

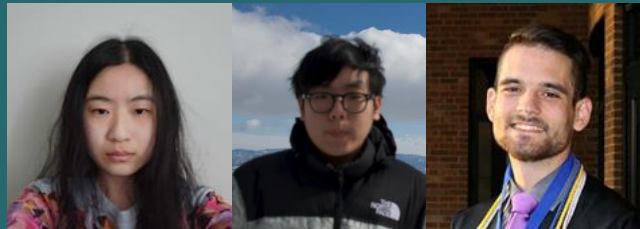


# Anomaly Detection: Hybrid Butterflies

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and

The Imageomics ML Challenge Team



Special thanks to Lisa Wu, Ziheng Zhang, David Carlyn

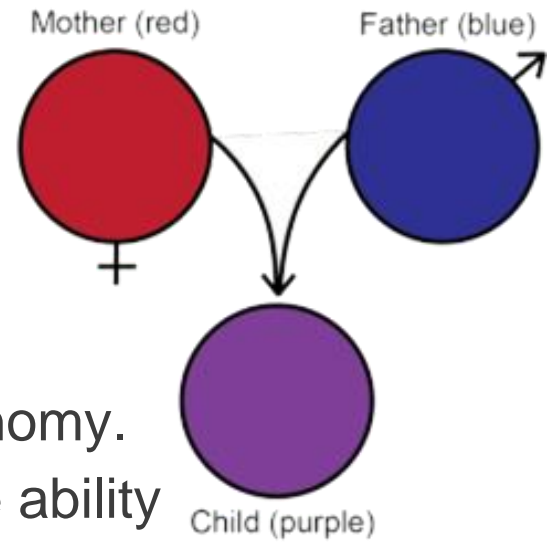


# Hybrid Detection

A brief history

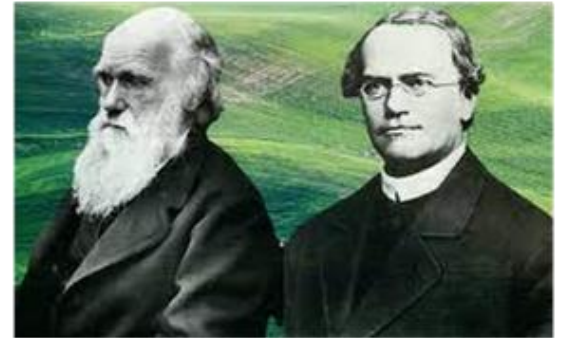
# Hybrid Detection

- Researchers have sought a means to detect hybrids since the creation of the field of taxonomy.
- Detecting hybrids would give taxonomists the ability to determine what constitutes a true *species* or *subspecies*.
- The question is **how?**
  - *How* do we recognize a hybrid?
  - What does a hybrid look like?



# Hybrid Detection: History

- Darwin first posed this question of “What does a hybrid look like?”
- Mendel answered with his pea plant experiment.



# Hybrid Detection: History

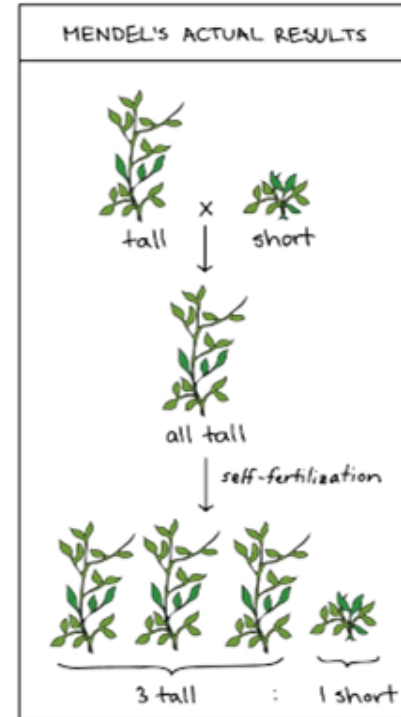
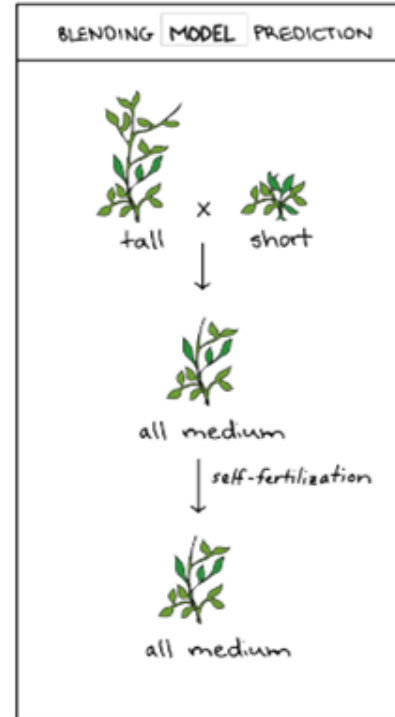
## Mendel's Hypothesis:

### Blending Inheritance

- Inheritance of traits is **continuous**.

