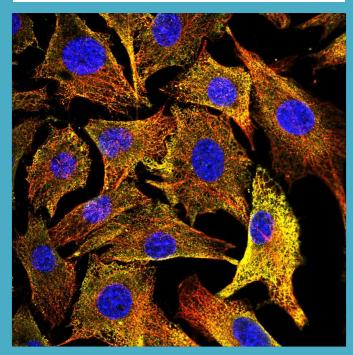


#### Human Protein Image Classification

Luke LaJoie June 16, 2020

#### THE HUMAN PROTEIN ATLAS



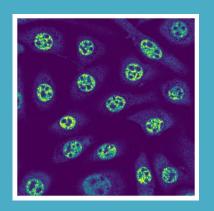
#### **Human Protein Classification**

- Classify different proteins in human cells
- Proteins do most of the work in the cell



#### **Human Protein Classification**

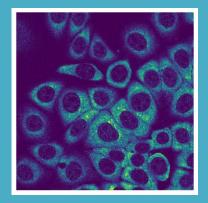
- Classify different proteins in human cells
- Proteins do most of the work in the cell
- High throughput confocal microscope



Nuclear speckles



- Classify two different proteins
  - Nuclear speckles
  - Cytosol



Cytosol

## Methodology



#### **Gather Data**

Data imagery provided by Human Protein Atlas and available via Kaggle

#### kaggle



#### **Process Data**

Images needed to be transformed in different ways to work with different models

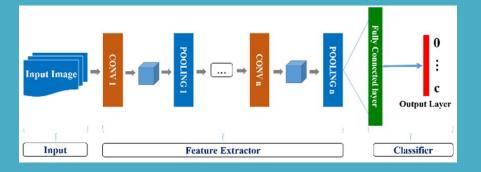


#### **Build and Run Models**

Random Forest and Convolutional Neural Network models were run and compared

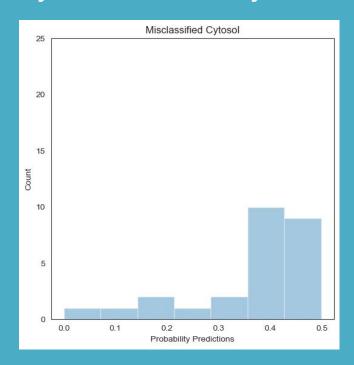


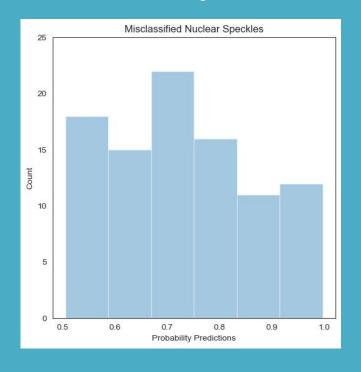
- CNN
  - 3 block VGG
  - Output: 0 to 1
    - NS: 0 to 0.5
    - C: 0.5 to 1

