Name:	
	Hanliang Jiang
Netid:	
	hj33

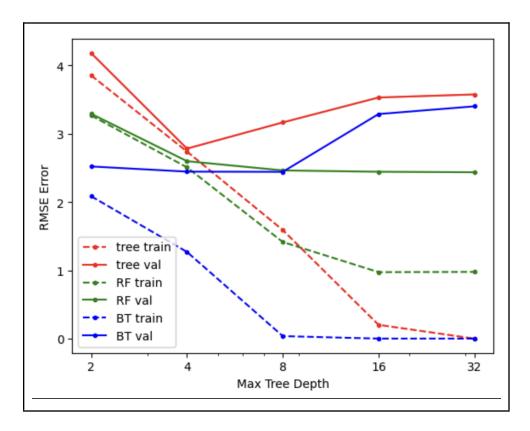
CS 441 - HW 4: Trees and MLPs

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

Total Points Claimed	[]/170
Model Complexity with Tree Regressors	
a. Depth vs. Error plot	[]/10
b. Analysis	[]/20
2. MLPs with MNIST	
a. Loss Curves	[]/20
b. Model Selection and Results	[]/20
3. Species Prediction	
a. Feature Analysis	[]/10
b. Simple Rule	[]/10
c. Model Design	[]/10
4. Stretch Goals	
 a. Improve MNIST classification 	[]/30
b. A second simple rule	[]/10
 c. Positional encoding of RGB Image 	[]/30

1. Model Complexity with Tree Regressors

a. Include your plot below.



b. Analyze your results:

1. 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

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

4

3. A model "overfits" when increasing the complexity increases the validation error. Which model is least prone to overfitting? Why?

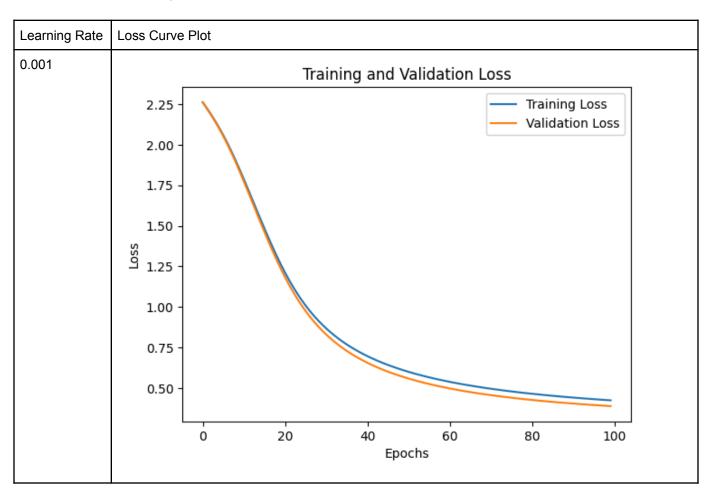
Average of many tree predictions reduces variance, thus reduce overfitting

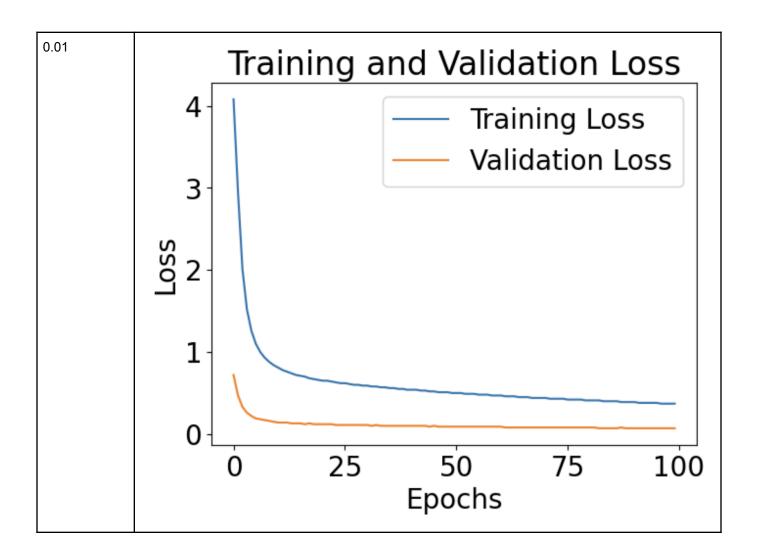
4. Do boosted trees seem to perform better with smaller or larger trees? Why?

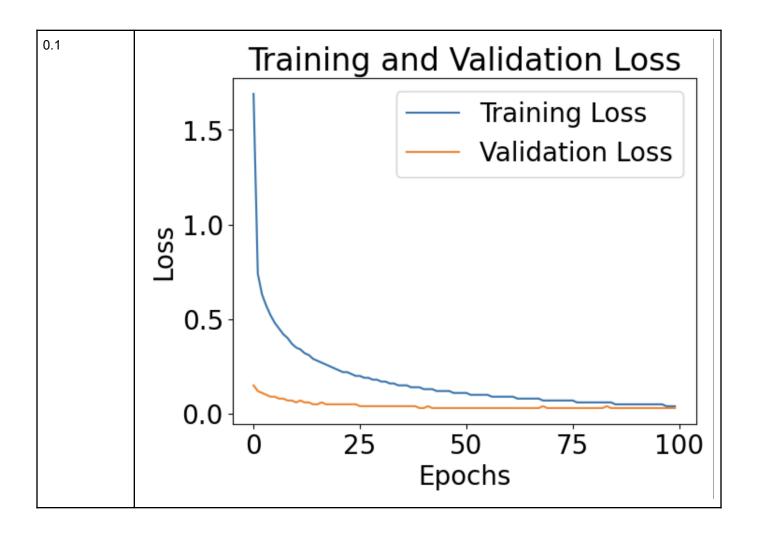
Smaller trees. Too large trees eliminates train error but increases test error

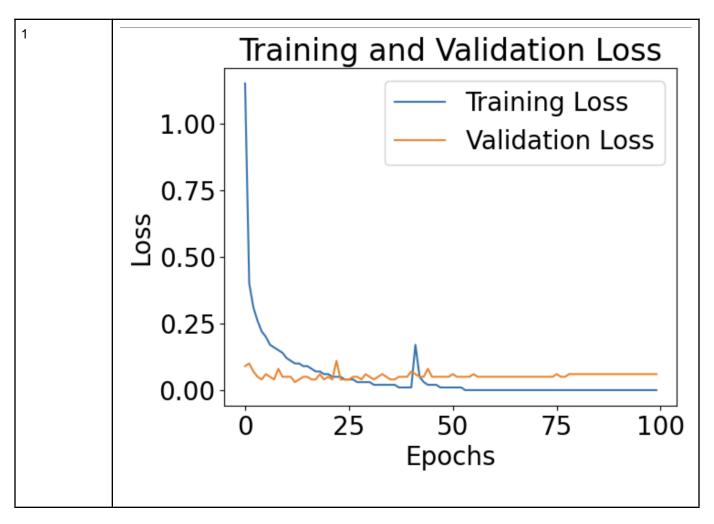
2. MLPs with MNIST

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.



