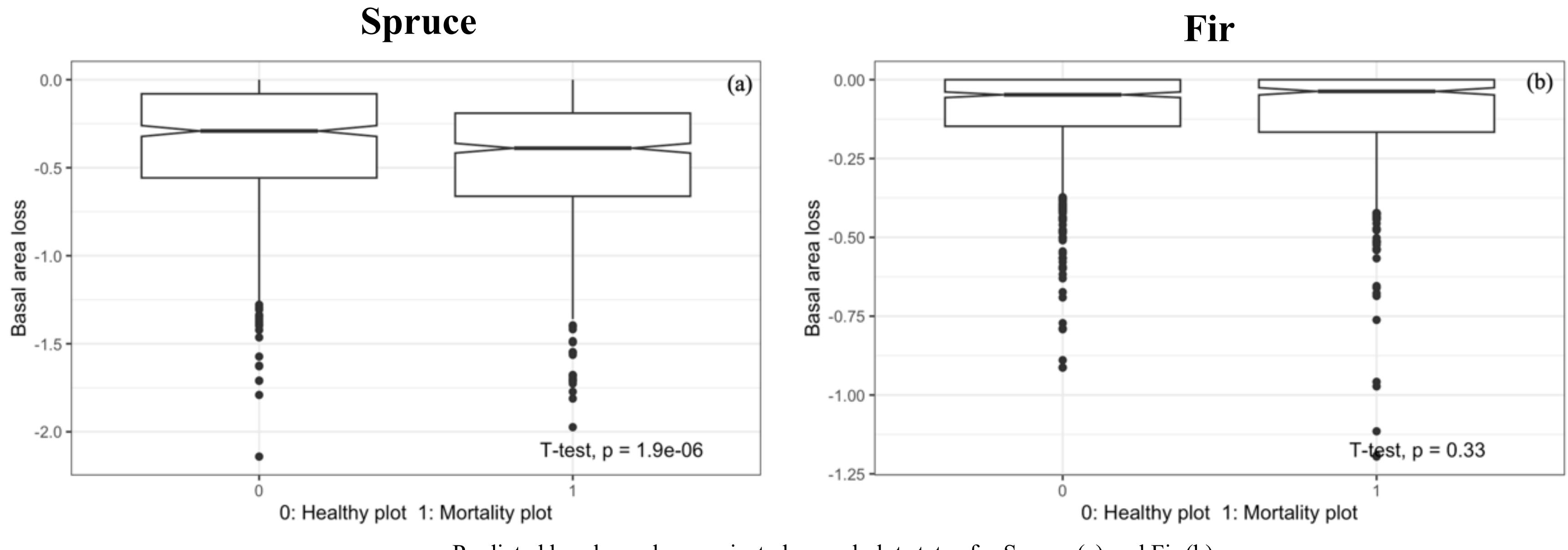
At tree scale



Monitor and log **unique tree ID**, noting their disappearance as an indication of death.

