

# AUTOMATING IDENTIFICATION OF VECTOR INSECTS

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# Chagas Disease

- The disease is caused by *Trypanosoma cruzi*
- Taxonomic impediment regarding vector species
- Most of the 150+ species of the family Triatominae feed on vertebrate blood
- Species identification is crucial because different species have different ecologies



<https://goo.gl/images/m7rfq8>



# VIRTUAL VECTOR LAB

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Spencer Art Museum, ITTC, Biodiversity Institute @ KU

Universidade de Brasília

Instituto Nacional de Salud Pública de Mexico



# Automated identification of insect vectors of Chagas disease in Brazil and Mexico: the Virtual Vector Lab

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## ABSTRACT

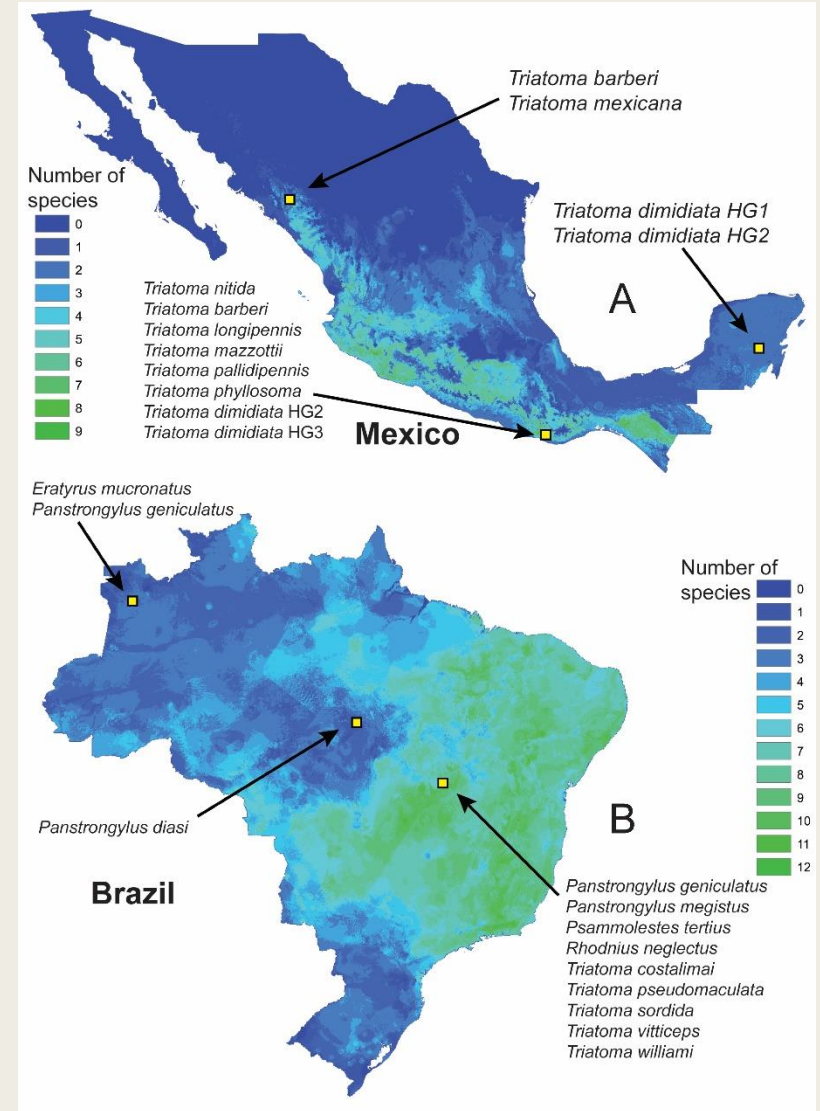
Identification of arthropods important in disease transmission is a crucial, yet difficult, task that can demand considerable training and experience. An important case in point is that of the 150+ species of Triatominae, vectors of *Trypanosoma cruzi*, causative agent of Chagas disease across the Americas. We present a fully automated system that is able to identify triatomine bugs from Mexico and Brazil with an accuracy consistently above 80%, and with considerable potential for further improvement. The system processes digital photographs from a photo apparatus into landmarks, and uses ratios of measurements among those landmarks, as well as (in a preliminary exploration) two measurements that approximate aspects of coloration, as the basis for classification. This project has thus produced a working prototype that achieves reasonably robust correct identification rates, although many more developments can and will be added, and—more broadly—the project illustrates the value of multidisciplinary collaborations in resolving difficult and complex challenges.

**Subjects** Entomology, Computational Science

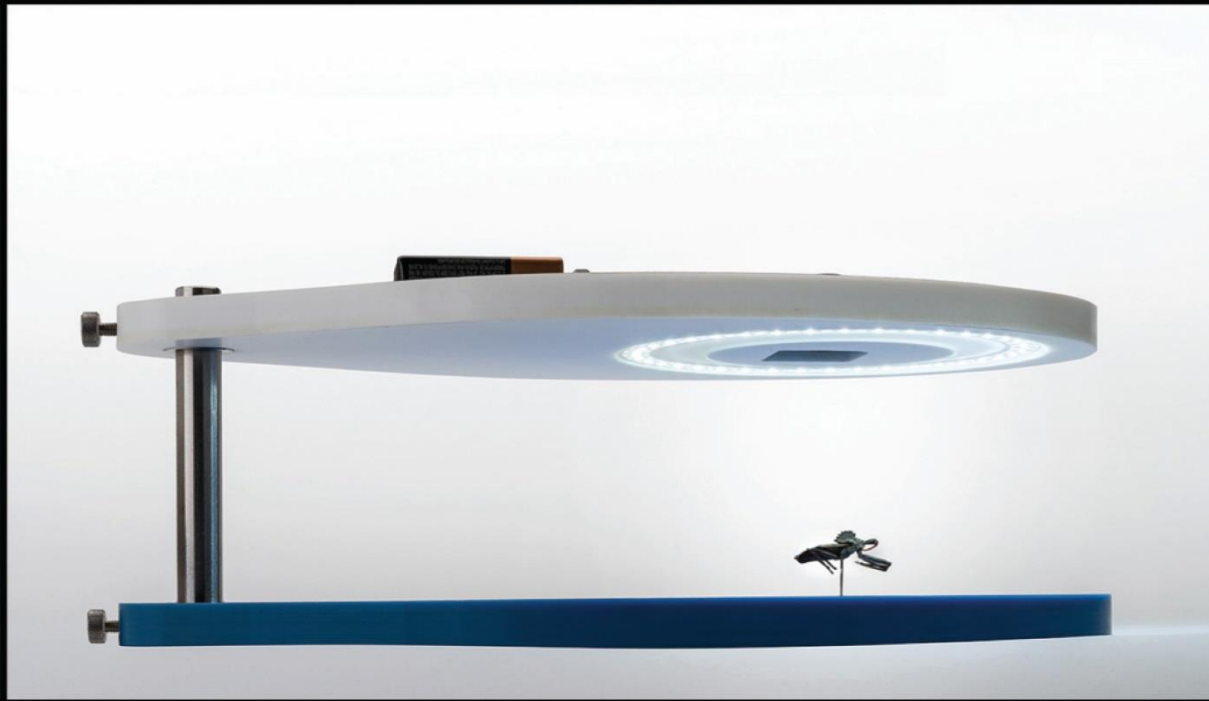
**Keywords** Identification, Chagas disease, Triatominae, Automation, Primary occurrence data

