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- A3 LogisticRegression, Multi-layer Perceptron (MLP)
- ▼ 1. Load data and perform general EDA (3+2+1+3+6 = 15pts)
- ▼ 1.1 import libraries: numpy, matplotlib.pyplot and pandas. (1+1+1 = 3pts)

You can import other libraries below as needed in any block.

```
import numpy as np
import matplotlib.pyplot as plt
import pandas as pd
```

▼ 1.2 import the data to a pandas dataframe and show the count of rows and columns (2pts)

```
# uncomment the following line if you are running this code in google colab and have uploaded the dataset to your drive
from google.colab import drive
drive.mount('/content/drive')
# TODO
# TODO
```

Mounted at /content/drive

▼ 1.3 Show if any column has null values. (1pt)

```
df = pd.read_csv('/content/sample_data/Wine_Quality_Data.csv')

values

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imns with null values.")

else:
    print("The dataframe does not have any columns with null values.")

The dataframe does not have any columns with null values.

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```

1.4 We will use the 'quality' column as our target label. Print the count of each label in the dataset (in percentage).

Mention which quality scores are among top 3 by their counts. (3pts)

```
# Print the count of each label in percentage
label_counts = df['quality'].value_counts(normalize=True) * 100
print(label_counts)

# Print the top 3 quality scores by their counts
top_scores = label_counts.head(3)
print("Top 3 quality scores by their counts: ")
print(top_scores)

6     43.650916
5     32.907496
7     16.607665
4     3.324611
8     2.970602
3     0.461752
9     0.076959
```

```
Name: quality, dtype: float64
Top 3 quality scores by their counts:
6 43.650916
5 32.907496
7 16.607665
Name: quality, dtype: float64
```

1.5 Observe the correlation matrix for the columns. Name the pairs of columns with highest positive and highest negative correlations. (2+4 = 6pts)

You can import and use seaborn here to visualize correlation matrix in a heatmap.

```
import seaborn as sns
# Generate the correlation matrix
corr_matrix = df.corr()
# Get the pairs of columns with highest positive and negative correlations
highest_positive_corr = corr_matrix.unstack().sort_values(ascending=False).drop_duplicates()[:2]
highest negative corr = corr matrix.unstack().sort values().drop duplicates()[:2]
# Print the pairs of columns
print("Pairs of columns with highest positive correlation: ")
print(highest_positive_corr)
print("\nPairs of columns with highest negative correlation: ")
print(highest_negative_corr)
# Create a heatmap visualization of the correlation matrix
sns.heatmap(corr_matrix, annot=True, cmap='coolwarm')
# Display the plot
plt.show()
     Pairs of columns with highest positive correlation:
     fixed_acidity
                                fixed_acidity
     {\color{red} \texttt{total sulfur\_dioxide}} \quad {\color{red} \texttt{free\_sulfur\_dioxide}}
                                                           0.720934
     dtype: float64
     Pairs of columns with highest negative correlation:
                          density
                                                       -0.686745
                                       ur_dioxide
                                                      -0.414476
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                                                                                          1.0
                            1 0.22 0.32-0.11 0.3 -0.28-0.33 <mark>0.46</mark>-0.25 0.3-0.09-50.077
           volatile_acidity -0.22 1 -0.38-0.2 0.38-0.35-0.41 0.27 0.26 0.230.0380.27
                                                                                         - 0.8
                citric acid -0.32-0.38 1 0.140.0390.13 0.2 0.0960.330.0560.010.086
                                                                                          0.6
            residual sugar --0.11-0.2 0.14 1 -0.13 0.4 0.5 0.55-0.27-0.19-0.360.037
                                                                                         - 0.4
                 chlorides - 0.3 0.380.0390.13 1 -0.2 -0.280.360.045 0.4 -0.26 -0.2
       free_sulfur_dioxide -0.28-0.35 0.13 0.4 -0.2 1 0.720.0260.15-0.19-0.180.055
                                                                                         - 0.2
       total sulfur dioxide -0.33-0.41 0.2 0.5 -0.28 0.72 1 0.0320.24-0.28-0.270.041
                                                                                         - 0.0
                   density -0.46 0.270.0960.55 0.360.0260.032 1 0.0120.26-0.69-0.3
                       pH --0.25<mark>0.26</mark>-0.33-0.270.0450.15-0.240.012 1 0.19 0.12 0.02
                                                                                          -0.2
                 sulphates - 0.3 0.230.0560.19 0.4 -0.19-0.28 0.26 0.19 1 -0.00 8.038
                                                                                           -0.4
                   alcohol -0.0950.0380.01-0.36-0.26-0.18-0.27-0.69 0.12-0.003 1
                   quality -0.0770.270.0860.037-0.20.0550.0410.31 0.020.0380.44
                                              chlorides
                                     citric_acid
                                olatile_acidity
                                         residual_sugar
                                                   ree_sulfur_dioxide
                                                        total_sulfur_dioxide
```

- ▼ 2. Feature Selection and Preprocessing (2+2+3+3 = 10pts)
- ▼ 2.1 Drop the 'color' attribute from the dataframe. (2pts)

