









Global Orchid Diversity in the Mediterranean Biome: identifying hotspots and exploring automated extinction risk assessment with deep neural networks

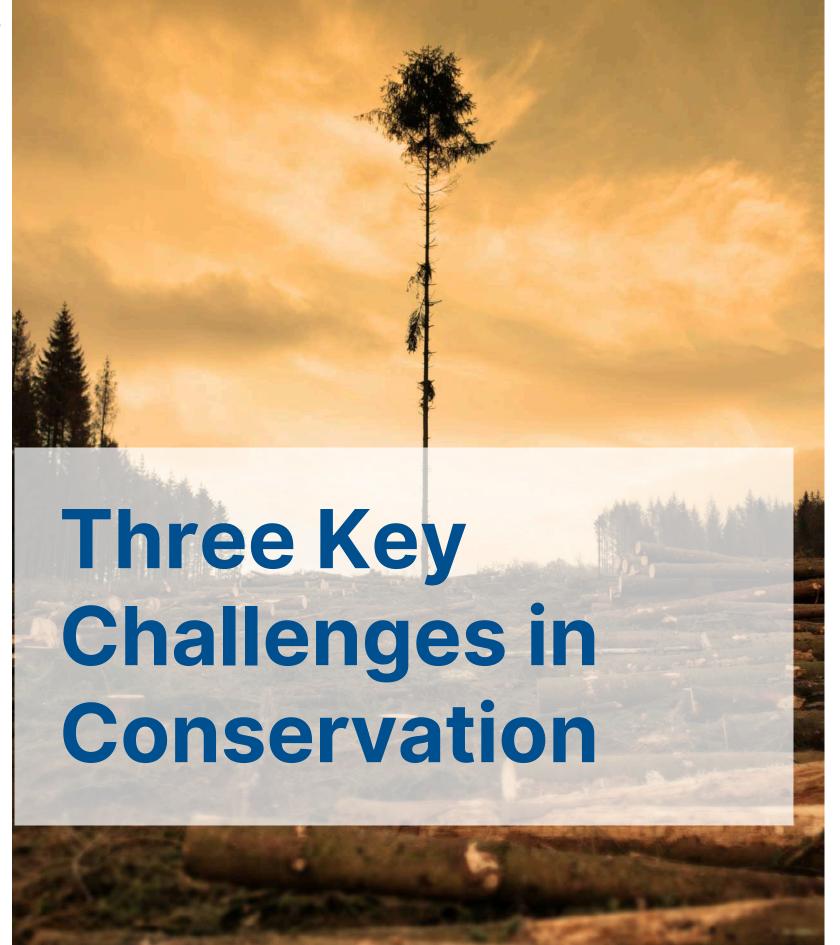
Lussu, Scaccabarozzi, Brundett, Santi, Silvestro, Chiarucci