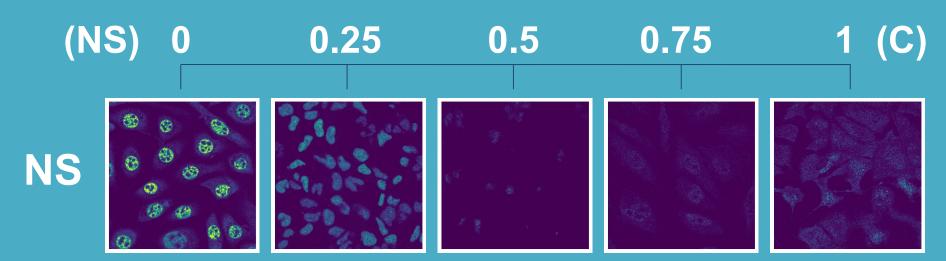


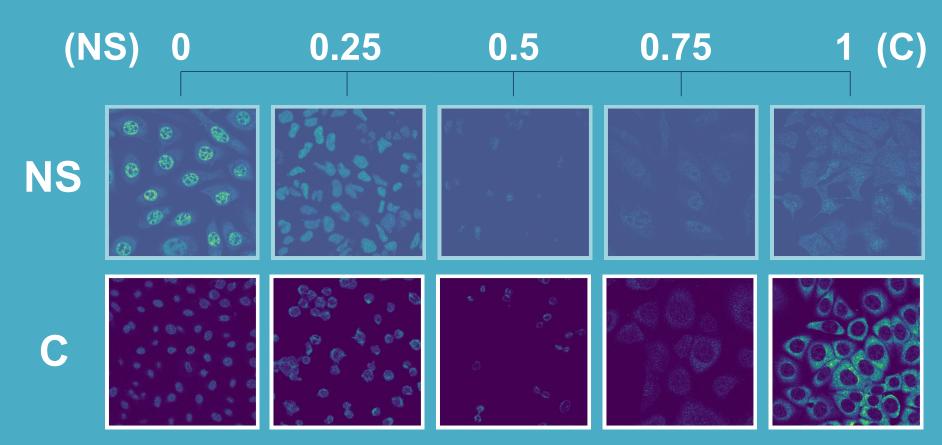
**Accuracy Score: 0.930** 

Cytosol less likely to be classified incorrectly











#### Conclusion

 CNN was a better classifier than RF for this image data

Accuracy of 93%

 Potentially can save time and resources by automating classification



#### **Credits**

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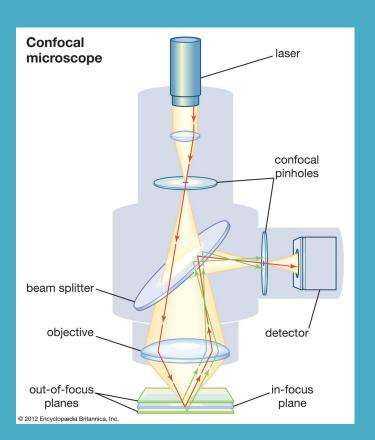
Illustrated images in this template were created by **GoogleSlidesppt.com** 

Inserted pictures in this template were created by <u>pixabay.com</u>

#### **Future Work**

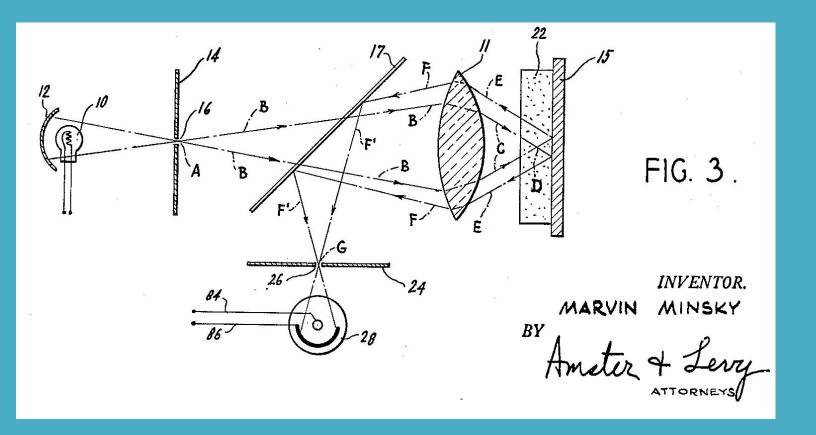
- Work with multiclass / multilabel data
- Try other pretrained models for transfer learning

## Appendix



## Human Protein Atlas Kaggle competition

- High throughput confocal microscopy
- Classify subcellular proteins in human cells







## Methodology

Insert the title of your subtitle Here

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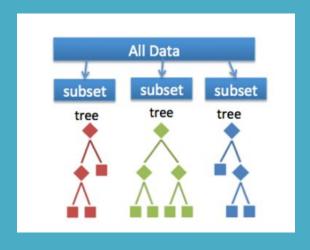
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You can simply impress your audience and add a unique zing and appeal to your Presentations.

