



BEeS

The LifeWatch ERIC Biodiversity & Ecosystem
eScience Conference



Heraklion, 30 June - 3 July 2025

30 June 2025 | 15:00



Session: Taxonomy: Identifying the units of diversity in life

1 July 2025 | 14:30-16:30



Waking up from a taxonomist's nightmare:
Ophrys species delimitation meets AI in the
phylogenomic era

Presenter: Elisa Addonizio

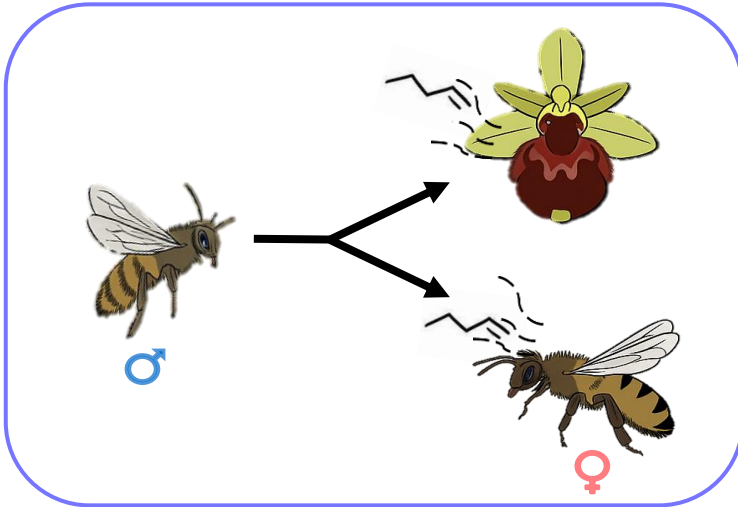


“Species are groups of actually or potentially interbreeding natural populations, which are reproductively isolated from other such groups”

– Ernst Mayr, 1942, *Systematics and the Origin of Species*



Case study: *Ophrys araneola*/sphegodes group



➤ Pollination in the orchid genus *Ophrys* works by **sexual deception**

➤ This genus generates deep controversy over species delimitation; estimates range from 9 to over 350 taxa, often based on subtle differences and presumed pollinator specificity



MAIN AIM

Develop and validate an automatic classification system based on deep learning to distinguish morphologically related species of the genus *Ophrys* from flower images

1

Can AI accurately discriminate morphologically similar *Ophrys* species from flower images?

2





How do model architecture, training strategy, and dataset composition influence classification performance?

3

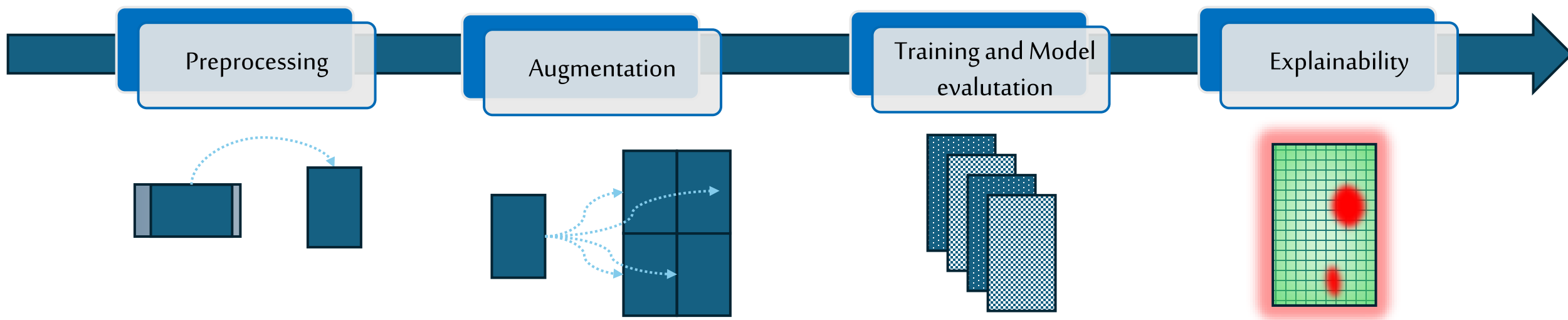
Is it possible to interpret and visualize the morphological features guiding the neural network's decisions?

Materials and methods

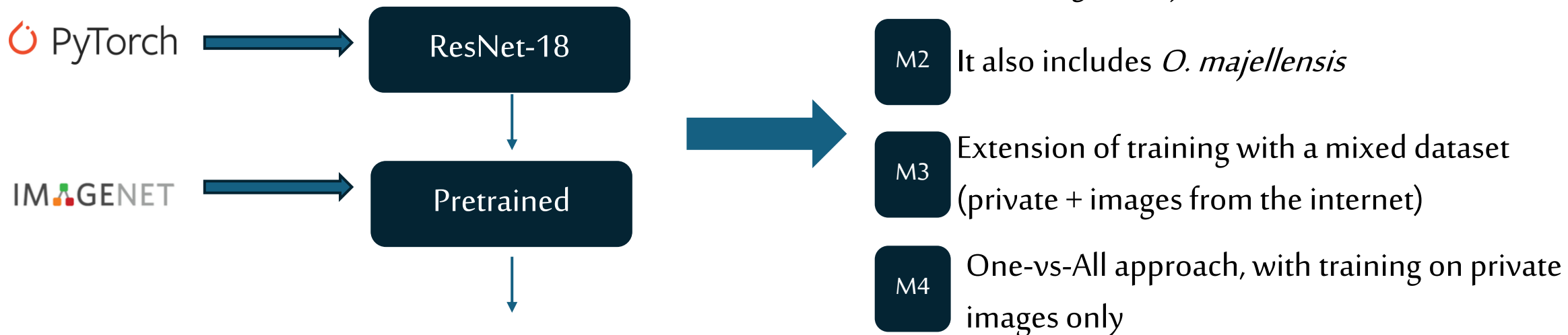
1. Dataset acquisition and image labelling

| <i>O. exaltata</i> | <i>O. garganica</i> | <i>O. incubacea</i> | <i>O. sphegodes</i> |
|--|---|--|--|
|  |  |  |  |
| 283 images | 251 images | 156 images | 100 images |
| 218 images | 319 images | 210 images | 218 images |