## Mendel's Results:

Inheritance is often **discrete**.



# Hybrid Detection: Butterflies

- Consider these two species:
- Hybridization may lead to a variety of resulting patterns.
- There are several [dominant] genes that control color pattern on wings.
  - Ex: red on hindwings is a dominant trait.
- Dominance: hybrids may look like one parent.
- In practice, identifying hybrids requires knowledge of their parent species/subspecies.



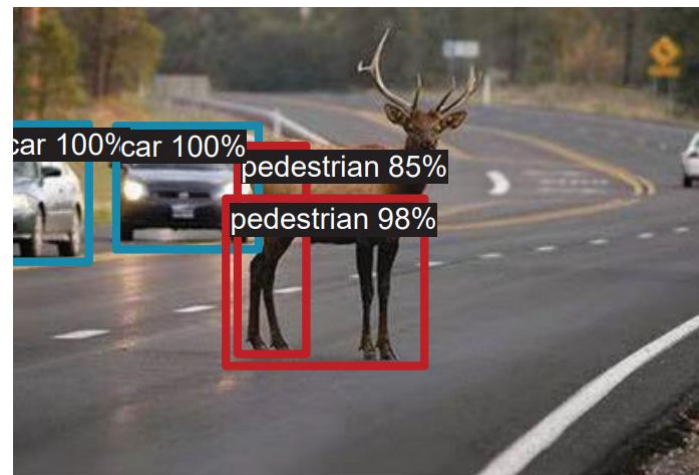


# Anomaly Detection

A brief history

# Anomaly Detection: History

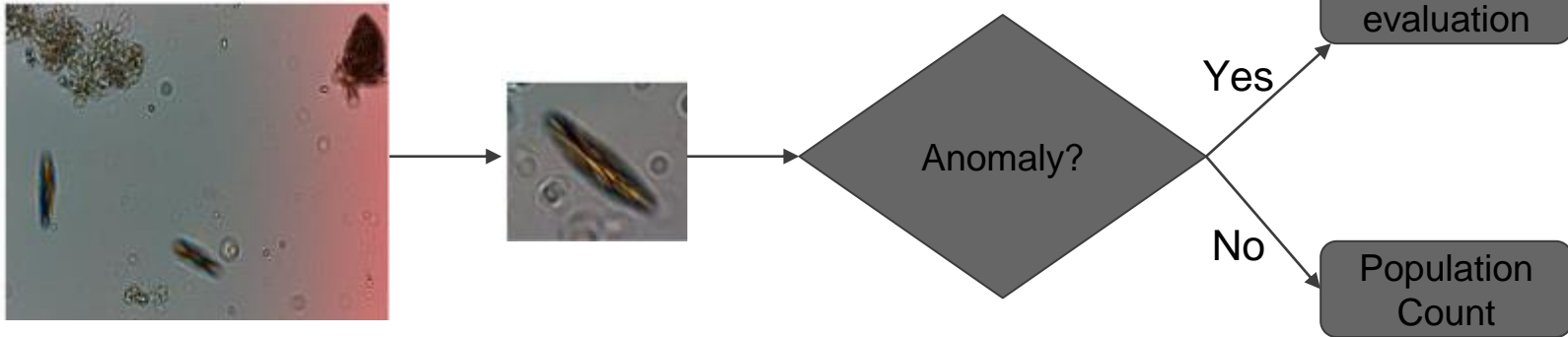
- Early topics include banking.
  - Detecting fraudulent or irregular spending or requests.
- In Machine Learning (ML), questions on classification:
  - Is the object a new one that the classifier has not seen?
- In Computer Vision (CV), questions for autonomous vehicles:
  - Is that a pedestrian or a deer that just ran into the road?





# Anomaly Detection: History

- In Biology, questions on:
  - Gene function identification [1]
    - What phenotype anomaly resulted from a gene knockout?
  - Ecosystem health monitoring [2]
    - Tracking plankton population.



[1] Ito, E. et al. (2022). Phenotype Anomaly Detection for Biological Dynamics Data Using a Deep Generative Model. In: Pimenidis, E., Angelov, P., Jayne, C., Papaleonidas, A., Aydin, M. (eds) Artificial Neural Networks and Machine Learning – ICANN 2022. ICANN 2022. Lecture Notes in Computer Science, vol 13530. Springer, Cham. [https://doi.org/10.1007/978-3-031-15931-2\\_36](https://doi.org/10.1007/978-3-031-15931-2_36)

[2] Pastore, V.P., Zimmerman, T.G., Biswas, S.K. et al. Annotation-free learning of plankton for classification and anomaly detection. Sci Rep 10, 12142 (2020). <https://doi.org/10.1038/s41598-020-68662-3>



