



The Nature Conservancy: Advancing Computer Vision for Invasive Species Detection



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The Nature Conservancy & Key Reserves

The Nature Conservancy (TNC) is a wide-reaching **environmental conservation organization** whose primary mission is to conserve “the lands and waters on which all life depends”



Santa Cruz Island is a TNC-managed island off the coast of California and leading example of successful island restoration. Through 40 years of work, the island is **completely free** of non-native animals and is home to over 1000 species, **12** of which are unique to the island.

The Jack and Laura Dangermond Preserve (JLDP) is a vast and diverse TNC-managed ecosystem in Southern California.

The Challenge

Invasive-species driven extinctions of birds, amphibians, mammals, and reptiles on islands make up **75% of all globally**.

Remedial actions and long-term effects of invasive species can be ecologically and economically devastating, costing some **\$1 trillion** in the last 50 years [1].

The low data availability for invasive species, the similarity between classes of species and poor transferability of models between biomes make for a **hard ML problem**.

The Solution

Build invasive species detectors using an established network of camera traps. In doing so, push forward the SOTA in camera trap computer vision.

Main Contribution: Proof of concept of a camera trap anomaly detection algorithm that learns representations of native species to identify non-native species in new images

Contribution 2: Learnings for future applications of CV to camera trap imagery

Anomaly Detection Algorithm

Two key elements in the algorithm:

SimCLR: Learns rich features from images by maximizing agreement between differently augmented views of the same data example via a contrastive loss in the latent space [2]

OneClassSVM/DBSCAN: Outlier detection comparing SimCLR representation of a new image with representation of native images
The workflow used in our proof-of-concept is below:

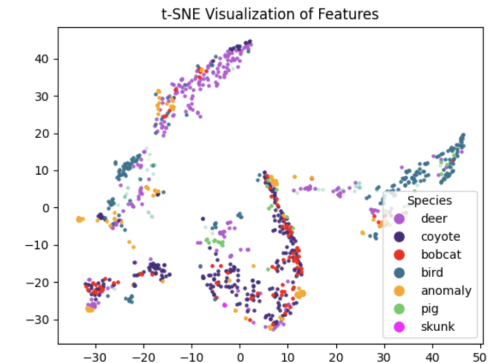
```
Data: 4.5K JLDP images,  $X_{tr}, y_{tr}, X_{te}, y_{te}$ 
 $f \leftarrow \text{ResNet50}$ ;
# Remove anomalies (bobcats) in train
 $X_{tr}, y_{tr} \leftarrow X_{tr} - X_{bobcat}, y_{tr} - y_{bobcat}$ ;
# Train for features & extractor
 $h_{tr}, f_{tr} = \text{SimCLR}(X_{tr}, y_{tr}, f)$ ;
if species_specific then
    for  $y_{tr}^i \in y_{tr}$  do
        # Train one-class SVM
         $1\text{CSVM}_i = 1\text{CSVM}(h_{tr}^i)$ ;
    end
end
# To test, run Yolo to crop images
 $X_{crop}, y_{pred} = \text{YOLO}(X_{te})$ ;
 $h_{pred} = f_{tr}(X_{crop})$ ; # Get SimCLR features
if species_specific then
    for  $y_{pred}^i \in y_{pred}$  do
        | Result:  $1\text{CSVM}_i(h_{pred}^i)$ 
    end
else
    | Result:  $\text{DBSCAN}(h_{pred})$ 
end
```

Notes:

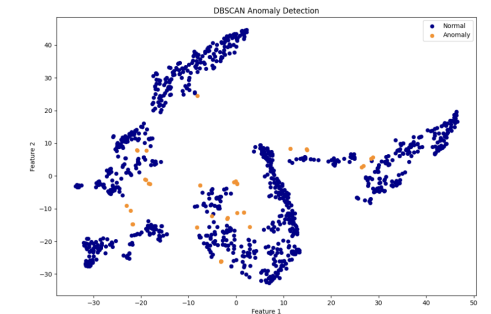
1. We created new anomalous images by swapping crops of bobcats onto different backgrounds
2. We ran both global anomaly detection and species specific anomaly detection
3. t-SNE was used for visualizations

Initial results

Result 1: At species level, model can identify truly anomalous data, but struggles when representations are similar (e.g. bobcat-coyote)



Result 2: Model can identify species not present in the train data as anomalous (e.g. skunks)



Further work

- New models, e.g. isolation forest or GMM's for outlier detection
- Higher quality generated anomalies with GAN's
- Consider low-confidence YOLO predictions as potential anomalies

References

1. R. D. Zenni, F. Essl, E. García-Berthou, and S. M. McDermott, 'The economic costs of biological invasions around the world', *NeoBiota*, vol. 67, pp. 1–9, Jul. 2021, doi: 10.3897/neobiota.67.69971.
2. T. Chen, S. Kornblith, M. Norouzi, and G. Hinton, 'A Simple Framework for Contrastive Learning of Visual Representations'. *arXiv*, Jun. 30, 2020. doi: 10.48550/arXiv.2002.05709.