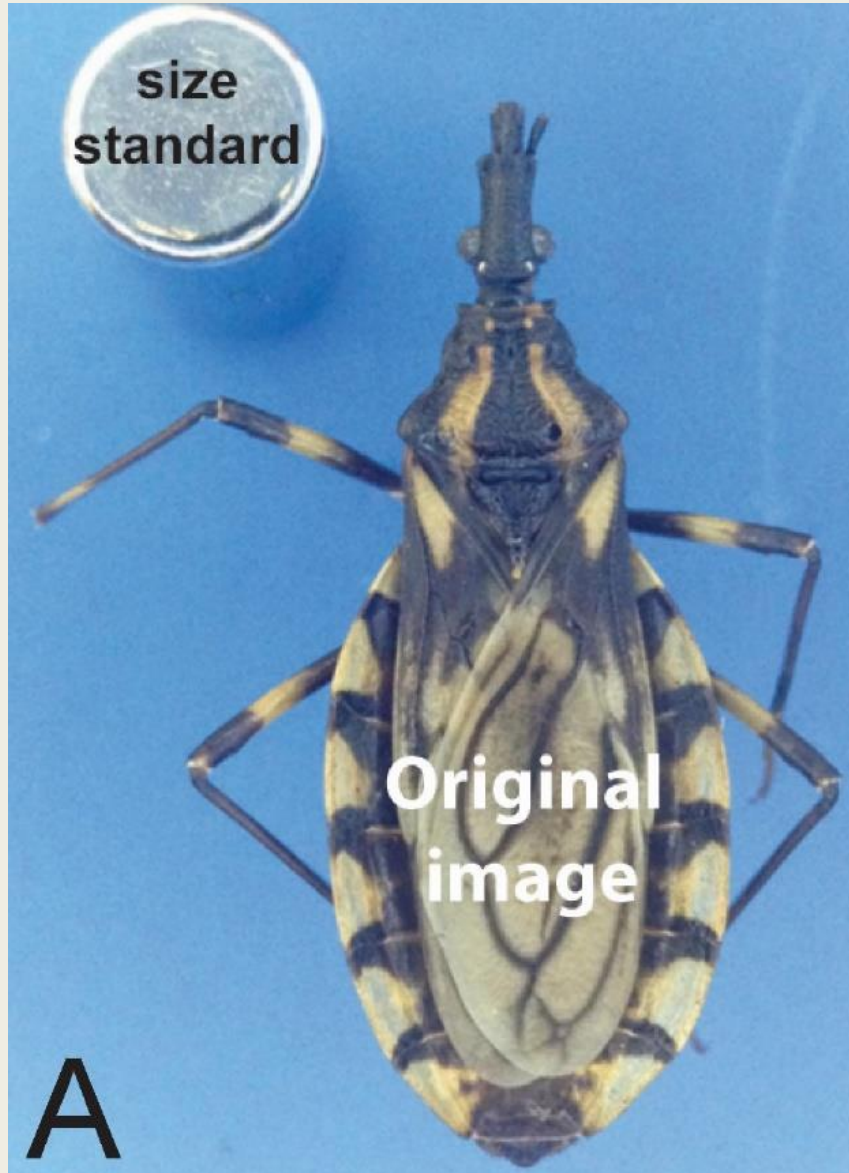
Submitted 26 September 2016  
Accepted 28 January 2017  
Published 18 April 2017