# Result

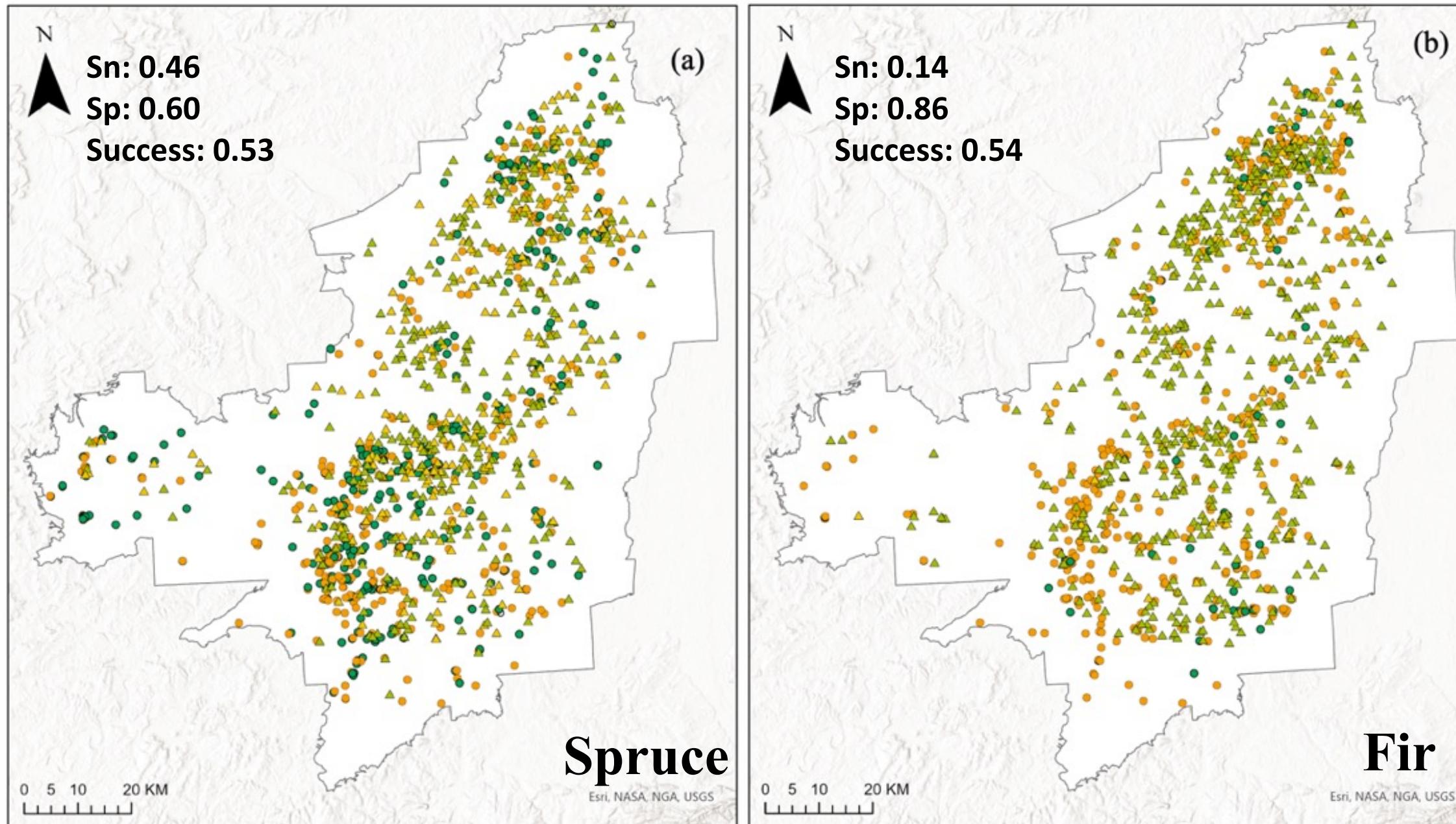
## Vosges: Mortality prediction vs observation at plot scale



- No significant difference between healthy and mortality plot.
- Difference of Spruce is slight obvious than Fir.

# Result

## Vosges: Mortality prediction at plot scale



Quality of prediction distribution for Spruce (a) and Fir (b) in Vosges

- Model has better mortality prediction for Spruce.
- Mortality is underestimated for both species, especially Fir.
- Model could partly reproduce mortality spatial pattern.

