

De Pl@ntNet à GeoPl@ntNet: nouvelles approches d'IA pour le monitoring de la biodiversité

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A citizen science platform that uses AI to help people identify plants with their mobile phones





25 Million users 200+ countries Up to 2M identifications per day

Personal Usage



Nature, walks





Phytotherapy



Professional Usage



Agro-ecology



Education, animation



Natural Areas Management



Tourism

Trade

Key concept of Pl@ntNet: Collaborative Al







Multi-head model trained on Jean Zay super-computer on a big dataset of 8M valid observations (5-6 days of training)

$$\sigma_{species}(f(x)) = 50 \text{K classes}$$

$$\sigma_{organs}(f(x))$$

$$\sigma_{genus}(f(x))$$

$$\sigma_{disease}(f(x))$$

Model = Vision transformer DinoV2

- Backbone pre-trained on 100M images using SSL (by Meta/Inria)
- Final multi-head model fine-tuned on 8M PI@ntNet images (by PI@ntNet team)

A difficult problem: uncertainty

Irreducible uncertainty

Species ambiguity

Model uncertainty Increased by long-tail distribution

Top1 Identification accuracy:

Common species	=	~90%
Average species	=	~70%
Rare species	=	~40%

Use of regional or thematic floras

Restricting the hypothesis space to a particular flora allows improving the identification accuracy

Pl@ntNet Similarity search

Deep neural network

Similarity search engine **9M images**

\rightarrow Sub-linear algorithm based on locality sensitive hashing

Joly, A., & Buisson, O. (2011, June). Random maximum margin hashing. In CVPR 2011 (pp. 873-880). IEEE.

User's contributions

Users can contribute their observations

User's revisions

Users can revise observations of other users.

← 4 Nom(s) commun(s) Français					
Larix decidua Mill.					
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Pin de Briançon	4 [≣]	1 6 Votes			
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Ajouter un nom					

Salsir l'espèce

Cooperative Learning algorithm

The most probable label of an observation is determined with a weighted majority voting rule:

 $U_i =$ Set of users who provided a label y_i^u for the observation x_i

Cooperative Learning algorithm

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The weight of a user in PI@ntNet is a function of the **estimated number of species** he is able to identify

940M raw observations (=queries)

Pl@ntNet Data visualisation tools

Bryonia cretica L.

Mag

سار دارو، فاشرا ,White bryony, Cretan bryony

Common name(s) White bryony Cretan bryony مار داروه فاشر ا View all / Edit &

Halling Hadden (Perty)

Pl@ntNet Data shared in GBIF

Top-5 data provider to GBIF (world's largest infrastructure for biodiversity data)

- Shared data = revised observations + trusted queries identified by the AI (AI score>0.95)
- Quality filters: potted & cultivated plants removal, region-based filtering (Kew POWO)

https://doi.org/10.15468/mma2ec

Objective: which species are present in a given location and why?

Raw species occurrence data needs to be interpolated in space and time:

Many plant occurrences at world scale

But very few locally for most species

Species Distribution Models (SDM)

Species Distribution Models (SDM)

Motivations

- Help conservation/ plans
- Invasive plant monitoring
- Simulation under climate change
- Learn about species preferences

Credits: "Introduction to species distribution modelling (SDM) in R", Damaris Zurell

Different types of SDMs

Niche models (e.g. GLM, MAXENT)

- Input: **low-dimensional** (e.g. temperature, precipitation)
- Purpose: interpretability, explicability

ML models (e.g. Random Forest, XGBoost)

- Input: high-dimensional vectors (e.g. 100 environmental variables)
- Purpose: **performance**, easy to use

Deep SDMs (e.g. CNNs, transformers)

- Input: **complex signals** (e.g. remote sensing images, time series)
- Purpose: performance on large number of species, very high resolution

Remote sensing based SDM

PLOS COMPUTATIONAL BIOLOGY

Convolutional neural networks improve species distribution modelling by capturing the spatial structure of the environment

Benjamin Deneu 🖪, Maximilien Servajean, Pierre Bonnet, Christophe Botella, François Munaz, Alexis Joly

Model Input = data cubes

Model Output = Suitability score of each species

From models to species mapping

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Inference/mapping phase

Different tasks vs. available data

-

Input data: ${\mathcal X}$

target: y

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12

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- Abundance data (very hard to produce) Task: predict $\hat{y} = f_{\theta}(x) \in \mathbb{R}^d$
- Presence / absence data (hard to produce) Task: predict $\hat{y} = f_{\theta}(x) \in [0,1]^d$
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- Presence only data (more data available)

