CS294 Midterm GitHub Link: https://github.com/angusYuhao/CS294-midterm 6.1:

a)

Binary case:

```
[22] num_exp = 20
     test_dimensions = [2, 4, 8]
     for dim in test_dimensions:
         n_avg = 0
         for i in range(num_exp):
            dataset = generate_random_dataset(dim, n_full, 2)
             X = dataset.drop('target', axis=1)
             y = dataset['target']
             X train = X.copy()
             y_train = y.copy()
             for idx in range(n_full):
                 X_train = X.drop(index=idx)
                 y_train = y.drop(index=idx)
                 knn = KNeighborsClassifier(n neighbors=1)
                 knn.fit(X_train, y_train)
                 y_pred = knn.predict(X)
                 different = False
                 for i in range(len(y)):
                     if y[i] != y_pred[i]:
                         different = True
                         break
                 if different:
                     n avg += 1
         n_avg /= num_exp
         print(f"d={dim}: n_full={n_full}, Avg. req. points for memorization n_avg={n_avg:.2f}, n_full/n_avg={n_full/n_avg}")
     d=4: n_full=16, Avg. req. points for memorization n_avg=8.20, n_full/n_avg=1.9512195121951221 d=8: n_full=256, Avg. req. points for memorization n_avg=129.15, n_full/n_avg=1.9821912504839334
```

b)

Multiclass case:

```
num_exp = 20
test_dimensions = [2, 4, 8]
for dim in test dimensions:
    n_full = 2 ** dim
    n_avg = 0
    for i in range(num_exp):
         dataset = generate_random_dataset(dim, n_full, 5)
         X = dataset.drop('target', axis=1)
         y = dataset['target']
         X_train = X.copy()
         y train = y.copy()
         for idx in range(n_full):
             X_train = X.drop(index=idx)
             y_train = y.drop(index=idx)
             knn = KNeighborsClassifier(n_neighbors=1)
             knn.fit(X_train, y_train)
             y_pred = knn.predict(X)
             different = False
             for i in range(len(y)):
                  if y[i] != y_pred[i]:
                      different = True
              if different:
                 n_avg += 1
    n_avg /= num_exp
    print(f"d={dim}: n_full={n_full}, Avg. req. points for memorization n_avg={n_avg:.2f}, n_full/n_avg={n_full/n_avg}")
d=2: n_full=4, Avg. req. points for memorization n_avg=3.45, n_full/n_avg=1.1594202898550725
d=4: n_full=16, Avg. req. points for memorization n_avg=12.45, n_full/n_avg=1.285140562248996 d=8: n_full=256, Avg. req. points for memorization n_avg=203.35, n_full/n_avg=1.2589132038357511
```

a)

Using the heart dataset:

```
data_raw = pd.read_csv('/content/heart.csv')
X = data_raw.drop('output', axis=1)
y = data_raw['output']
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=0)
X_train.head()
                                                                                        age sex cp trtbps chol fbs restecg thalachh exng oldpeak slp caa thall
 74
      43
                     122
                           213
                                                   165
                                                                 0.2
                                                                                        ılı
153
      66
               2
                     146
                           278
                                  0
                                           0
                                                  152
                                                          0
                                                                  0.0
                                                                                   2
            0
 64
      58
                     140
                           211
                                                  165
                                                          0
                                                                  0.0
                                                                        2
                                                                             0
      63
               0
                     124
                           197
                                                                             0
                                                                                   2
296
            0
                                  0
                                                  136
                                                                  0.0
                     154
                           232
                                  0
                                           0
                                                  164
                                                          0
                                                                 0.0
                                                                        2
                                                                                   2
287
      57
```

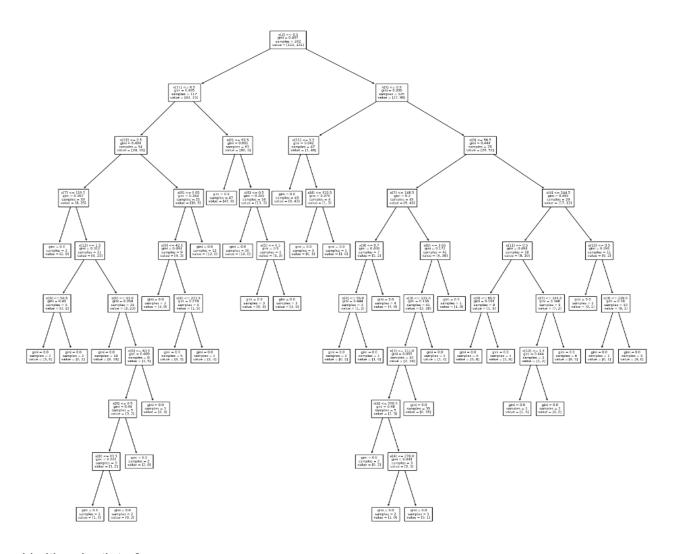
Initial decision tree:

```
dt1 = tree.DecisionTreeClassifier()

dt1.fit(X_train, y_train)

y_pred = dt1.predict(X_test)
accuracy = accuracy_score(y_test, y_pred)
print("Accuracy:", accuracy)

Accuracy: 0.7868852459016393
```