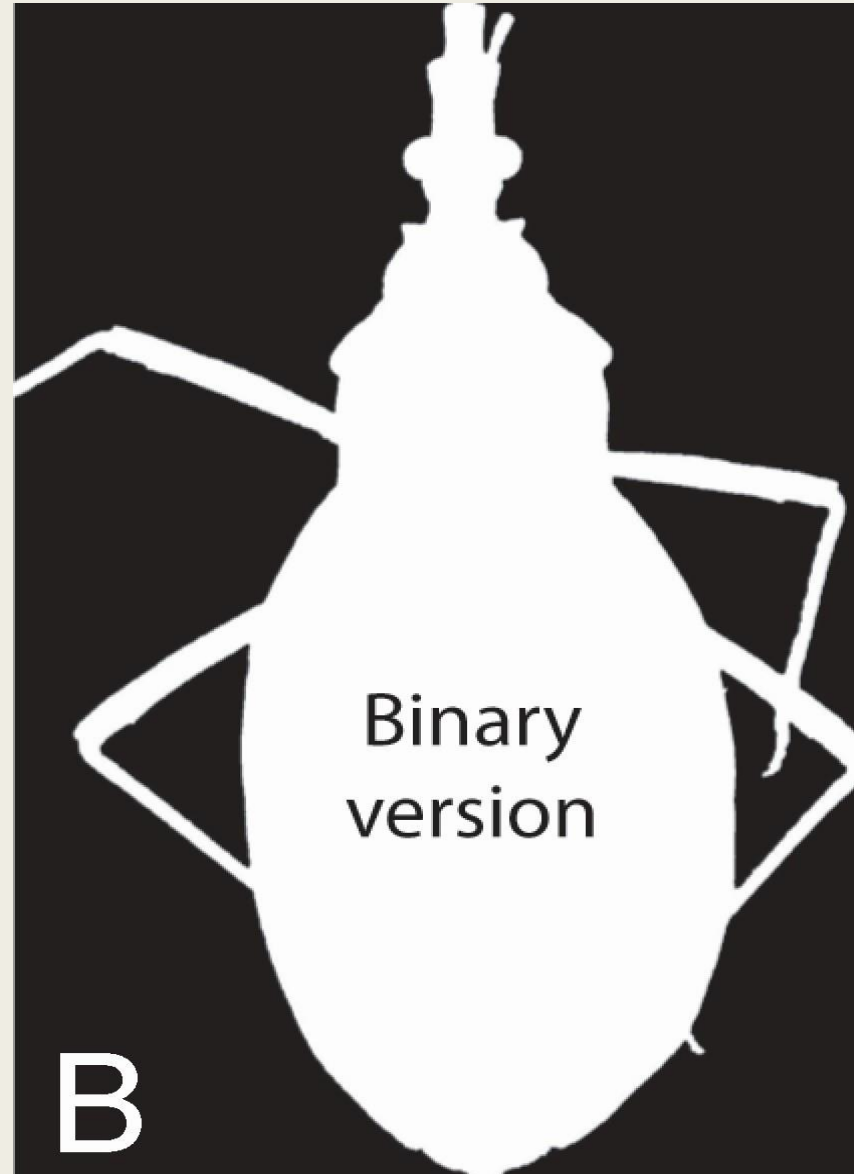
Corresponding author  
A Townsend Peterson, town@ku.edu