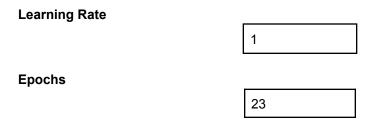






b. Model selection and results

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



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

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

Training Loss	Validation Loss	
3.3E-2	1.3E-1	

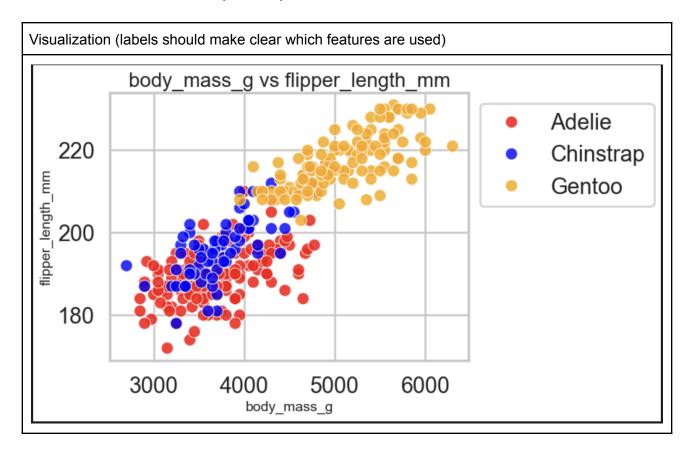
Show two decimal places for percent

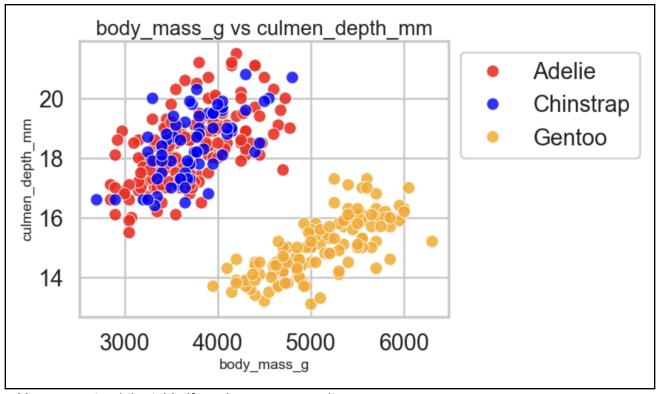
Training Error (%)	Validation Error (%)	Test Error (%)
1.1E-2	3.2E-2	3.3E-2

3. Species Prediction

a. Visualization of Features

Include at least two scatterplots of pairs of features.





You may extend the table if you have more results

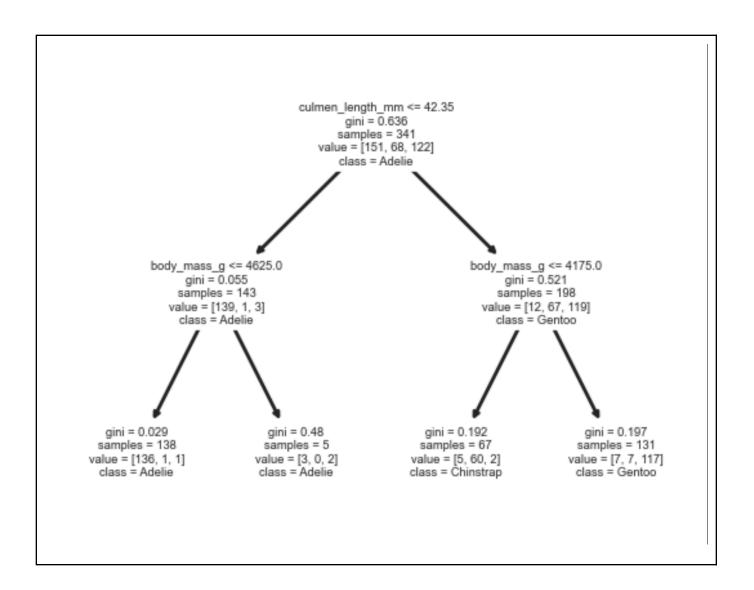
Of these three options, which two features (by themselves) are best able to classify the penguin species?

- 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 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)

117/341

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

117 /122

c. Model Design

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

Random forest model achieves the best accuricy

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 higner error rate

Validation Error (%)	Test Error (%)
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 <= 206.5

and

Culmen length <= 40.85

then species is Gentoo.

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

121/341

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

121/122

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 n	etwork is pos_enc(x, y)
Input to n	etwork is pos_enc(x, y)
Input to n	etwork is pos_enc(x, y)
Input to n	etwork is pos_enc(x, y)
Input to n	etwork is pos_enc(x, y)
Input to n	etwork is pos_enc(x, y)
Input to n	etwork is pos_enc(x, y)
Input to n	etwork is pos_enc(x, y)
Input to n	etwork is pos_enc(x, y)
Input to n	etwork is pos_enc(x, y)
Input to n	etwork is pos_enc(x, y)
Input to n	etwork is pos_enc(x, y)
Input to n	etwork is pos_enc(x, y)

Acknowledgments / Attribution

None

CS441: Applied ML - HW 4

Part 1: Model Complexity and Tree-based Regressors

One measure of a tree's complexity is the maximum tree depth. Train tree, random forest, and boosted tree regressors on the temperature regression task, using all default parameters except:

- max_depth={2,4,8,16,32}
- random state=0
- For random forest: max_features=1/3

Measure train and val RMSE for each and plot them all on the same plot using the provided plot_depth_error function. You should have six lines (train/val for each model type), each with 5 data points (one for each max depth value). Include the plot and answer the analysis questions in the report.