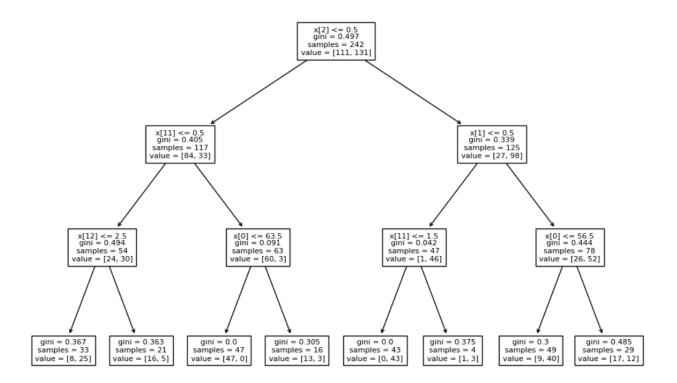


Limiting depth to 3:

```
dt2 = tree.DecisionTreeClassifier(max_depth=3)
dt2.fit(X_train, y_train)

y_pred = dt2.predict(X_test)
accuracy = accuracy_score(y_test, y_pred)
print("Accuracy:", accuracy)

Accuracy: 0.819672131147541
```



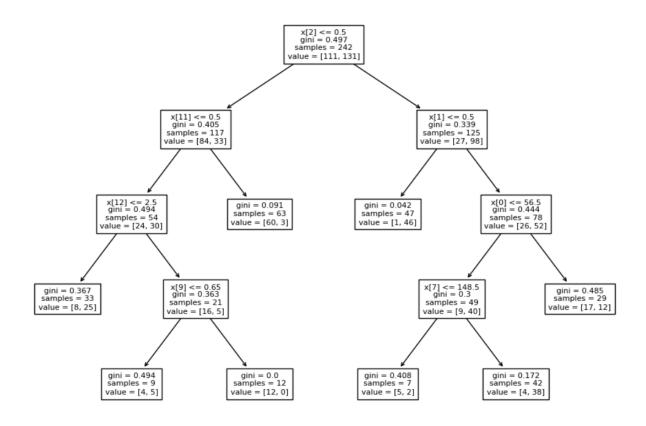
Limiting leaf nodes to 8:

```
dt3 = tree.DecisionTreeClassifier(max_leaf_nodes=8)

dt3.fit(X_train, y_train)

y_pred = dt3.predict(X_test)
accuracy = accuracy_score(y_test, y_pred)
print("Accuracy:", accuracy)

Accuracy: 0.7540983606557377
```



b) Using the stroke dataset:

<pre>data_raw = pd.read_csv('/content/healthcare-dataset-stroke-data.csv') data_raw = data_raw.dropna() X = data_raw.drop('stroke', axis=1) y = data_raw['stroke'] X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=0) X_train.head()</pre>											
	id	gender	age	hypertension	heart_disease	ever_married	work_type	Residence_type	avg_glucose_level	bmi	smoking_status
5036	57159	Male	56.0								
			30.0		0	Yes	Self-employed	Rural	125.87	24.6	never smoked
3179	23893	Male		0	0		Self-employed Private	Rural Urban	125.87 103.45		never smoked smokes
3179 4149		Male Female	24.0			Yes				25.1	
	57080		24.0 81.0			Yes Yes	Private	Urban	103.45	25.1 20.7	smokes

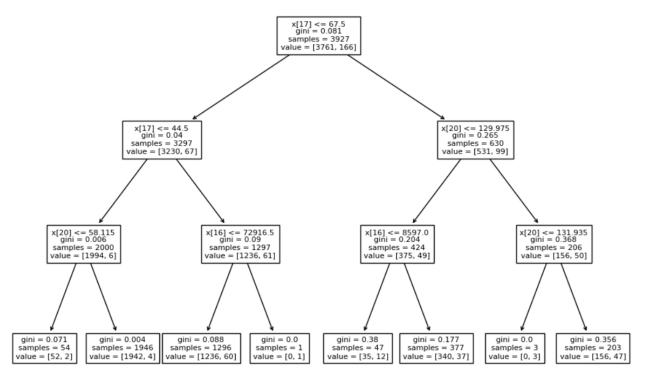
Limiting depth to 3:

```
categorical_cols = ['gender', 'ever_married', 'work_type', 'Residence_type', 'smoking_status']
preprocessor = ColumnTransformer(
    transformers=[
        ('cat', OneHotEncoder(), categorical_cols)
    ],
    remainder='passthrough'
)

pipeline = Pipeline([
        ('preprocessor', preprocessor),
        ('classifier', tree.DecisionTreeClassifier(max_depth=3))
])

pipeline.fit(X_train, y_train)
y_pred = pipeline.predict(X_test)
accuracy = accuracy_score(y_test, y_pred)
print("Accuracy:", accuracy)

Accuracy: 0.9541751527494908
```