2. Design of a complete classification pipeline



3. Model training



Metrics for the model (private dataset)



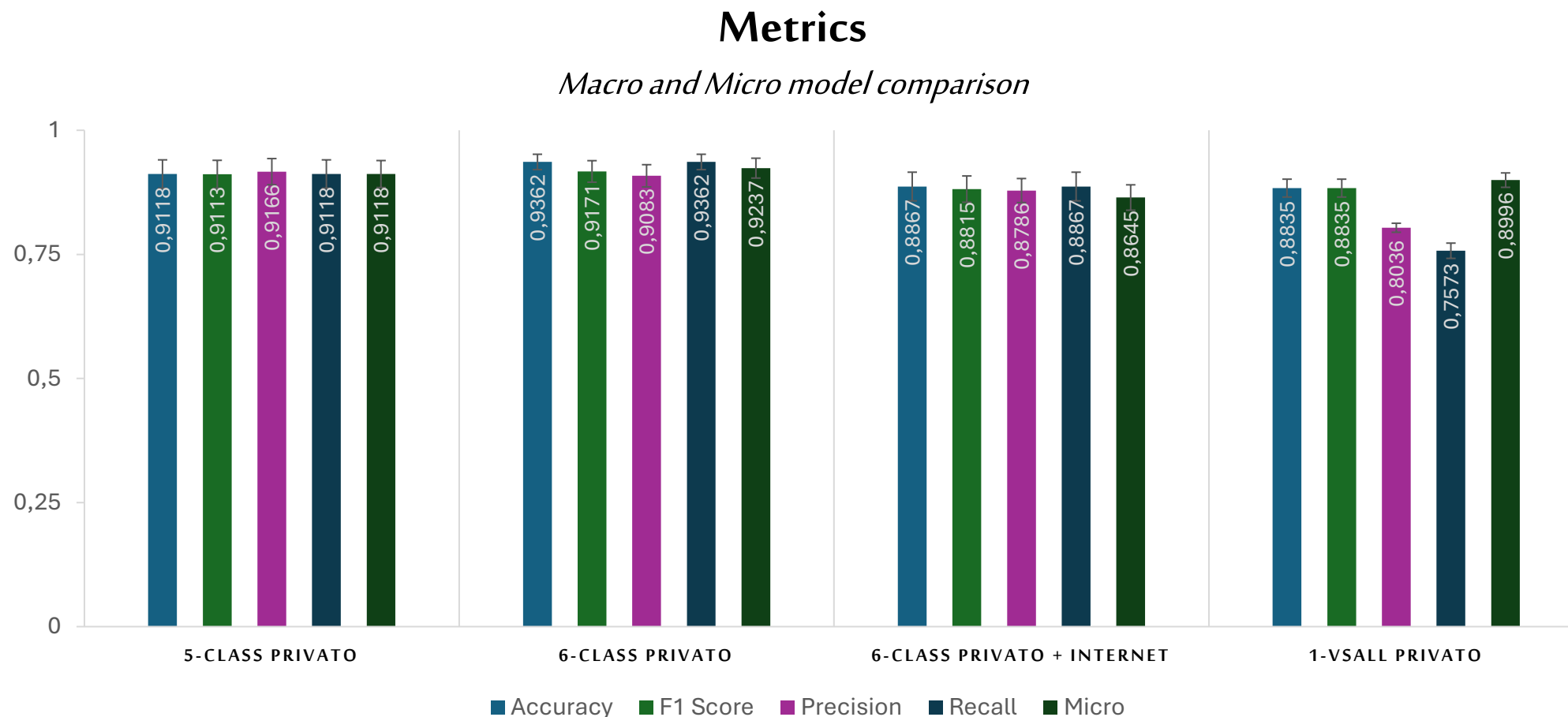
➤ This model achieves high performance, confirming the effectiveness of the automated morphological approach in distinguishing *Ophrys* species

Confusion Matrix for model (private dataset)

| Label\Predicted | <i>O. exaltata</i> | <i>O. garganica</i> | <i>O. incubacea</i> | <i>O. sphegodes</i> |
|---------------------|--------------------|---------------------|---------------------|---------------------|
| <i>O. exaltata</i> | 48 | 0 | 0 | 8 |
| <i>O. garganica</i> | 0 | 45 | 3 | 0 |
| <i>O. incubacea</i> | 0 | 1 | 30 | 0 |
| <i>O. sphegodes</i> | 0 | 0 | 0 | 20 |

➤ The results confirm strong class separation and high accuracy across all categories

