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## CS 441 - HW 4: Trees and MLPs

Complete the sections below. You do not need to fill out the checklist. **Do select all relevant pages in Gradescope.**

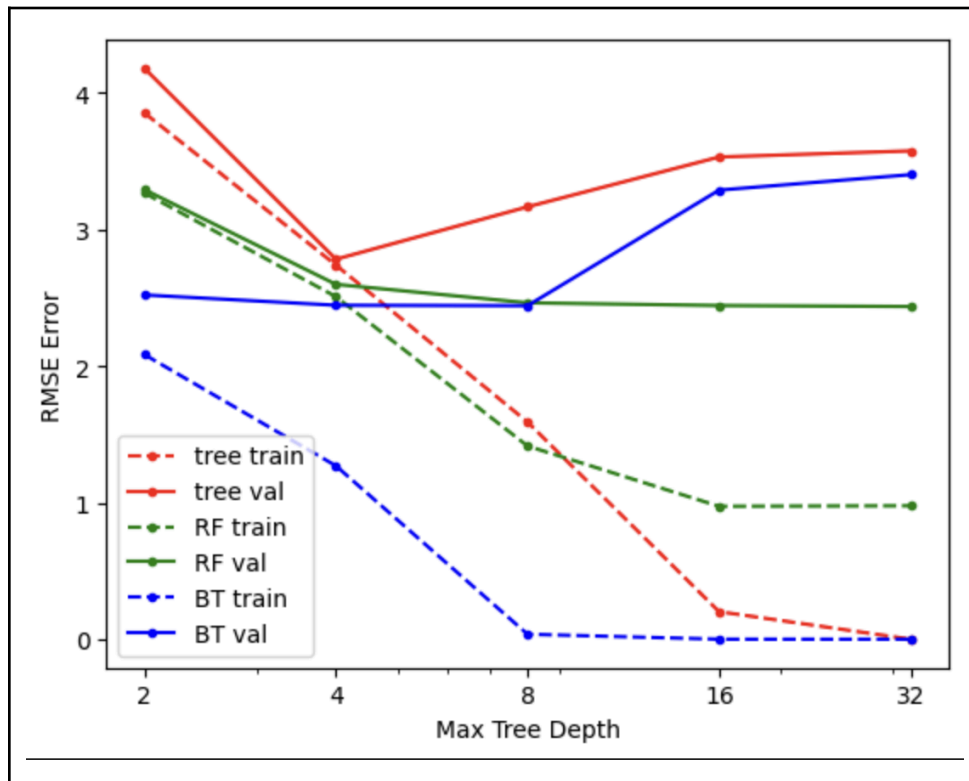
Total Points Claimed

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1. Model Complexity with Tree Regressors
  - a. Depth vs. Error plot [ ] / 10
  - b. Analysis [ ] / 20
2. MLPs with MNIST
  - a. Loss Curves [ ] / 20
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3. Species Prediction
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  - b. Simple Rule [ ] / 10
  - c. Model Design [ ] / 10
4. Stretch Goals
  - a. Improve MNIST classification [ ] / 30
  - b. A second simple rule [ ] / 10
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### 1. Model Complexity with Tree Regressors

- a. Include your plot below.



**b. Analyze your results:**

- For a given max tree depth, which of regressor model (single tree, random forest, boosted tree) has the lowest bias (or most powerful)?

Boosted tree

- For single regression trees, what tree depth achieves minimum validation error?

4

- A model “overfits” when increasing the complexity increases the validation error. Which model is least prone to overfitting? Why?