Task: predict
$$\hat{y} = f_{\theta}(x) \in \{1, ..., d\}$$

Limitations of models trained on presence-only data

Sensitive to taxonomic reporting bias

Observation probability ≠ Presence probability

The threshold λ over the estimated probabilities is **hard to set** (we don't know how many species there are)

The probability of each species is **relative** to the others and depends on the **number** of species present somewhere \rightarrow this is not appropriate for mapping each species individually GeoLifeCLEF challenge 2023 & 2024

GeoLifeCLEF challenge 2023 - results

GeoLifeCLEF challenge 2023 - best approach

Leverage Samples with Single Positive Labels to Train CNN-based Models For Multi-label Plant Species Prediction Huy Quang Ung, Ryoichi Kojima, Shinya Wada

Architecture

GeoLifeCLEF challenge 2023 - best approach

Leverage Samples with Single Positive Labels to Train CNN-based Models For Multi-label Plant Species Prediction Huy Quang Ung, Ryoichi Kojima, Shinya Wada

PA = Presence/Absence data (with Binary Cross Entropy loss) PO = Presence only data (with Cross Entropy loss)

GeoLifeCLEF challenge 2023 - best approach

Leverage Samples with Single Positive Labels to Train CNN-based Models For Multi-label Plant Species Prediction Huy Quang Ung, Ryoichi Kojima, Shinya Wada

1280 x 1280 m 10 m **10K** species resolution 10 m 0,3 1 km resolution 128 x 128 resolution pixels **IOK** species Sentinel RGB-IR CNN model FI-score 0.2 64x64 km **Ensemble model 10K** species 0,1 64 x 64 **pixels** Environmental CNN model 0.0 MAMBO ensemble model MAMBO RGB-IR model MAMBO environmental model

Architecture

Contribution of modalities

Integration in PI@ntNet for EU-scale species mapping

Greece

Species predictions Model: GPN_RGBI_2 Grid: greece Resolution: 50 m Species threshold: 50

Fraxinus ornus L. (#6353)

WMS opacity:

Fraxinus ornus Family: Oleaceae Genus: Fraxinus Common Name: Manna Model prediction score: 13.32

Species predictions Model: GPN_RGBI_2 Grid: cbnmed Resolution: 50 m Species threshold: 50

Thymus vulgaris L. (#5404)

WMS opacity:

Thymus vulgaris Family: Lamiaceae Genus: Thymus Common Name: Garden thyme Model prediction score: 13.67

Anthemis maritima

Family: Asteraceae

Genus: Anthemis

Common Name:

Seaside Chamomile

GeoPl@ntNet Barcelo	ne
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Biodiversity indicators	-
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	4216	Salicornia fruticosa (!:J1.,.				83.82	D62
	9752	Phragmites australis (Cav.) Trin. ex Steud.	LC			82.70	-
	3878	<u>Trii:iolium</u> i;iannonicum <u>(Jacg.)</u> Dobrocz.				82.56	
	7137	Juncus maritimus Lam.				81.70	Contraction of Method
3 La.	2212	Limonium <u>vulgare Mill</u> .				77.42	
	8238	Juncus acutus L.	LC			72.80	
	4012	Anthemis maritima L.				70.72	
	7590	Suaeda maritima (bl Dumort.				67.99	
de Carrie	9411	Artemisia caerulescens L.				67.39	
- di	4846	Suaeda vera Forssk, ex J.F.Gmel.				67.19	
1 and	7950	Schoenus nigricans L.	LC			63.67	
Ch.	8456	Elaeagnus angustifolia L.	LC			63.62	

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Mapping biodiversity conservation indicators

From the species assemblage predicted at each point

$$S_{\lambda}(x) := \{k \in \mathcal{Y} : \hat{\eta}_k(x) > \lambda\}$$

We can compute indicators such as:

- The number of endangered species (e.g. on IUCN red list)
- The proportion of woody species (carbon capture)
- The diversity of species (e.g. Shanon index)
- The number or rare species
- The EUNIS habitat (using a species-to-habitat model)

We can construct maps of such indicators at very high resolution by computing $S_{\lambda}(x)$ for all x_i on a dense spatial grid

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CBN Med

Biodiversity indicators

Specialization

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Biodiversity indicators

EU directive

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Discover plant biodiversity close to home and help protect it better

Thank you

INRAC agropolis fondation