# Our Challenge

How **you** can contribute to answering this important biological question

## Hybrid

Species A subspecies I



Species A subspecies II



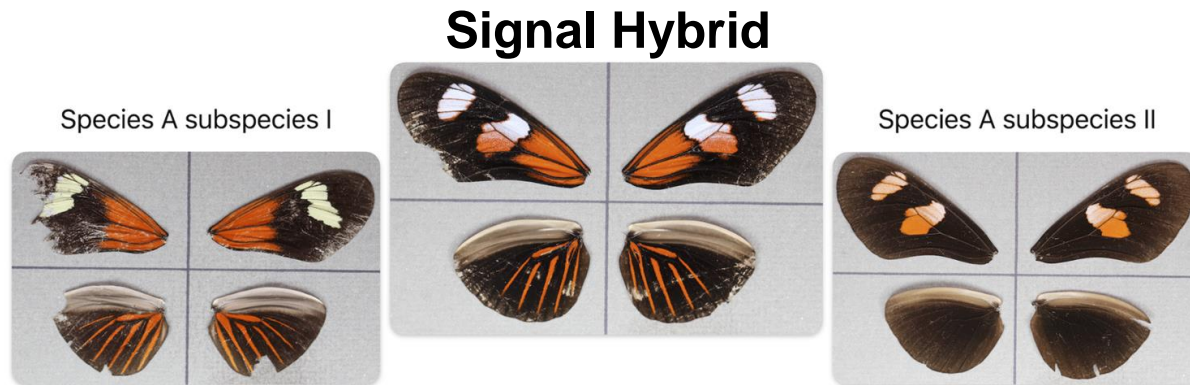
Images are from Zenodo  
records [2764332](#),  
[2088062](#), [2672849](#), and  
[2604621](#), considered  
images are from Zenodo  
records [2714636](#) and  
[252608](#).  
Hybrid graphic generated  
using Canva Magic Media  
CC-BY 4.0, annually edited.



Training Data

# Our Challenge: Training Data

- ~2200 images of Species A:
  - Multiple *subspecies*.
  - Selected signal hybrids of two *subspecies*.



# Our Challenge: Dev & Test Data

- Includes:
  - All Species A subspecies.
  - Signal hybrids from training data.
- Further introduces:
  - Other Species A hybrids (non-signal).
  - Species B: Mimics of Species A signal hybrid parents (& their hybrids).
- The numbers:
  - Validation Data (Dev): ~1100 images
  - Test Data: ~2200 images

# The Challenge: Find the Hybrids

- Among Species A & B, can your algorithm find...
  - Species A signal hybrids?
  - Species A non-signal hybrids?
  - Species B hybrids (mimics of Species A signal hybrids)?

Species A subspecies I



Species A subspecies II



Species A subspecies III



Species A subspecies IV



Species B subspecies II



Species B subspecies I



# Sample Submissions Repository



Files

feature/notebook

Go to file

- > BioCLIP\_code\_submission
- > BioCLIP\_train
- > DINO\_SGD\_code\_submission
- > DINO\_train
- .gitignore
- LICENSE
- README.md
- butterfly\_anomaly.bib
- butterfly\_sample\_notebook.ipynb
- requirements.txt

HDR-anomaly-challenge-sample / butterfly\_sample\_notebook.ipynb

Preview

Code

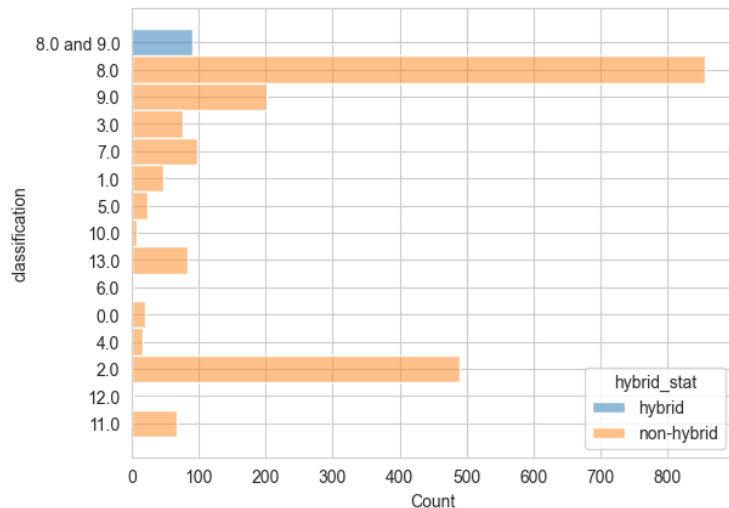
Blame

606 lines (606 loc) · 53.3 KB

## Get distribution of images by subspecies (colored by hybrid status)

```
In [15]: sns.histplot(df, y = "classification", hue = "hybrid_stat")
```

```
Out[15]: <Axes: xlabel='Count', ylabel='classification'>
```





Join the  
Challenge!

# Thank you!

Questions?