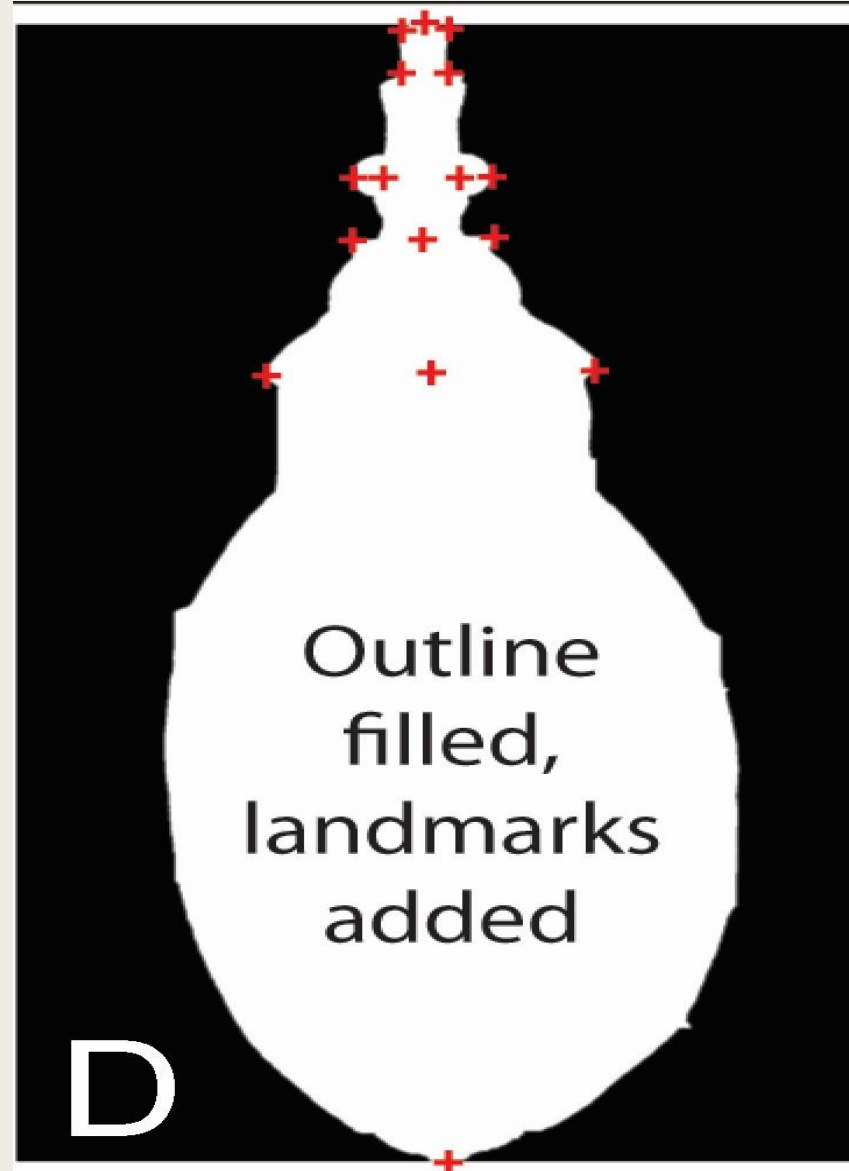


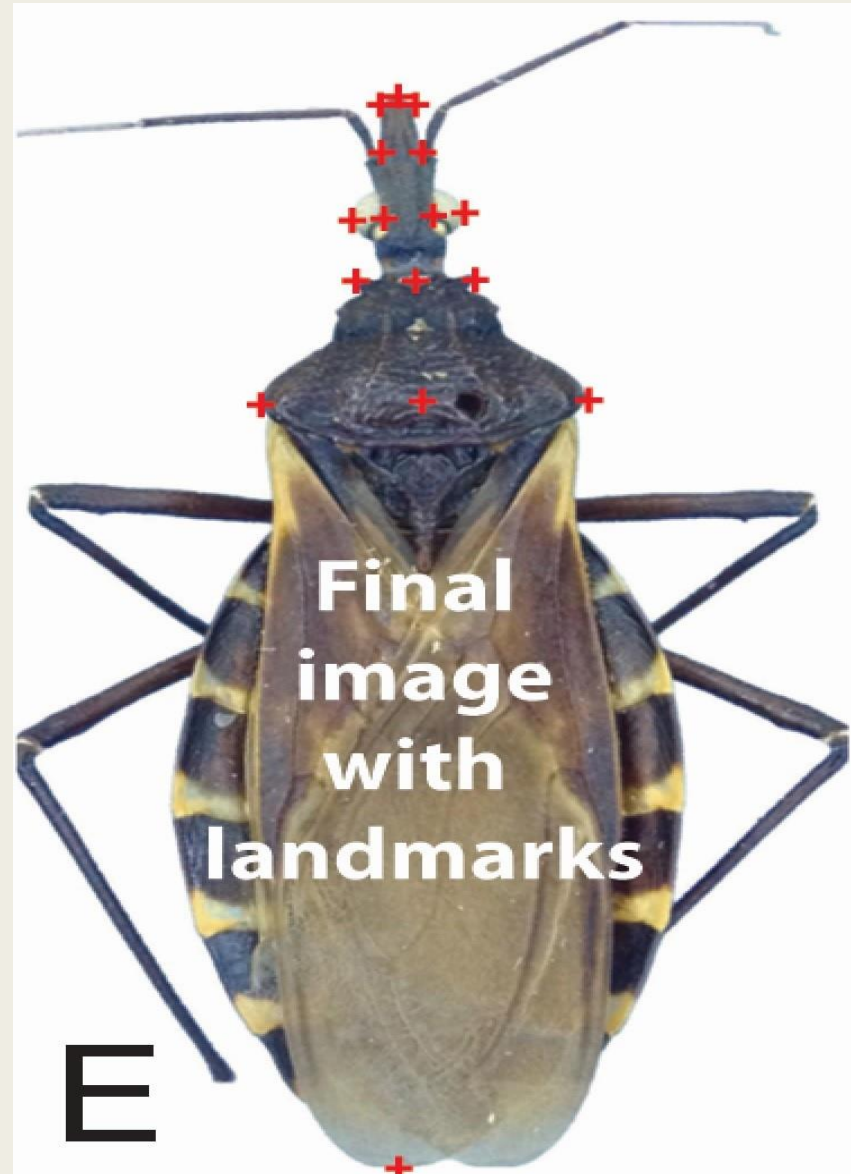






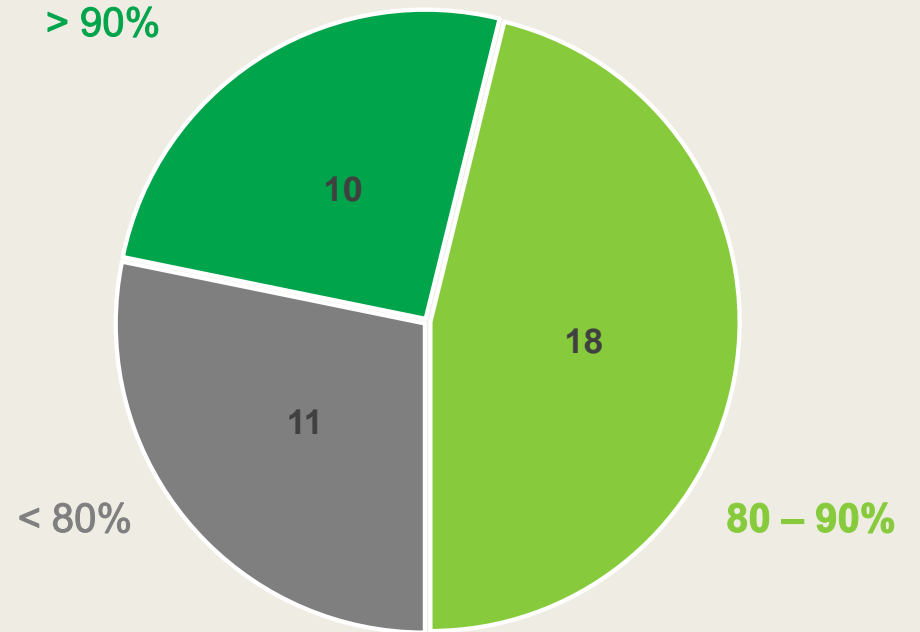






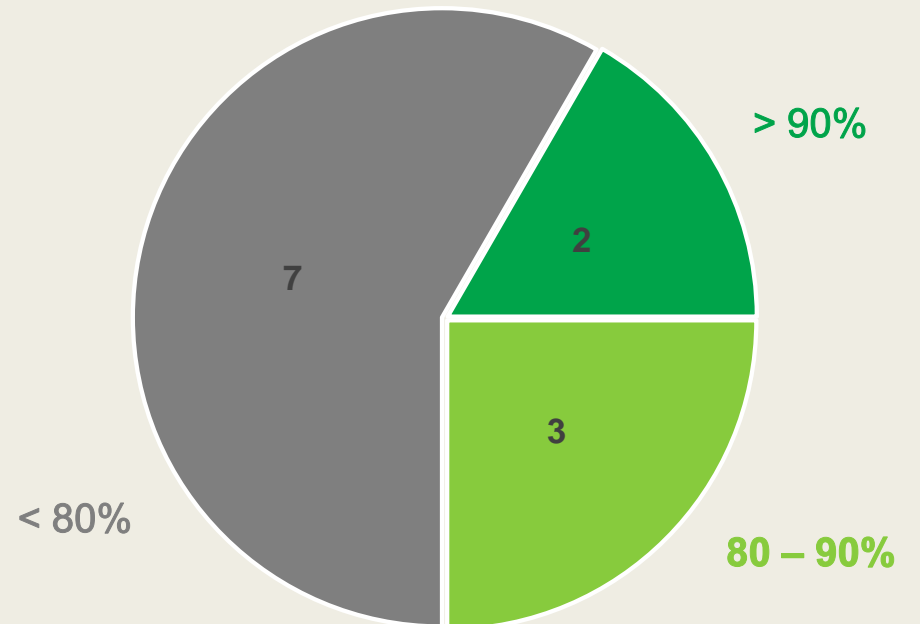
# Previous Results

39 Brazilian Species → Overall Rate: 83.3%



Results achieved using diverse statistical classifiers and simple neural networks