Presenter: Michele Lussu







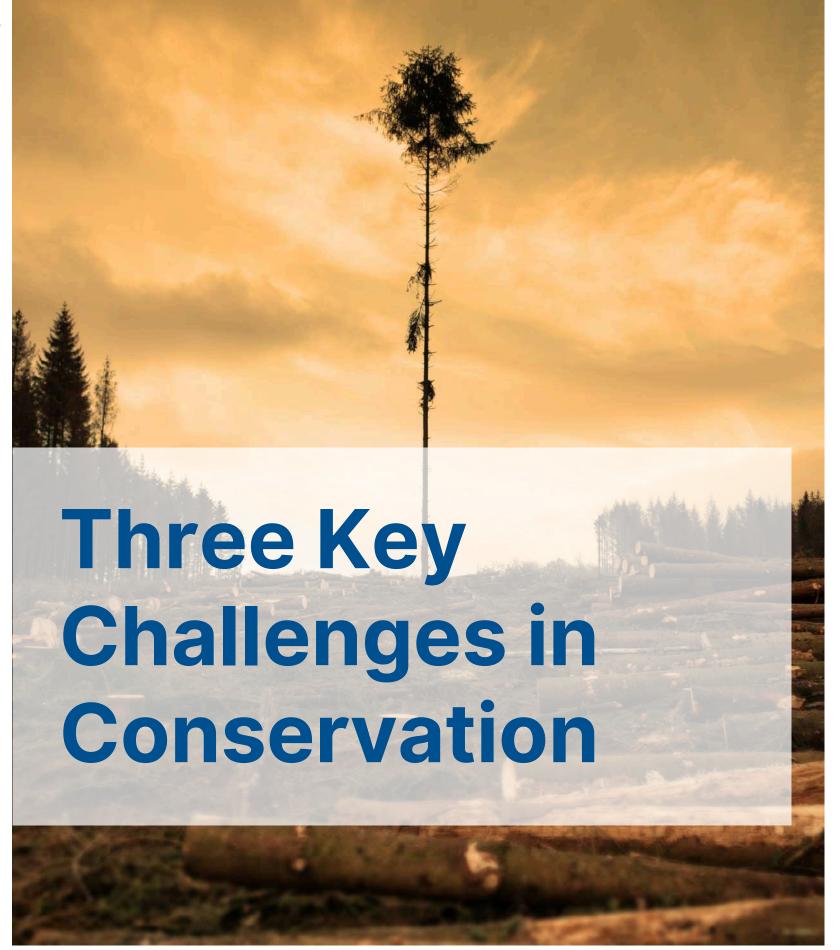


Biodiversity is complex and multidimensional: species richness, functional and phylogenetic diversity

Collecting and analyzing biodiversity data is time-consuming and resource-intensive.

The biodiversity crisis is urgent, requiring faster and more effective action



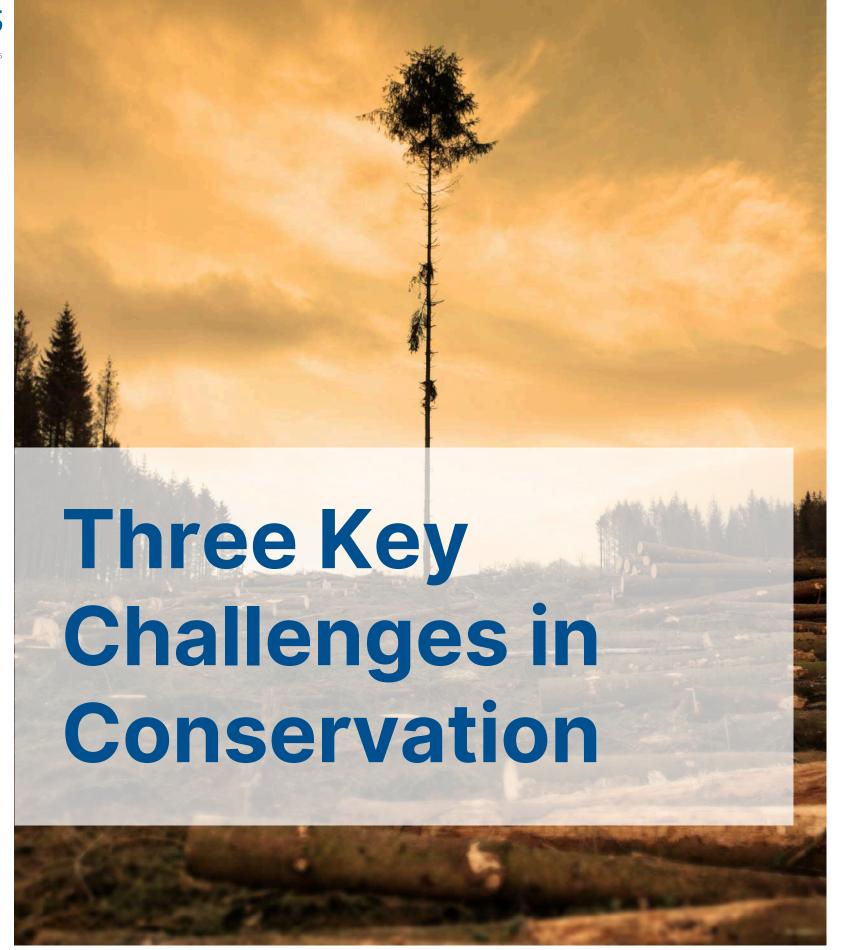


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ARTIFICIAL INTELLIGENCE

The potential for AI to revolutionize conservation: a horizon scan

Trends in Ecology & Evolution



Revie

Species delimitation 4.0: integrative taxonomy meets artificial intelligence*

Kevin Karbstein ^{1,*}, Lara Kösters ¹, Ladislav Hodač ¹, Martin Hofmann ², Elvira Hörandl ³, Salvatore Tomasello ³, Natascha D. Wagner ³, Brent C. Emerson ⁴, Dirk C. Albach ⁵, Stefan Scheu ^{6,7}, Sven Bradler ⁶, Jan de Vries ^{8,9,10}, Iker Irisarri ¹¹, He Li ¹², Pamela Soltis ¹³, Patrick Mäder ^{2,14,15}, and Jana Wäldchen ^{1,14}