c)
Using random dataset:

```
random_dataset = generate_random_dataset(5, 1000, 2)
X = random_dataset.drop('target', axis=1)
y = random_dataset['target']
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=0)
X_train.head()
                                                     翩
    feature_1 feature_2 feature_3 feature_4 feature_5
     687
                                                     ıl.
500 -1.438866 0.579128 -1.788463 -0.692195 -0.315527
     0.626235 -0.353869 1.370535 0.073198 0.322181
332
979 -0.330763 -0.291903 0.119736 -0.115354 -0.255583
817
     1.635701 0.703540 -0.277989 2.043371 -0.870054
```

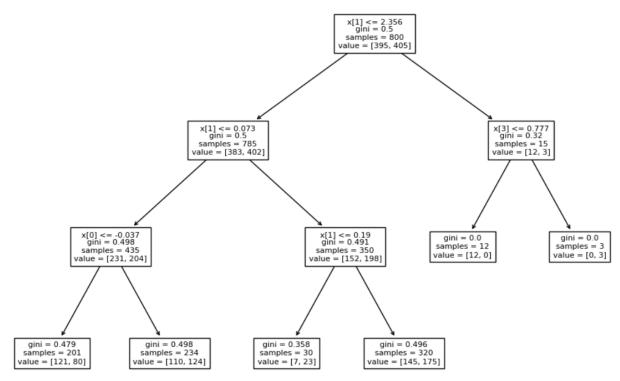
Limiting depth to 3:

```
dt5 = tree.DecisionTreeClassifier(max_depth=3)

dt5.fit(X_train, y_train)

y_pred = dt5.predict(X_test)
accuracy = accuracy_score(y_test, y_pred)
print("Accuracy:", accuracy)

Accuracy: 0.48
```



6.3:

```
[ ] original_string = generate_random_string(200)
    compressed_data = zlib.compress(original_string.encode())

    original_size = sys.getsizeof(original_string)
    compressed_size = sys.getsizeof(compressed_data)

    print("Original String:", original_string)
    print("Original String Size:", original_size)
    print("Compressed Data:", compressed_data)
    print("Compressed Data Size:", compressed_size)
    print("Compression Ratio:", compressed_size / original_size)

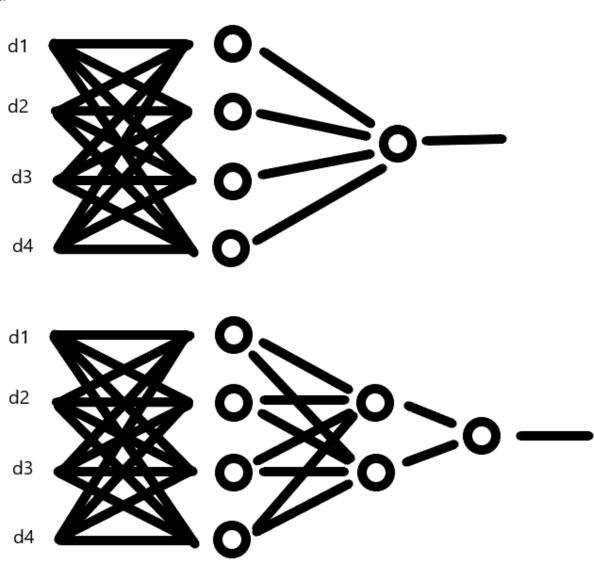
Original String: kp3ijzSQruGn4ucezwwsgvyW8vNdOYqgGZj1CHbBctSNZH7FrbyBjvbzb52YuPUBzkqY76ENvcaE0d14UDOfQ5UD2J8m8HJThyRT6Ah
    Original String Size: 249
    Compressed Data: b'x\x9c\x05\xc1\xc9\x16C0\x14\x00\xd0_2\x86m\r\x95\xe3\x10UT\xd3\x1d1\x87h\xd5P\xef\xeb{/\x7f\xab\xfd\x
Compressed Data Size: 215
    Compression Ratio: 0.8634538152610441

4
```

We see that the compression ratio is not very good. This is because the string is random, and there are no patterns/redundency for the compression algorithm to exploit. In a completely random string, the expected compression ratio should be close to 1:1.