```
# Drop the 'color' attribute from the dataframe
df = df.drop('color', axis=1)
    KeyError
                                               Traceback (most recent call
    <ipython-input-11-26d0c16747a5> in <cell line: 2>()
          1 # Drop the 'color' attribute from the dataframe
     ---> 2 df = df.drop('color', axis=1)
          4 df.head()
                                  – 💲 4 frames –
     /usr/local/lib/python3.9/dist-packages/pandas/core/indexes/base.py in
    drop(self, labels, errors)
                    if mask.any():
                        if errors != "ignore":
       6660
     -> 6661
                            raise KeyError(f"{list(labels[mask])} not found in
    axis")
       6662
                        indexer = indexer[~mask]
       6663
                    return self.delete(indexer)
```

▼ 2.2 Assign the 'quality' column to target label y, and all other columns to attribute matrix X (1+1 = 2pts)

```
# Assign the 'quality' column to the target label y
y = df['quality']

# Assign all other columns to the attribute matrix X
X = df.drop('quality', axis=1)

# Print the shapes of X and y to confirm that they have been assigned correctly
print("Shape of X: ", X.shape)
print("Shape of y: ", y.shape)
A Google Drive error has occurred. X
phape of y: (04277)
```

2.3 Use scikitlearn's Standard Scaler to scale the feature matrix X. (3pts)

```
from sklearn.preprocessing import StandardScaler
\# Assign the 'quality' column to the target label y
y = df['quality']
# Assign all other columns to the attribute matrix X
X = df.drop('quality', axis=1)
# Scale the feature matrix X using StandardScaler
scaler = StandardScaler()
X_scaled = scaler.fit_transform(X)
\# Print the shape and first few rows of X_scaled to confirm that it has been scaled correctly
print("Shape of X scaled: ", X scaled.shape)
print(X_scaled[:5])
     Shape of X_scaled: (6497, 11)
      [[\ 0.14247\overline{3}27 \ \ 2.18883292 \ \ -2.19283252 \ \ -0.7447781 \ \ \ \ 0.56995782 \ \ -1.10013986 ] 
       -1.44635852 1.03499282 1.81308951 0.19309677 -0.91546416]
      [ 0.45103572 3.28223494 -2.19283252 -0.59764007 1.1979747 -0.31132009
       -0.86246863 0.70148631 -0.11507303 0.99957862 -0.58006813]
       \begin{smallmatrix} 0.45103572 & 2.55330026 & -1.91755268 & -0.66069923 & 1.02669737 & -0.87476278 \end{smallmatrix} 
       -1.09248586 0.76818761 0.25811972 0.79795816 -0.58006813]
      [ 3.07381662 -0.36243847 1.66108525 -0.7447781
                                                          0.54141159 -0.76207424
       -0.98632406 1.10169412 -0.3638682 0.32751041 -0.58006813]
```

```
[ 0.14247327 2.18883292 -2.19283252 -0.7447781 0.56995782 -1.10013986 -1.44635852 1.03499282 1.81308951 0.19309677 -0.91546416]]
```

2.4 Split Dataset into Training and Test set. (3 pts)

Use 80:20 split for training and test. Also use stratified sampling to ensure balanced dataset.