2 How do model architecture, training strategy, and dataset composition influence classification performance?



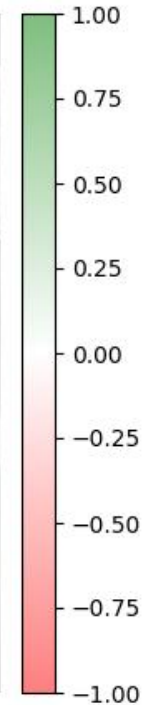
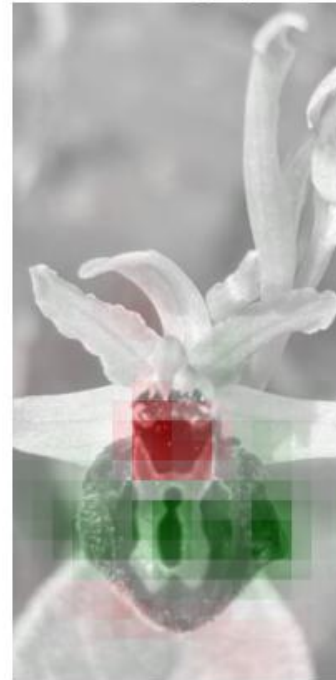
- All models show high performance; expanding to 6 classes and adding online data increases variability and generalization, while One-vs-All allows detection of unknown classes

Explainability

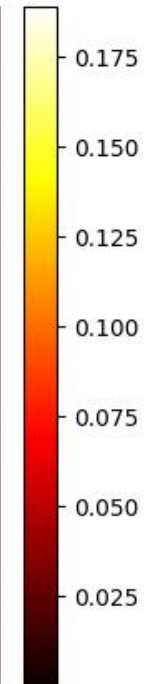
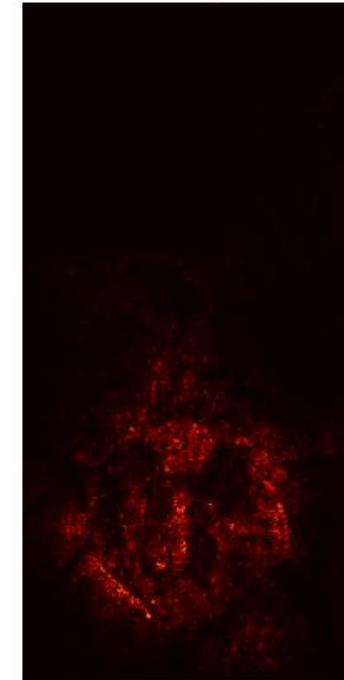
Predicted: *O. garganica*
Label: *O. garganica*



