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Close-range drone imagery for mapping tree canopy seed fertility in European temperate forest

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2025 Senckenberg

Background

- Variation in seed production among adult trees controls their genetic contribution to the next generation, shaping effective population size, gene flow and forest resilience.
- Tree-level seed counts over whole stands are logistically demanding, species-dependent and rarely feasible beyond small plots.
- Close-range RGB imagery from low-cost drones offers a scalable way to turn crown observations into spatial fertility maps for genetic monitoring and sampling design.

Hypothesis

Under favourable weather and flight conditions, highresolution drone RGB imagery, analysed with Machine Learning (ML) or visual scoring, yields crown fertility classes reliable enough to map seed fertility from individual crowns to stand scales.

Eight Genetic Conservation Units (1 - 8), four different European temperate tree species GBR00001 · Pinus 3 - FRA00051 · Pinus 4 - ITA00019 · Pinu 6 - ESP00395 · Populus

Step 3 – Image Processing & Seed database by tree

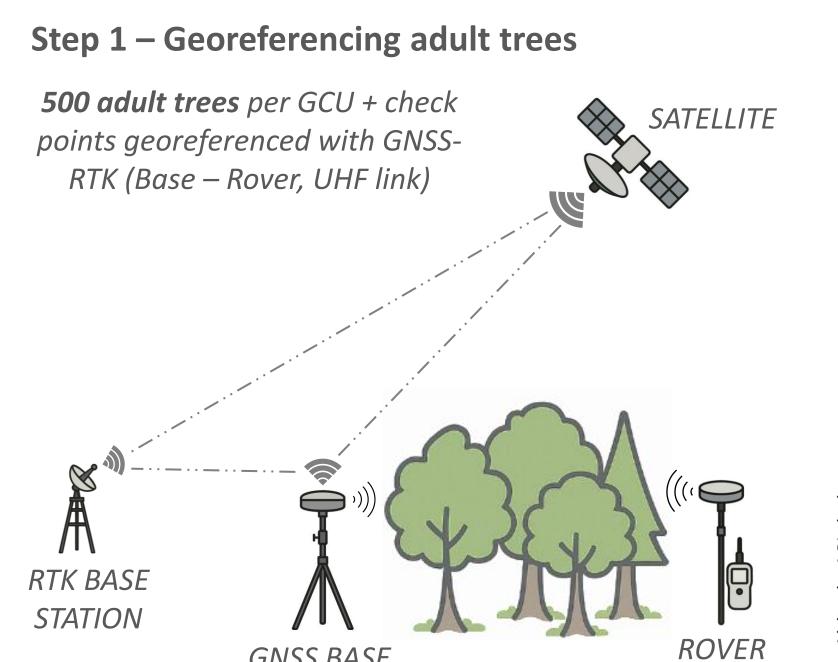
Study sites

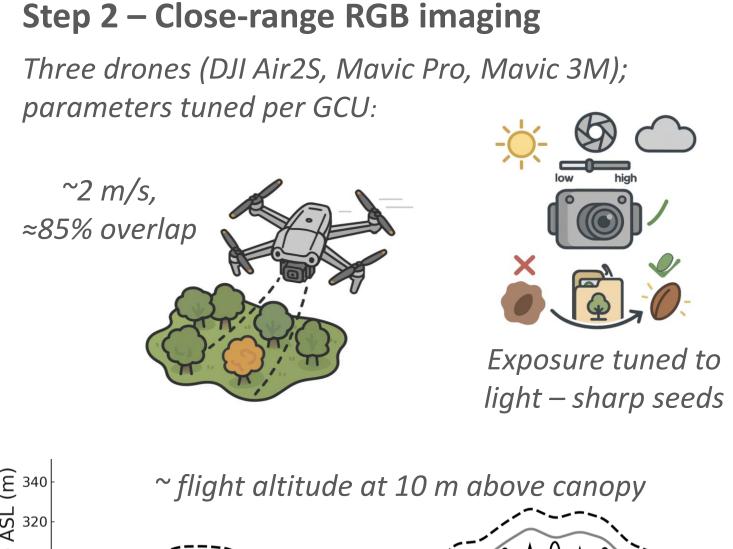
Aggregated forest map showing the dominant species at 1x1 km [1]:

O Pinus sylvestris O Populus nigra | • UMR (Philips-Universität Marburg) O Pinus pinaster O Fagus sylvatica

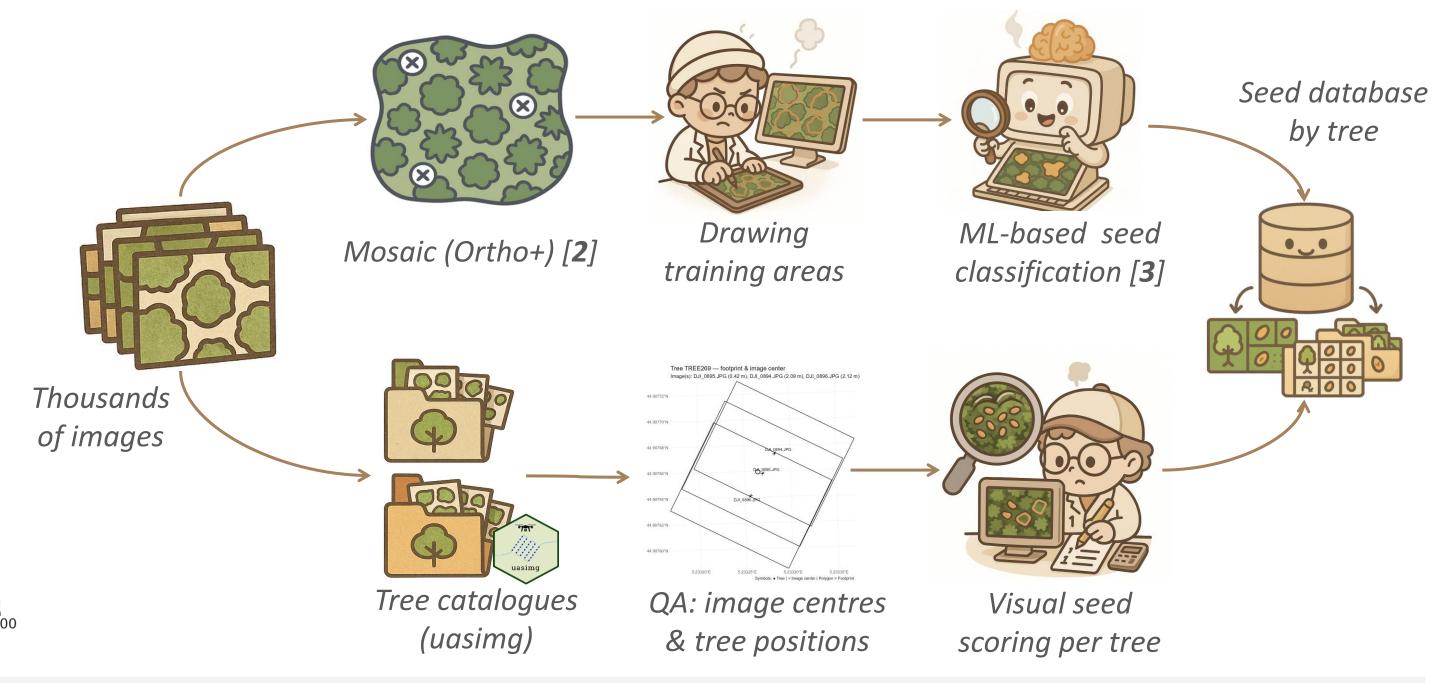
8 - FRA00045 · Fagus 7- SVN00047 · Fagus

Study design





Distance / time (relative)



