

# AI as a Catalyst in Entomological Research by Simplifying Species Identification

**Hossein Shirali**

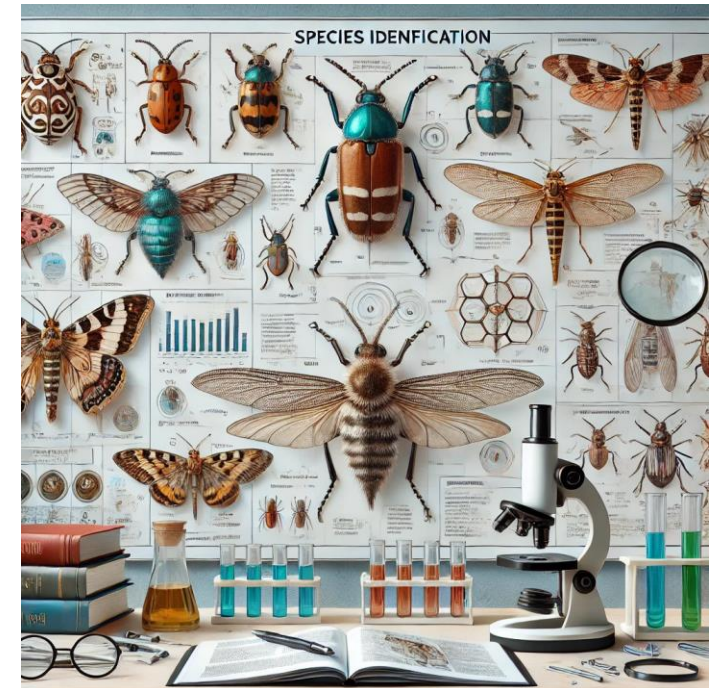
Karlsruhe Institute of Technology



# Introduction

## Challenges in Species Identification<sup>[1]</sup> :

- High diversity and morphological similarity among species.
- Manual identification prone to human error.
- Time-consuming and labor-intensive processes.
- Requires specialized expertise and extensive training.
- Limited scalability for large-scale studies.





# Introduction

## Opportunities of AI<sup>[2]</sup>

- AI enhances the efficiency and accuracy of rapid species identification.
- Automated image recognition and classification.
- Machine learning algorithms tailored for entomological data.
- Enhanced data collection and analysis.
- Facilitates large-scale biodiversity studies.



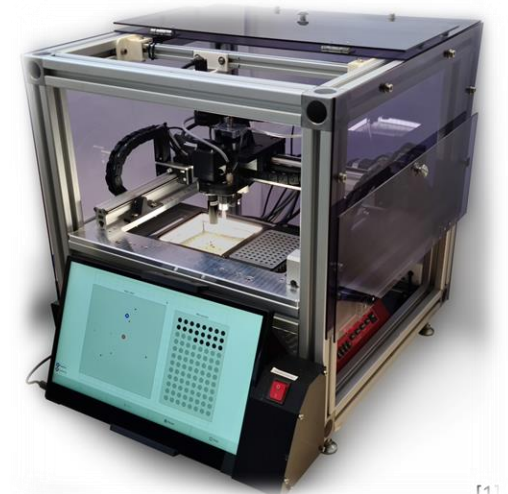
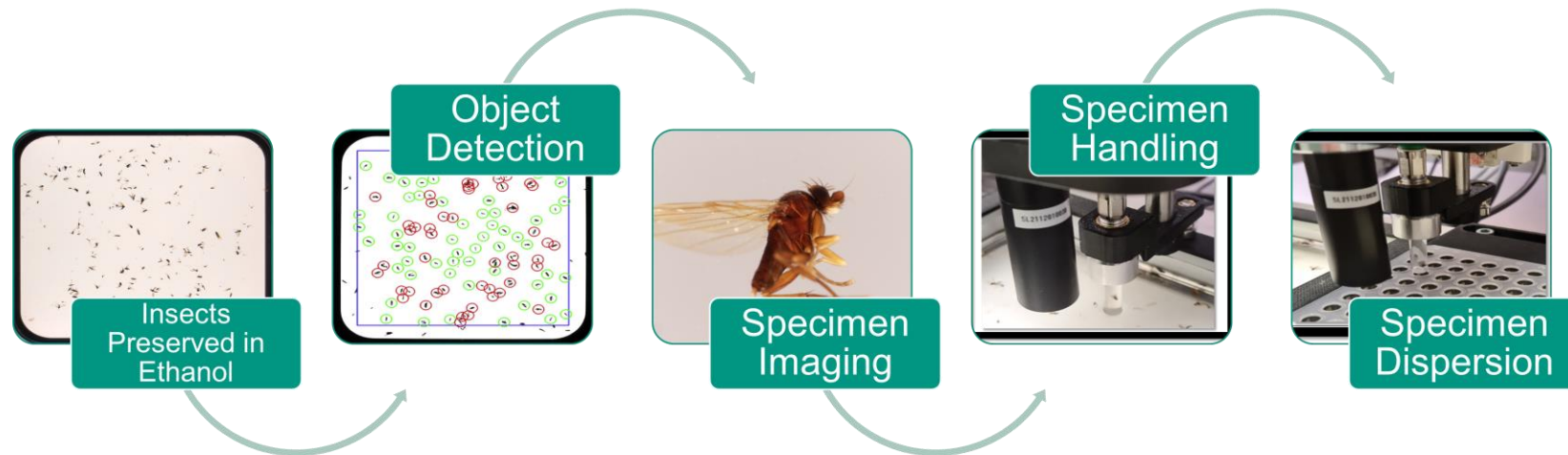
# AI in Entomological Research