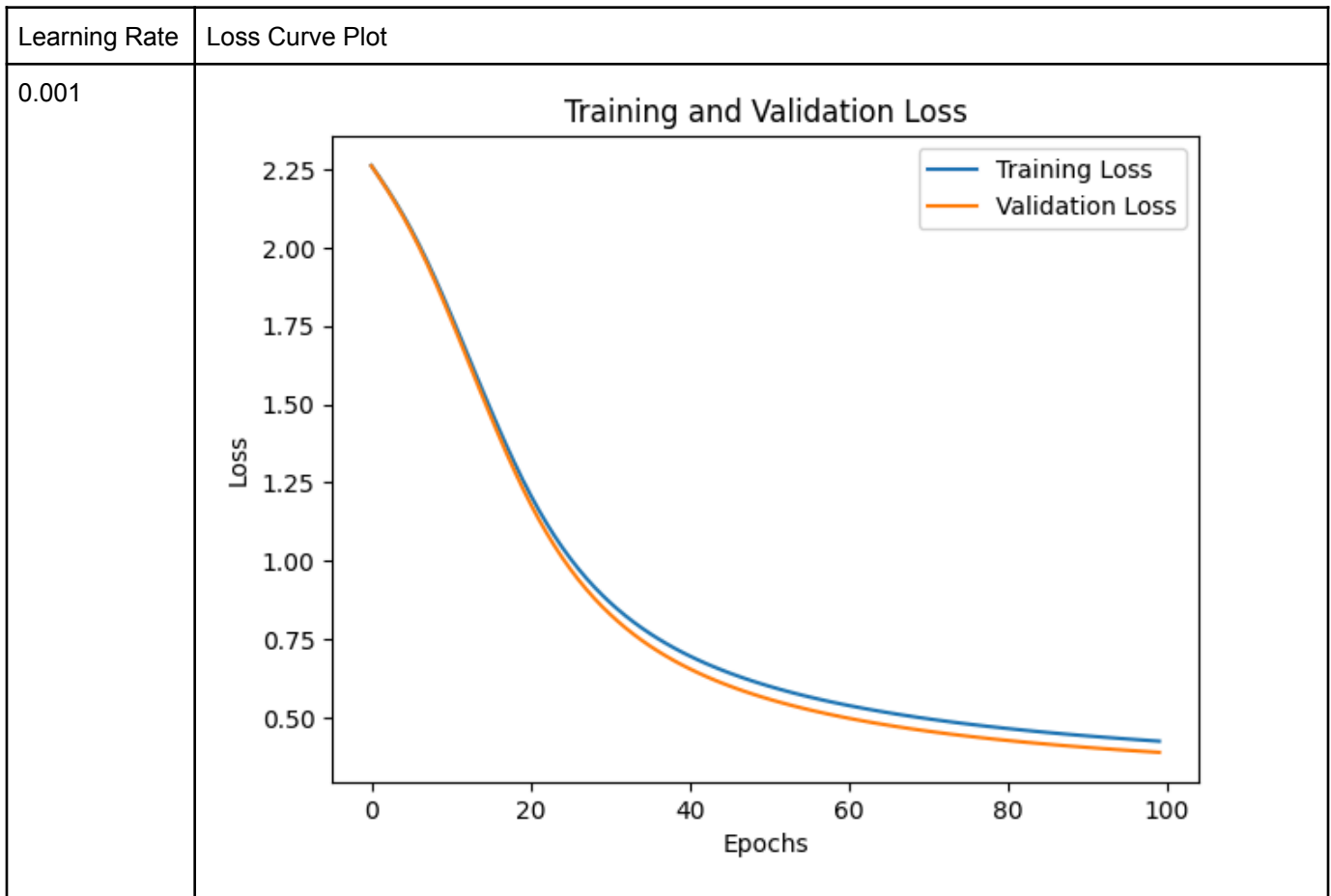
Average of many tree predictions reduces variance, thus reduce overfitting

- Do boosted trees seem to perform better with smaller or larger trees? Why?