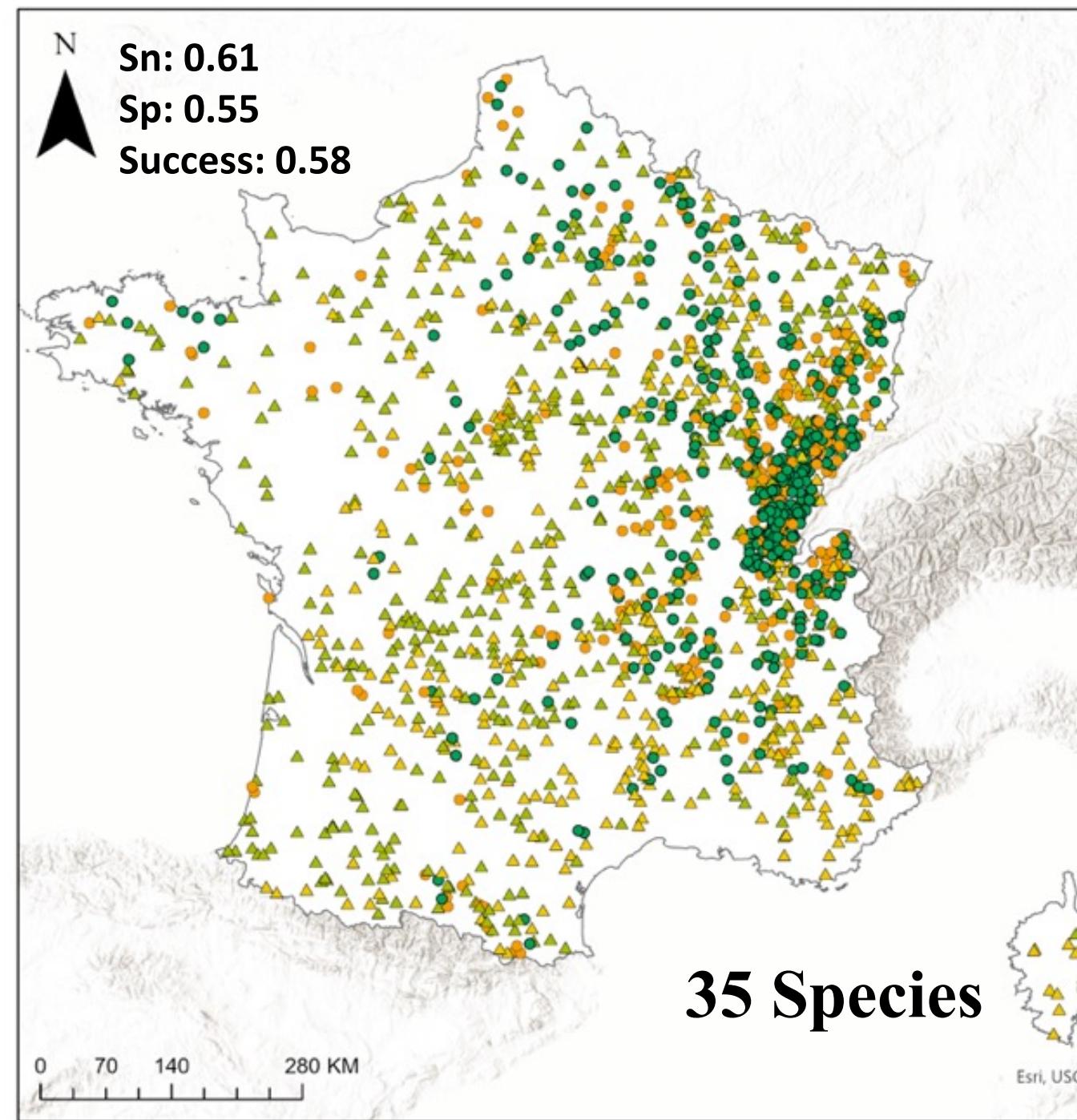
### Ratio comparison

Item	Spruce	Fir
Observed mortality frequency	0.55	0.44
Predicted mortality frequency	0.43	0.14

Observation Prediction	Mortality plot	Healthy plot
Mortality plot	● True Positive	▲ False Positive
Healthy plot	● False Negative	▲ True Negative

# Result

## NFI: Mortality prediction at plot scale



Quality of prediction distribution for all species for NFI dataset

### Ratio comparison

Item	NFI
Observed mortality frequency	0.71
Predicted mortality frequency	1.07

Observation Prediction	Mortality plot	Healthy plot
Mortality plot	● True Positive	▲ False Positive
Healthy plot	● False Negative	▲ True Negative

- Accurate mortality predictions are concentrated in the Jura mountains.
- Mortality is overestimated for NFI dataset, especially in south of France.

