# **AutoML Modeling Report**



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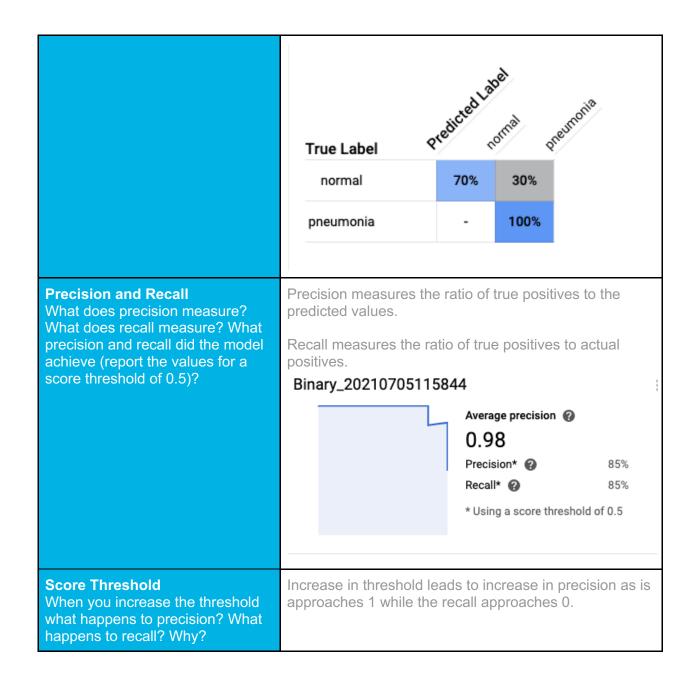
## Binary Classifier with Clean/Balanced Data

### **Train/Test Split** Labels Images How much data was used for training? How much data was used normal for testing? pneumonia Train Validation Test 80 10 10 80 10 10 **Confusion Matrix**

What do each of the cells in the confusion matrix describe? What values did you observe (include a screenshot)? What is the true positive rate for the "pneumonia" class? What is the false positive rate for the "normal" class?

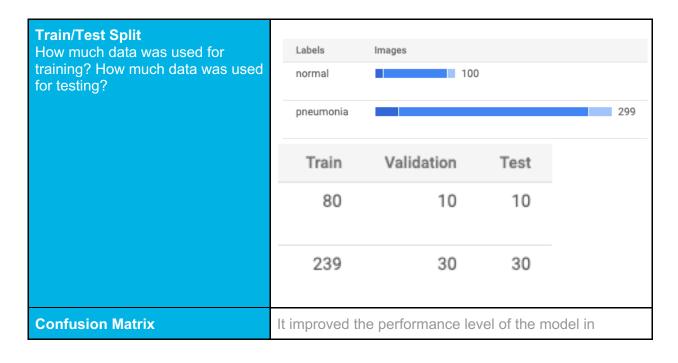
True positives are the correct classification of pneumonia while false positive is the incorrect classification of pneumonia.

In the confusion matrix, it shows the model was better at classifying pneumonia diseased lungs than healthy lungs.





# Binary Classifier with Clean/Unbalanced Data



How has the confusion matrix been affected by the unbalanced data? Include a screenshot of the new confusion matrix. classifying the healthy lungs in the confusion matrix with no effect on the pneumonia classification.

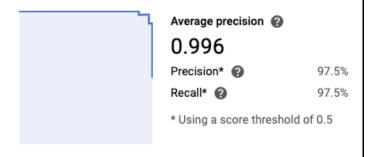


#### **Precision and Recall**

How have the model's precision and recall been affected by the unbalanced data (report the values for a score threshold of 0.5)?

There was significant improvement on the precision and recall of the model.