Sam A. Reynolds (1) 1,*, Sara Beery (2) Neil Burgess (3) Mark Burgman 5, Stuart H.M. Butchart (6) 6,7, Steven J. Cooke (6) 8, David Coomes (6) 9, Finn Danielsen (6) 10, Enrico Di Minin (6) 11,12, América Paz Durán (6) 13, Francis Gassert 14, Amy Hinsley (6) 15, Sadiq Jaffer (6) 16, Julia P.G. Jones (6) 17, Binbin V. Li 18,19,



AI IN CONSERVATION BIOLOGY

KEY APPLICATIONS

- Extinction risk prediction
- Trait imputation, data gap filling
- Prioritization of protected area
- Trend detection

PRO

- Handles complex datasets
- Accelerates decisionmaking

CONS

- Dependent on data quality
- Requires validation
- Cannot replace expert judgment

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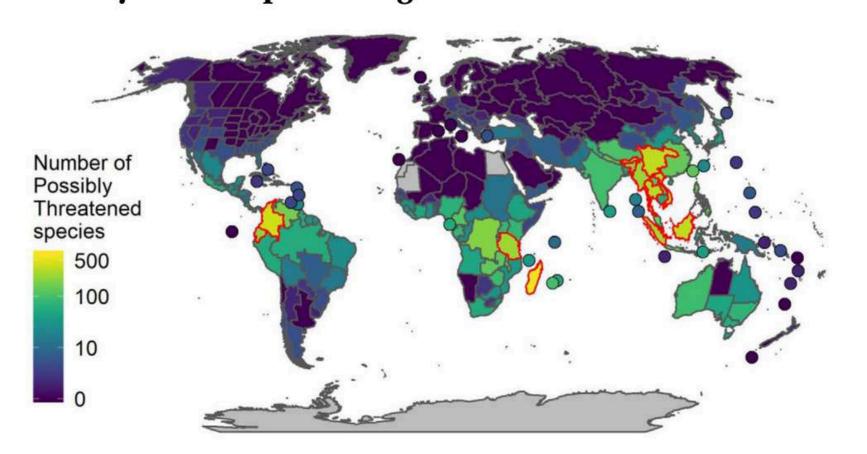
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Conservation Biology



Contributed Paper

Automated conservation assessment of the orchid family with deep learning



automated methods can significantly improve the speed and scope of extinction risk assessments for highly diverse groups like orchids