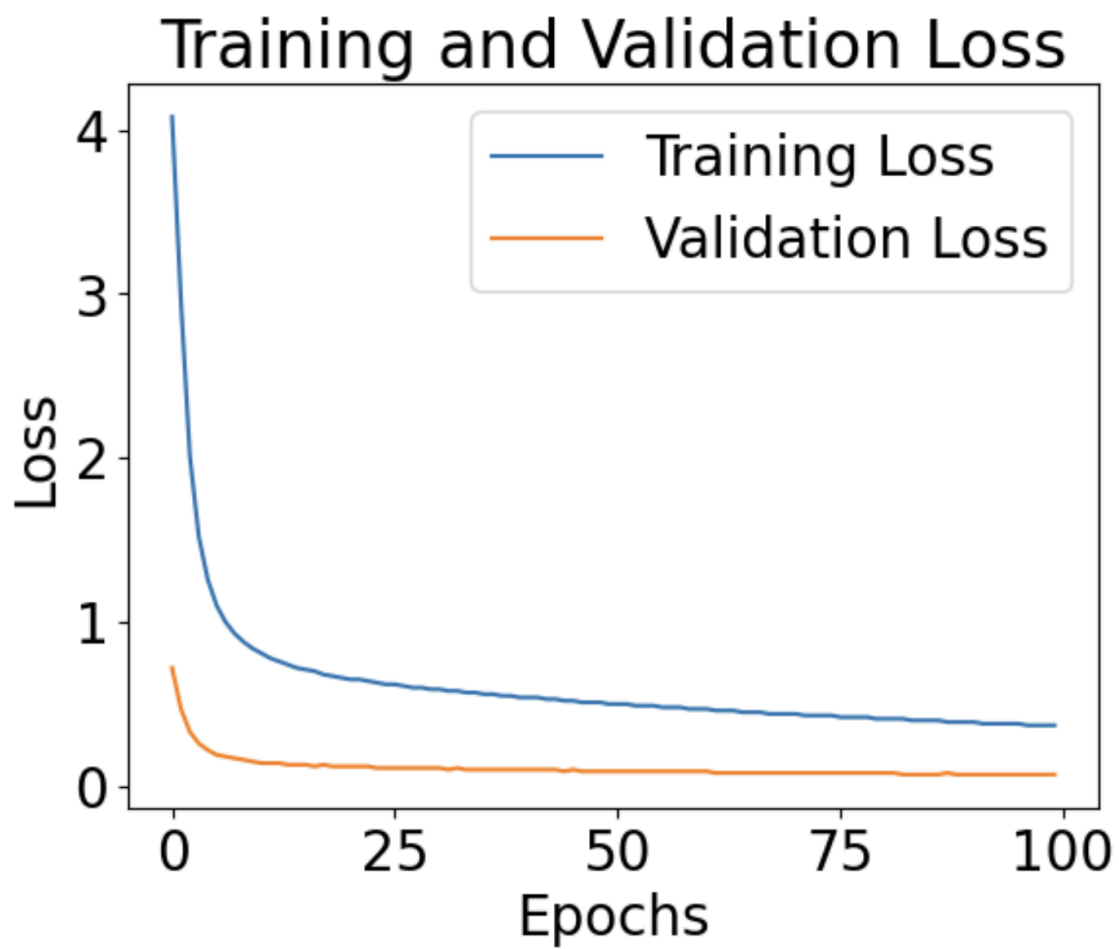
Smaller trees. Too large trees eliminates train error but increases test error

## 2. MLPs with MNIST

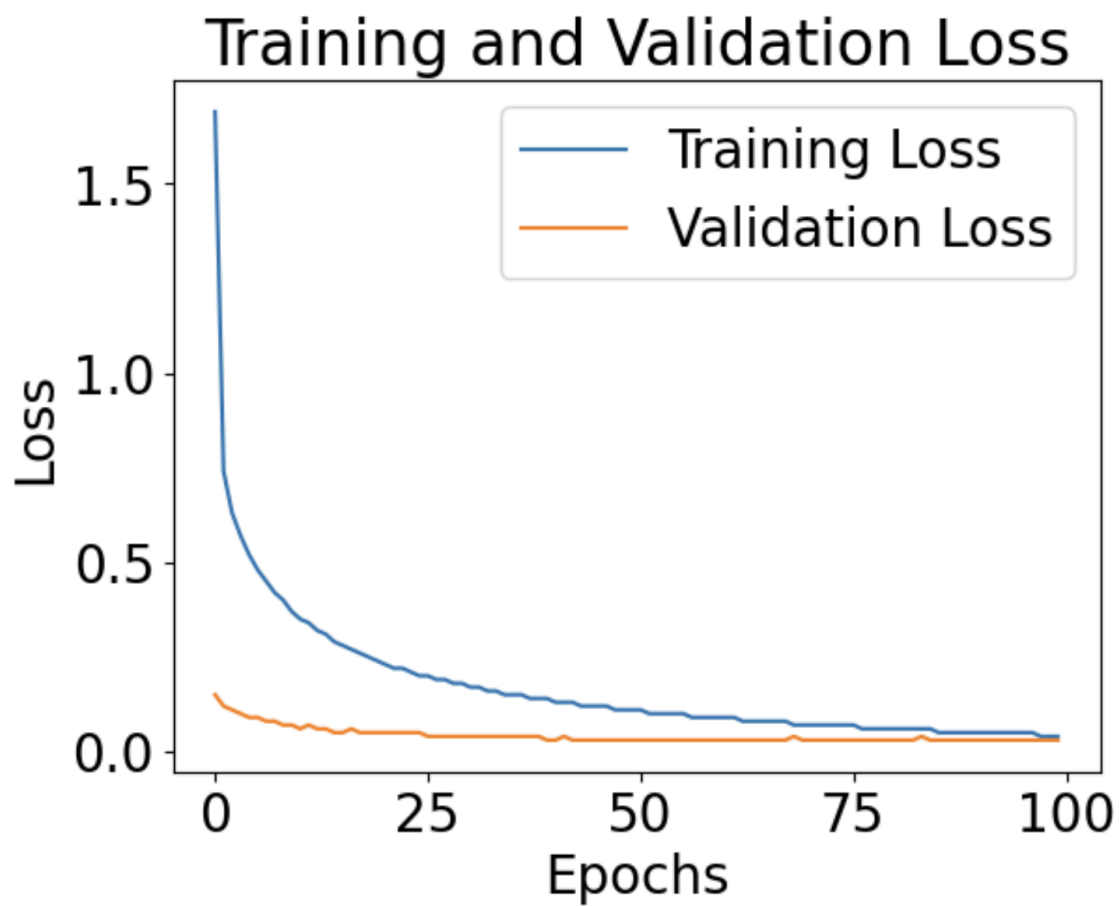
- a. **Show the loss curves** for 3 learning rates (1E-2, 1E-1, 1E1) training for 100 epochs.  
An example of the loss curves is shown for LR=0.001.