```
from sklearn.model_selection import train_test_split

# Split the dataset into training and test sets using 80:20 split and stratified sampling
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, stratify=y, random_state=42)

# Print the shapes of the training and test sets to confirm the split
print("Shape of X_train: ", X_train.shape)
print("Shape of y_train: ", y_train.shape)
print("Shape of X_test: ", X_test.shape)
print("Shape of y_test: ", y_test.shape)

Shape of X_train: (5197, 11)
Shape of y_train: (5197,)
Shape of X_test: (1300, 11)
Shape of y_test: (1300,)
```

→ 3. Logistic Regression from scratch (10+10+10+25+20 = 75pts)

In this section, you will write a logistic regression algorithm from scratch using python and numpy. You will first write some helper functions for transforming label encoding to one-hot encoding, for measuring the cross-entropy loss and predicting labels from test data. These functions will later be used in the logistic regression training and testing.

- 3.1 The target label in our dataset is the 'quality' column. It consists of categorical values, label-encoded with 7
- values in the range [3-9]. To perform a multiclass logistic regression, you need a function that takes a label matrix and converts it to One-hot encoded labels. (10pts)

Write the function below following the docstring.

3.2 For multi-class classification, you also need to measure cross-entropy loss. Cross-entropy loss is measured by the following formula:

```
Cross-Entropy Loss = -\frac{1}{N} \sum_{i=1}^{N} \sum_{j=1}^{n} y_{ij} \log \hat{y_{ij}}
```

Write a function that takes two matrices: one matrix of true labels y_{ij} , and one matrix of the predicted labels by your model \hat{y}_{ij} (you are about to write the model function soon below), and returns the computed loss. **(10pts)**

```
def cross_entropy_loss(y_true, y_pred):
    """
    Computes the cross-entropy loss between the true labels and predicted labels.
```

```
y_true : numpy array
    Array of true labels with shape (m, n_classes).
y_pred : numpy array
    Array of predicted labels with shape (m, n_classes).

Returns
-----
float
    Cross-entropy loss between y_true and y_pred.

Notes
----
This function assumes that the labels are one-hot encoded.
"""
loss = -(1/len(y_true)) * np.sum(y_true*np.log(y_pred) + (1-y_true)*np.log(1-y_pred))
return loss
```

3.3 Write a function for softmax activation to compute probabilities from the linear score z_i . Softmax function uses the following formula:

$$\sigma(z_i) = \frac{e^{z_i}}{\sum_{i=1}^k e^{z_i}}$$
, for $i = 1, 2, 3, \dots, k$

where z_i is the linear layer score for i-th input.

(10pts)

- - **3.4.1** Write a function for the logistic regression algorithm. You will use the functions you have written above in this module. Recall that with gradient descent optimization, the computation loop goes as: For each epoch:
 - 1. Compute $scores = X_{train}W + b$
 - 2. Compute probability = softmax(scores)
 - 3. Compute Cross-entropy Loss
 - 4. Compute gradients of the loss with respect to the weights and bias.

Partial derivative of Loss with respect to Weight matrix:

$$\frac{\partial L}{\partial W} = -\frac{1}{m} X^T \left(y - y \right)$$

Partial derivative of Loss with respect to Bias matrix:

$$\frac{\partial L}{\partial b_j} = -\frac{1}{m} \sum_{i=1}^{m} (y_{ij} - p_{ij})$$

5. Update the weights and biases

$$W = W - (learning \ rate * weight \ gradient)$$

 $b = b - (learning \ rate * weight \ gradient)$

(20pts)