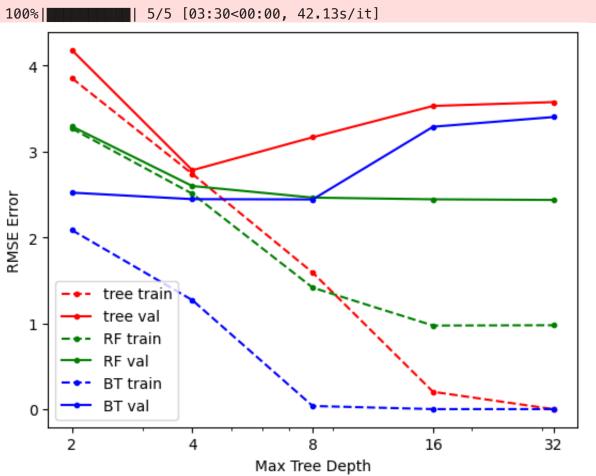
```
In [ ]: import numpy as np
        from matplotlib import pyplot as plt
        # load data (modify to match your data directory or comment)
        def load temp data():
          datadir = "./"
          T = np.load(datadir + 'temperature data.npz')
          x_train, y_train, x_val, y_val, x_test, y_test, dates_train, dates_val, dates_val
          T['x_train'], T['y_train'], T['x_val'], T['y_val'], T['x_test'], T['y_test
          return (x_train, y_train, x_val, y_val, x_test, y_test, dates_train, dates
        # plot one data point for listed cities and target temperature
        def plot_temps(x, y, cities, feature_to_city, feature_to_day, target_date):
          nc = len(cities)
          ndays = 5
          xplot = np.array([-5, -4, -3, -2, -1])
          yplot = np.zeros((nc,ndays))
          for f in np.arange(len(x)):
            for c in np.arange(nc):
              if cities[c] == feature to city[f]:
                yplot[feature_to_day[f]+ndays,c] = x[f]
          plt.plot(xplot,yplot)
          plt.legend(cities)
          plt.plot(0, y, 'b*', markersize=10)
          plt.title('Predict Temp for Cleveland on ' + target_date)
          plt.xlabel('Day')
```

```
plt.ylabel('Avg Temp (C)')
          plt.show()
        # load data
        (x_train, y_train, x_val, y_val, x_test, y_test, dates_train, dates_val, dat
In [ ]: | # to plot the errors
        def plot_depth_error(max_depths, tree_train_err, tree_val_err, rf_train_err,
          plt.figure()
          plt.semilogx(max_depths, tree_train_err, 'r.--', label='tree train')
          plt.semilogx(max_depths, tree_val_err, 'r.-', label='tree val')
          plt.semilogx(max_depths, rf_train_err, 'g.--', label='RF train')
          plt.semilogx(max_depths, rf_val_err, 'g.-', label='RF val')
          plt.semilogx(max_depths, bt_train_err, 'b.--',label='BT train')
          plt.semilogx(max_depths, bt_val_err, 'b.-', label='BT val')
          plt.ylabel('RMSE Error')
          plt.xlabel('Max Tree Depth')
          plt.xticks(max depths, max depths)
          plt.legend()
          plt.rcParams.update({'font.size': 20})
          plt.show()
In [ ]: from sklearn import tree
        from sklearn.tree import DecisionTreeRegressor
        from sklearn.ensemble import RandomForestRegressor
        from sklearn.ensemble import GradientBoostingRegressor
        from tqdm import tqdm
        max_depths = [2,4,8,16,32]
        rmse1 = []
        rmse2 = []
        rmse3 = []
        rmse4 = []
        rmse5 = []
        rmse6 = []
        for depth in tqdm(max_depths):
          model1 = DecisionTreeRegressor(random_state=0, max_depth=depth)
          model2 = RandomForestRegressor(random_state=0, max_depth=depth, max_featur
          model3 = GradientBoostingRegressor(random state=0, max depth=depth)
          model1.fit(x_train, y_train)
          y_pred1 = model1.predict(x_train)
          rmse1.append(np.sqrt(np.mean((y_train-y_pred1)**2)))
          y_pred2 = model1.predict(x_val)
          rmse2.append(np.sqrt(np.mean((y_val-y_pred2)**2)))
          model2.fit(x train, y train)
          y_pred3 = model2.predict(x_train)
          rmse3.append(np.sqrt(np.mean((y_train-y_pred3)**2)))
```

```
y_pred4 = model2.predict(x_val)
rmse4.append(np.sqrt(np.mean((y_val-y_pred4)**2)))

model3.fit(x_train, y_train)
y_pred5 = model3.predict(x_train)
rmse5.append(np.sqrt(np.mean((y_train-y_pred5)**2)))
y_pred6 = model3.predict(x_val)
rmse6.append(np.sqrt(np.mean((y_val-y_pred6)**2)))

plot_depth_error(max_depths, rmse1, rmse2, rmse3, rmse4, rmse5, rmse6)
```



Part 2: MLPs with MNIST

For this part, you will want to use a GPU to improve runtime. Google Colab provides limited free GPU acceleration to all users. Go to Runtime and change Runtime Type to GPU. This will reset your compute node, so do it before starting to run other cells.

See Tips for detailed guidance on this problem.

First, use PyTorch to implement a Multilayer Perceptron network with one hidden layer (size 64) with ReLU activation. Set the network to minimize cross-entropy loss, which is

the negative log probability of the training labels given the training features. This objective function takes unnormalized logits as inputs.

Do not use MLP in sklearn for this HW - use Torch.

```
In [ ]: # initialization code
        import numpy as np
         from keras.datasets import mnist
         from matplotlib import pyplot as plt
        from scipy import stats
         import torch
         import torch.nn as nn
        def load mnist():
           Loads, reshapes, and normalizes the data
           (x_train, y_train), (x_test, y_test) = mnist.load_data() # loads MNIST dat
          x train = np.reshape(x train, (len(x train), 28*28)) # reformat to 768-d
          x_{\text{test}} = \text{np.reshape}(x_{\text{test}}, (\text{len}(x_{\text{test}}), 28*28))
          maxval = x train.max()
          x_train = x_train/maxval # normalize values to range from 0 to 1
          x_{test} = x_{test/maxval}
           return (x_train, y_train), (x_test, y_test)
        def display_mnist(x, subplot_rows=1, subplot_cols=1):
          Displays one or more examples in a row or a grid
           if subplot_rows>1 or subplot_cols>1:
             fig, ax = plt.subplots(subplot_rows, subplot_cols, figsize=(15,15))
             for i in np.arange(len(x)):
               ax[i].imshow(np.reshape(x[i], (28,28)), cmap='gray')
               ax[i].axis('off')
          else:
               plt.imshow(np.reshape(x, (28,28)), cmap='gray')
               plt.axis('off')
           plt.show()
```

```
In []: # Sets device to "cuda" if a GPU is available (in Colabs, enable GPU by Edi
device = "cuda" if torch.cuda.is_available() else 'cpu'
print(device) # make sure you're using GPU instance
```