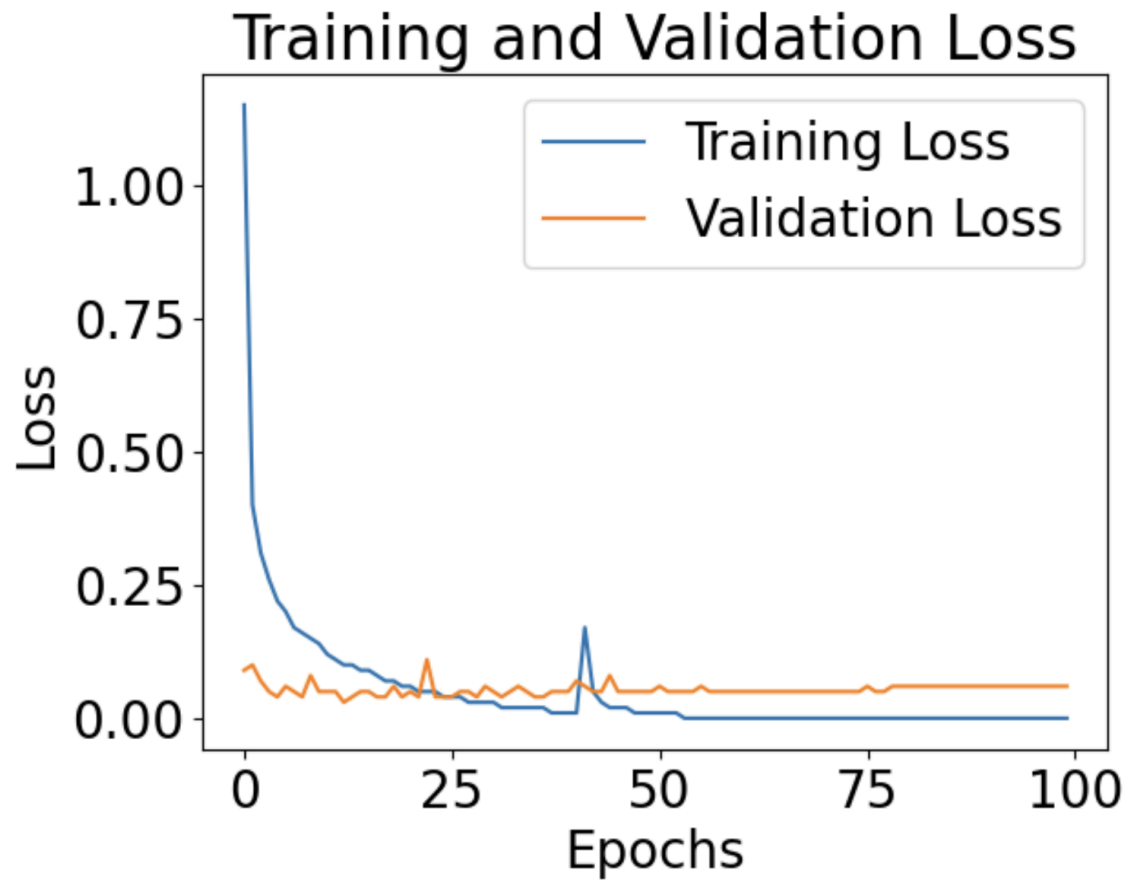
0.01



0.1



1



#### b. Model selection and results

**Select the best hyperparameters** (learning rate and number of epochs up to 100) based on minimizing the validation loss.

**Learning Rate**

**Epochs**

**Report the losses and errors for the model trained with these hyperparameters:**

Use scientific notation with one decimal place, e. 1.5E-3

Training <b>Loss</b>	Validation <b>Loss</b>	
3.3E-2	1.3E-1	

Show two decimal places for percent

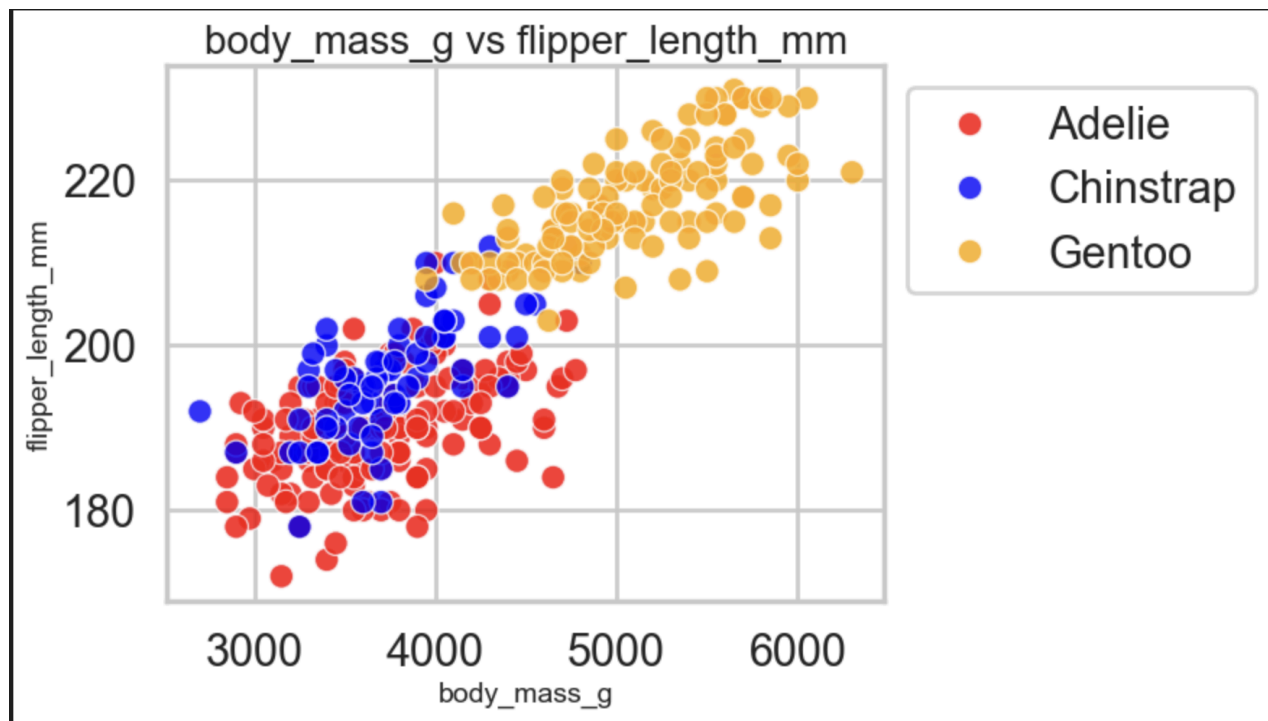
Training <b>Error</b> (%)	Validation <b>Error</b> (%)	Test <b>Error</b> (%)
1.1E-2	3.2E-2	3.3E-2