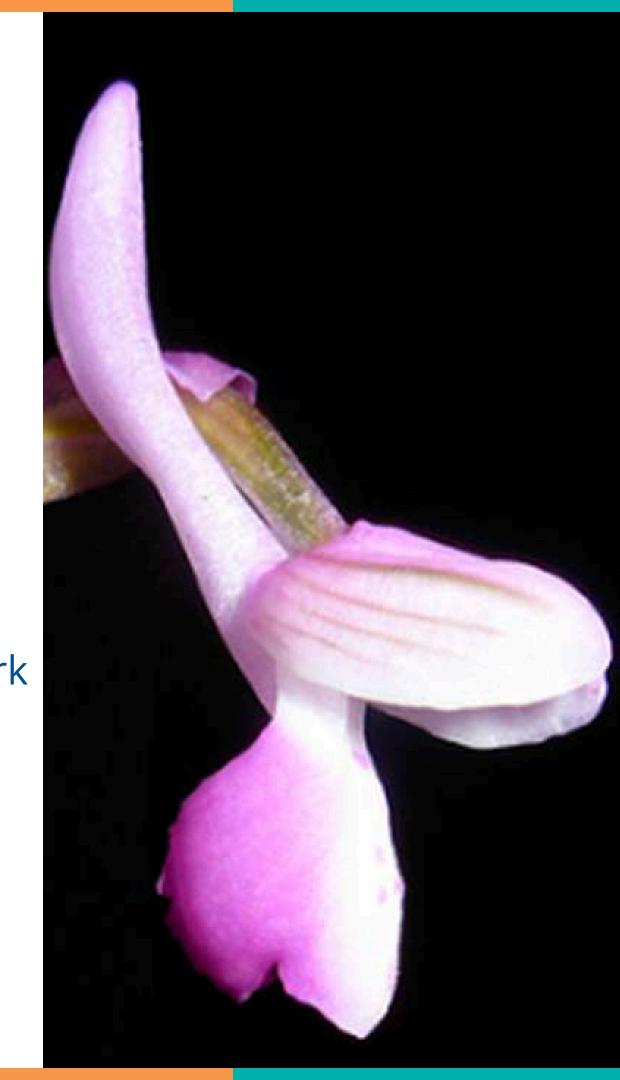
AIMS

01

Identify orchid hotspots, combining species richness, functional, and phylogenetic diversity accross Mediterranean Biome

Predict extinction risk of unassessed species using neural network

Identify conservation priority areas in Mediterranean Biome





THE MEDITERRANEAN BIOME

Hotspot of plant diversity

2% of the world surface is covered by the Mediterranean biome, it harbours more than 20% of the plant diversity

An orchidologically well-explored biome

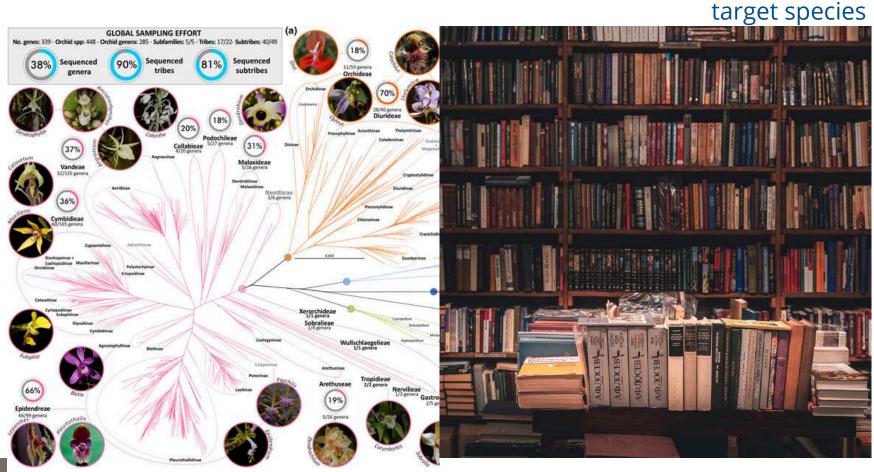
The Mediterranean biome is among the most botanically explored, with extensive floristic and taxonomic research

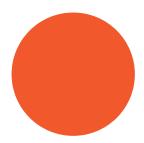




WHY ORCHIDS?







28.000 species world wide



Extensive literature



Recent phylogeny.

Perez-Escobar 2024



DATA WORKFLOW

- **Download species occurrence**s from GBIF for the Mediterranean Biome.
- Standardize taxonomy according to Plants of the World Online (POWO).
- Retrieve official IUCN Red List assessments via iucnredlist.org.
- Compile functional traits from published literature and expert sources.

Diversity metrics

- Occurrence data linked to TDWG Level 4. (species richness)
- **FD** computed using Gower distance (traits used: trophism, pollination strategy, growth form, fire response)
- Phylogenetic tree pruned to target species (picante) (**FD**)







The need of a rigorous global standard for assessing species extinction risk to train Al.

Key Strengths:

- Scientifically robust
- Globally recognized
- Essential for conservation planning and policy

Critical Gaps:

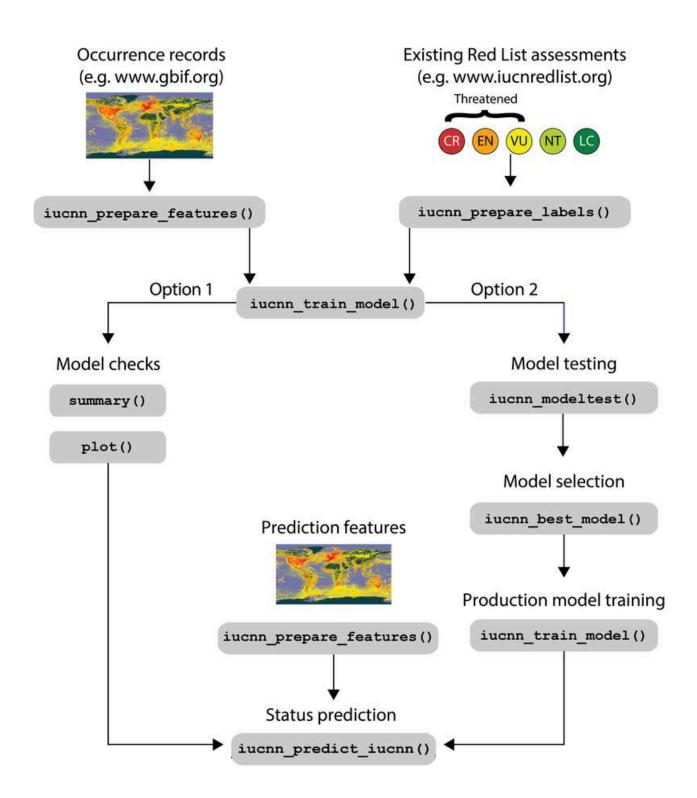
- <10% of species assessed globally
- Taxonomic bias (focus on vertebrates)
- Geographic bias (underrepresentation of biodiversity hotspots)
- Time- and data-intensive



AI-BASED RISK PREDICTION

Data Workflow

- Used only species classified in CR EN & VU categories
- Features: range size, climatic niche, human footprint, biome occupancy
- Retained only high-confidence predictions using Monte Carlo dropout
- Final selection: species flagged as "at risk" by ≥3 models (elbow threshold)