# Result

## NFI: Mortality prediction at tree scale

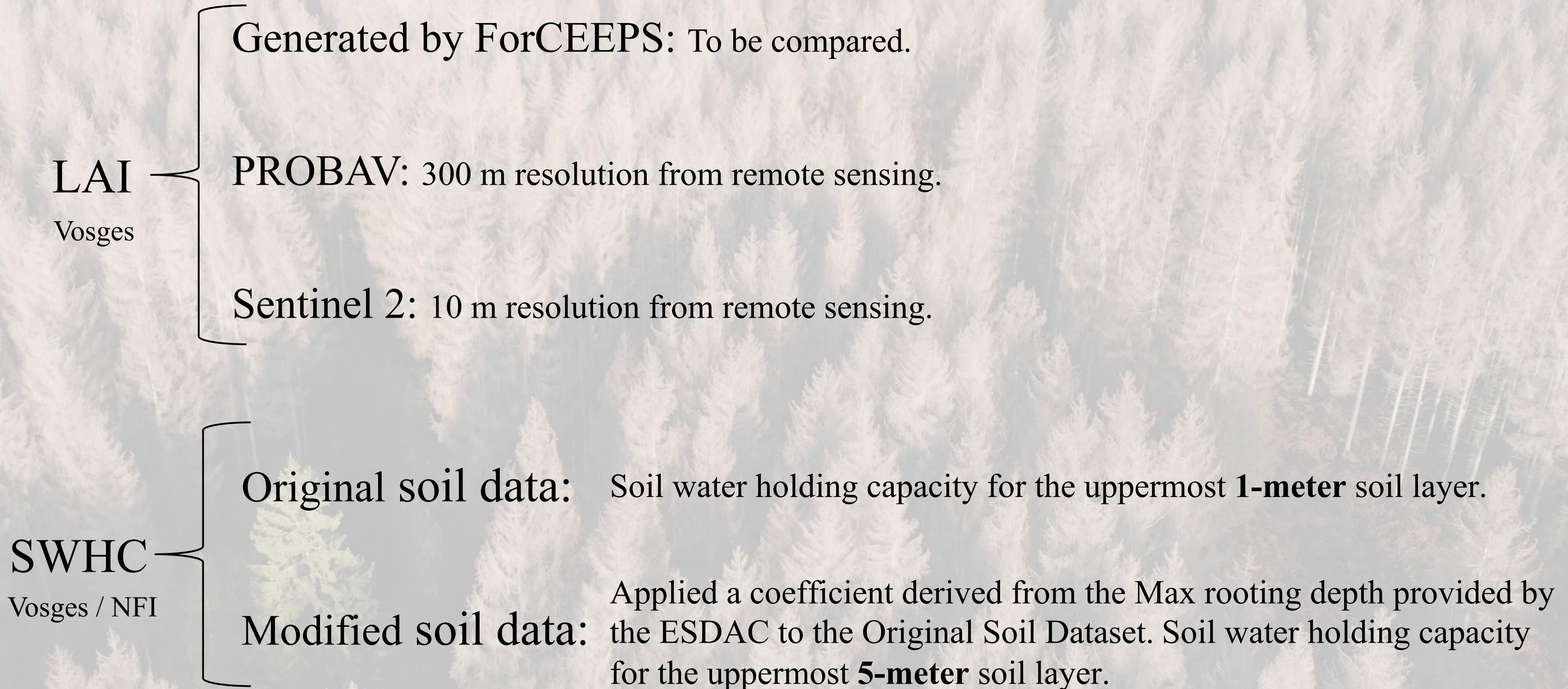
Comparison of mortality frequency

Species	Observed frequency	Predicted frequency
<i>Castanea sativa</i>	0.51	0.63
<i>Picea abies</i>	0.49	0.55
<i>Fraxinus excelsior</i>	0.49	0.51
<i>Corylus avellana</i>	<b>0.37</b>	<b>0.28</b>
<i>Pinus sylvestris</i>	0.34	0.55
<i>Populus tremula</i>	0.32	0.38
<i>Abies alba</i>	0.32	0.45
<i>Betula pendula</i>	0.27	0.37
<i>Carpinus betulus</i>	<b>0.27</b>	<b>0.69</b>

- Model predicted similar or slightly higher mortality frequency for almost all the listed species except *Corylus avellana*, and *Carpinus betulus* has a much higher predicted frequency.