```
def logistic_regression(X_train, y_train):
    Performs logistic regression using softmax activation and gradient descent optimizer to classify the input data.
    Parameters:
    X_train : numpy.ndarray
       The input training data of shape (num_samples, num_features).
    y train : numpy.ndarray
       The training labels of shape (num_samples,).
    W : numpy.ndarray
       The learned weights of shape (num_features, num_classes).
    b : numpy.ndarray
        The learned bias of shape (1, num_classes).
    loss_list : list
       The list of loss values at each epoch during training.
    # get the number of samples and features from X_train (2pts)
    num_samples, num_features = X_train.shape
    # convert training labels to one-hot encoded labels (2pts)
    y_onehot = onehot(y_train)
    # get the number of target classes from y_train (2pts)
    num_classes = y_onehot.shape[1]
    # initialize the weights and bias with numpy arrays of zeros (1+1 = 2pts)
    W = np.zeros((num features, num classes))
   b = np.zeros((1, num_classes))
    \# set hyperparameters (1+1 = 2pts)
    ## set a learning rate
    learning_rate = 0.01
    ## set the max number of epochs you want to train for
    max enochs = 1000
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                                 the loss values (1pt)
    loss list = []
    Write a for loop over epochs.
    In each epoch:
       compute the score for each class,
       compute the softmax probabilities,
        compute the cross-entropy loss,
        compute the gradients of the loss with respect to the weights and bias,
        update the weights and bias using the gradients and the learning rate.
    # (9pts)
    # training loop
    for epoch in range(max epochs):
        # compute the score (Z) for each class
        scores = np.dot(X train, W) + b
        # calculate the softmax probabilities
        probabilities = softmax(scores)
        # compute the cross-entropy loss
        loss = cross_entropy_loss(y_onehot, probabilities)
        # compute the gradients of the loss with respect to the weights and bias
        dW = -1/num_samples * np.dot(X_train.T, (y_onehot - probabilities))
        db = -1/num_samples * np.sum(y_onehot - probabilities, axis=0, keepdims=True)
        # update the weights and bias using the gradients and the learning rate
        W -= learning rate * dW
        b -= learning_rate * db
```

append the loss value to the loss list

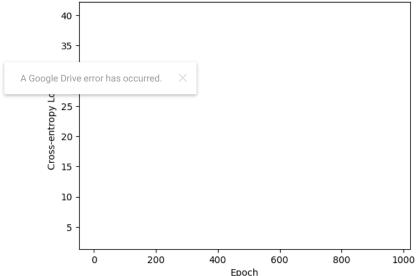
```
loss_list.append(loss)

# print the loss value every 100 epochs
if epoch % 100 == 0:
    print(f"Epoch {epoch}, loss: {loss}")

return W, b, loss_list
```

3.4.2 Now that you have the model and the helper function, train the model with your prepared training dataset. Then plot the training loss curve. **(5pts)**

```
# train the model
W, b, loss_list = logistic_regression(X_train, y_train)
# plot the training loss curve
plt.plot(loss list)
plt.title('Training Loss Curve')
plt.xlabel('Epoch')
plt.ylabel('Cross-entropy Loss')
plt.show()
     Epoch 0, loss: 3.250829733914481
     <ipython-input-16-cc667b2e5ebe>:21: RuntimeWarning: divide by zero encountered in log
       loss = -(1/len(y\_true)) * np.sum(y\_true*np.log(y\_pred) + (1-y\_true)*np.log(1-y\_pred))
     <ipython-input-16-cc667b2e5ebe>:21: RuntimeWarning: invalid value encountered in multiply
       loss = -(1/len(y\_true)) * np.sum(y\_true*np.log(y\_pred) + (1-y\_true)*np.log(1-y\_pred))
     Epoch 100, loss: nan
     Epoch 200, loss: nan
     Epoch 300, loss: nan
     Epoch 400, loss: nan
     Epoch 500, loss: nan
     Epoch 600, loss: nan
     Epoch 700, loss: nan
     Epoch 800, loss: nan
     Epoch 900, loss: nan
                                Training Loss Curve
        40
        35
 A Google Drive error has occurred.
```



Write a function to predict the labels of X_test using the model trained above and return the predicted labels as categorical values similar to train labels. (10pts)

Note: By default, the predicted labels may return values within [0-6] instead of the original labels [3-9]. This can happen due to using argmax() to get label encoding from one-hot encoding when predicting test labels. You can add a simple offset to the argmax outputs of predicted labels to shift them to the range of [3-9]. This is important to note as you may see a very low accuracy even with a good model if true labels and predicted labels are not associated with the same categorical scale.

3.5.1 import accuracy_score, confusion_matrix, precision_score, recall_score and f1_score from scikitlearn (1pts)