cpu

2a

Using the train/val split provided in the starter code, train your network for 100 epochs

with learning rates of 0.01, 0.1, and 1. Use a batch size of 256 and the SGD optimizer. After each epoch, record the mean training and validation loss and compute the validation error of the final model. The mean validation loss should be computed after the epoch is complete. The mean training loss can either be computed after the epoch is complete, or, for efficiency, computed using the losses accumulated during the training of the epoch. Plot the training and validation losses using the display_error_curves function.

```
In [ ]: (x_train, y_train), (x_test, y_test) = load_mnist()
        # create train/val split
        ntrain = 50000
        x_val = x_train[ntrain:].copy()
        y_val = y_train[ntrain:].copy()
        x_train = x_train[:ntrain]
        y train = y train[:ntrain]
In [ ]: def display_error_curves(training_losses, validation_losses):
          Plots the training and validation loss curves
          training losses and validation losses should be lists or arrays of the same
          num epochs = len(training losses)
          plt.plot(range(num_epochs), training_losses, label="Training Loss")
          plt.plot(range(num_epochs), validation_losses, label="Validation Loss")
          # Add in a title and axes labels
          plt.title('Training and Validation Loss')
          plt.xlabel('Epochs')
          plt.ylabel('Loss')
          # Display the plot
          plt.legend(loc='best')
          plt.show()
In [ ]: # Define the model
        class MLP(nn.Module):
            def __init__(self, input_size, hidden_size, output_size):
                super(MLP, self).__init__()
                # Needs code here
                self.fc1 = nn.Linear(input_size, hidden_size)
                self.relu = nn.ReLU()
                self.fc2 = nn.Linear(hidden_size, output_size)
            def forward(self, x):
              # Needs code here
```

```
x = self.fc1(x)
x = self.relu(x)
x = self.fc2(x)
return x
```

```
In [ ]: # This is a possible function definition for training MLP, but feel free to
        # You may also want to create helper functions, e.g. for computing loss or p
        import torch
        import torch.nn as nn
        import torch.optim as optim
        def train_MLP_mnist(train_loader, val_loader, lr=1e-1, num_epochs=100):
          \mathbf{I}_{-}\mathbf{I}_{-}\mathbf{I}_{-}
          Train a MLP
          Input: train_loader and val_loader are dataloaders for the training and
          val data, respectively. Ir is the learning rate, and the network will
          be trained for num epochs epochs.
          Output: return a trained MLP
          # TODO: fill in all code
          input_size = 784
          hidden size = 64
          output_size = 10
          # Instantiate the model
          mlp = MLP(input size, hidden size, output size)
          optimizer = optim.SGD(mlp.parameters(), lr=lr)
          training_losses, validation_losses = [], []
          loss_function = torch.nn.CrossEntropyLoss()
          # Train the model, compute and store train/val loss at each epoch
          for epoch in tqdm(range(0, num_epochs), desc=f"learning rate = {lr}"):
            # Iterate over the DataLoader for training data
            sum train = 0
            sum_val = 0
            for i, data in enumerate(train_loader, 0):
               inputs, targets = data # Get inputs
               optimizer.zero_grad() # Zero the gradients
               outputs = mlp(inputs) # Compute logit scores for current batch
               loss = loss function(outputs, targets) # Compute loss
               loss.backward() # Backprop loss
               optimizer.step() # Update weights
               sum_train += loss
            for i, data in enumerate(val_loader, 0):
               inputs, targets = data # Get inputs
               optimizer.zero_grad() # Zero the gradients
```

```
outputs = mlp(inputs) # Compute logit scores for current batch
      loss = loss_function(outputs, targets) # Compute loss
      sum_val += loss
    # print(type(sum_train))
    training losses.append(int(sum train))
    validation_losses.append(int(sum_val))
  training_losses = np.array(training_losses)/num_epochs
  validation_losses = np.array(validation_losses)/num_epochs
 # Display Loss Curves
 display_error_curves(training_losses, validation_losses)
  return mlp, validation losses
def evaluate_MLP(mlp, loader):
  ''' Computes loss and error rate given your mlp model and data loader'''
 N = 0
 acc = 0
  loss = 0
  loss_function = torch.nn.CrossEntropyLoss()
 with torch.set_grad_enabled(False):
    for i, data in enumerate(loader, 0):
     # Get inputs
      inputs, targets = data
     N += len(targets)
      # Perform forward pass
      outputs = mlp(inputs.to(device))
      # Compute sum of correct labels
      y_pred = np.argmax(outputs.cpu().numpy(), axis=1)
      y_gt = np.argmax(targets.numpy(), axis=1)
      acc += np.sum(y_pred==y_gt)
      # Compute loss
      loss += loss_function(outputs, targets.to(device)).item()*len(targets)
  loss /= N
  acc /= N
  return loss, 1-acc
```

```
In []: # Code for running experiments

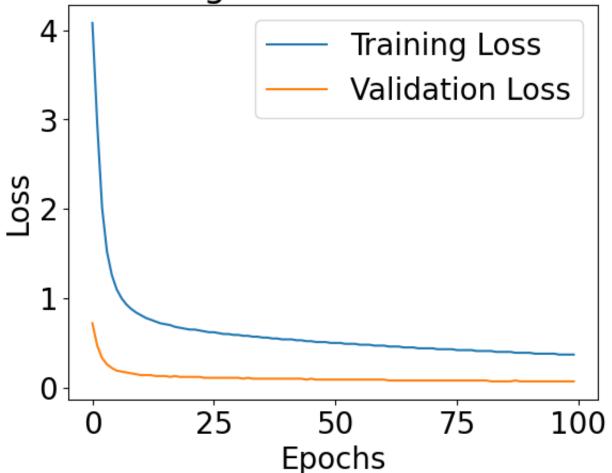
print(device) # make sure you're using GPU instance
torch.manual_seed(0) # to avoid randomness, but if you wanted to create an e
```

```
# TODO (set up dataloaders, and call training function)
trainset = torch.utils.data.TensorDataset(torch.Tensor(x_train), torch.Tensot
train_loader = torch.utils.data.DataLoader(trainset, batch_size=256, shuffle)
valset = torch.utils.data.TensorDataset(torch.Tensor(x_val), torch.Tensor(np.val_loader = torch.utils.data.DataLoader(valset, batch_size=256, shuffle=Tru.)
mlp1, loss1 = train_MLP_mnist(train_loader, val_loader, lr=0.01, num_epochs=mlp2, loss2 = train_MLP_mnist(train_loader, val_loader, lr=0.1, num_epochs=1mlp3, loss3 = train_MLP_mnist(train_loader, val_loader, lr=1, num_epochs=100)
```