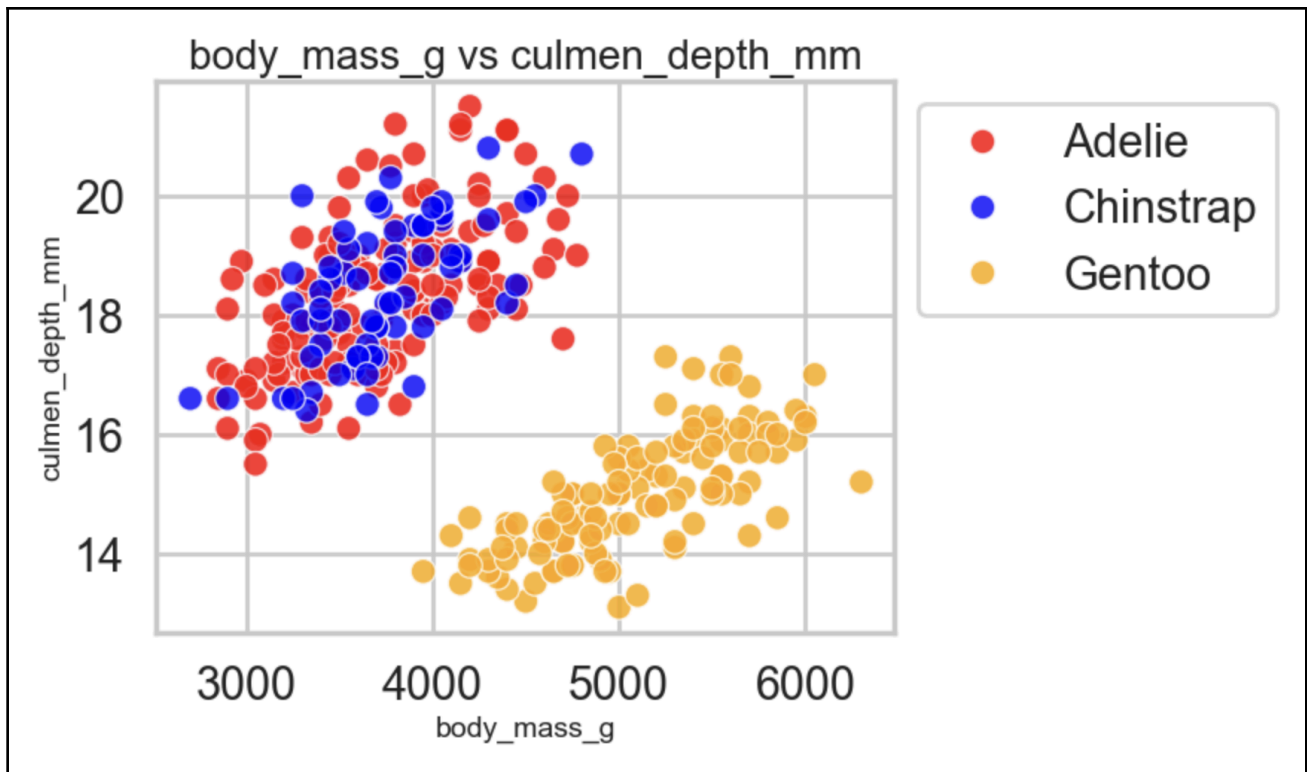
### 3. Species Prediction

#### a. Visualization of Features

Include at least two scatterplots of pairs of features.

Visualization (labels should make clear which features are used)