Occlusion attributions
for class *O. garganica*

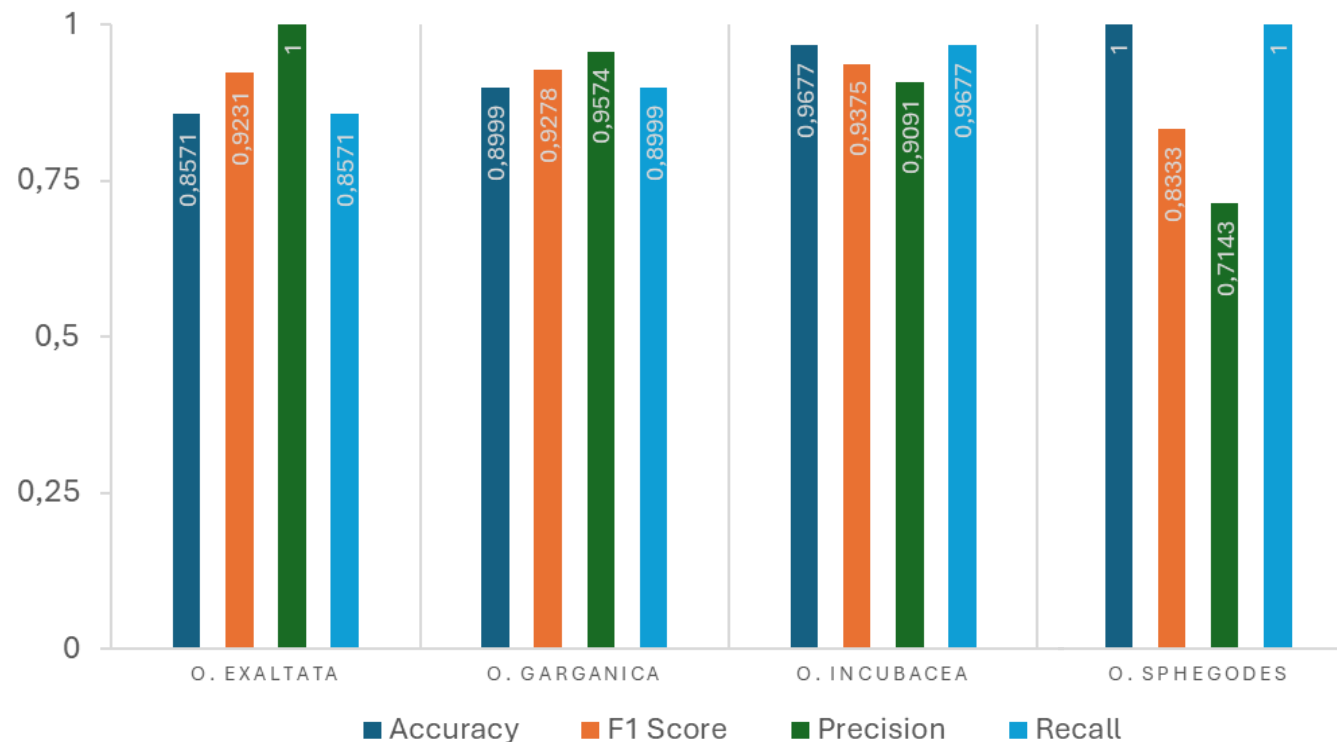


IntegratedGradients attributions
for class *O. garganica*

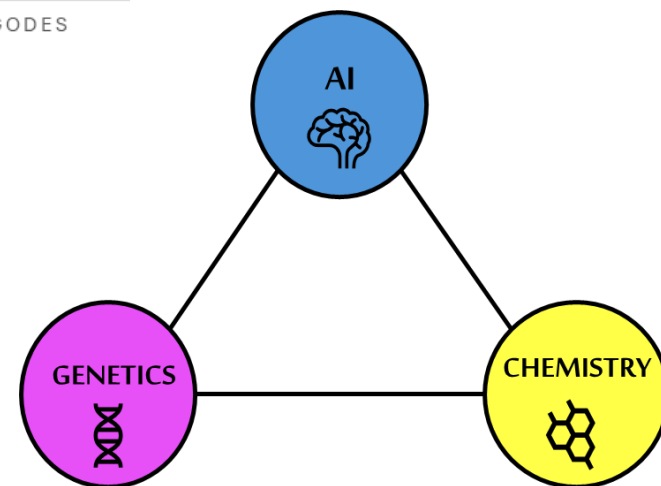


- Explainability techniques such as Occlusion and Integrated Gradients reveal that, across all analysed images, the most informative regions for classification consistently correspond to the labellum, confirming that the model relies on key morphological traits consistent with the biology of the *Ophrys* genus

The AI model successfully distinguished different *Ophrys* species, with several F1-scores above 0.93



Does the model's high classification accuracy reflect underlying
genetic and **chemical** differences?



Case study application: *O. sphegodes* and *O. majellensis* in Majella National Park

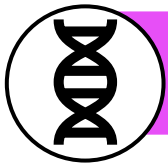


Ophrys sphegodes
❖ Flowering period: **April**



Ophrys majellensis
❖ Flowering period: **June**

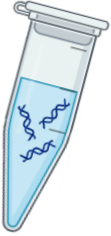
They share the lawn, but flower at different times