## Image-based species identification

1. Tools and Data Collection
2. Data Labeling and Cleaning
3. Machine Learning Models for Species Identification
4. Automation in specimen classification and sorting

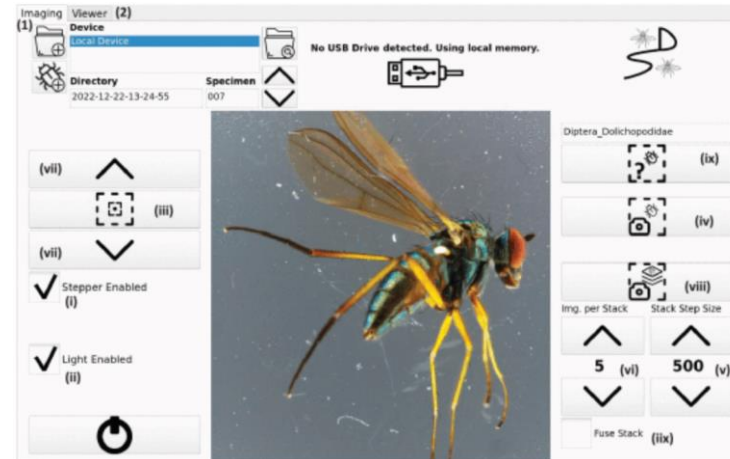
# Tools and Data Collection

- **DiversityScanner** [3]:
  - Automatically Capture high-resolution images and sort specimens.



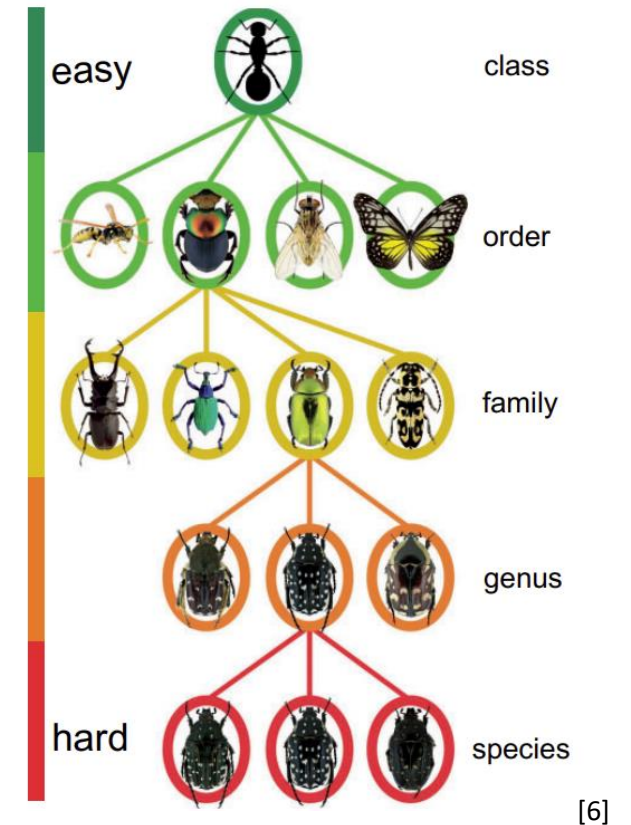
## Tools and Data Collection

- **Entomoscope** [4]:
  - A DIY photomicroscope for imaging invertebrates preserved in ethanol.
- **Workflow Integration:**
  - Streamlines data collection processes.
  - Significantly improves research efficiency.
  - Reduces time spent on data collection.