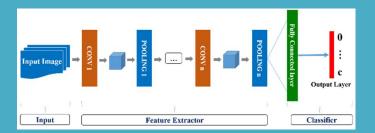
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## Methodology

- PCA
- Random Forest



- CNN
  - o 3 block VGG

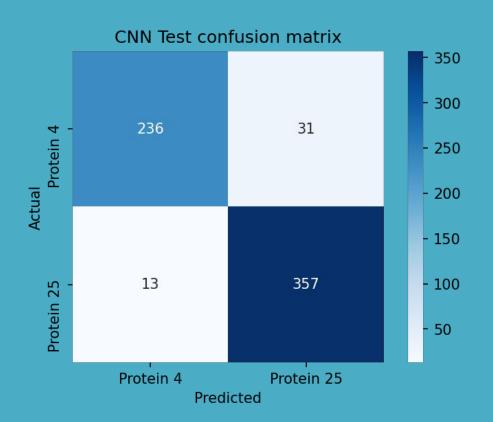


#### CNN

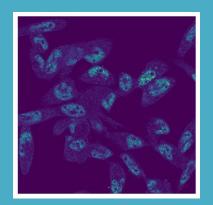
Accuracy Score: 0.930 Precision Score: 0.920

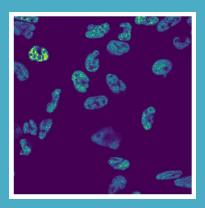
Recall Score: 0.964

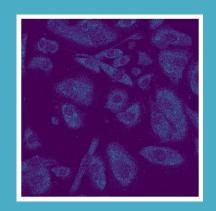
F1 Score: 0.941

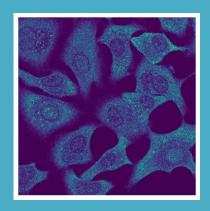


### **Protein 4 Protein 25**

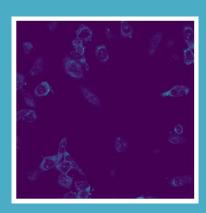






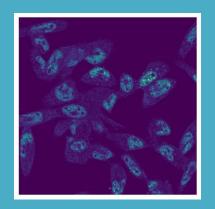


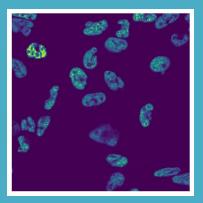


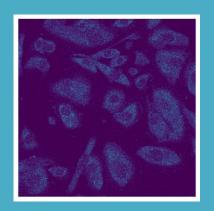


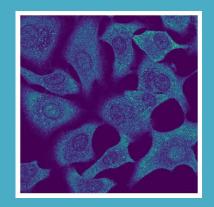


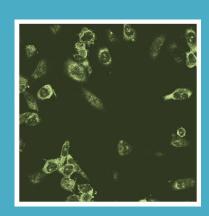
### **Protein 4 Protein 25**





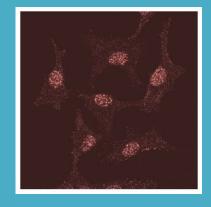






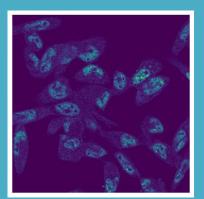
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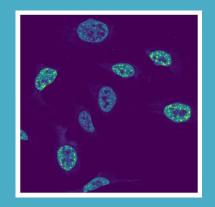
0.528