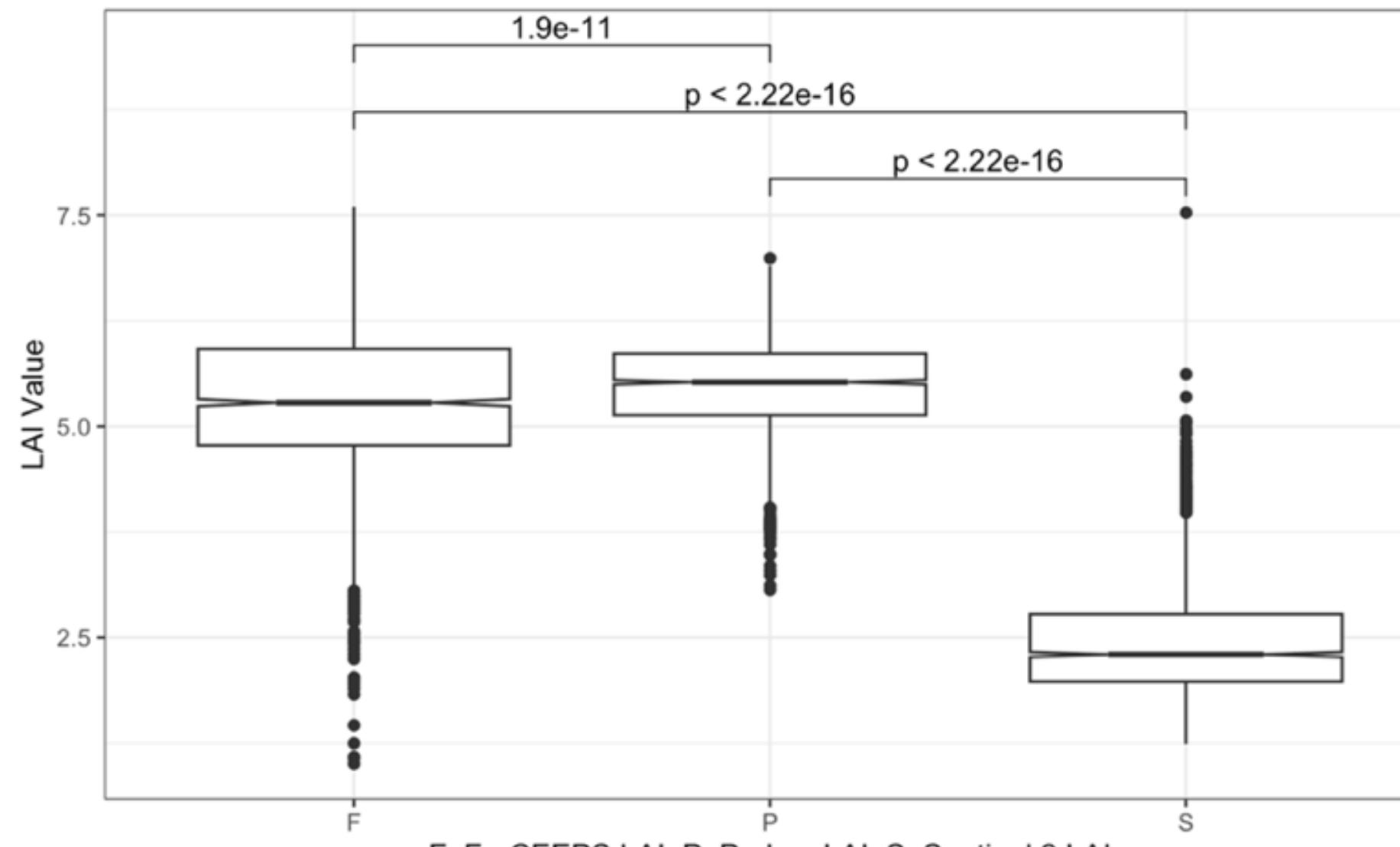
# Method

## 3 LAI datasets and 2 SWHC datasets

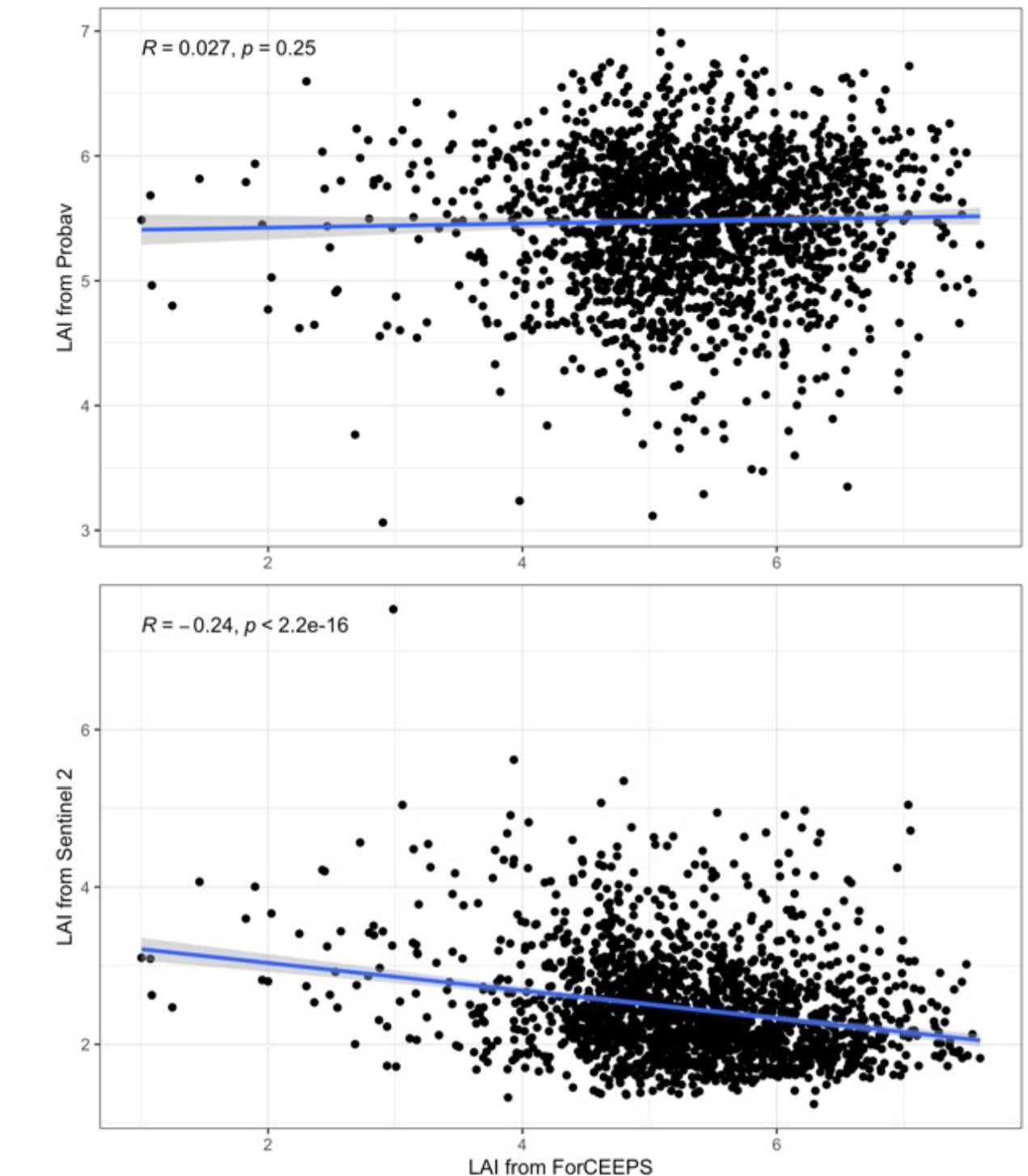


# Result

## LAI comparison



LAI values from ForCEEPS, Probav and Sentinel 2

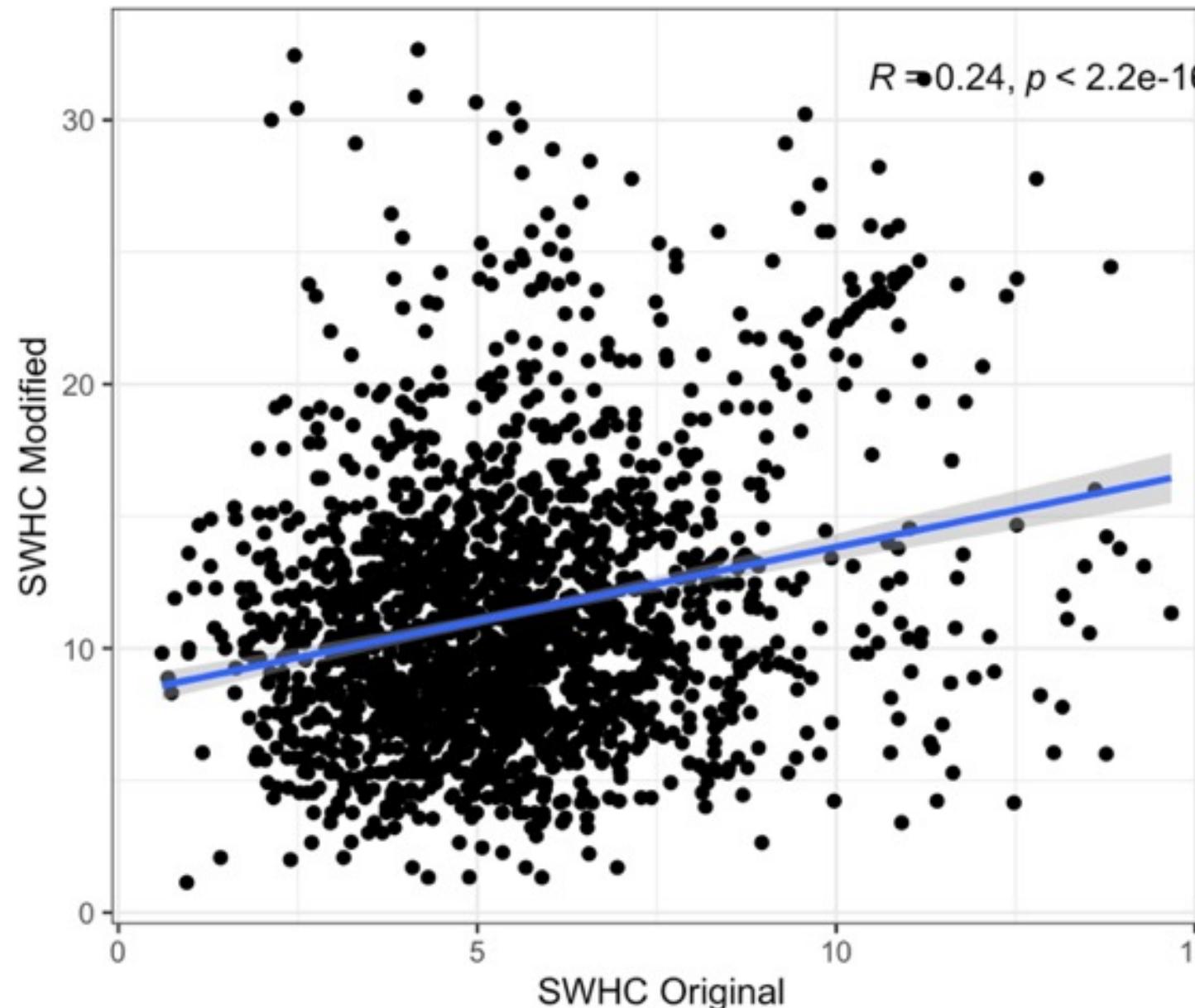


LAI ForCEEPS against Probav and Sentinel 2

- ForCEEPS LAI has a weak positive correlation with Probav and a strange negative correlation with Sentinel 2.
- Three different LAI has small similarity and with a large uncertainty.