# Data Labeling and Cleaning

- **Expert Labeling & DNA Analysis:**
  - Accurate labeling by experts.
  - Morphologically or using DNA analysis.
  - Depending on the project's goal, labeling can range from class to species level.
- **Importance of Clean Data:**
  - Clean data is crucial for training effective machine learning models.
  - Ensures accurate predictions and reliable insights.

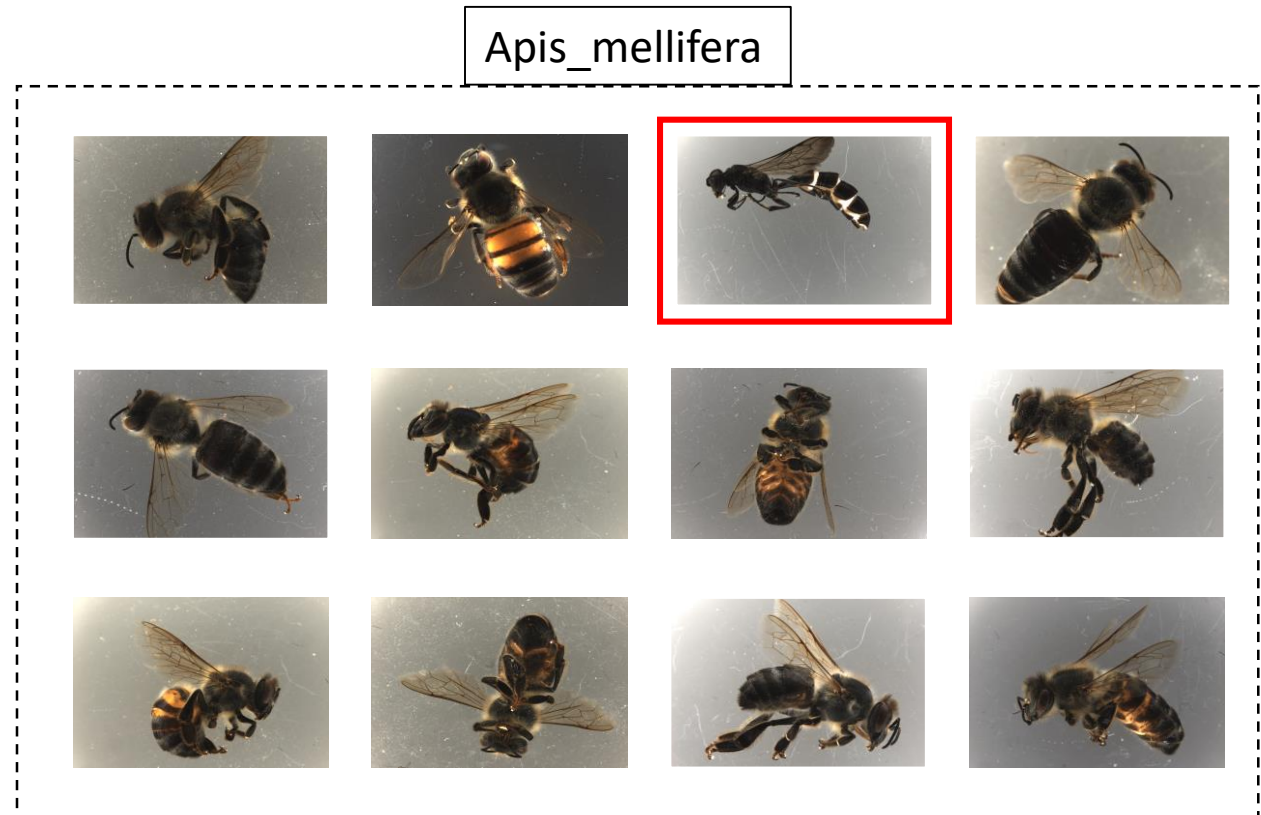




# Data Labeling and Cleaning

## Outlier Detection Model:

- Identify contaminations using unsupervised methods
- Flagging Potential Issues
- Enhancing Dataset Integrity
- Efficiency in Quality Control