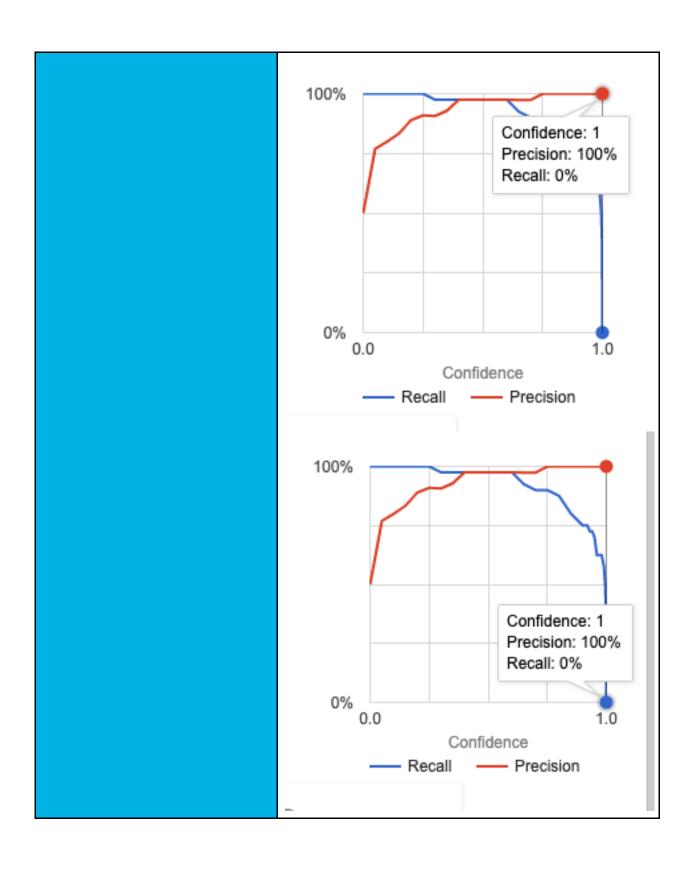
#### Unbalanced\_20210706124043



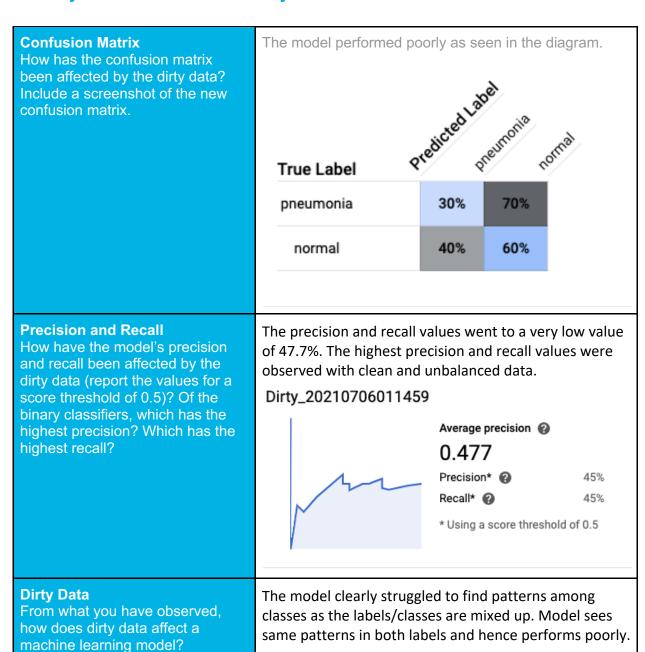
#### **Unbalanced Classes**

From what you have observed, how do unbalanced classed affect a machine learning model?

Unbalanced data introduces bias which impacts the accuracy of the model. With the increase in accuracy, the recall on both didn't change.



### Binary Classifier with Dirty/Balanced Data



### 3-Class Model

Confusion	Mat	rix	
Summarize	the	3-class	confusion

By intuition, the bacterial and viral classes are likely to be confused. But the model was able to classify the viral

matrix. Which classes is the model pneumonia class accurately with false positives in bacterial class and the normal class. most likely to confuse? Which class(es) is the model most likely to get right? Why might you do to To remedy this, I would change the labelling to exclude try to remedy the model's the pneumonia metadata. "confusion"? Include a screenshot of the new confusion matrix. True Label bacterial 40% viral 100% normal 10% 10% 80% **Precision and Recall** What are the model's precision Class\_20210706015017 and recall? How are these values calculated (report the values for a Average precision (2) score threshold of 0.5)? 0.905 Precision\* (2) 85.19% Recall\* (2) 76.67% \* Using a score threshold of 0.5 F1 Score F1 = 80.701% What is this model's F1 score?