Results

How seeds look in the images

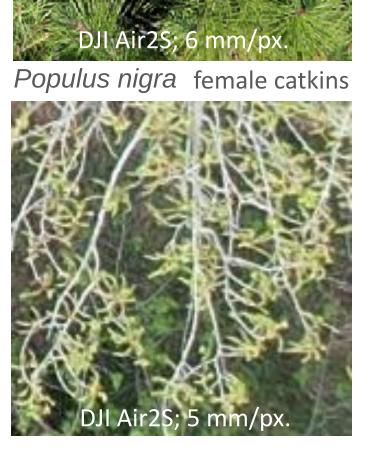
GNSS BASE

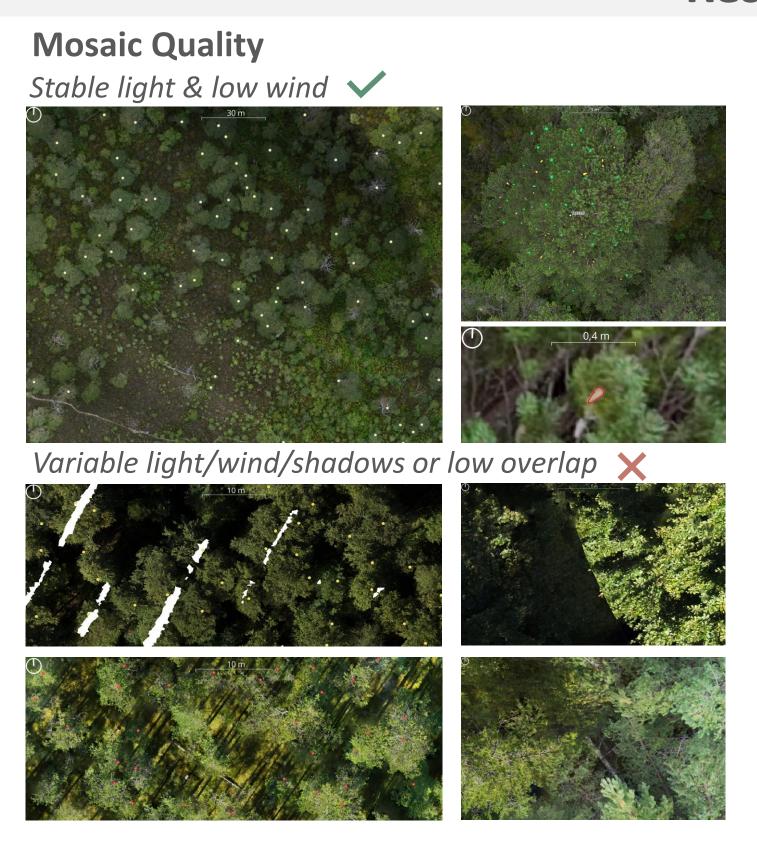
STATION



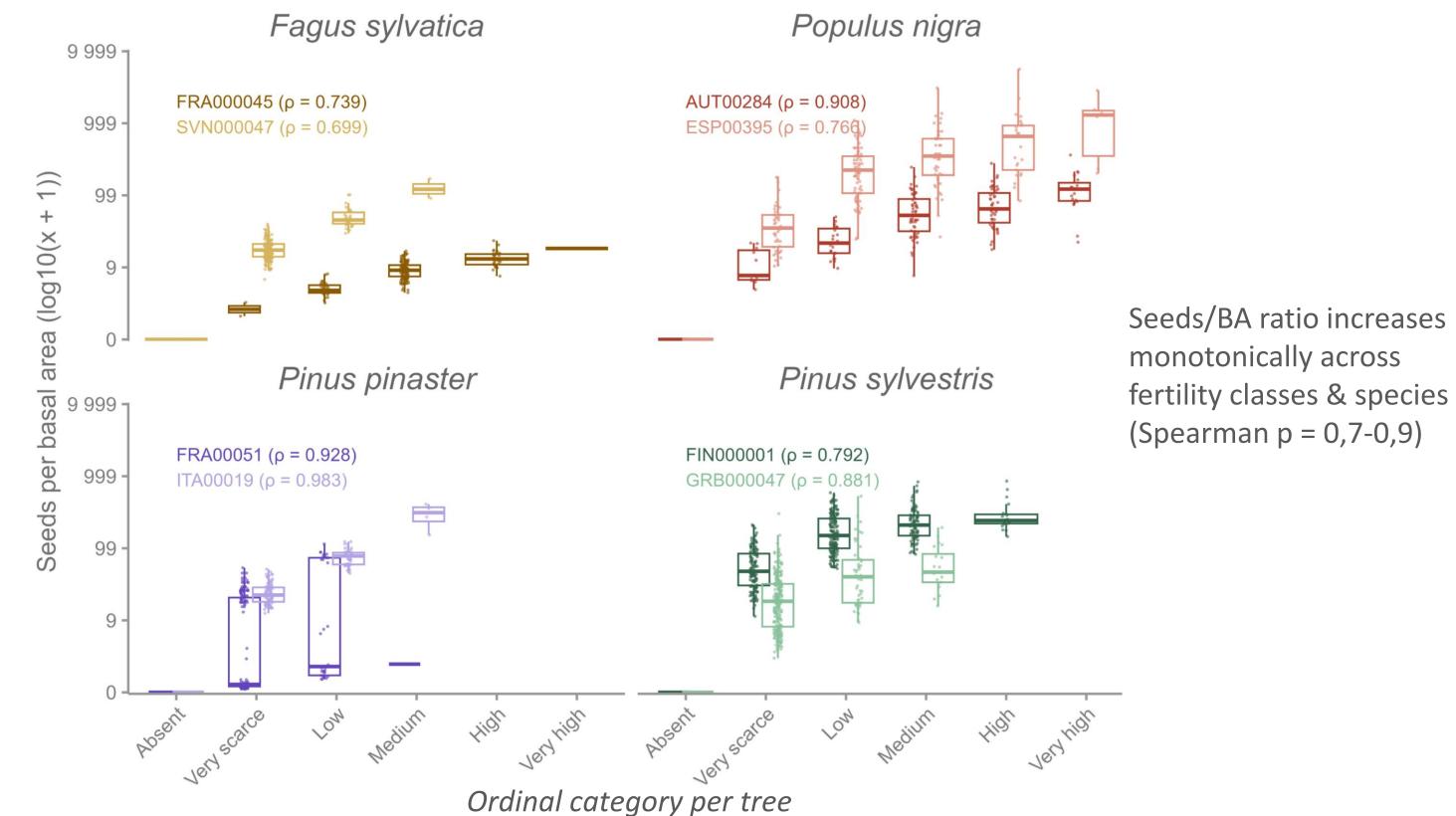
Fagus sylvatica Beechnut



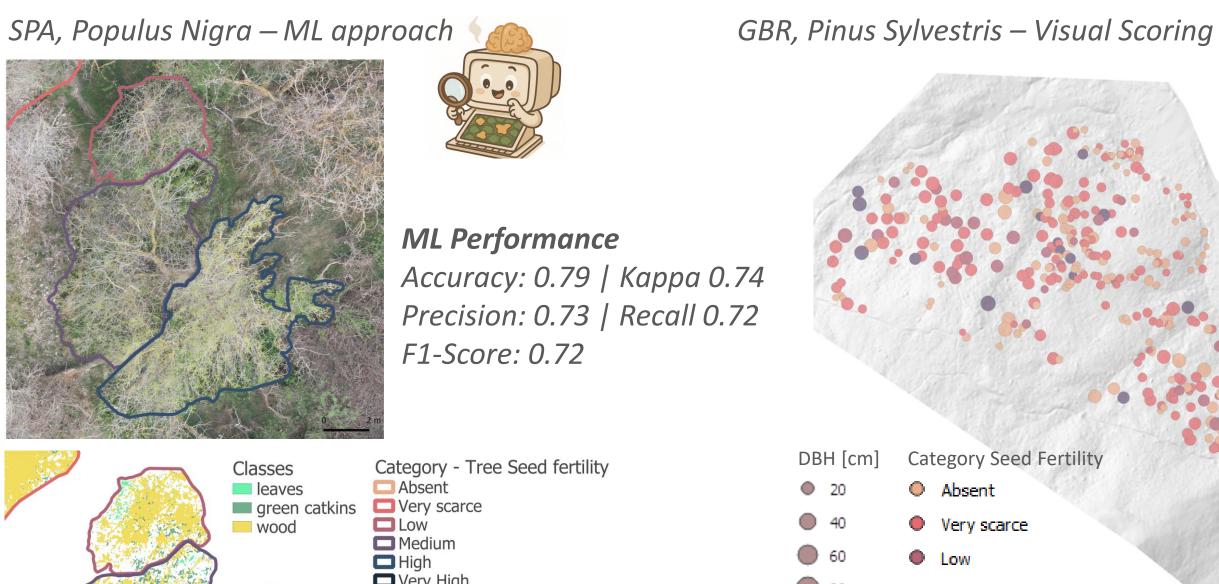


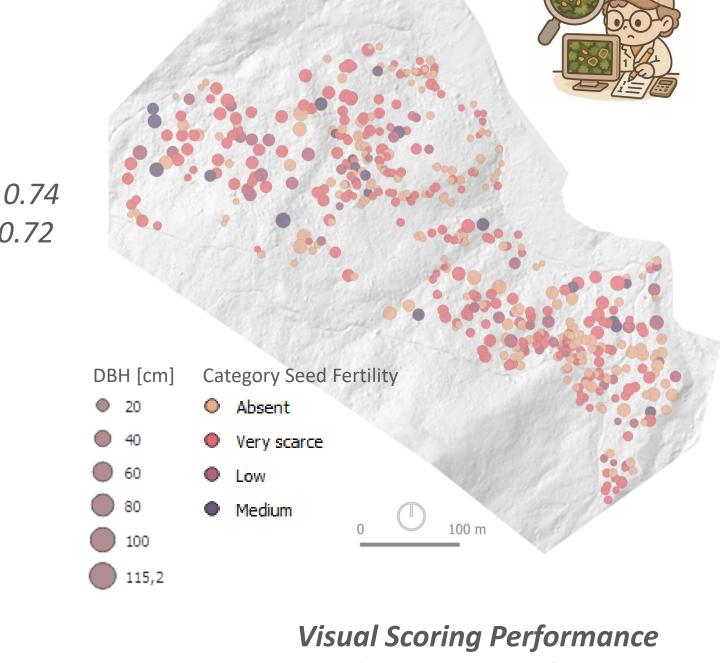


QA – do image-based fertility classes match seed/BA ratio? Fagus sylvatica 9 999



Seed-fertility maps from crown to stand scale – basis for spatial sampling and genetic interpretation





Fertility categories from semantic interpretation – visual ground truth; subjective by definition