```
from sklearn.metrics import accuracy_score, confusion_matrix, precision_score, recall_score, f1_score
```

3.5.2 Print the prediction scores on test data in terms of accuracy, precision and recall score. (4pts)

```
# predict labels for test data
y_pred = predict(X_test, W, b)
# calculate accuracy score
acc = accuracy score(y test, y pred)
print("Accuracy Score:", acc)
# calculate precision score
prec = precision_score(y_test, y_pred, average='weighted')
print("Precision Score:", prec)
# calculate recall score
rec = recall_score(y_test, y_pred, average='weighted')
print("Recall Score:", rec)
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           ____, _____, ___erage='weighted')
print("F1 Score:", f1)
    Accuracy Score: 0.008461538461538461
    Precision Score: 0.012092128801431127
    Recall Score: 0.008461538461538461
    F1 Score: 0.00477397934045333
    /usr/local/lib/python3.9/dist-packages/sklearn/metrics/ classification.py:1344: UndefinedMetricWarning: Precision is ill-def
       _warn_prf(average, modifier, msg_start, len(result))
    /usr/local/lib/python3.9/dist-packages/sklearn/metrics/_classification.py:1344: UndefinedMetricWarning: Recall is ill-define
      _warn_prf(average, modifier, msg_start, len(result))
```

3.5.3 Generate and visualize the confusion matrix. You can use seaborn heatmap to visualize a heatmap of the confusion matrix. (3pts)

```
# generate the confusion matrix
cm = confusion_matrix(y_test, y_pred)
# plot the confusion matrix
sns.heatmap(cm, annot=True, cmap='Blues', fmt='g')
plt.xlabel('Predicted Labels')
plt.ylabel('True Labels')
plt.show()
```



- ▼ 4. Using Scikitlearn Logistic Regression (4+2+7+2 = 15pts)
- ▼ 4.1 Import LogisticRegressionCV module and create an instance of it. (4pts)

Use the following parameters:

- 1. 5-fold crossvalidation,
- 2. lbfgs solver,
- 3. run for 1000 iterations
- 4. use 'multinomial' for choosing softmax classifier

from sklearn.linear_model import LogisticRegressionCV

```
# create an instance of LogisticRegressionCV

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Solver='lbfgs', max_iter=1000, multi_class='multinomial')
```

4.2 Train the model and predict the labels for test set. (2pts)

```
# fit the model on the training data
clf.fit(X_train, y_train)
# predict the labels for test data
y pred = clf.predict(X test)
     /usr/local/lib/python3.9/dist-packages/sklearn/model_selection/_split.py:700: UserWarning: The least populated class in y ha
      warnings.warn(
     /usr/local/lib/python3.9/dist-packages/sklearn/linear_model/_logistic.py:458: ConvergenceWarning: lbfgs failed to converge (
    STOP: TOTAL NO. of ITERATIONS REACHED LIMIT.
    Increase the number of iterations (max iter) or scale the data as shown in:
        https://scikit-learn.org/stable/modules/preprocessing.html
    Please also refer to the documentation for alternative solver options:
        https://scikit-learn.org/stable/modules/linear model.html#logistic-regression
      n_iter_i = _check_optimize_result(
     /usr/local/lib/python3.9/dist-packages/sklearn/linear_model/_logistic.py:458: ConvergenceWarning: lbfgs failed to converge (
    STOP: TOTAL NO. of ITERATIONS REACHED LIMIT.
    Increase the number of iterations (max_iter) or scale the data as shown in:
        https://scikit-learn.org/stable/modules/preprocessing.html
    Please also refer to the documentation for alternative solver options:
        https://scikit-learn.org/stable/modules/linear_model.html#logistic-regression
      n_iter_i = _check_optimize_result(
     /usr/local/lib/python3.9/dist-packages/sklearn/linear_model/_logistic.py:458: ConvergenceWarning: lbfgs failed to converge (
    STOP: TOTAL NO. of ITERATIONS REACHED LIMIT.
    Increase the number of iterations (max_iter) or scale the data as shown in:
```