12 Mexican Species → Overall Rate: 74.9%





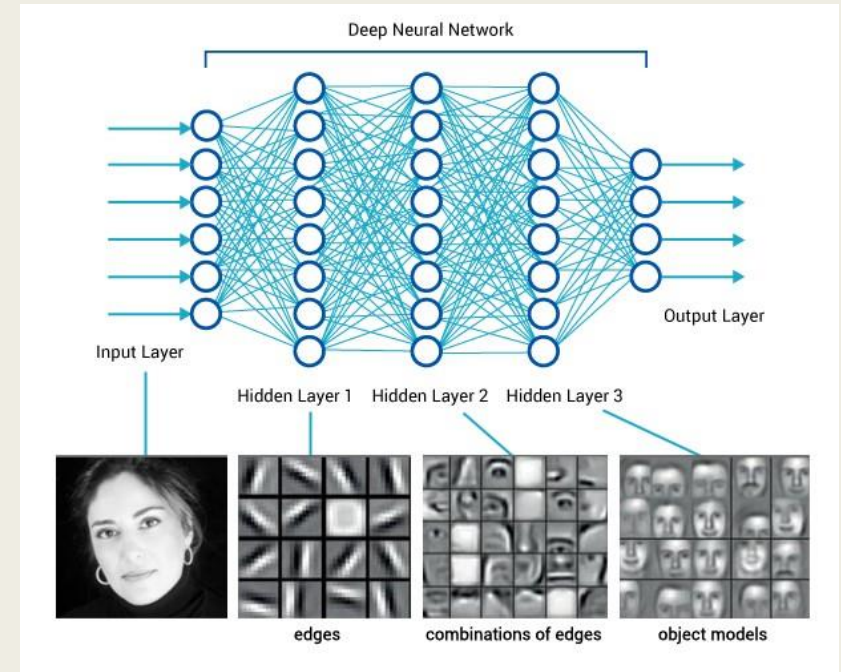
TensorFlow



# Google Brain Team

Make machines intelligent. Improve people's lives.

- Open source software
- Released in November 2015
- Provides a platform for deep learning technique



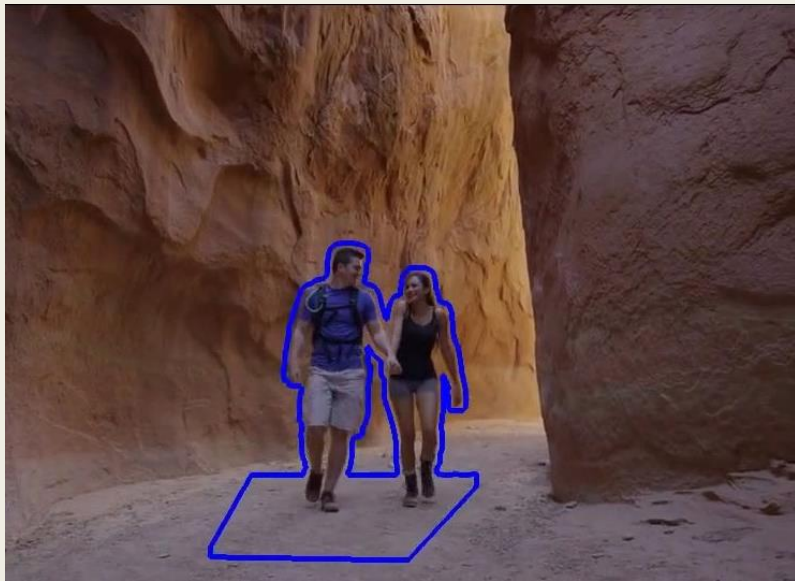
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**Before**