# Result

## Sensitivity of SWHC



Model performance for two different soil input for Vosges dataset

Soil input	Spruce			Fir		
	Sn	Sp	Success	Sn	Sp	Success
Modified soil data	0.46	0.60	<b>0.53</b>	0.14	0.86	<b>0.54</b>
Original soil data	0.11	0.90	0.43	0.10	0.88	0.51

Model performance for two different soil input for NFI dataset

Soil input	Sn	Sp	Success
Modified soil data	0.61	0.55	<b>0.58</b>
Original soil data	0.85	0.27	0.51

- A weak positive correlation has been observed in two soil data.
- Modified soil data has better model performance compared to Original soil data for both datasets.

# Discussion

## Answer to research objective

- This coupled model could give the consistent results but the performance is low (0.53-0.58) at plot scale. And it can differentiate species and partly reproduce mortality spatial pattern.
- LAI generated by this model is not similar with external LAI which still needs to be enhanced. And the model is sensitive to SWHC calculation method.

## Ways to improve

- Test new mortality dataset with detailed stand characteristics and mortality variation.
- Based on ecological resource balance theory, a new method to collect soil water holding capacity data is going on.
- Achieving equilibrium within each plot before drought simulation is pivotal.

# Conclusion

## Future perspective

The first time to couple three different process-based model together to simulate forest dynamics and predict drought-induced mortality on regional and national scale.

Answer theoretical questions regarding tree mortality

Drought tolerance of each species?

Tree size effects mortality?

Growth with mortality?

PLC threshold?

Answer practical questions in forest sector

Species choice?

Optimize ecosystem services?

The effect of mixed forest?

Assistant migration?



Thank you for listening

Q&A

# Result

## LAI Sentinel 2 and mortality

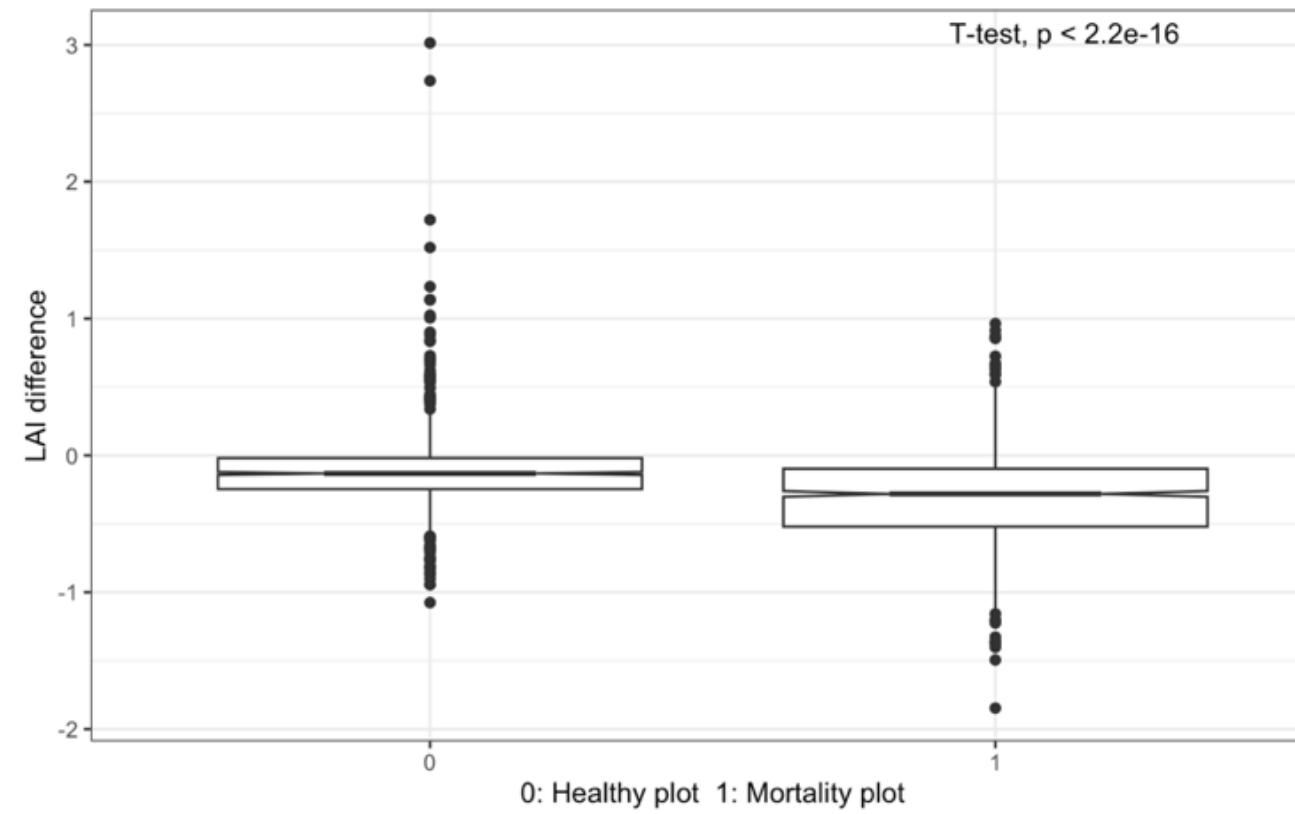


Fig 12. LAI sentinel 2 difference and plot status.

