1)	
	"I certify that all solutions in this document are entirely my own
	"I certify that all solutions in this document are entirely my own and that I have not looked at anyone elsis solution. I have
	given credit to cell external source I consulter."
	U
	XMU

Went to office hours, helped/got help from TAs and various students.

$$E[\frac{1}{n}\sum_{i=0}^{i\leq n}Y_i]=\frac{1}{n}\sum_{i=0}^{i\leq n}E[Y_i]=u$$

$$Vor\left(\frac{1}{n}\sum_{i=0}^{i\in n}Y_i\right)=\frac{1}{n^2}\sum_{i=0}^{i\in n}Vor(Y_i)=\frac{\sigma^2}{n}$$

$$Vor(\frac{1}{n}\sum_{i=0}^{i n} Z_i) = \frac{\sigma^2}{n} \cdot \rho n = \sigma^2 \rho$$

If p is very large then that means the correlation between Zi is very controlling and coveraging will do less. If p is very small, then overaging will be effective and will cost closer to uncorrelated variable.

Middling is in the middle

$$1-e^{-1}$$
 is prob. of gratting chorn $= 0.63$

117	Cross-Nalidation. Run at different n'value and
	ison at different in value and
	see which one delivers the best
	performance

3) a) If
$$K=T$$
, then $\alpha=\gamma=0$ minimizes $\alpha_1=1$

b) h contains all 1

Normal Equation

a e orgmin 11 Ka-y112

Xxn= x, 1 1h 0=1

Kisall ls so a=0 to minimize

in order to get y=0 a=0 so K(0) = 0.

h(2)= | due to all 14 equalling 0.

c)
$$K = \begin{bmatrix} 1 + \frac{X_1 X_1}{2\sigma^2} & 1 + \frac{X_1 X_2}{2\sigma^2} \\ 1 + \frac{X_2 X_1}{2\sigma^2} & 1 + \frac{X_2 X_2}{2\sigma^2} \end{bmatrix}$$

or & organin 11 Ka-y112 if 1/2)

Y has the seeme property as parts

Normal Equation

KT Ka=KTY

 $a = (K^{1}K)K^{1}y = \begin{bmatrix} -\sigma^{2} \\ -\sigma^{2} \end{bmatrix}$ $h(2) = Sign\left(\sum_{i=1}^{n} \alpha_{i} \frac{x_{i}^{2}}{2\sigma^{2}} + 1\right) = \frac{1}{2\sigma^{2}}$

 $o^{2}\frac{|(1)|}{2\sigma^{2}}$ $-o^{2}\frac{(-1)(-1)}{2\sigma^{2}}$

h62)=1

```
oredict(self, X):
collect = [self.traverse(i, self.tree) for i in X]
return no.array(collect)
    rapy(self, y):
_af_ys = Counter(y)
                in x_cols:
hat_to_gain = self.info(y, X_column, 1)
if what_to_gain = best:
best = what_to_gain
index = i
thresh = i
ddex, thresh
       n left_index, right_index
```

4.2

- 1. How did you deal with categorical features and missing values?
 - a. Substituted for categorical features the most frequent occurrence, based on the idea that most likely, the missing features would most likely be the most common feature. For the missing values, replaced them with median value.
 - b. I completed them by using scikit pipeline coupled with simpleimputer, onehotencoder, and ColumnTransformer.
- 2. What was your stopping criterion?
 - a. When depth was hit, it is a tunable hyperparameter.
- 3. How did you implement random forests?
 - a. Leveraged the decision tree. The random forest implementation is basically just a decision tree in disguise.
- 4. Did you do anything special to speed up training?
 - a. Nothing special other than changing the hyperparameters (slowly and painfully)
- 5. Anything else cool you implemented?
 - a. I'm proud of my pre-processing, took me a very long time to figure it out and a lot of reading python libraries commands.

4.4

Titanic Decision Tree Training Accuracy 0.8314745972738538
Titanic Decision Tree Validation Accuracy 0.801980198019802
Titanic Random Forest Training Accuracy 0.781908302354399
Titanic Random Forest Training Accuracy 0.792079207921

Spam Data Decision tree Validation: 0.8419182948490231 Spam Data Decision tree Training: 0.8247834776815456 Spam Data RandomForest Validation: 0.7655417406749556 Spam Data RandomForest Training: 0.737508327781479

Spam Kaggle: 0.817 Titanic Kaggle: 0.822

```
def print_tree(node, depth = 1):
    if node.is_leaf():
        print("\t" * depth + "Leaf: ", node.feature)
    else:
        print("\t" * depth + "Node: ", node.feature, " = ", node.threshold)
        if node.right is not None:
            print("\t" * depth + "Right: ")
            print_tree(node.right, depth + 1)

        if node.left is not None:
            print("\t" * depth + "Left: ")
            print_tree(node.left, depth + 1)

decision_tree = DecisionTree(2)
    decision_tree.fit(X_train_np, y_train_np)

print_tree(decision_tree.tree)
```

```
import matplotlib.pyplot as plt
     depths = range(1, 40)
     accuracies = []
     X_train_np = preprocessor.transform(X_train)
     y_train_np = y_train
     X_val_np = preprocessor.transform(X_val)
     y_val_np = y_val
     for depth in depths:
         model.fit(X_train_np, y_train_np)
         y_pred = model.predict(X_val_np)
         accuracy = accuracy_score(y_val, y_pred)
         accuracies.append(accuracy)
     plt.figure(figsize=(5, 5))
     plt.plot(depths, accuracies)
     plt.xlabel('Depth')
     plt.ylabel('Accuracy')
     plt.show()
  0.806
  0.804
Accuracy
  0.802
  0.800
  0.798
```

Depth 10, 20, 27, 34, 37 gives the highest accuracy. This is broken. It should not be like this. I do not know why, I swear it was working earlier.

15

20

Depth

25

30

35

40

5

10

```
def print_tree(node, depth = 1):
    if node.is_leaf():
        print("\t" * depth + "Leaf: ", node.feature)

else:
        print("\t" * depth + "Node: ", node.feature, " = ", node.threshold)

        if node.right is not None:
            print("\t" * depth + "Right: ")
            print_tree(node.right, depth + 1)

        if node.left is not None:
            print("\t" * depth + "Left: ")
            print_tree(node.left, depth + 1)

decision_tree = DecisionTree(2)
decision_tree.fit(X_train_np, y_train_np)

print_tree(decision_tree.tree)
```

References:

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