8.1:

8.2:

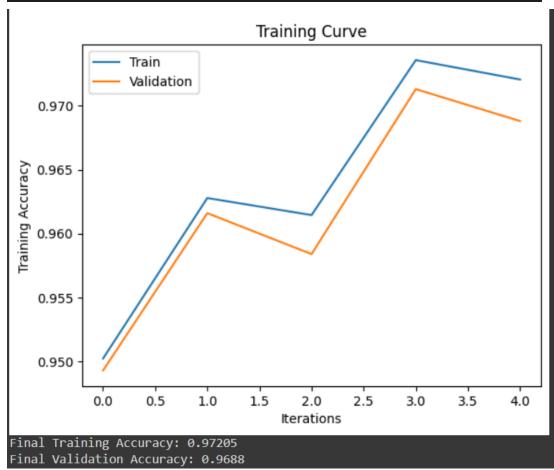


```
My brain's capacity: 10^11 x 1000 x 2 = 2E14
Total visual information in bits:
   Assume my brain can process 30 images per second, and each image is RGB (3 x 8 = 24 bits) and full HD (1920 x 1080 =
    2073600 pixels) and I am awake 16 hours per day
    Total visual information in bits = 23(years) x 365(days) x 16(hours) x 3600(seconds) x 30(images) x 2073600(pixels)
    x 24(bits) = 7.2E17
Total audio information in bits:
    Assume my brain can process 80 Kbps audio, this is 2.88E8 bits/hour
    Total audio information in bits = 23(years) x 365(days) x 16(hours) x 2.88E8(bits) = 1.1E13
Audio information size is negligible compared to visual information, and the combined size is a lot larger than my
brain's capacity. This makes sense because I don't remember all of this information.
Total bits in Shakespeare's works:
    There are 884647 words in Shakespeare's works, each word consists of 6.47 letters, and each letter consists of 1.6 bits.
    Total bits in Shakespeare = 884647 x 6.47 x 1.6 = 9.2E6
This is a lot less than the capacity of my brain.
We can use the same algorithm to handle non-binary classification cases, since we are simply counting the number of
times the label changes
We can simply define a small value 'e', in the second for loop of the algorithm we check if the next class is at least
increment threshold
```

9.1: Initial implementation using MNIST dataset:

```
class SmallNet(nn.Module):
    def __init__(self):
        super(SmallNet, self).__init__()
        self.conv = nn.Conv2d(1, 3, 3, 1)
        self.pool = nn.MaxPool2d(2, 2)
        self.fc = nn.Linear(3 * 13 * 13, 10)

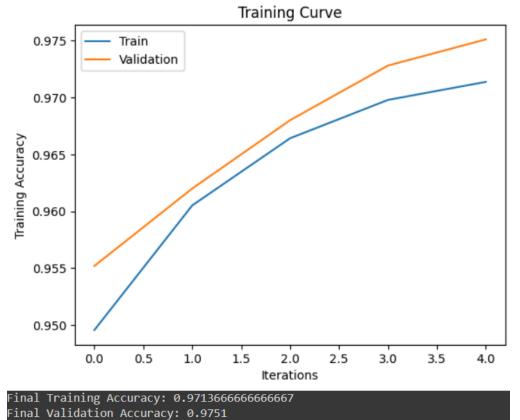
    def forward(self, x):
        x = self.pool(F.relu(self.conv(x)))
        x = x.view(-1, 3 * 13 * 13)
        x = self.fc(x)
        x = x.squeeze(1) # Flatten to [batch_size]
        return x
```



In the above case, the MEC of the fully connected layer is 3x13x13x10 = 5070, which is very large. We can make the MEC of the fully connected layer smaller by compressing more information in the CNN layers:

```
class SmallerNet(nn.Module):
    def __init__(self):
        super(SmallerNet, self).__init__()
        self.conv1 = nn.Conv2d(1, 3, 5, 1)
        self.pool1 = nn.MaxPool2d(2, 2)
        self.conv2 = nn.Conv2d(3, 3, 3, 1)
        self.pool2 = nn.MaxPool2d(2, 2)
        self.fc = nn.Linear(3 * 5 * 5, 10)

def forward(self, x):
        x = self.pool1(F.relu(self.conv1(x)))
        x = self.pool2(F.relu(self.conv2(x)))
        x = x.view(-1, 3 * 5 * 5)
        x = self.fc(x)
        x = x.squeeze(1) # Flatten to [batch_size]
        return x
```



This model has a smaller fully connected layer with MEC of 3x5x5x10=750, but it can classify just as well as the previous model.