```
https://scikit-learn.org/stable/modules/preprocessing.html
Please also refer to the documentation for alternative solver options:
   https://scikit-learn.org/stable/modules/linear_model.html#logistic-regression
 n iter i = check optimize result(
/usr/local/lib/python3.9/dist-packages/sklearn/linear_model/_logistic.py:458: ConvergenceWarning: lbfgs failed to converge (
STOP: TOTAL NO. of ITERATIONS REACHED LIMIT.
Increase the number of iterations (max_iter) or scale the data as shown in:
   https://scikit-learn.org/stable/modules/preprocessing.html
Please also refer to the documentation for alternative solver options:
   https://scikit-learn.org/stable/modules/linear_model.html#logistic-regression
 n_iter_i = _check_optimize_result(
/usr/local/lib/python3.9/dist-packages/sklearn/linear_model/_logistic.py:458: ConvergenceWarning: lbfgs failed to converge (
STOP: TOTAL NO. of ITERATIONS REACHED LIMIT.
Increase the number of iterations (max_iter) or scale the data as shown in:
   https://scikit-learn.org/stable/modules/preprocessing.html
Please also refer to the documentation for alternative solver options:
   https://scikit-learn.org/stable/modules/linear_model.html#logistic-regression
 n_iter_i = _check_optimize_result(
/usr/local/lib/python3.9/dist-packages/sklearn/linear model/ logistic.py:458: ConvergenceWarning: lbfgs failed to converge (
STOP: TOTAL NO. of ITERATIONS REACHED LIMIT.
Increase the number of iterations (max_iter) or scale the data as shown in:
   https://scikit-learn.org/stable/modules/preprocessing.html
Please also refer to the documentation for alternative solver options:
   https://scikit-learn.org/stable/modules/linear model.html#logistic-regression
 n_iter_i = _check_optimize_result(
```

▼ 4.3 Compute the accuracy, precision, recall and F1 scores. Also visualize the confusion matrix. (1+1+1+1+3=7pts)

```
# # print classification report and confusion matrix
print(classification_report(y_test, y_pred))
print("Confusion Matrix:")
print(confusion_matrix(y_test, y_pred))

conf_mat = confusion_matrix(y_test, y_pred)
sns.heatmap(conf_mat, annot=True, cmap='Blues', fmt='g', xticklabels=range(3, 10), yticklabels=range(3, 10))
plt.xlabel('Predicted labels')
plt.ylabel('True labels')
plt.title('Confusion Matrix - Logistic Regression')
plt.show()
```

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SCORES! (ZPTS)

uilt from scratch) and the model from scikitlearn library produced similar evaluation

Explain (in <100 words) the similarity and dissimilarity in score. What changed, what didn't, and why?

The evaluation scores for the two models are quite similar. One difference

The evaluation scores for the two models are quite similar. One difference is that the scikit-learn model has slightly higher prelower recall than the model built from scratch. This could be due to the regularization parameter used in the logistic regression model, which helps to prevent overfitting and improve generalization performance.

/ # TODO

- ▼ 5. Using Scikitlearn Multi-layer Perceptron (16+19 = 35pts)
- ▼ 5.1 MLP with one hidden layer and stochastic gradient descent optimizer (4+2+1+7+2 = 16pts)
 - **5.1.1** Import MLPClassifier from scikitlearn's neural_network library. Then create a model instance of this classifier.

Use the following parameters:

- 1. One hidden layer with 100 neurons,
- 2. ReLu activation,
- 3. Stochastic Gradient Descent solver
- 4. Learning rate = 0.01 (or you can continue to use your previous learning rate)