cpu

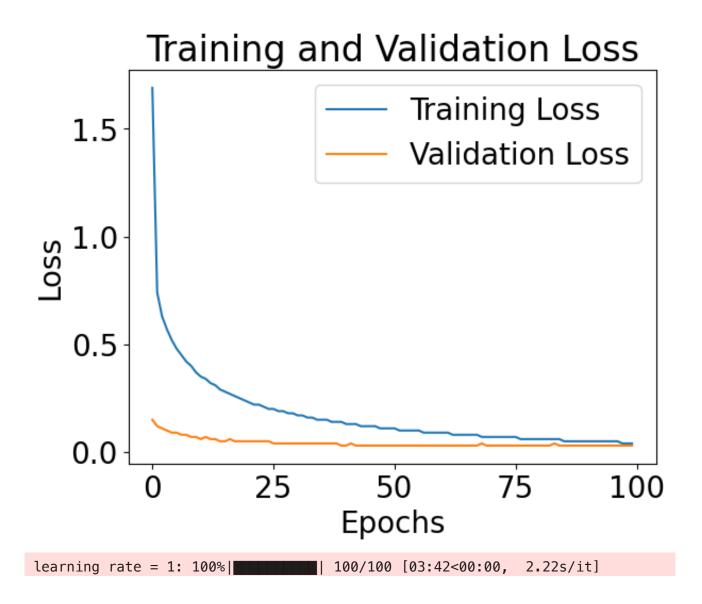
learning rate = 0.01: 100%| 100/100 [03:42<00:00, 2.23s/it]

Training and Validation Loss

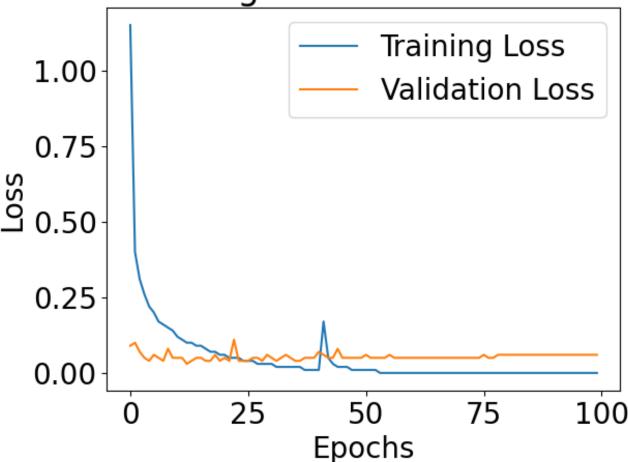


learning rate = 0.1: 100%||

| 100/100 [03:39<00:00, 2.20s/it







```
In []: ind1 = np.argmax(loss1)
    print(f"max loss is {loss1[ind1]} for index = {ind1}")
    ind2 = np.argmax(loss2)
    print(f"max loss is {loss2[ind2]} for index = {ind2}")
    ind3 = np.argmax(loss3)
    print(f"max loss is {loss3[ind3]} for index = {ind3}")

max loss is 0.72 for index = 0
    max loss is 0.15 for index = 0
    max loss is 0.11 for index = 22
```

Based on the loss curves, select the learning rate and number of epochs that minimizes the validation loss. Retrain that model (if it's not stored), and report training loss, validation loss, training error, validation error, and test error.

```
In []: # TO DO (retrain if needed, and evaluate model on train, val, and test sets)
  testset = torch.utils.data.TensorDataset(torch.Tensor(x_test), torch.Tensor(
  test_loader = torch.utils.data.DataLoader(testset, batch_size=256, shuffle=1
```

```
mlp, _ = train_MLP_mnist(train_loader, val_loader, lr=1, num_epochs=23)
val_loss, val_err = evaluate_MLP(mlp, val_loader)
test_loss, test_err = evaluate_MLP(mlp, test_loader)
train_loss, train_err = evaluate_MLP(mlp, train_loader)
print(f"validation loss: {val_loss}, validation error: {val_err}")
print(f"test loss: {test_loss}, test error: {test_err}")
print(f"train loss: {train_loss}, train error: {train_err}")
```

learning rate = 1: 100%| 23/23 [00:50<00:00, 2.19s/it]

Part 3: Predicting Penguin Species

Include all your code for part 3 in this section.

```
In [ ]: import numpy as np
from matplotlib import pyplot as plt
```

```
import pandas as pd
import seaborn as sns
#styling preferences for sns
sns.set_style('whitegrid')
sns.set_context('poster')
datadir = "./"
df_penguins = pd.read_csv(datadir + 'penguins_size.csv')
df penguins.head(10)
# convert features with multiple string values to binary features so they ca
def get_penguin_xy(df_penguins):
 data = np.array(df_penguins[['island', 'culmen_length_mm', 'culmen_depth_m'
 y = df penguins['species']
 ui = np.unique(data[:,0]) # unique island
  us = np.unique(data[:,-1]) # unique sex
 X = np.zeros((len(y), 10))
  for i in range(len(y)):
    f = 0
    for j in range(len(ui)):
      if data[i, f]==ui[j]:
        X[i, f+j] = 1
    f = f + len(ui)
   X[i, f:(f+4)] = data[i, 1:5]
    f=f+4
    for j in range(len(us)):
      if data[i, 5]==us[j]:
        X[i, f+j] = 1
  feature_names = ['island_biscoe', 'island_dream', 'island_torgersen', 'cul
 X = pd.DataFrame(X, columns=feature names)
  return(X, y, feature_names, np.unique(y))
```

```
In [ ]: get_penguin_xy(df_penguins)[0].to_csv("X.csv")
```