**After**



<https://goo.gl/images/aAWKjE>





# VVL & TENSORFLOW

## Previous Results

12 Mexican Species → Overall Rate: 74.9%

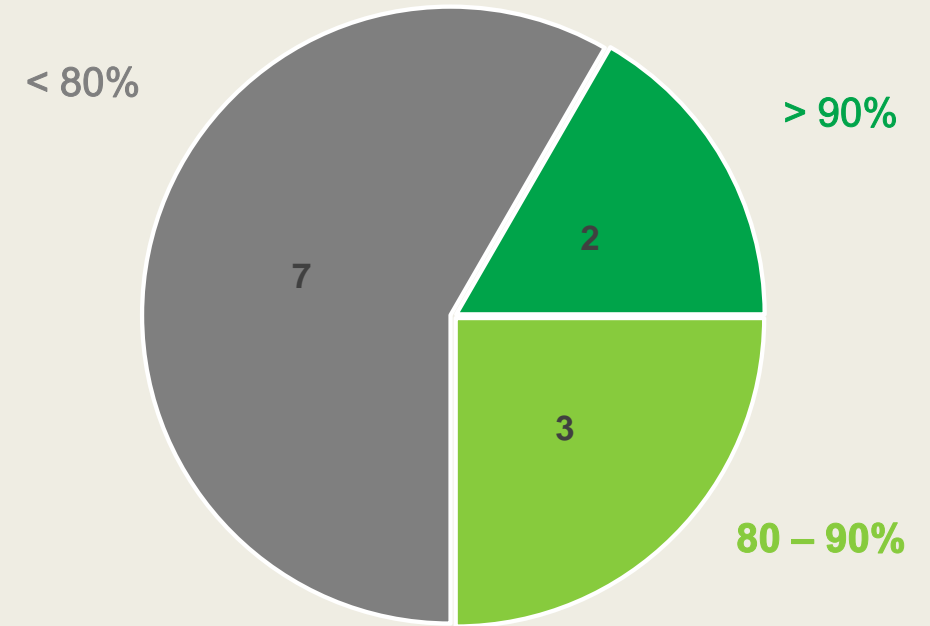
## TensorFlow

12 Mexican Species → Overall Rate: 83.0%



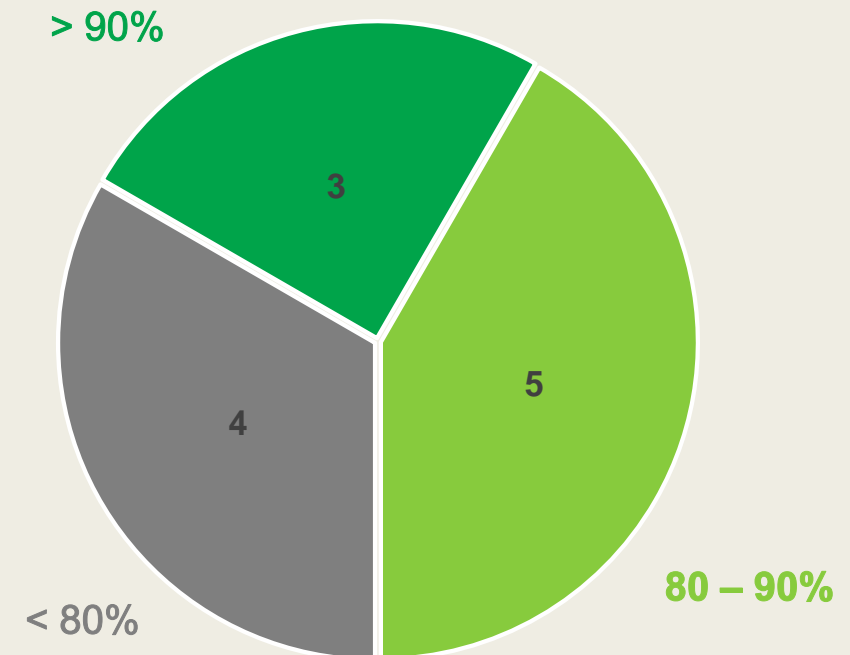
## Previous Results

12 Mexican Species → Overall Rate: **74.9%**



## TensorFlow

12 Mexican Species → Overall Rate: **83.0%**



Translation: an 8.1% improvement in overall correct ID rate over our previous VVL results

## Previous Results

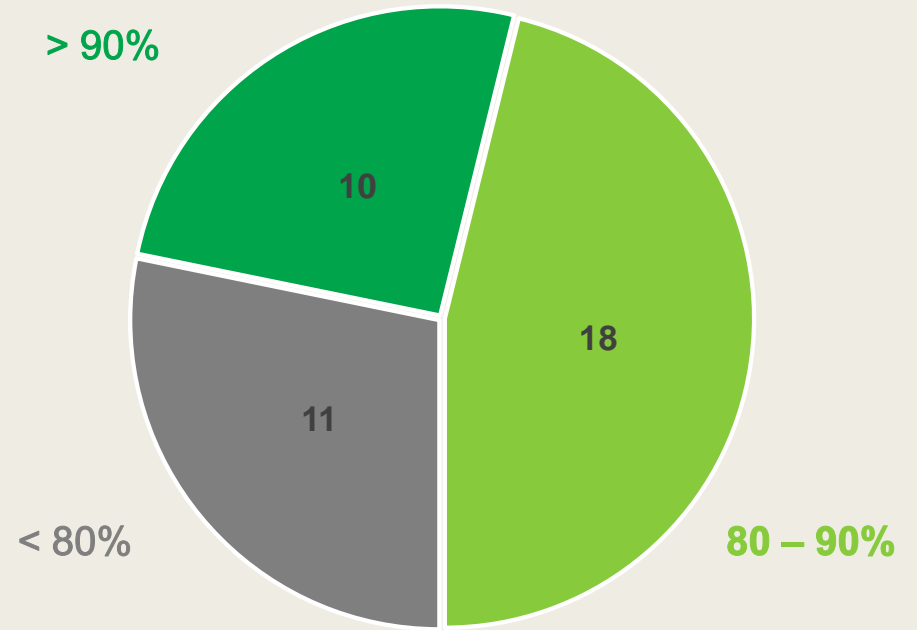
39 Brazilian Species → Overall Rate: **83.3%**