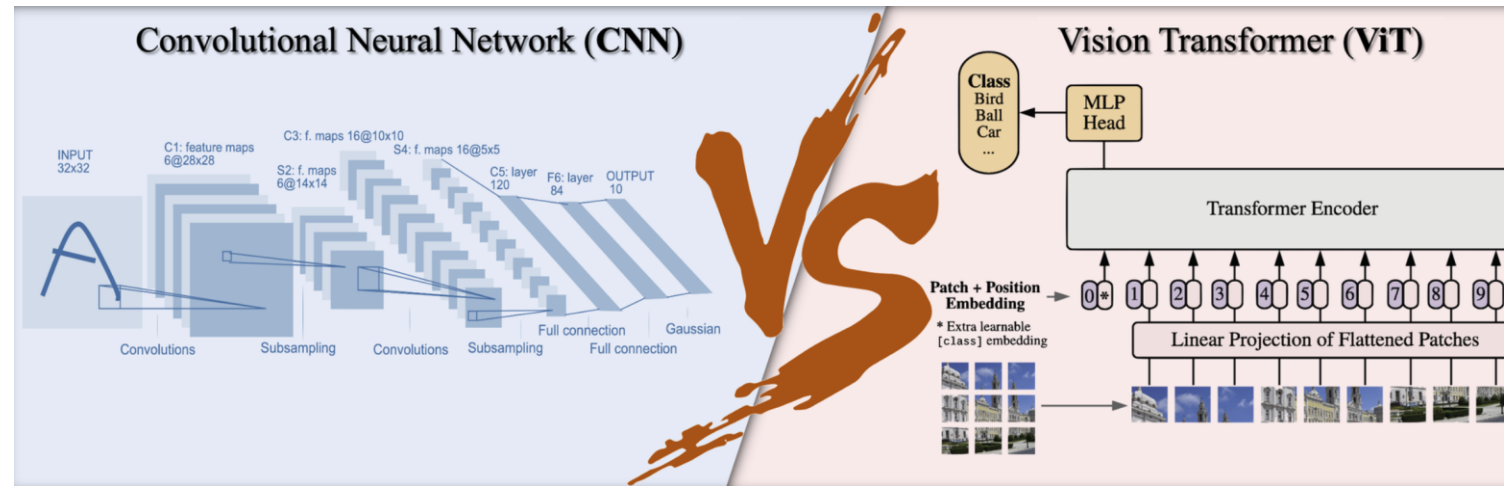




# Machine Learning Models for Species Identification

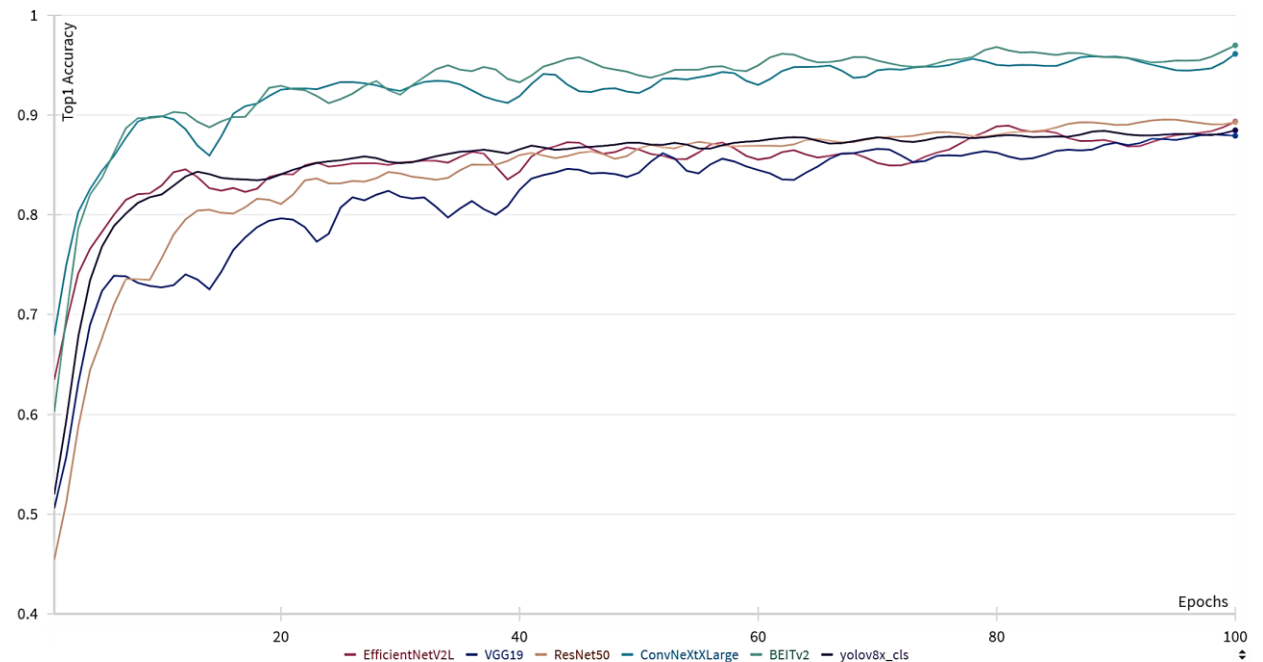
## Core Models:

- Convolutional Neural Networks (CNNs): Highly effective for extracting and processing spatial features in image data.
- Vision Transformers (ViTs): Advanced architecture designed for capturing complex patterns and global relationships in images.



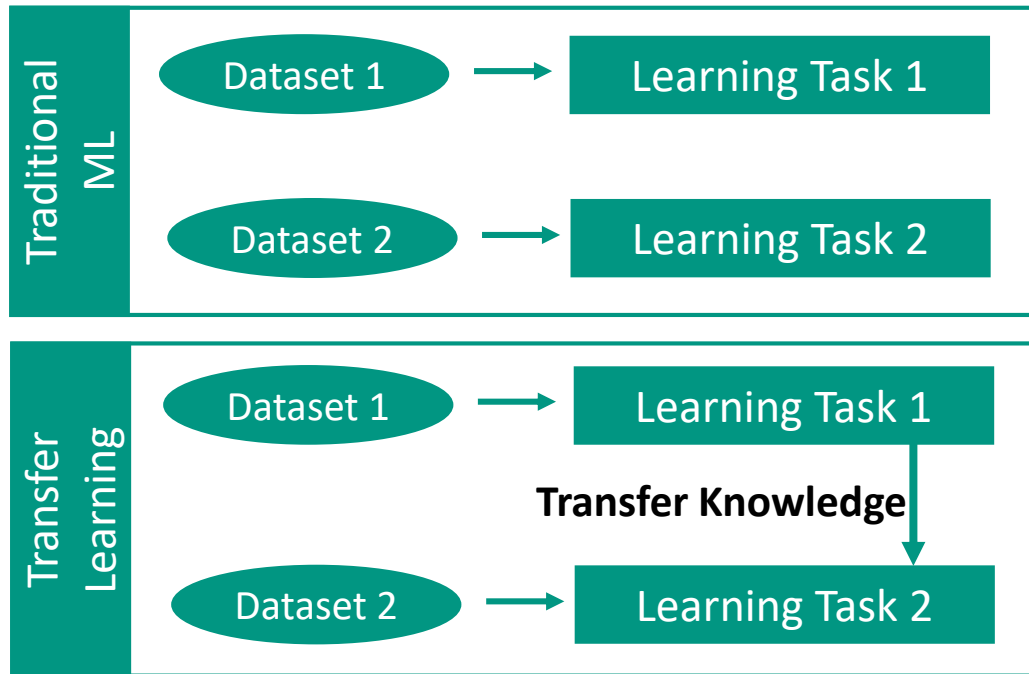
# Model Selection and Evaluation

- Model Benchmarking:
  - Systematic evaluation of multiple models to identify the most suitable one for our dataset.
- Evaluation Metrics:
  - Using metrics such as accuracy, precision, recall, and F1-score and Confusion matrix to rigorously assess model performance and reliability

