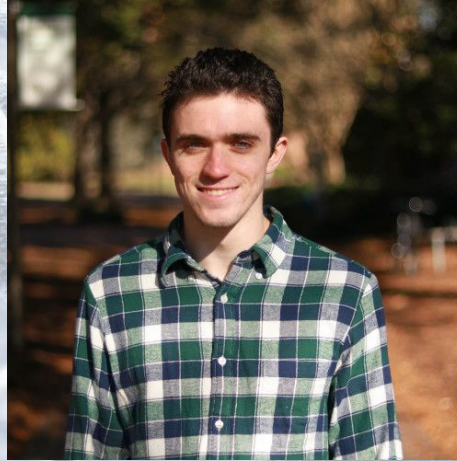




Classifying Zoonotic Viruses

8/24/23

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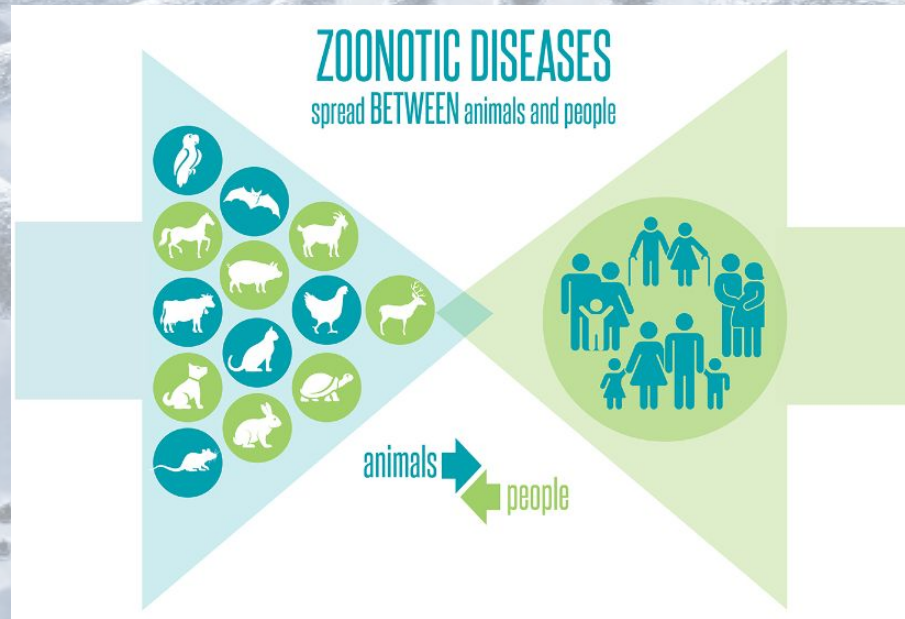
James McLaughlin

[LinkedIn](#)

[GitHub](#)

What are zoonotic viruses?

- **Caused by germs that spread between animals and people**
- **Can spread through:**
 - **Direct contact**
 - **Indirect contact**
 - **Vectors**
 - **Food**
 - **Water**



Bottom line

- **Model can correctly identify associated animal about 95% of the time**
- **Quick way to make predictions for a large number of genetic sequences**
- **Useful supplement to [BLAST](#)**

Why it matters

- **Identifying the animal origins of COVID is an ongoing search**
- **3 out of 4 emerging infectious diseases come from animals**
- **Key for vaccine development and preventing future outbreaks**

Data Overview

- Data comes from [ZOVER](#)
- ~50,000 unique viruses associated with bats, rodents, mosquitos, and ticks
- DNA and protein sequences



Data Limitations



- **Incomplete sequences**
- **No numeric data**
- **Limited to 4 animals**

Model

- **Voting classifier model that combines XGBoost and Neural Networks**
- **XGBoost has higher accuracy, less overfitting**
- **Neural Networks identify hidden features in data**

Model metrics

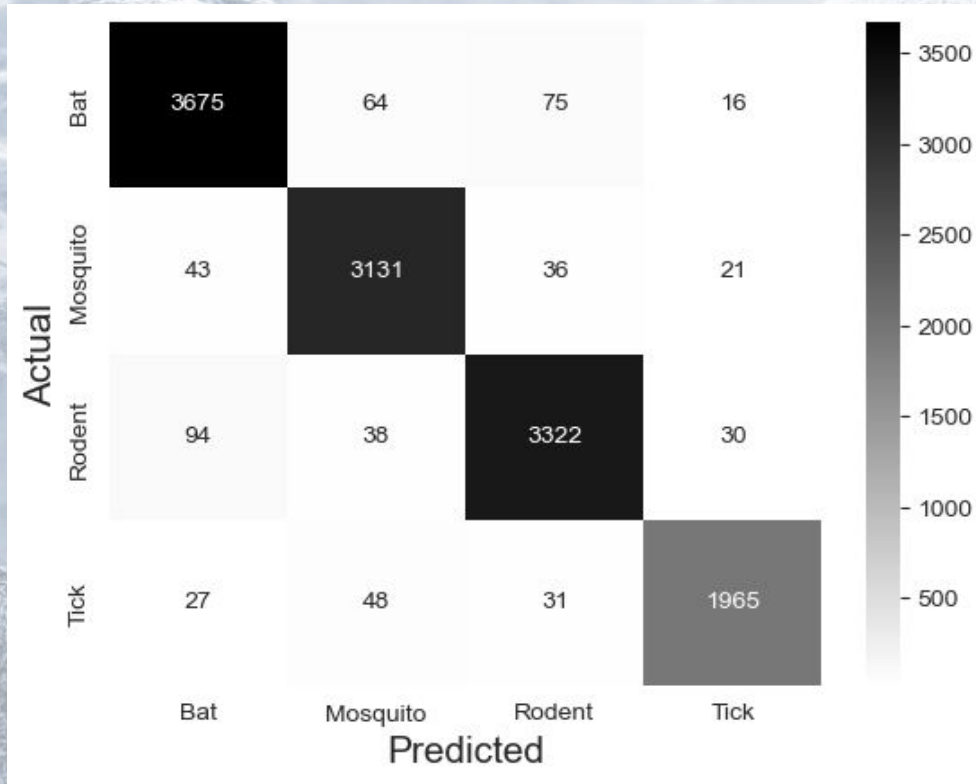
Accuracy

95.4%

ROC-AUC Score

0.97

Model metrics



Sources

<https://www.cdc.gov/onehealth/basics/zoonotic-diseases.html>

<https://www.theatlantic.com/science/archive/2023/03/covid-origins-research-raccoon-dogs-wuhan-market-lab-leak/673390/>