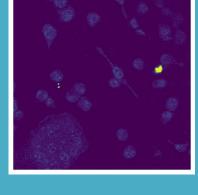


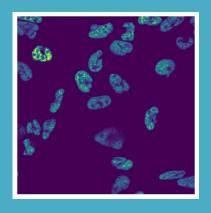
**4** 0.550

## Results: Nuclear Speckles

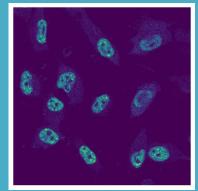




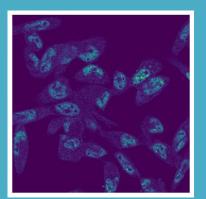


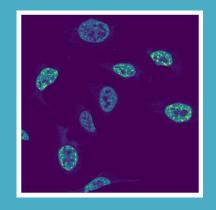


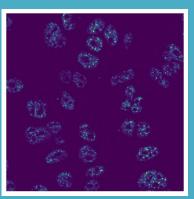


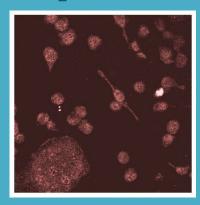


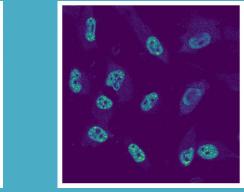
## Results: Nuclear Speckles



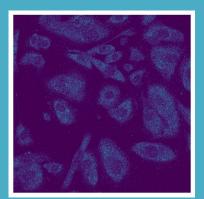


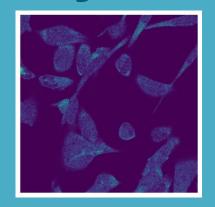


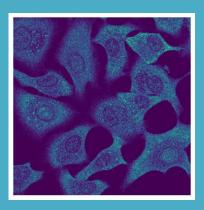


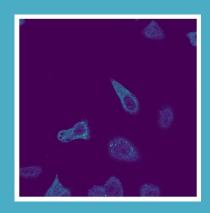


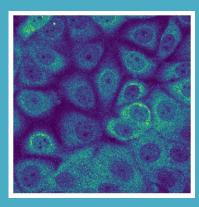
## Results: Cytosol



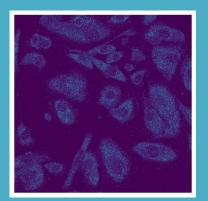


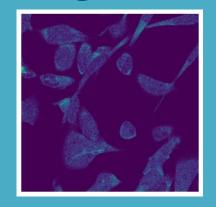


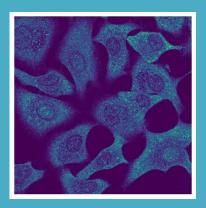


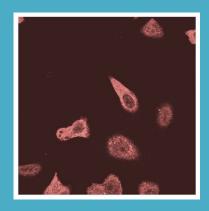


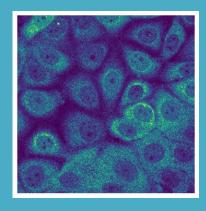
## Results: Cytosol

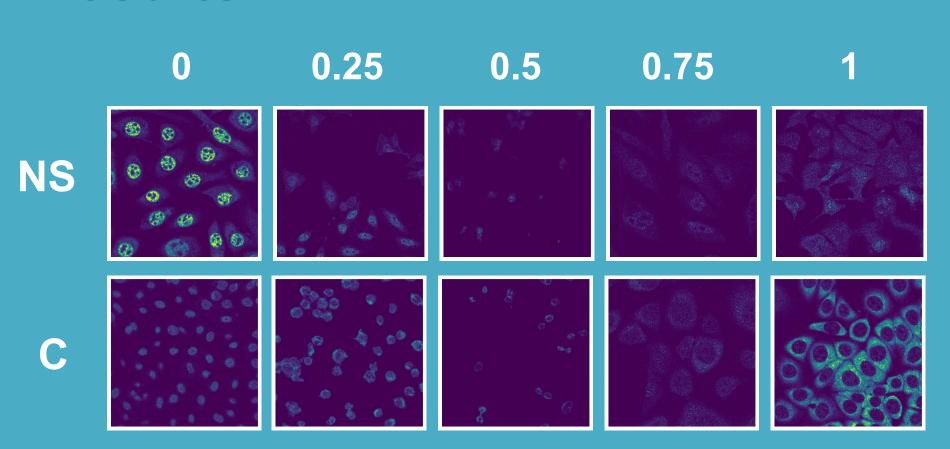




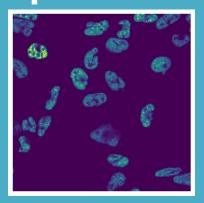




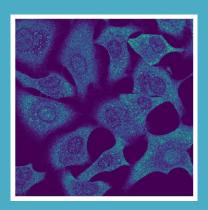




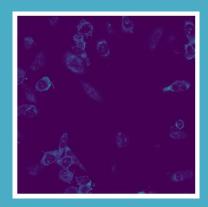
# Nuclear speckles



#### Cytosol

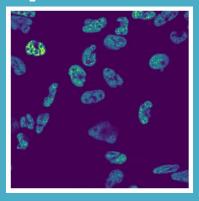


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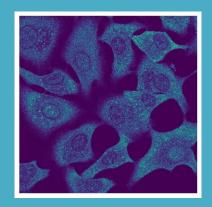