You may extend the table if you have more results

**Of these three options, which two features (by themselves) are best able to classify the penguin species?**

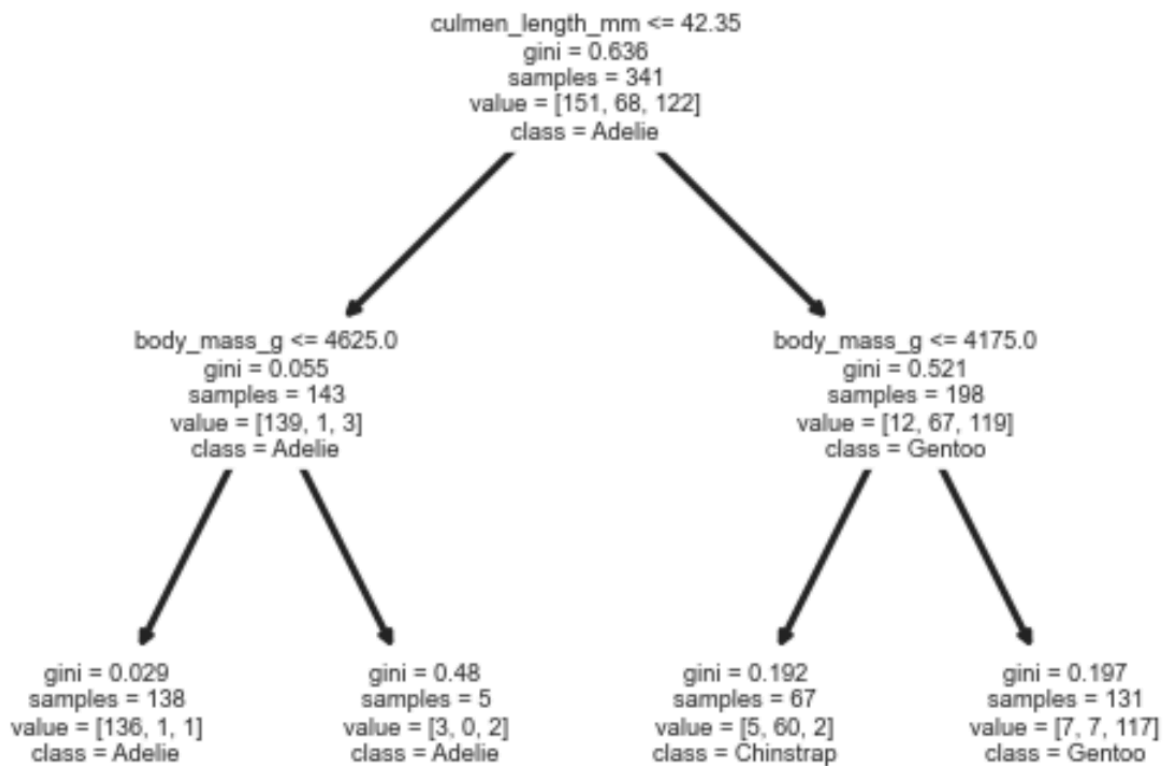
1. Culmen Depth + Flipper Length
2. Flipper Length + Culmen Length
3. Flipper Length + Body Mass

Flipper Length + Culmen Length

**b. Simple rule to identify Gentoo**

**Display your decision tree with labeled features and classes.**





**Write down the simple two-part rule to identify Gentoo.** For example, the format should be “If Mass > 3000 and Culmen Depth < 17, then species is Gentoo”.

If...

If Mass <= 4175.0

and

Culmen Depth <= 42.35

then species is **Gentoo**.

**Rule precision:** fraction of penguins that satisfy this rule that are Gentoos (# gentoo predicted / # predicted)

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**Rule recall:** fraction of all Gentoo penguins that are identified as Gentoo using this rule (# gentoo predicted / # gentoo)

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### c. Model Design

Describe the model that achieves best 5-fold cross-validation accuracy:

Random forest model achieves the best accuracy

5-fold Cross-Validation Accuracy: (xx.x%)

99.1%

## 3. Stretch Goals

### a. Improve MNIST Classification Performance using MLPs

Report the classification val and test errors and details of your best method. Describe your approach and parameters. Feel free to change the MLP batch size, optimizer (e.g. try Adam), learning rate, number of epochs, hidden layer size, activation layer, or anything else.

#### Description and key parameters

Optimizer = Adam  
Hidden layer(s) = 128,64,32  
Learning rate = 0.01  
Number of epochs = 84

Any other details: attempts different hidden layer structure, found that a structure more complex than this one may end up higher error rate

Validation <b>Error</b> (%)	Test <b>Error</b> (%)
2.2E-2	2.2E-2

**b. Find a second simple rule to identify Gentoo**

Provide the second two-part rule here (that is substantially different from your first rule).