## TensorFlow

39 Brazilian Species → Overall Rate: **86.7%**

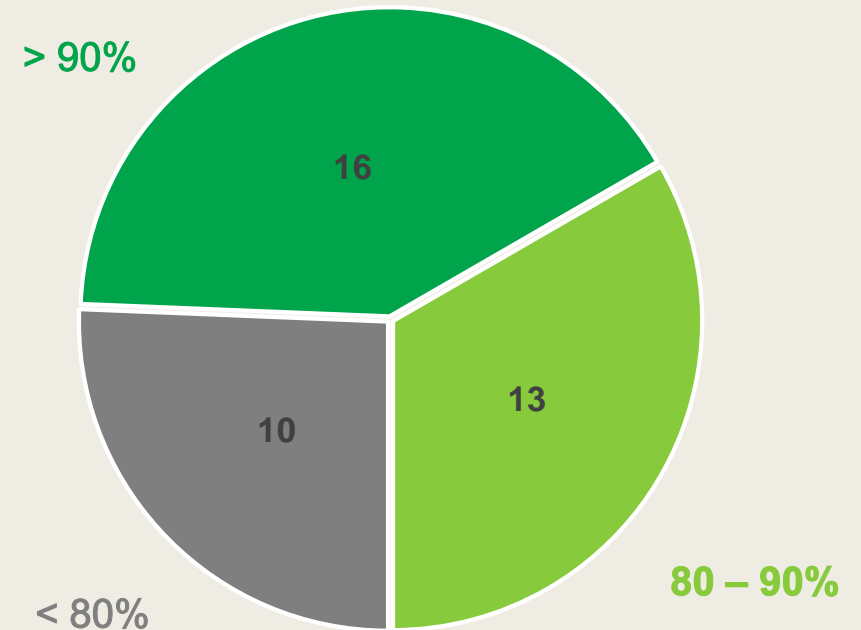
## Previous Results

39 Brazilian Species → Overall Rate: **83.3%**

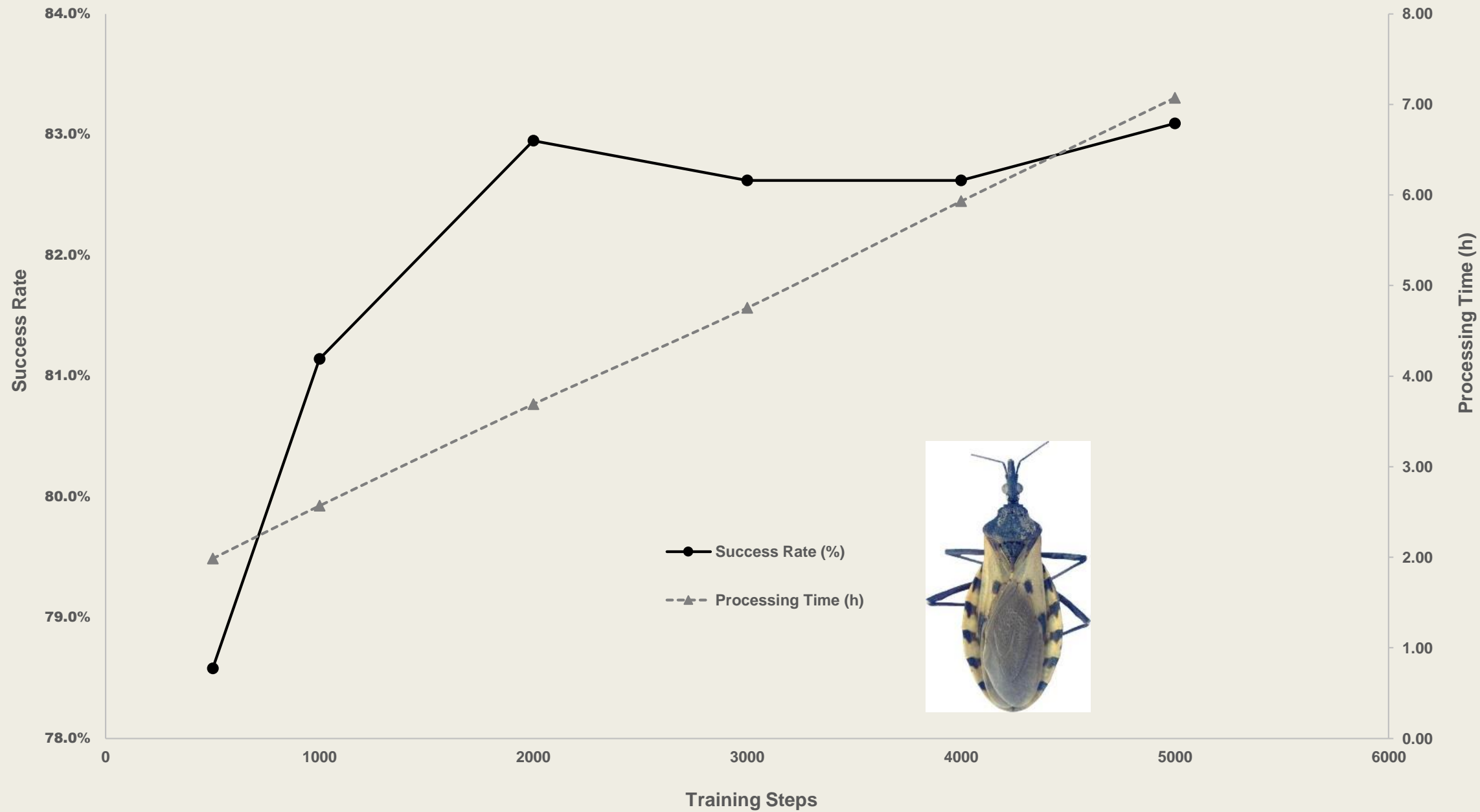


## TensorFlow

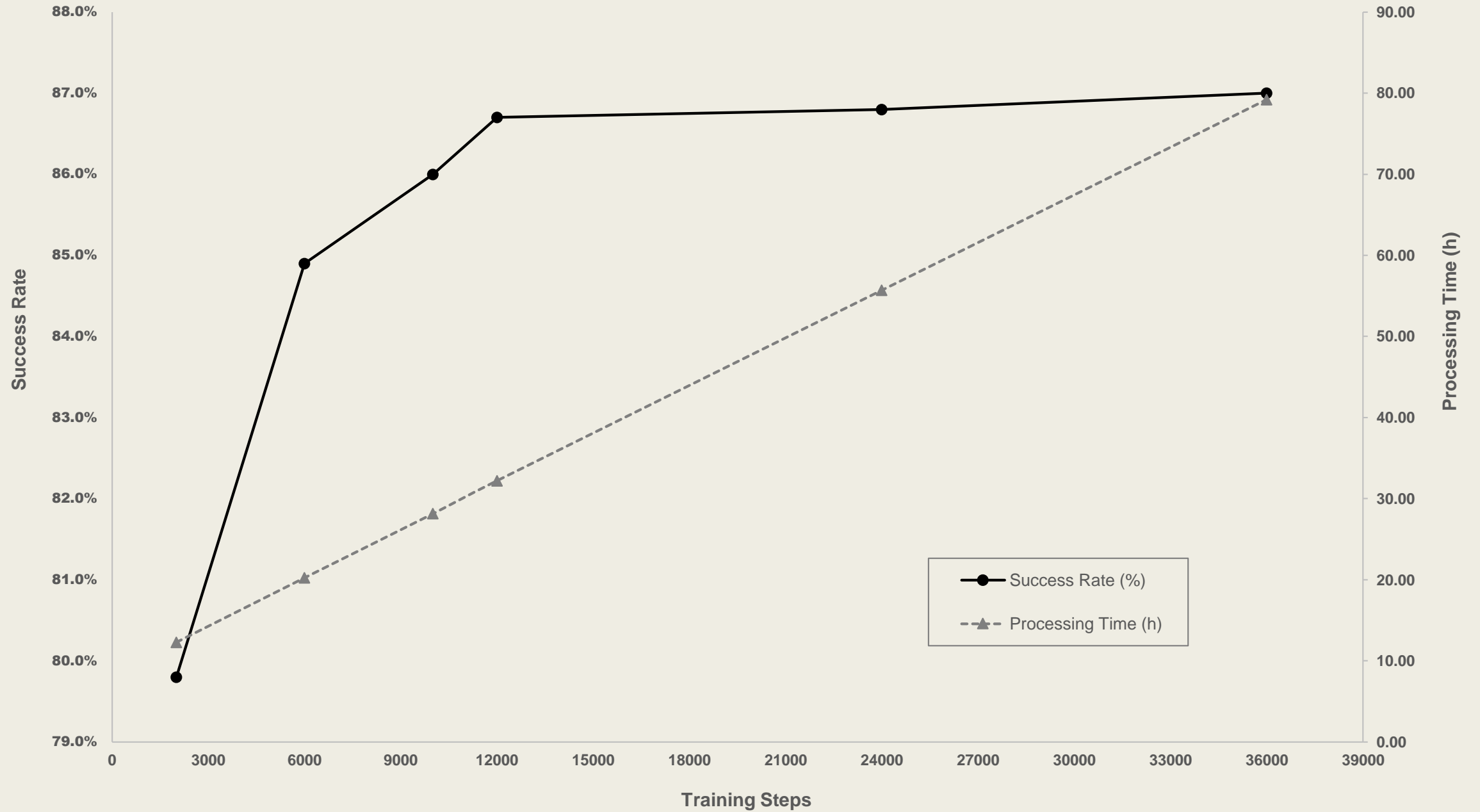
39 Brazilian Species → Overall Rate: **86.7%**