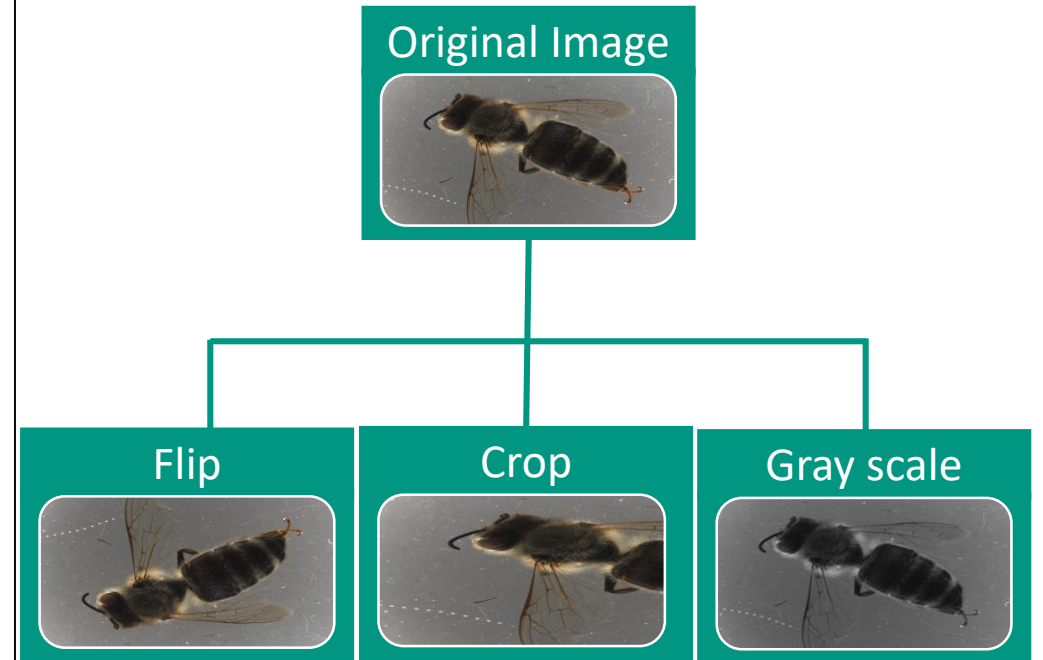


# Advanced Techniques

## Transfer Learning

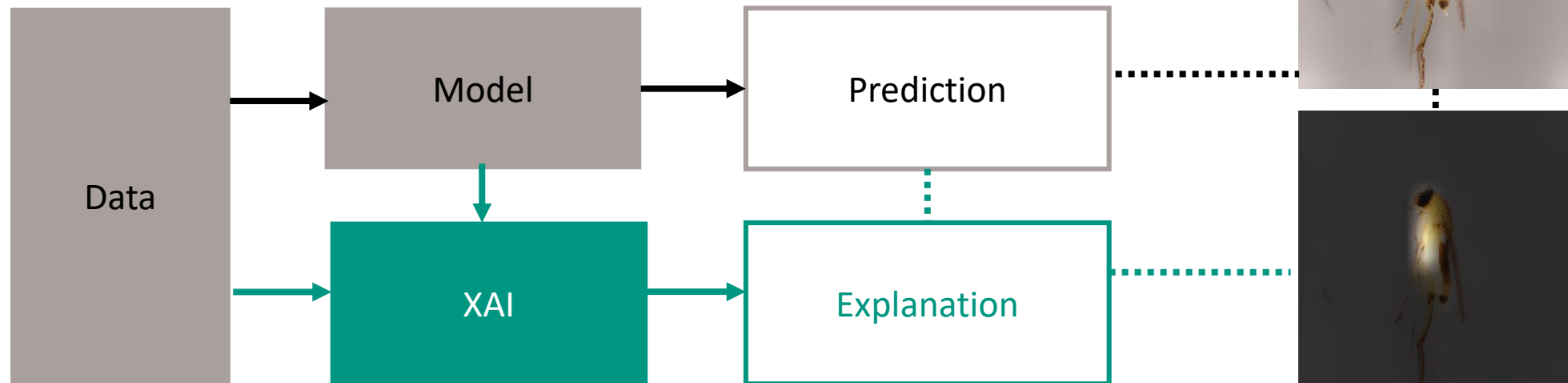


## Data Augmentation



# Explainable AI (XAI)

- Increased trust in AI models.
- Easier debugging and model improvement.
- Enhanced collaboration between AI and domain experts.
- Verification of results with experts.