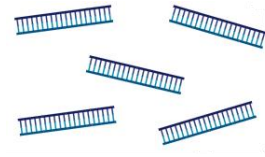
GENETICS

METHODS

DNA extraction

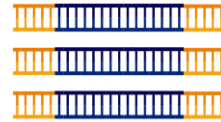


Digestion by
Double Restriction
enzymes



DNA fragments

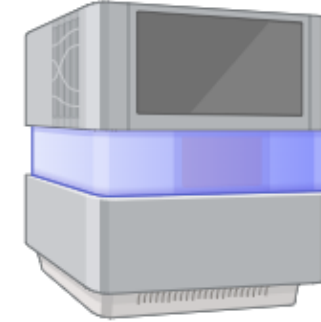
ddRADSeq



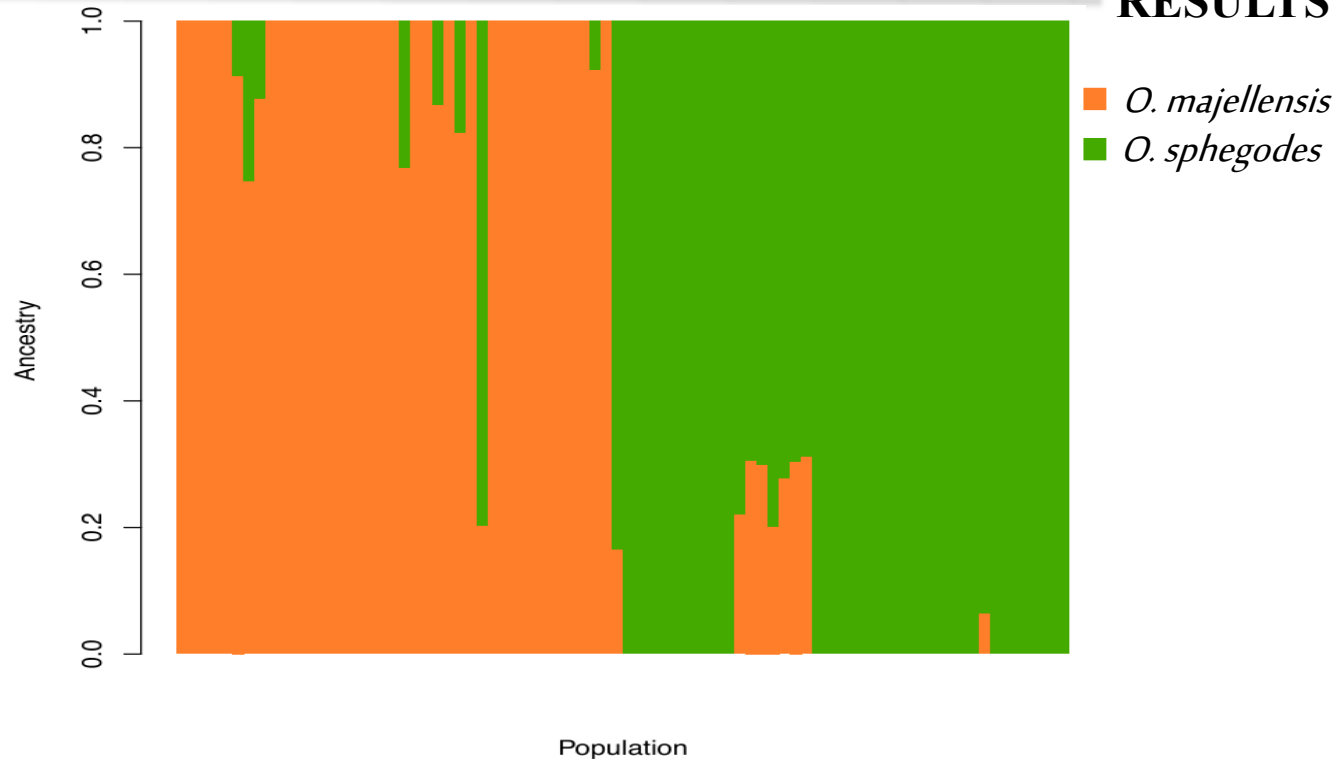
Adapter ligation



Sequencing



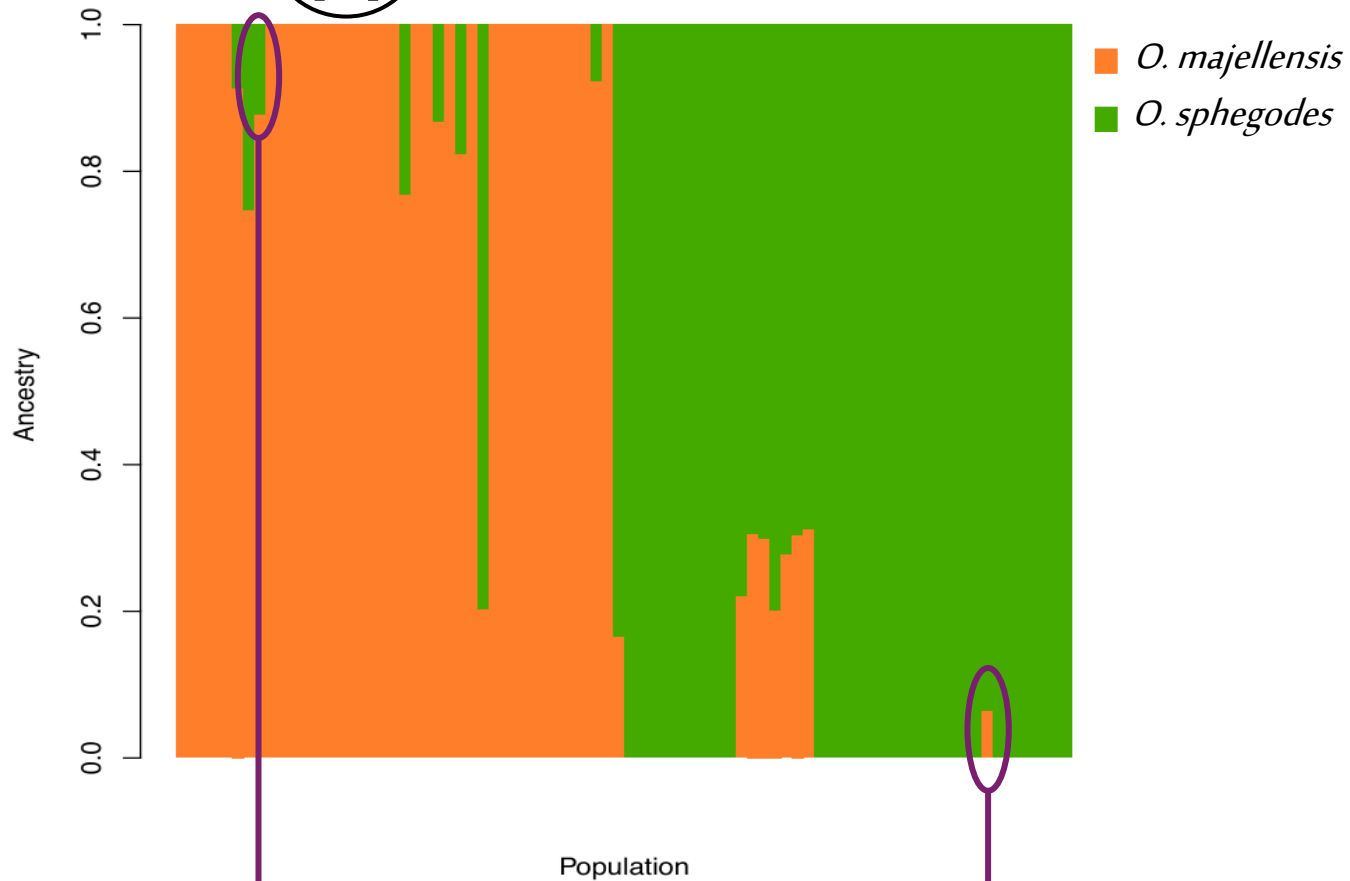
RESULTS



Some individuals show intermediate genetic assignment: in these cases, what do the morphological and chemical data indicate?



GENETICS



INDIVIDUAL: «**DM32**»
Ophrys majellensis

INDIVIDUAL: «**SC31**»
Ophrys sphegodes

100%
90%
80%
70%
60%
50%
40%
30%
20%
10%
0%

Genetic assignment

Genetic assignment for **DM32**:

- 87% *O. majellensis*
- 13% *O. sphegodes*

94%
93%
92%
91%
90%
89%
88%
87%
86%
85%
84%

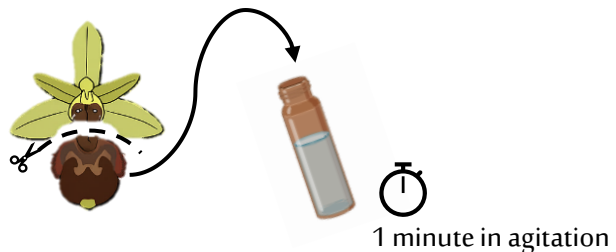
Genetic assignment

Genetic assignment for **SC31**:

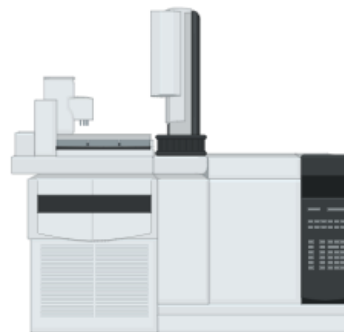
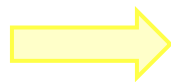
- 93% *O. majellensis*
- 7% *O. sphegodes*



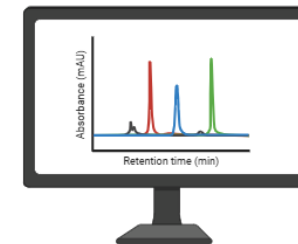
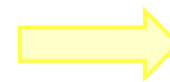
METHODS



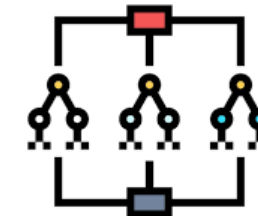
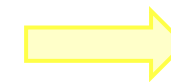
Sampling of labella in hexane



GC-MS analysis



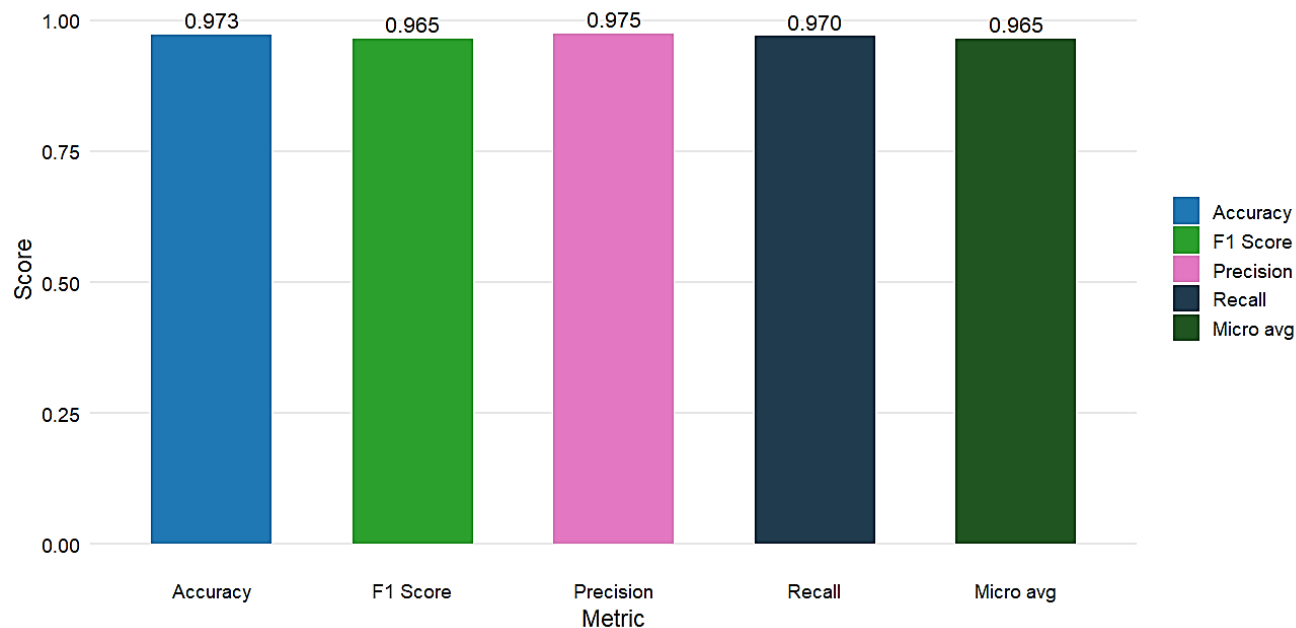
Chromatogram analysis and
peak assignment



Species classification using
VOCs and Random Forest

RESULTS

General Performance of the Model



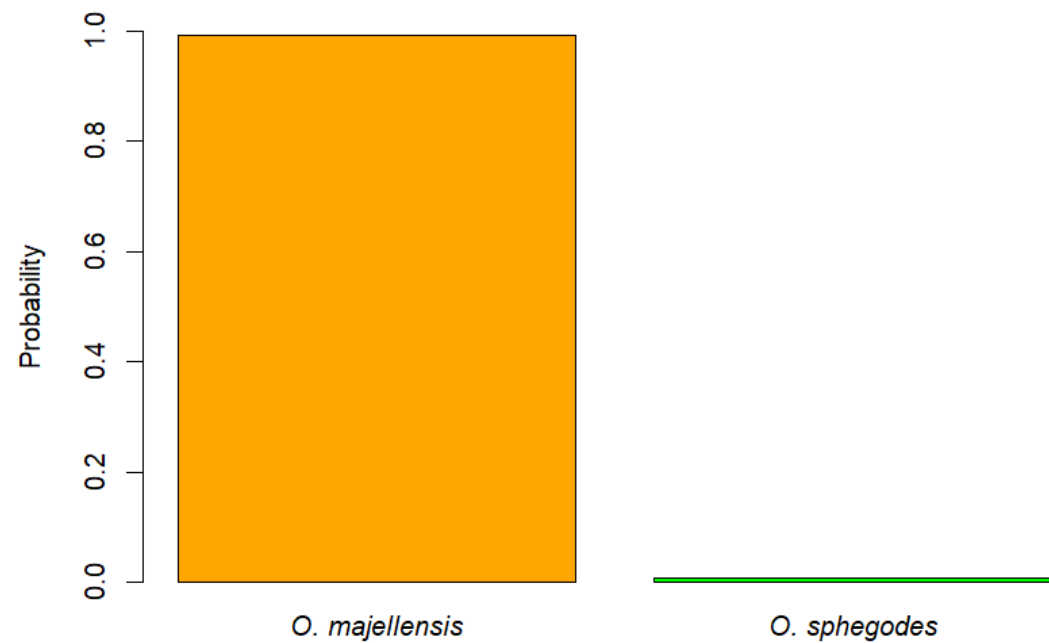
| Label\Predicted | <i>O. majellensis</i> | <i>O. sphegodes</i> |
|-----------------------|-----------------------|---------------------|
| <i>O. majellensis</i> | 38 | 2 |
| <i>O. sphegodes</i> | 0 | 33 |

