A decorative graphic on the left side of the slide consisting of two overlapping parallelograms. The front one is blue and the back one is a light greenish-blue. They are positioned diagonally, with the blue one partially covering the green one.

# Analysis of Diabetic Retinopathy Fundus Imagery using Convolutional Neural Networks

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# Objectives

- Learn more about the Data Analytics process
- Learn more about the development of convolutional neural networks
- Develop a CNN model that correctly classifies the presence of diabetic retinopathy in fundus images as either being present or not present

# Implemented Models





# Constants

- All models share the following components:
  - Constructed using Keras' built-in Sequential structure
  - Compiled with the Adam optimizer
  - Utilize the binary cross entropy loss function
  - Contain a Flatten layer and a Dense layer



# Base Model

The Base Model's layer architecture is as follows:

1. 2D Convolution Layer
  - a. 64 nodes
  - b. Kernel Size =  $3 \times 3$
  - c. ReLU activation function
2. 2D Convolution Layer
  - a. 32 nodes
  - b. Kernel Size =  $3 \times 3$
  - c. ReLU activation function
3. Flatten Layer
4. Dense Layer
  - a. Sigmoid activation function



# 1 Convolution Layer

The 1 Convolution Layer's architecture is as follows:

1. 2D Convolution Layer
  - a. 64 nodes
  - b. Kernel Size =  $3 \times 3$
  - c. ReLU activation function
2. Flatten Layer
3. Dense
  - a. Sigmoid activation function



# Max Pooling and Kernel Adjustments

The Max Pooling Model's architecture is as follows:

1. 2D Convolution Layer
  - a. 64 nodes
  - b. Kernel Size =  $5 \times 5$
  - c. ReLU activation function
2. Max Pooling 2D Layer
  - a. Pool Size =  $3 \times 3$
  - b. Padding = "same"
3. 2D Convolution Layer
  - a. 32 nodes
  - b. Kernel Size =  $3 \times 3$
  - c. ReLU activation function
4. Flatten Layer
5. Dense Layer
  - a. Sigmoid activation function

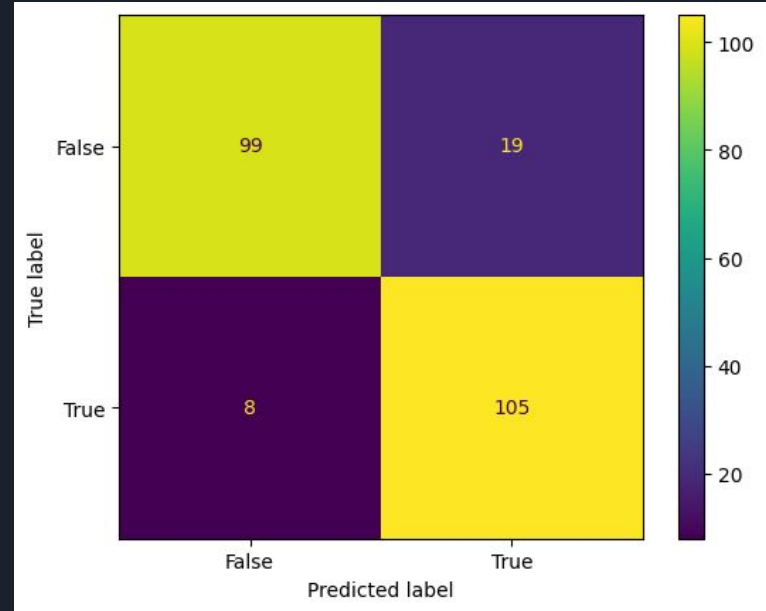
# Results and Analysis





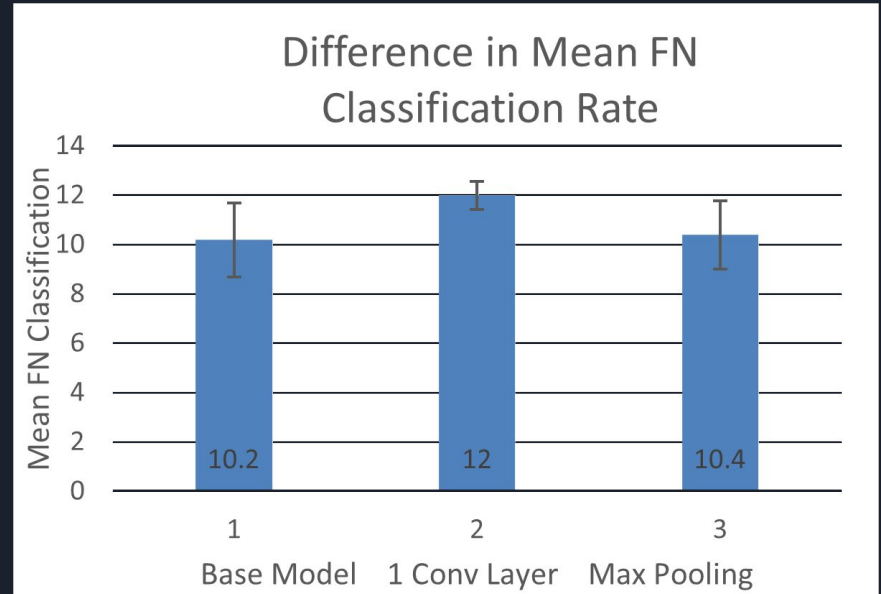
# Results and Analysis

- Success Criteria = Minimal False Negatives (FN)
- Difference of Means T-Test
- Comparison of Other Relevant Metrics
  - Precision
  - Recall
  - F1 Score



# Difference of Means T-Test

- Conducted comparing the two models achieving the lowest average FN counts across five runs: Base Model (10.2), and Max Pooling Model (10.4)
- P-value of 0.815 = insignificant statistical difference
- \*A t-test was also conducted comparing the Base Model to the 1 Convolution Layer Model, which indicated a statistically significant difference in performance





# Further Comparisons

The following are averages achieved by each model in the corresponding metrics

	Base Model	Max Pooling Model
Testing Accuracy	88.75%	89.78%
Precision	86.77%	88.64%
Recall	90.97%	90.80%
F1 Score	88.79%	89.69%

Future Scope





# Future Scope

- Further examination of hyperparameters
  - Learning Rate
- Experimentation with model architecture
  - Node counts
  - Activation and loss functions
  - Additional layers; deeper network
- Additional image preprocessing
  - Data augmentation
    - Rotation
    - Skew
    - Shear
    - Flipping

# Reflections and Conclusion





# Reflections and Conclusion

- False negative classification rate (~4.50%) at present is still higher than what's required for practical implementation
- Gained a greater appreciation for the Data Analytics Process
- Obtained exposure to the development of CNNs
- Better accuracy likely could've been achieved with adjustments to learning rate, amongst other parameters

Thank you!