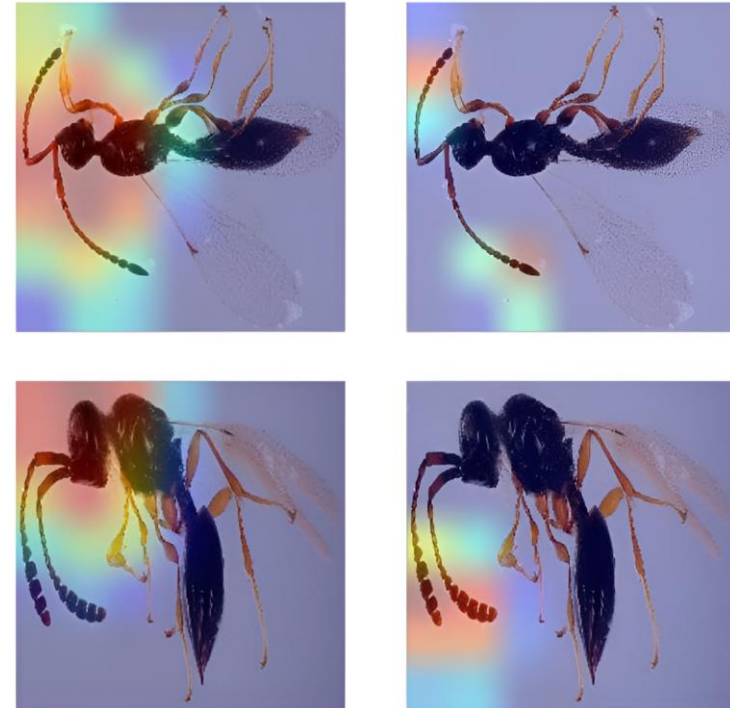
# Image-based recognition of parasitoid wasps using advanced neural networks <sup>[5]</sup>

- Prepared a dataset of 2,257 images.
- Labeled using DNA barcoding and expert morphological analysis.
- Taxonomically identified to the genus and gender levels:
  - 11 diaprid genera.
  - An additional mixed group of mixed 'other Hymenoptera'.



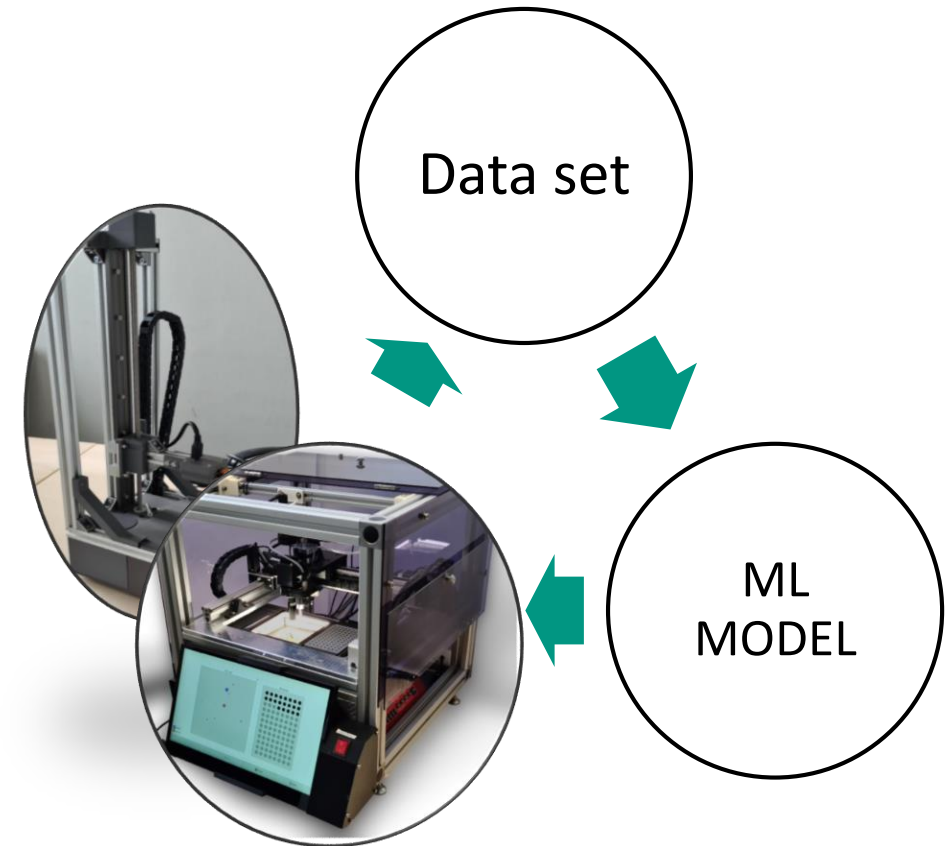
# Image-based recognition of parasitoid wasps using advanced neural networks <sup>[5]</sup>