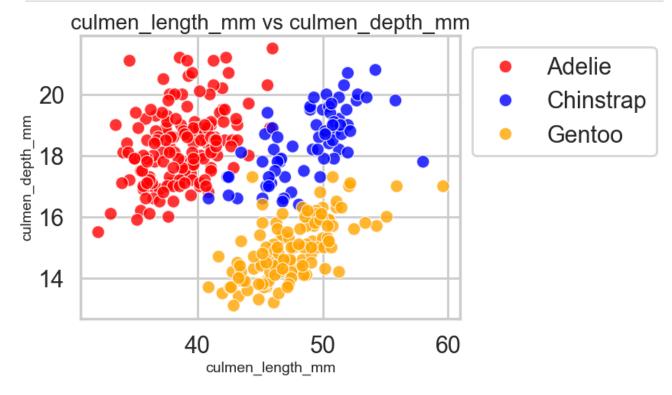
3a

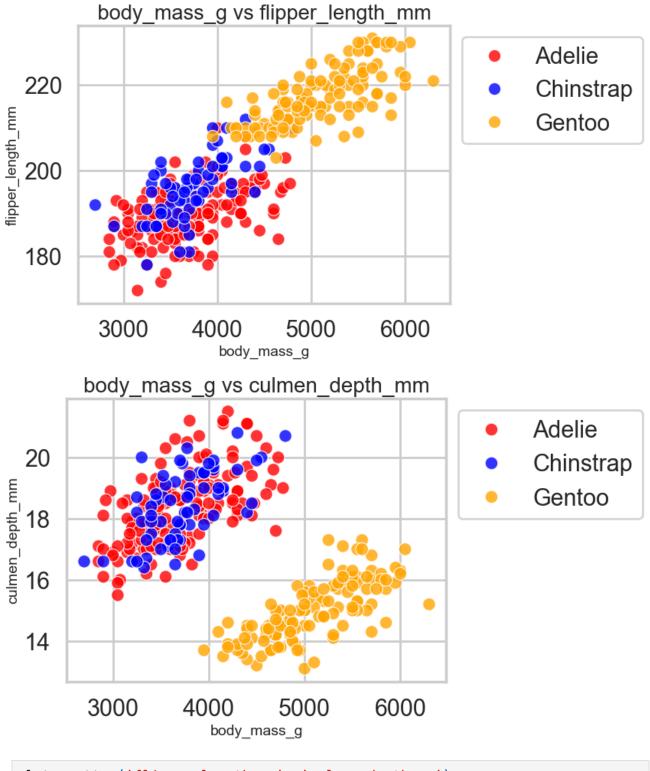
Spend some time to visualize different pairs of features and their relationships to the species. We've done one for you. Include in your report at least two other visualizations.

```
hue = 'species', palette=palette, alpha=0.8)
# Doc: https://seaborn.pydata.org/generated/seaborn.scatterplot.html

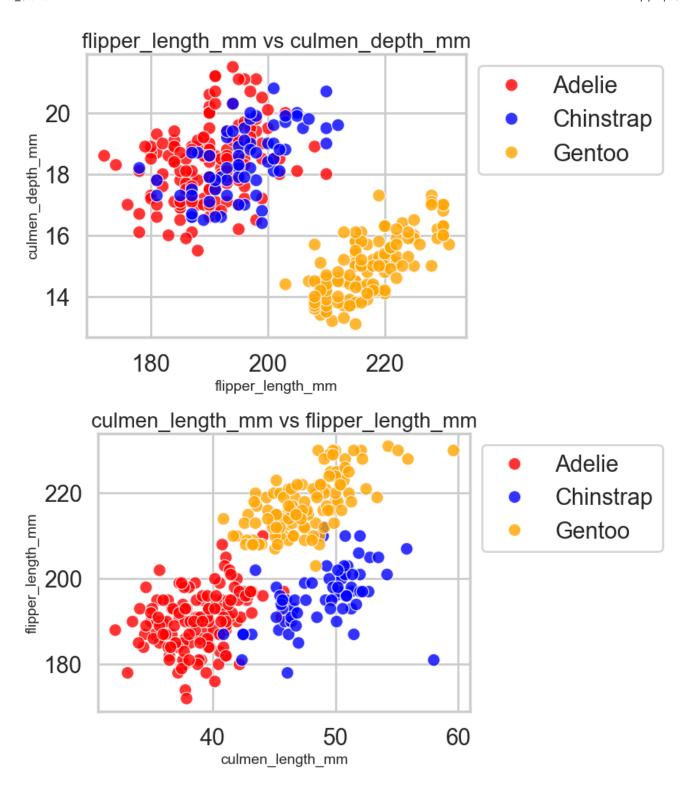
plt.xlabel(feature1, fontsize=14)
plt.ylabel(feature2, fontsize=14)
plt.title(feature1 + ' vs ' + feature2, fontsize=20)
plt.legend(bbox_to_anchor=(1.0, 1.0), loc='upper left')
plt.show()

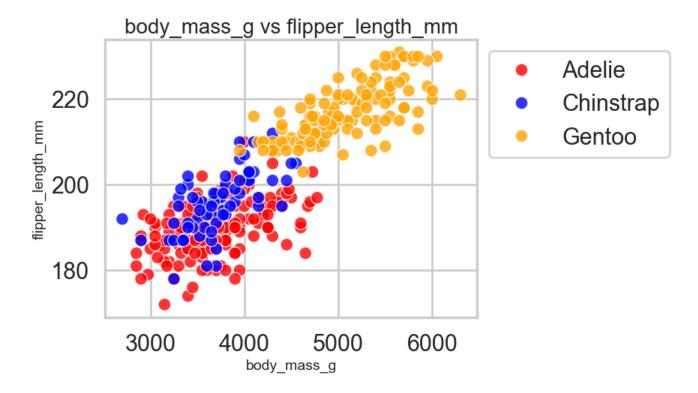
# TO DO call plot_scatter with different feature pairs to create some visual
plot_scatter('culmen_length_mm', 'culmen_depth_mm')
plot_scatter('body_mass_g', 'flipper_length_mm')
plot_scatter('body_mass_g', 'culmen_depth_mm')
```





```
In []: plot_scatter('flipper_length_mm', 'culmen_depth_mm')
    plot_scatter('culmen_length_mm', 'flipper_length_mm')
    plot_scatter('body_mass_g', 'flipper_length_mm')
```



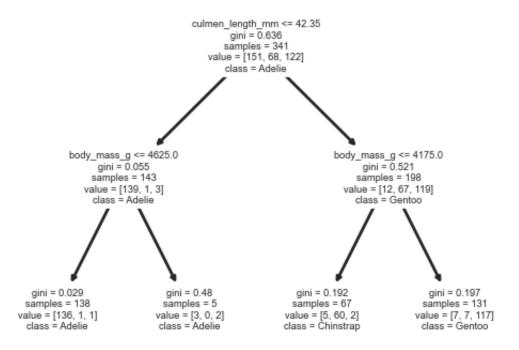


3b

Suppose you want to be able to identify the Gentoo species with a simple rule with very high accuracy. Use a decision tree classifier to figure out such a rule that has only two checks (e.g. "mass greater than 4000 g, and culmen length less than 40 mm is Gentoo; otherwise, not"). You can use the library DecisionTreeClassifier with either 'gini' or 'entropy' criterion. Use sklearn.tree.plot_tree with feature_names and class_names arguments to visualize the decision tree. Include the tree that you used to find the rule in your report and the rule.

```
In []: # TO DO (Train a short tree to identify a good rule, plot the tree, report t
from sklearn.tree import DecisionTreeClassifier
from sklearn.tree import plot_tree

classify = DecisionTreeClassifier(max_depth=2)
features = df_penguins[["body_mass_g", "culmen_length_mm"]]
target = df_penguins["species"]
classify.fit(features, target)
plot_tree(classify, feature_names=["body_mass_g", "culmen_length_mm"], class
```



```
In []: pred = classify.predict(features)

print(np.sum((pred=="Gentoo")&(target=="Gentoo")))
print(pred.shape[0])
print(np.sum((pred=="Gentoo")&(target=="Gentoo")))
print(np.sum(target=="Gentoo"))
```