VOC-based Random Forest classification supports a distinct chemical profile for *O. majellensis* and *O. sphegodes*



Species assignment based on volatile compound profiles

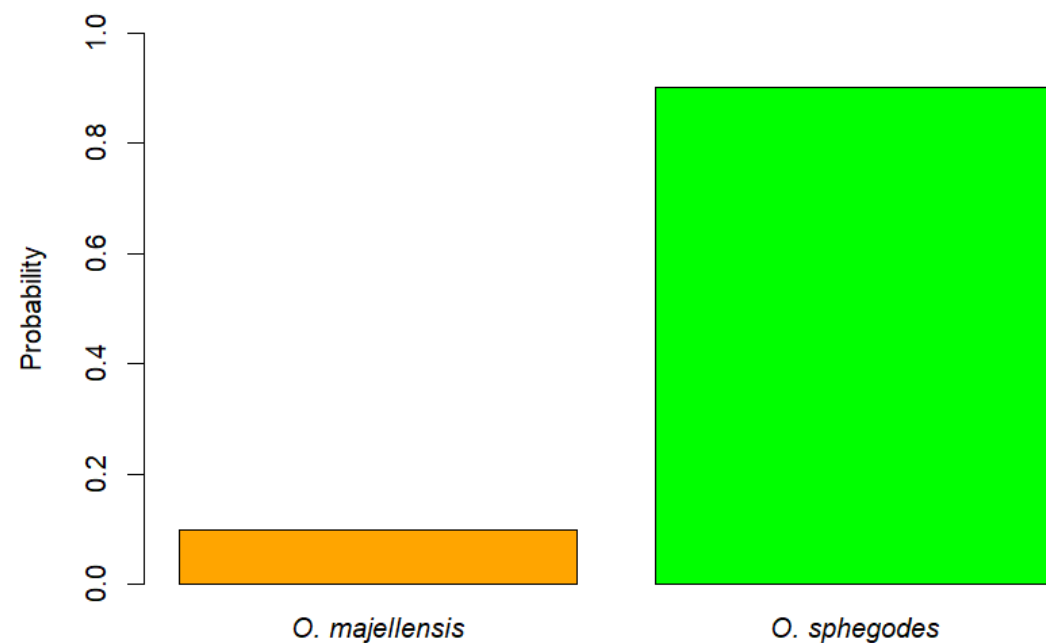
Assignment probability for DM32



O. majellensis: 99.3%

O. sphegodes: 0.007

Assignment probability for SC31



O. majellensis: 9.9%

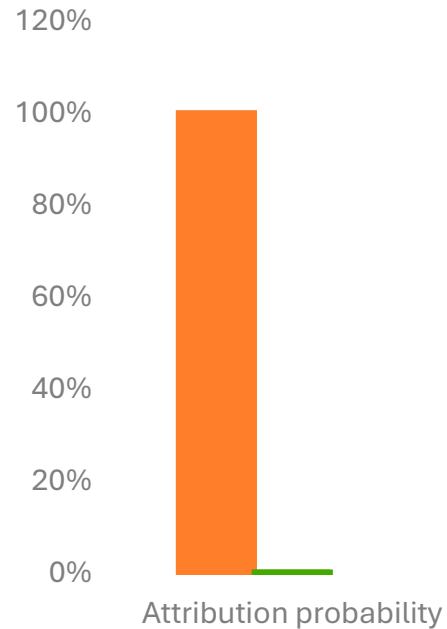
O. sphegodes: 90.1%

The percentages represent classification probabilities generated by a Random Forest model



ARTIFICIAL INTELLIGENCE

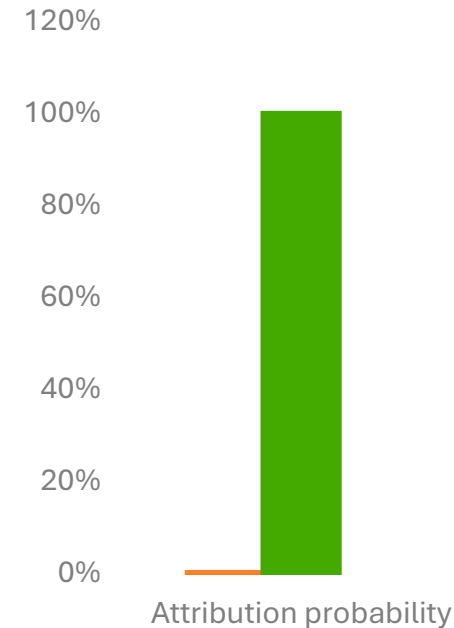
Species attribution obtained through automatic classification of the flower image using a convolutional neural network



Predicted class probability for «**DM32**»:

O. majellensis: **99.98%**

O. sphegodes: 0.02%



Predicted class probability for«**SC31**»:

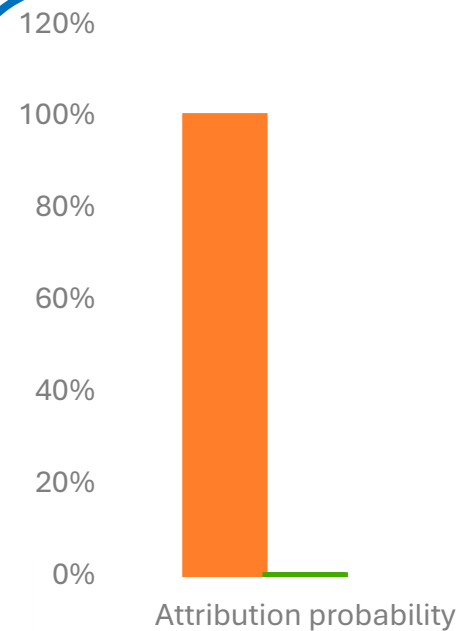
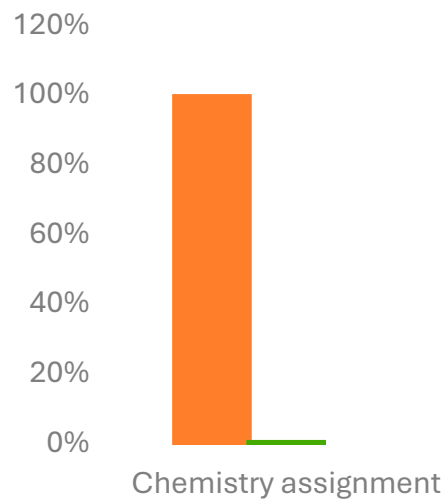
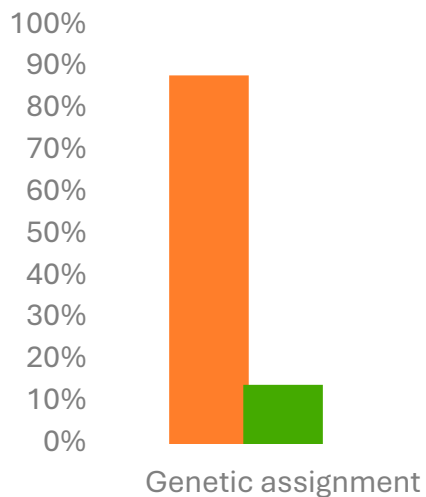
O. majellensis: 0.01%

O. sphegodes: **99.99%**

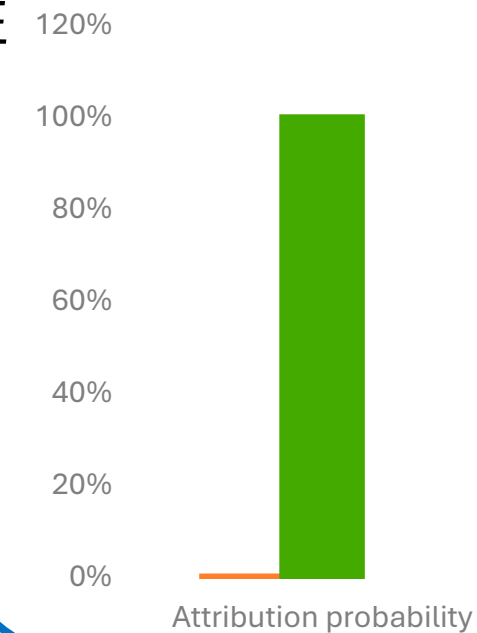
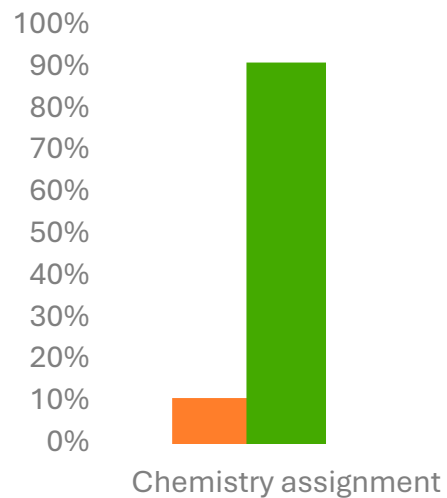
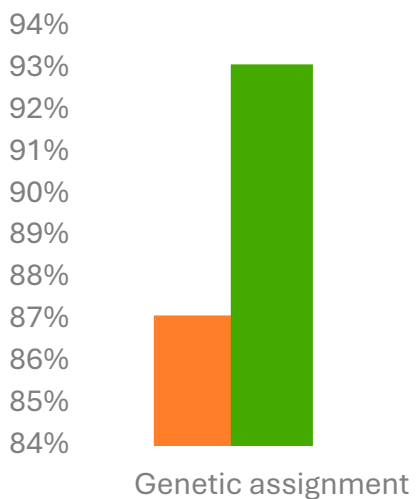
The percentages represent the probabilities of assignment predicted by the model for each species

CONCLUSION

DM32



SC31



ARTIFICIAL
INTELLIGENCE



GENETIC



CHEMISTRY

CONCLUSION

- ❖ The results highlight how AI techniques achieve high accuracy rates
- ❖ Samples with less defined genetic assignments show high assignment with chemical data and AI
- ❖ AI-based classification system is able to capture complex morphological and phenotypic nuances that escape genetic analysis alone, thus offering a promising and more comprehensive tool for the classification of species in *Ophrys*

Thank you!

Questions?

elisa.addonizio@unina.it



BEeS

The LifeWatch ERIC Biodiversity & Ecosystem
eScience Conference



Heraklion, 30 June - 3 July 2025