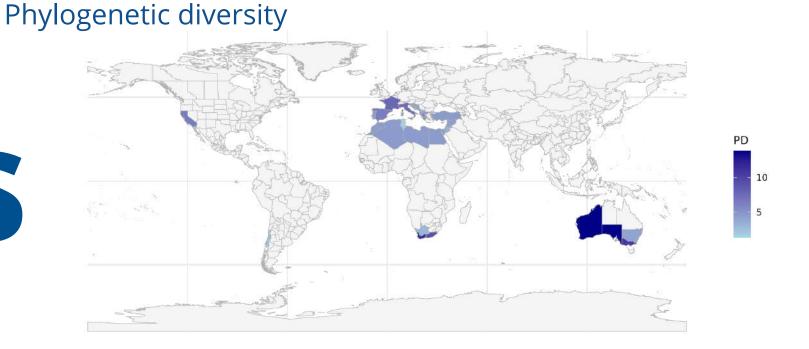


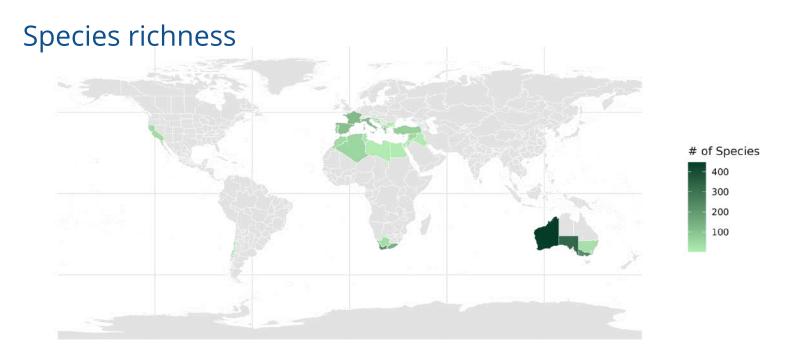
RESULTS

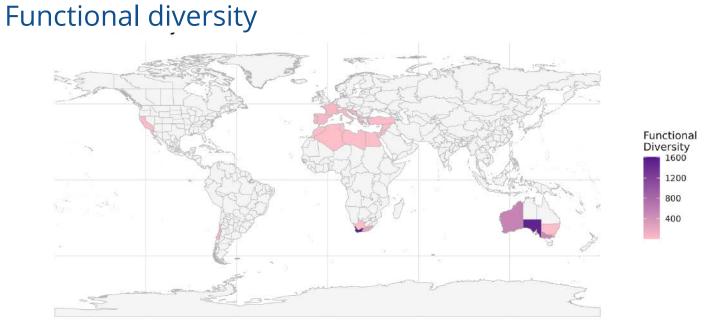




1264 species







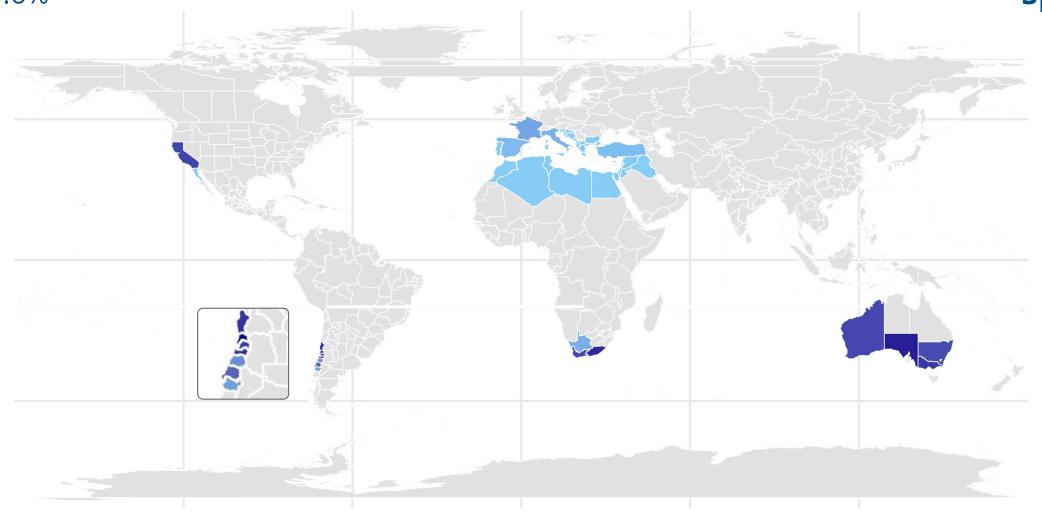
There is a highly significant difference between biodiversity metrics. This result shows that the distributions of PD, FD, and SR within MB are not equivalent.



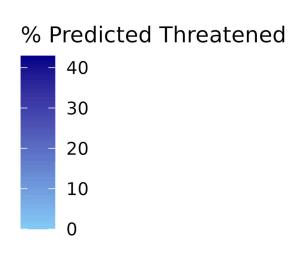
DL-BASED RISK PREDICTION

IUCN assessment: NA 92%, LC 4.6%, NT 1.1%, VU 0.7%, EN

0.7%, CR 0.6%



1160 species evaluated by DL



438 (34.6% of total) species predicted as threatened



Discussions

- SR, FD and PD shows different patterns
- 34.6% of orchid species are predicted as threatened VS 2,4% already assessed
- The % of predicted threatened species varies accross Mediterranean botanical countries



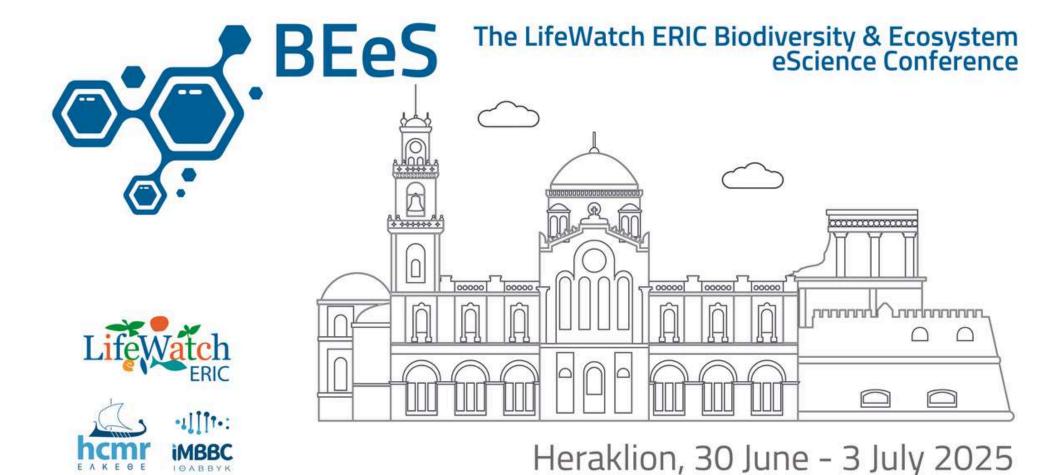


Take-Home Messages

- Integrating taxonomic, functional, and phylogenetic perspectives is crucial for effective conservation.
- Deep learning emerges as a powerful ally for conservation biology, **enabling fast, scalable, to support** assessments that traditional methods alone cannot achieve.

Thank you!

Questions? michele.lussu@unibo.it



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