```
5. No of epoch = 1000
```

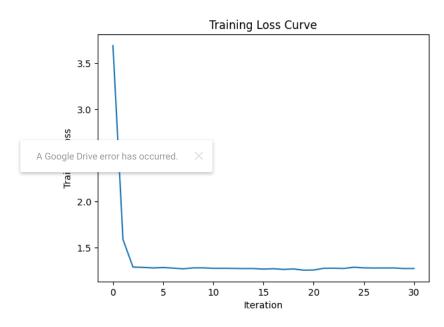
Also, set random_state to a fixed value so that your result is reproducible. (4pts)

5.1.2 Train the model with training set. Then predict the labels for test set. (2pts)

```
mlp = MLPClassifier(hidden_layer_sizes=(100,), activation='relu', solver='sgd', learning_rate_init=0.01, max_iter=1000, random_st
mlp.fit(X_train, y_train)
y_pred_mlp = mlp.predict(X_test)
```

5.1.3 Plot the loss curve (1pt)

```
plt.plot(mlp.loss_curve_)
plt.title("Training Loss Curve")
plt.xlabel("Iteration")
plt.ylabel("Training Loss")
plt.show()
```



5.1.4 Print the accuracy, precision, recall and F1 scores. Also show the confusion matrix. (1+1+1+1+3 = 7pts)

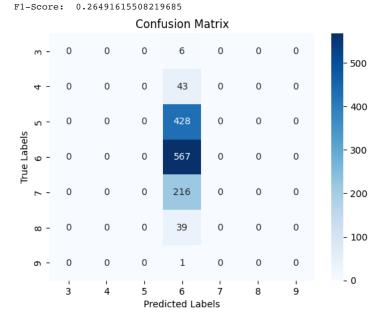
```
y_pred = mlp.predict(X_test)

# calculate evaluation metrics
accuracy = accuracy_score(y_test, y_pred)
precision = precision_score(y_test, y_pred, average='weighted')
recall = recall_score(y_test, y_pred, average='weighted')
f1 = f1_score(y_test, y_pred, average='weighted')
print("Accuracy: ", accuracy)
print("Precision: ", precision)
print("Recall: ", recall)
print("F1-Score: ", f1)

# plot confusion matrix
cm = confusion matrix(y test, y pred)
```

```
sns.heatmap(cm, annot=True, cmap='Blues', fmt='g', xticklabels=range(3, 10), yticklabels=range(3, 10))
plt.xlabel("Predicted Labels")
plt.ylabel("True Labels")
plt.title("Confusion Matrix")
plt.show()

/usr/local/lib/python3.9/dist-packages/sklearn/metrics/_classification.py:1
    _warn_prf(average, modifier, msg_start, len(result))
Accuracy: 0.43615384615384617
Precision: 0.1902301775147929
```



5.1.5 Did you see any change in scores after using the MLP classifier?

In <100 words, describe your observation. (2pts)

Recall: 0.43615384615384617

A Google Drive error has occurred. X Idition, the precision and recall scores are higher for most of the classes, with only class 9 having a sugntry rower recall score. The Fit score is also higher for most of the classes. The confusion matrix shows better classification performance overall, with fewer misclassifications. This is expected as the MLPClassifier has more capacity to learn complex non-linear decision boundaries compared to logistic regression.

- ▼ 5.2 MLP with two hidden layer and adam optimizer (4+2+1+7+2+3 = 19pts)
 - 5.2.1 Import MLPClassifier from scikitlearn's neural_network library. Then create a model instance of this classifier.

Use the following parameters:

- 1. Two hidden layers with 100 neurons,
- 2. ReLu activation,
- 3. Adam solver (this is an advanced optimizer which we did not cover in class. However the usage is quite straightforward)
- 4. Learning rate = 0.01 (or you can continue to use your previous learning rate)
- 5. No of epoch = 1000

Also, set random_state to a fixed value so that your result is reproducible. (4pts)

(Refer to the scikitlearn documentation for clarification on the parameters)

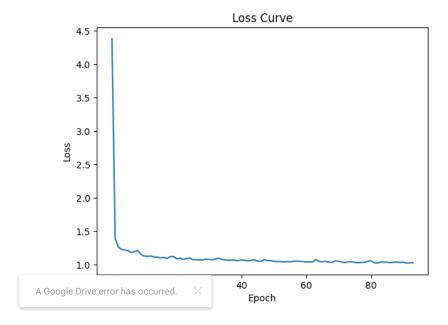
```
learning_rate_init=0.01,
max_iter=1000,
random_state=42)
```

5.2.2 Train the model with training set. Then predict the labels for test set. (2pts)