117

341

117

122

3c

Use any method at your disposal to achieve maximum 5-fold cross-validation accuracy on this problem. To keep it simple, we will use sklearn.model_selection to perform the cross-validation for us. Report your model design and 5-fold accuracy. It is possible to get more than 99% accuracy.

```
In []: # design a classification model, import libraries as needed
        from sklearn.model_selection import cross_val_score
        from sklearn.linear_model import LogisticRegression
        import warnings
        # Ignore all warnings
        warnings.filterwarnings("ignore")
        X, y, feature_names, class_names = get_penguin_xy(df_penguins)
        # TO DO -- choose some model and fit the data
        model = LogisticRegression()
        scores = cross_val_score(model, np.array(X), np.array(y), cv=5)
        print('CV Accuracy: {}'.format(scores.mean()))
        from sklearn.tree import DecisionTreeClassifier
        model = DecisionTreeClassifier()
        scores = cross_val_score(model, np.array(X), np.array(y), cv=5)
        print('CV Accuracy: {}'.format(scores.mean()))
        from sklearn.ensemble import RandomForestClassifier
        model = RandomForestClassifier()
        scores = cross_val_score(model, np.array(X), np.array(y), cv=5)
        print('CV Accuracy: {}'.format(scores.mean()))
        from sklearn.svm import SVC
        model = SVC()
        scores = cross_val_score(model, np.array(X), np.array(y), cv=5)
        print('CV Accuracy: {}'.format(scores.mean()))
        from sklearn.neighbors import KNeighborsClassifier
        model = KNeighborsClassifier()
        scores = cross_val_score(model, np.array(X), np.array(y), cv=5)
        print('CV Accuracy: {}'.format(scores.mean()))
       CV Accuracy: 0.9883205456095482
       CV Accuracy: 0.9764705882352942
       CV Accuracy: 0.9911764705882353
       CV Accuracy: 0.7331628303495311
```