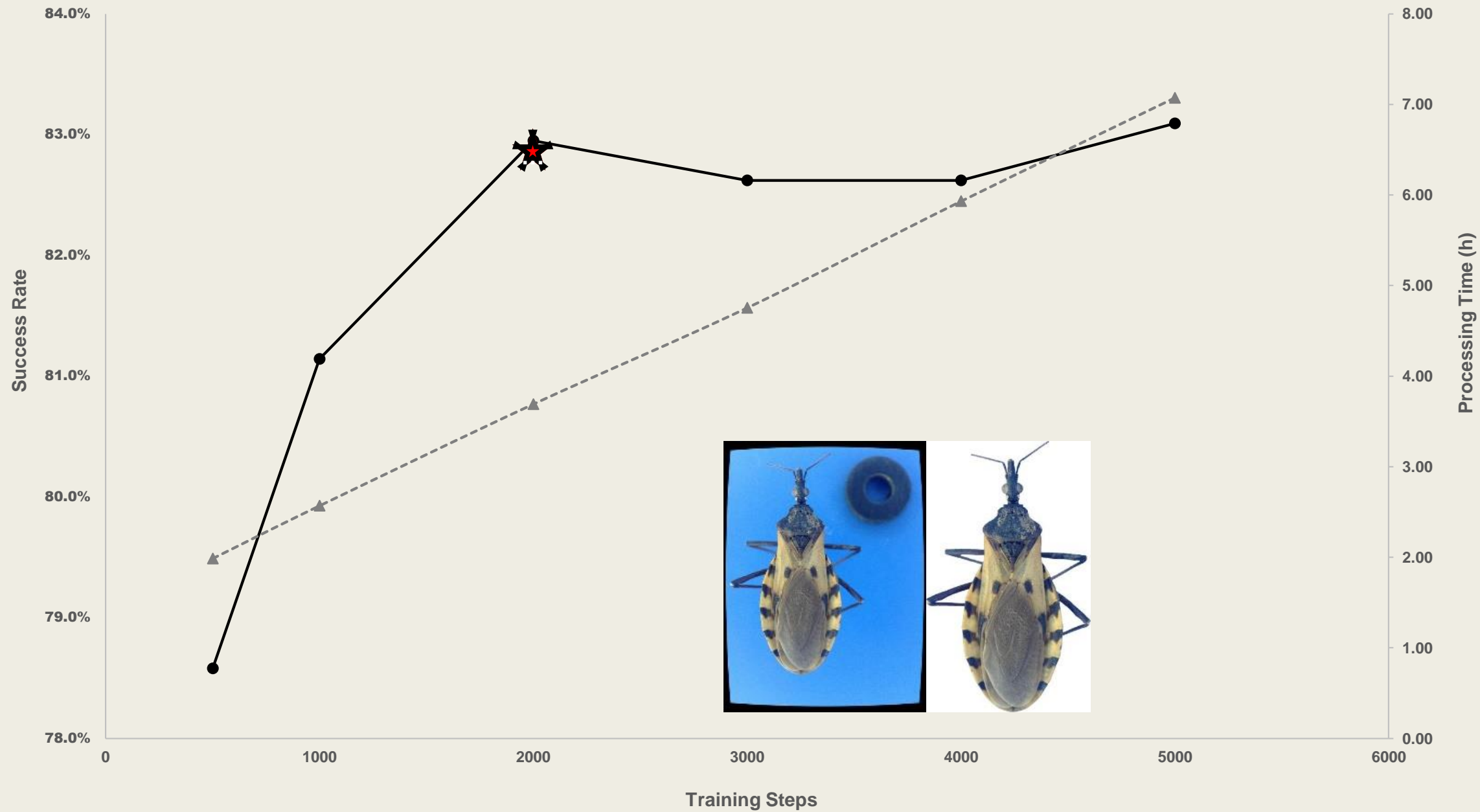


Translation: a 3.4% improvement in overall correct ID rate over our previous VVL results











<https://goo.gl/images/Bno76q>

# Conclusion

- Improved overall classification of triatomine species markedly over our previous results, particularly the total number of species with ID rate of above 80%
- Opened the possibility of using raw images in classification, without a need for pre-processing
- Confidence rate information provided by TensorFlow can inform us about when identification is uncertain
- Technical support from Google will update and improve versions, and offer web-based implementations

# Thank you!

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