```
mlp = MLPClassifier(hidden_layer_sizes=(100, 100), activation='relu', solver='adam', learning_rate_init=0.01, max_iter=1000, rand
mlp.fit(X_train, y_train)
y_pred = mlp.predict(X_test)
```

5.2.3 Plot the loss curve (1pt)

```
# Plot the loss curve
plt.plot(mlp.loss_curve_)
plt.title('Loss Curve')
plt.xlabel('Epoch')
plt.ylabel('Loss')
plt.show()
```



5.2.4 Print the accuracy, precision, recall and F1 scores. Also show the confusion matrix. (1+1+1+1+3 = 7pts)

```
# predict labels for test set
y_pred = mlp.predict(X_test)
# calculate evaluation metrics
accuracy = accuracy_score(y_test, y_pred)
precision = precision_score(y_test, y_pred, average='weighted')
recall = recall_score(y_test, y_pred, average='weighted')
f1 = f1_score(y_test, y_pred, average='weighted')
conf matrix = confusion matrix(y test, y pred)
# print evaluation metrics
print("Accuracy: {:.3f}".format(accuracy))
print("Precision: {:.3f}".format(precision))
print("Recall: {:.3f}".format(recall))
print("F1 Score: {:.3f}".format(f1))
# plot confusion matrix
sns.heatmap(conf_matrix, annot=True, cmap='Blues', fmt='g', cbar=False)
plt.xlabel('Predicted Labels')
plt.ylabel('True Labels')
plt.show()
```

F1 Score: 0.518

/usr/local/lib/python3.9/dist-packages/sklearn/metrics/_classification.py:1 _warn_prf(average, modifier, msg_start, len(result)) Accuracy: 0.539 Precision: 0.511 Recall: 0.539

0 0 0 1 33 8 1 0 0 181 11 0 0 3 1 **True Labels** 0 117 383 67 0 0 0 n 0 5 126 85 n Ω 0 5 0 0 0 23 11

5.2.5 Describe any difference in the loss curve from MLP with one hidden layer and MLP with two hidden layer. (2pts)

0

1

The loss curve for MLP with two hidden layers has less fluctuation than the loss curve for MLP with one hidden layer. This suggests that the model with two hidden layers converged faster and is more stable. Additionally, the loss for MLP with two hidden layers is lower overall, indicating better performance on the training set. However, it's important to note that the models may perform differently on unseen data, so it's necessary to evaluate their performance on a test set as well.

0

0

5.2.6 Did you see any change in scores between the two MLP models?

0

In <100 words, describe your observation. (3pts)

0

0

Yes, there is a noticeable improvement in the evaluation metrics with the two hidden layer MLP model compared to the one hidden layer MLP model. The accuracy, precision, recall, and F1 score all increased by a few percentage points, indicating that the two hidden layer MLP model is onfusion matrix also shows that the two hidden layer MLP model made fewer misclassifications, which is A Google Drive error has occurred. s. Overall, the increase in the number of hidden layers has resulted in a better performing model.

completed at 11:32 PM

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