Architectures	Genus accuracy	Sex accuracy
BEITV2	0.96	0.97
ConvNeXt XLarge	0.94	0.95
YOLOv8	0.89	0.94



# Automation in Specimen Classification and Sorting

- Integration of trained models with DiversityScanner and Entomoscope:
- Benefits:
  - Fast and high-accuracy identification of species.
  - Facilitates the identification of unknown species.
  - Enables scaling of models through active learning.
  - Enhances model performance with iterative expert feedback and retraining.



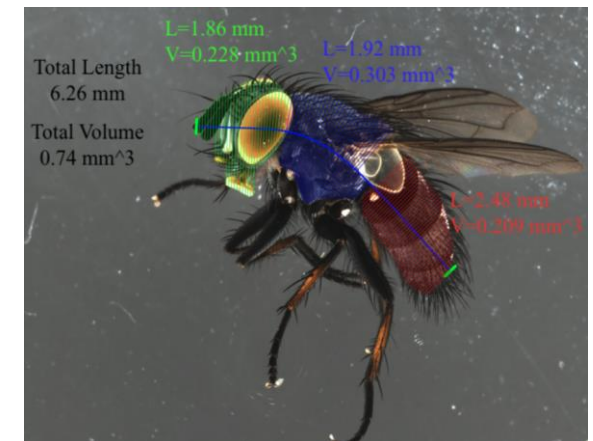
## Future Directions and Impact

- Scaling Up Classification:

- Enhancing model capabilities and training with diverse datasets.
- Goal to identify the most common species captured in a Malaise trap.

- Biomass Estimation from 2D Images:

- Developing new techniques to estimate biomass (length and volume) from 2D images.
- Integrating image analysis with ecological data for comprehensive insights.





Thank you for your attention!



Let's connect !



## References

- [1] Guerrero, M. S., & Cayabyab, B. F. (2024). DNA barcoding of *Locusta migratoria manilensis* (Orthoptera: Acrididae) reveals insights into the species and subspecies differentiation. *Journal of Entomological Science*, 59, 125-132.
- [2] Sharma, R. P., et al. (2023). Exploring the significance of insects in ecosystems: A comprehensive examination of entomological studies. *International Journal of Environment and Climate Change*. <https://doi.org/...>
- [3] Wührl, L., Pylatiuk, C., Giersch, M., Lapp, F., Balke, M., Schmidt, S., Cerretti, P., & Meier, R. (2022). DiversityScanner: Robotic handling of small invertebrates with machine learning methods. *Molecular Ecology Resources*, 22(4), 1626-1638. <https://doi.org/10.1111/1755-0998.13567>

## References

[4] Wührl, L., Rettenberger, L., Meier, R., Hartop, E., Graf, J., & Pylatiuk, C. (2024). Entomoscope: An open-source photomicroscope for biodiversity discovery. *IEEE Access*, 12, 11785-11794.

<https://doi.org/10.1109/ACCESS.2024.3355272>

[5] Shirali, H., Hübner, J., Both, R., Raupach, M., Reischl, M., Schmidt, S., & Pylatiuk, C. (2024). Image-based recognition of parasitoid wasps using advanced neural networks. *Invertebrate Systematics*, 38, IS24011.

[6] Miroslav Valan, Karoly Makonyi, Atsuto Maki, Dominik Vondráček, Fredrik Ronquist, Automated Taxonomic Identification of Insects with Expert-Level Accuracy Using Effective Feature Transfer from Convolutional Networks, *Systematic Biology*, Volume 68, Issue 6, November 2019, Pages 876–895, <https://doi.org/10.1093/sysbio/syz014>