Table 5. Statistical comparison of LAI Sentinel 2 for four years.

Index	2017	2018	2019	2020
Min	0.46	0.67	0.72	0.23
Max	6.41	6.30	6.69	5.92
Median	2.00	1.93	1.84	1.74
Mean	<b>2.12</b>	<b>2.07</b>	<b>1.94</b>	<b>1.81</b>
Std	0.59	0.59	0.64	0.66

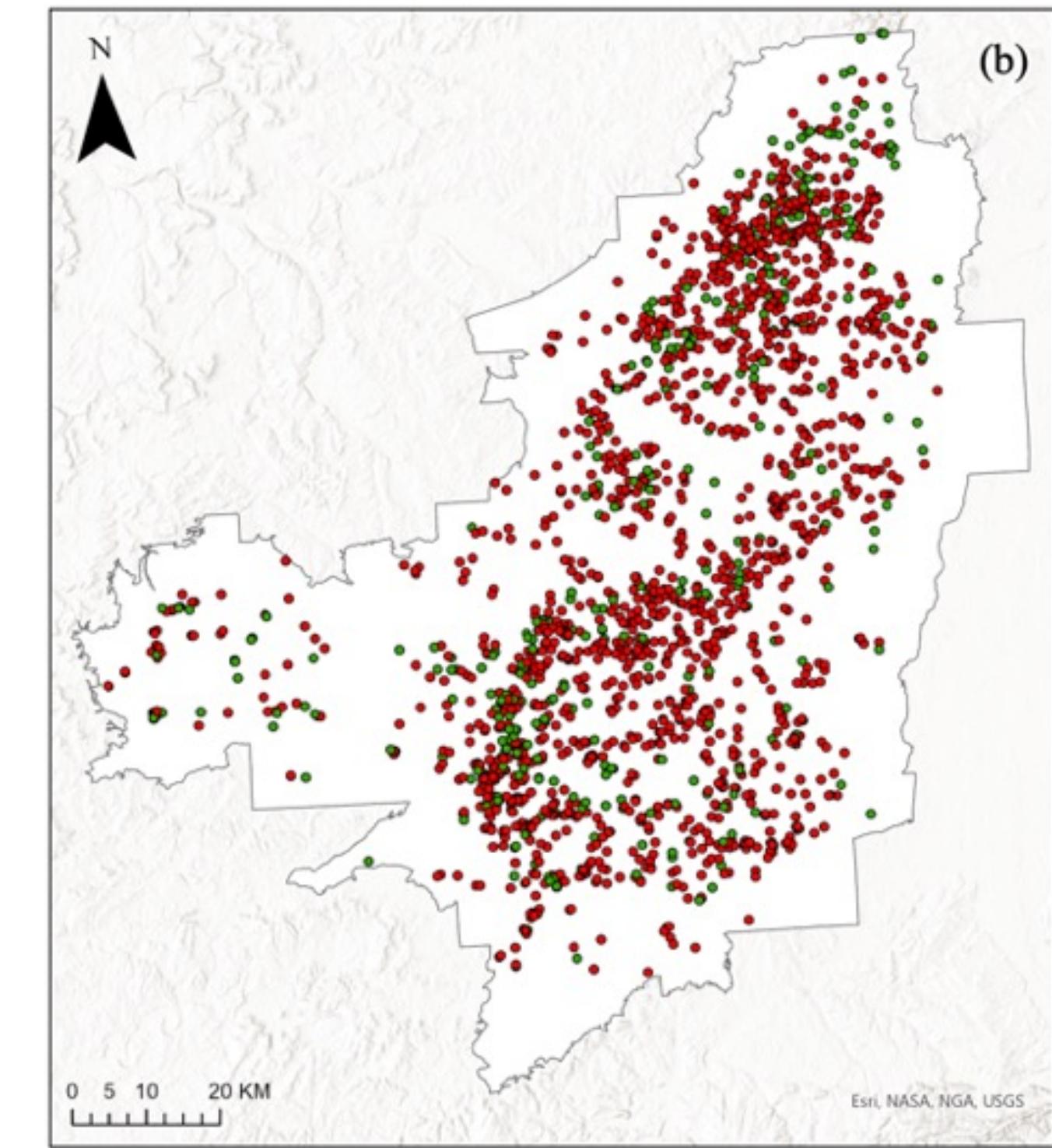


Fig 13. LAI difference between 2017-2018 and 2019-2020.

- Mean value of LAI decreased from 2017 to 2020.
- LAI decrease pattern is consistent as the distribution of observed mortality.