CV Accuracy: 0.7890451832907076

Part 4: Stretch Goals

Include any new code needed for Part 4 here

```
In []: # TO DO (optional)
        # TO DO (optional)
        import numpy as np
         from keras.datasets import mnist
        from matplotlib import pyplot as plt
         from scipy import stats
         import torch
         import torch.nn as nn
        def load mnist():
           \mathbf{I} \cdot \mathbf{I} \cdot \mathbf{I}
           Loads, reshapes, and normalizes the data
           (x_train, y_train), (x_test, y_test) = mnist.load_data() # loads MNIST dat
          x_train = np.reshape(x_train, (len(x_train), 28*28)) # reformat to 768-d
          x_{test} = np.reshape(x_{test}, (len(x_{test}), 28*28))
          maxval = x train.max()
          x train = x train/maxval # normalize values to range from 0 to 1
          x_{test} = x_{test/maxval}
           return (x_train, y_train), (x_test, y_test)
         (x_train, y_train), (x_test, y_test) = load_mnist()
        # create train/val split
         ntrain = 50000
        x_val = x_train[ntrain:].copy()
        y_val = y_train[ntrain:].copy()
        x_train = x_train[:ntrain]
        y_train = y_train[:ntrain]
        def display_error_curves(training_losses, validation_losses):
           Plots the training and validation loss curves
           training_losses and validation_losses should be lists or arrays of the sam
           num_epochs = len(training_losses)
           plt.plot(range(num_epochs), training_losses, label="Training Loss")
           plt.plot(range(num epochs), validation losses, label="Validation Loss")
           # Add in a title and axes labels
           plt.title('Training and Validation Loss')
           plt.xlabel('Epochs')
           plt.ylabel('Loss')
```

```
# Display the plot
  plt.legend(loc='best')
  plt.show()
# Define the model
class MLP(nn.Module):
    def __init__(self, input_size, hidden_size, output_size):
        super(MLP, self).__init__()
        self.fc1 = nn.Linear(input_size, 2*hidden_size)
        self.relu1 = nn.ReLU()
        self.fc2 = nn.Linear(2*hidden_size, hidden_size)
        self.relu2 = nn.ReLU()
        self.fc3 = nn.Linear(hidden size, hidden size//2)
        self.relu3 = nn.ReLU()
        self.fc4 = nn.Linear(hidden_size//2, output_size)
    def forward(self, x):
        x = self.fc1(x)
        x = self.relu1(x)
        x = self.fc2(x)
        x = self.relu2(x)
        x = self.fc3(x)
        x = self.relu3(x)
        x = self.fc4(x)
        return x
# This is a possible function definition for training MLP, but feel free to
# You may also want to create helper functions, e.g. for computing loss or p
import torch
from tqdm import tqdm
import torch.nn as nn
import torch.optim as optim
def train_MLP_mnist(train_loader, val_loader, lr=1e-1, num_epochs=100):
 Train a MLP
  Input: train_loader and val_loader are dataloaders for the training and
 val data, respectively. Ir is the learning rate, and the network will
  be trained for num_epochs epochs.
  Output: return a trained MLP
 # TODO: fill in all code
  input size = 784
  hidden size = 64
  output_size = 10
  # Instantiate the model
  mlp = MLP(input_size, hidden_size, output_size)
```

```
optimizer = optim.Adam(mlp.parameters(), lr=lr)
  training_losses, validation_losses = [], []
  loss_function = torch.nn.CrossEntropyLoss()
 # Train the model, compute and store train/val loss at each epoch
  for epoch in tgdm(range(0, num epochs), desc=f"learning rate = {lr}"):
    # Iterate over the DataLoader for training data
    sum_train = 0
    sum_val = 0
    for i, data in enumerate(train_loader, 0):
      inputs, targets = data # Get inputs
      optimizer.zero grad() # Zero the gradients
      outputs = mlp(inputs) # Compute logit scores for current batch
      loss = loss_function(outputs, targets) # Compute loss
      loss.backward() # Backprop loss
      optimizer.step() # Update weights
      sum_train += loss
    for i, data in enumerate(val_loader, 0):
      inputs, targets = data # Get inputs
      optimizer.zero_grad() # Zero the gradients
      outputs = mlp(inputs) # Compute logit scores for current batch
      loss = loss_function(outputs, targets) # Compute loss
      sum_val += loss
    # print(type(sum_train))
    training losses.append(int(sum train))
    validation_losses.append(int(sum_val))
 # Display Loss Curves
 # display_error_curves(training_losses, validation_losses)
  return mlp, validation_losses
def evaluate_MLP(mlp, loader):
  ''' Computes loss and error rate given your mlp model and data loader'''
 N = 0
  acc = 0
  loss = 0
  loss_function = torch.nn.CrossEntropyLoss()
 with torch.set_grad_enabled(False):
    for i, data in enumerate(loader, 0):
      # Get inputs
      inputs, targets = data
     N += len(targets)
     # Perform forward pass
```

```
outputs = mlp(inputs.to(device))
      # Compute sum of correct labels
      y_pred = np.argmax(outputs.cpu().numpy(), axis=1)
      y_gt = np.argmax(targets.numpy(), axis=1)
      acc += np.sum(y pred==y qt)
      # Compute loss
      loss += loss_function(outputs, targets.to(device)).item()*len(targets)
  loss /= N
  acc /= N
  return loss, 1-acc
# Code for running experiments
# print(device) # make sure you're using GPU instance
torch.manual_seed(0) # to avoid randomness, but if you wanted to create an e
# TODO (set up dataloaders, and call training function)
trainset = torch.utils.data.TensorDataset(torch.Tensor(x_train), torch.Tenso
train_loader = torch.utils.data.DataLoader(trainset, batch_size=256, shuffle
valset = torch.utils.data.TensorDataset(torch.Tensor(x_val), torch.Tensor(ng
val_loader = torch.utils.data.DataLoader(valset, batch_size=256, shuffle=Tru
mlp1, loss1 = train_MLP_mnist(train_loader, val_loader, lr=0.001, num_epochs
mlp2, loss2 = train_MLP_mnist(train_loader, val_loader, lr=0.01, num_epochs=
mlp3, loss3 = train_MLP_mnist(train_loader, val_loader, lr=0.1, num_epochs=1
ind1 = np.argmax(loss1)
print(f"max loss is {loss1[ind1]} for index = {ind1}")
ind2 = np.argmax(loss2)
print(f"max loss is {loss2[ind2]} for index = {ind2}")
ind3 = np.argmax(loss3)
print(f"max loss is {loss3[ind3]} for index = {ind3}")
# TO DO (retrain if needed, and evaluate model on train, val, and test sets)
# Sets device to "cuda" if a GPU is available (in Colabs, enable GPU by Edi
device = "cuda" if torch.cuda.is_available() else 'cpu'
print(device) # make sure you're using GPU instance
testset = torch.utils.data.TensorDataset(torch.Tensor(x_test), torch.Tensor(
test loader = torch.utils.data.DataLoader(testset, batch size=256, shuffle=1
if loss1[ind1]>loss2[ind2]:
  if loss1[ind1]>loss3[ind3]:
    lr=0.001
    num epochs=ind1
  else:
    lr=0.1
```

```
num_epochs=ind3
                 else:
                     if loss2[ind2]>loss3[ind3]:
                         1r=0.01
                         num_epochs=ind2
                     else:
                         lr=0.1
                         num epochs=ind3
                 mlp, _ = train_MLP_mnist(train_loader, val_loader, lr=lr, num_epochs=num_epochs
                 val_loss, val_err = evaluate_MLP(mlp, val_loader)
                 test_loss, test_err = evaluate_MLP(mlp, test_loader)
                 train_loss, train_err = evaluate_MLP(mlp, train_loader)
                 print(f"validation loss: {val loss}, validation error: {val err}")
                 print(f"test loss: {test loss}, test error: {test err}")
                 print(f"train loss: {train_loss}, train error: {train_err}")
              learning rate = 0.001: 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100\% | 100
                                                                         100/100 [07:56<00:00, 4.76s/it]
              learning rate = 0.1: 100%
              \max loss is 11 for index = 0
              max loss is 11 for index = 83
              max loss is 81 for index = 18
              cpu
              learning rate = 0.1: 100% | 18/18 [01:27<00:00, 4.85s/it]
              validation loss: 0.7412997485160827, validation error: 0.2164000000000004
              test loss: 0.7280186690330506, test error: 0.2221999999999995
              train loss: 0.7102787209129333, train error: 0.21914
In [ ]: import numpy as np
                 from matplotlib import pyplot as plt
                 import pandas as pd
                 import seaborn as sns
                 #styling preferences for sns
                 sns.set style('whitegrid')
                 sns.set context('poster')
                 datadir = "./"
                 df_penguins = pd.read_csv(datadir + 'penguins_size.csv')
                 df_penguins.head(10)
                 # convert features with multiple string values to binary features so they ca
                 def get_penguin_xy(df_penguins):
                     data = np.array(df_penguins[['island', 'culmen_length_mm', 'culmen_depth_m
                     y = df_penguins['species']
                     ui = np.unique(data[:,0]) # unique island
                     us = np.unique(data[:,-1]) # unique sex
                    X = np.zeros((len(y), 10))
                     for i in range(len(y)):
                         f = 0
                         for j in range(len(ui)):
                             if data[i, f]==ui[j]:
```

```
X[i, f+j] = 1
    f = f + len(ui)
   X[i, f:(f+4)] = data[i, 1:5]
   f=f+4
   for j in range(len(us)):
      if data[i, 5]==us[j]:
        X[i, f+j] = 1
  feature_names = ['island_biscoe', 'island_dream', 'island_torgersen', 'cul
 X = pd.DataFrame(X, columns=feature_names)
  return(X, y, feature_names, np.unique(y))
# TO DO (Train a short tree to identify a good rule, plot the tree, report t
from sklearn.tree import DecisionTreeClassifier
from sklearn.tree import plot tree
classify = DecisionTreeClassifier(max depth=2)
features = df_penguins[["flipper_length_mm", "culmen_length_mm"]]
target = df_penguins["species"]
classify.fit(features, target)
plot_tree(classify, feature_names=["flipper_length_mm", "culmen_length_mm"],
pred = classify.predict(features)
print(np.sum((pred=="Gentoo")&(target=="Gentoo")))
print(pred.shape[0])
print(np.sum((pred=="Gentoo")&(target=="Gentoo")))
print(np.sum(target=="Gentoo"))
```

121341

121

122