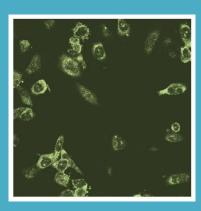


# Nuclear speckles



#### Cytosol





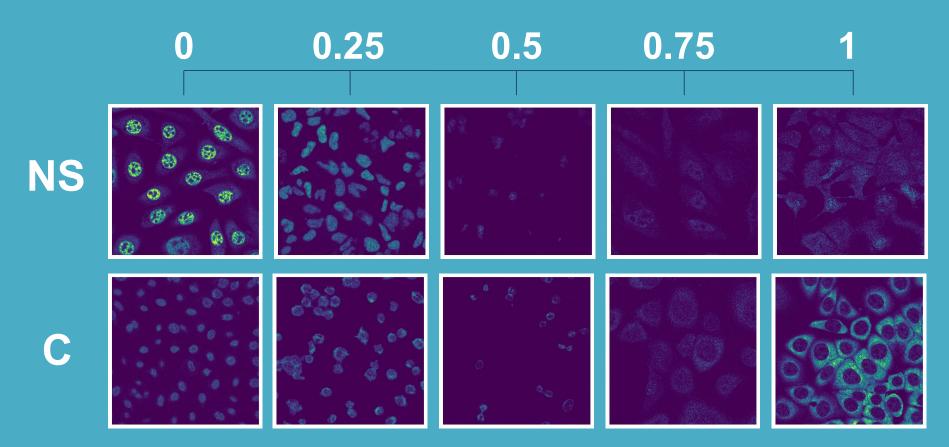
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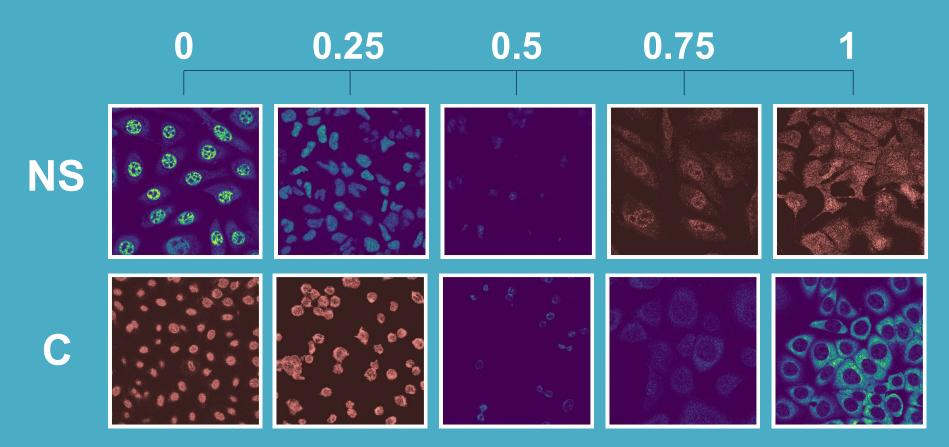
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NS

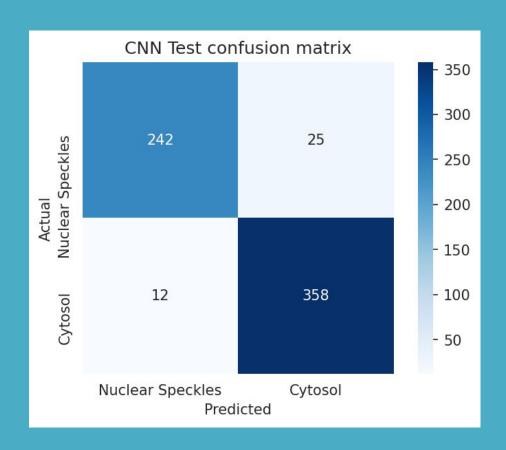
0.550





- CNN
  - 3 block VGG
  - Output: 0 to 1
    - Nuc Spec: < 0.5
    - Cytosol: > 0.5

Accuracy Score: 0.930





#### Conclusion

- CNN was a better classifier than RF for this image data
- Instances of CNN model "uncertainty" may require human intervention



Cytosol less likely to be classified incorrectly