**If...**

Flipper length  $\leq 206.5$

**and**

Culmen length  $\leq 40.85$

**then species is Gentoo.**

**Rule precision:** fraction of penguins that satisfy this rule that are Gentoos (# gentoo predicted / # predicted)

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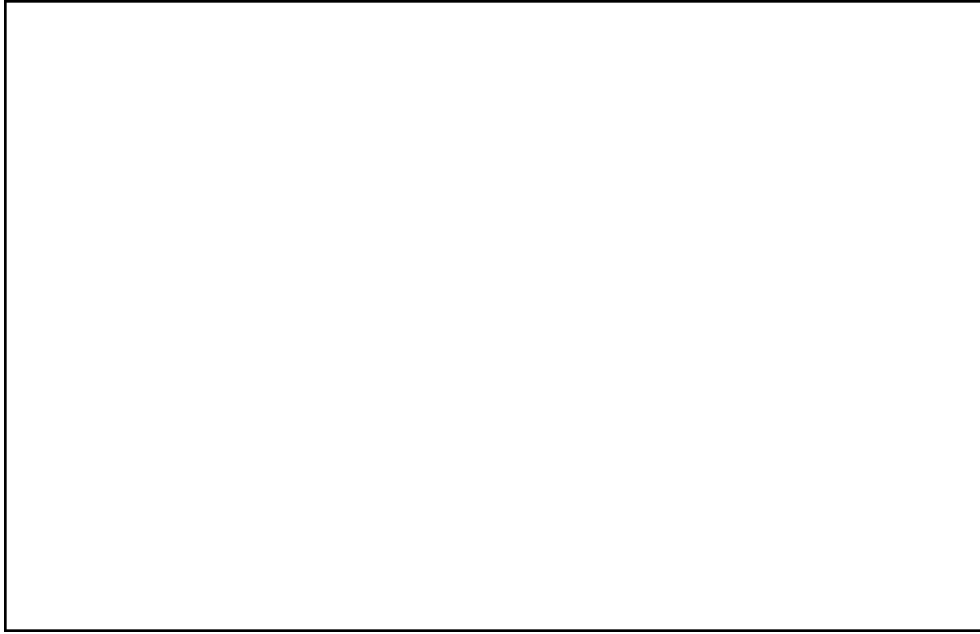
**Rule recall:** fraction of all Gentoo penguins that are identified as Gentoo using this rule (# gentoo predicted / # gentoo)

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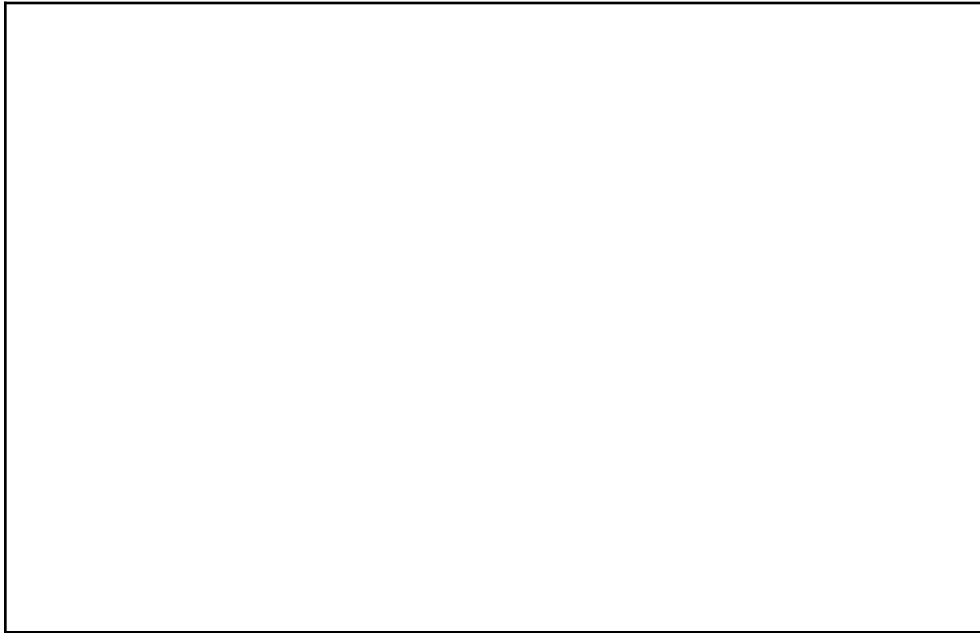
**c. Positional encoding**

Show the RGB image obtained by predicting directly from (x,y) and the image obtained by predicting from the positional encoding.

**Input to network is (x,y)**



**Input to network is  $\text{pos\_enc}(x, y)$**



**Acknowledgments / Attribution**

None