Outlook Tree decision-path Artifacts? Drone seed mapping is powerful but fragile – it only works in a narrow "good condition" window

Fertility classes are consistent with seeds counts – higher class means more seeds per basal area across species Multiscale seed-fertility maps reveal intra-stand aggregation and provide spatial priors for genetic analysis

References

[1] Schuck, A., Van Brusselen, J., Päivinen, R., Häme, T., Kennedy, P., Folving, S., 2002. Compilation of a calibrated European forest map derived from NOAA-AVHRR data. EFI Technical Report 13, European Forest Institute.

Multi-scale seed

autocorrelation?

[2] Reudenbach, C. (2022). MetashapeTools: Scripts, utilities and optimized workflows for Agisoft Metashape Software. GitHub. Available at: https://github.com/gisma/MetashapeTools [3] Meyer, H., & Pebesma, E. (2021). Predicting into unknown space? Estimating the area of applicability of spatial prediction models. Methods in Ecology and Evolution, 12(9), 1620-1633. https://doi.org/10.1111/2041-210X.13650

All cartoon-style illustrations were edited using Bilbatree Art Designer

Two pinecone photographs used in the Study Site figure were obtained from iNaturalist.

Aerial images of the study site figure were acquired with our drones, except for site FRA0051, which is a Google Earth Pro screenshot









Pinus Pinaster



