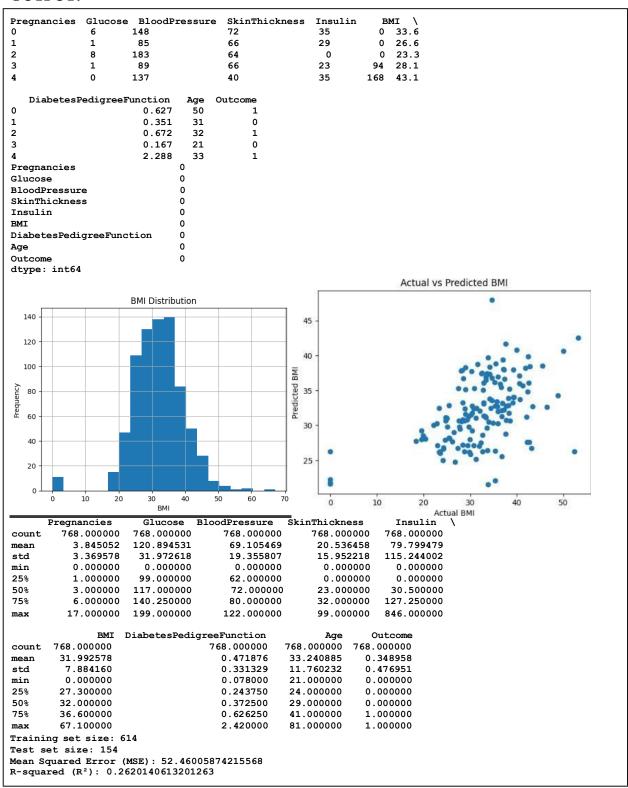
Aim: Diabetes dataset for linear regression practical

```
import pandas as pd
import numpy as np
from sklearn.model selection import train test split
from sklearn.linear model import LinearRegression
from sklearn.metrics import mean squared error, r2 score
import matplotlib.pyplot as plt
# Dataset URL
url = "https://raw.githubusercontent.com/jbrownlee/Datasets/master/pima-
indians-diabetes.data.csv"
# Column names based on the dataset documentation
column names = ['Pregnancies', 'Glucose', 'BloodPressure',
'SkinThickness', 'Insulin', 'BMI', 'DiabetesPedigreeFunction', 'Age',
'Outcome']
# Load the dataset into a pandas DataFrame
data = pd.read csv(url, names=column_names)
# Check the first few rows
print(data.head())
# Check for missing values
print(data.isnull().sum())
# Visualize the distribution of BMI (target variable)
data['BMI'].hist(bins=20)
plt.title('BMI Distribution')
plt.xlabel('BMI')
plt.ylabel('Frequency')
plt.show()
# Statistical summary
print(data.describe())
```

```
# Features (X) and target (y)
X = data.drop('BMI', axis=1) # All columns except BMI
y = data['BMI'] # BMI column as target
# Split the data into training and testing sets (80% train, 20% test)
X train, X test, y train, y test = train test split(X, y, test size=0.2,
random state=42)
print(f"Training set size: {X_train.shape[0]}")
print(f"Test set size: {X_test.shape[0]}")
# Initialize the linear regression model
model = LinearRegression()
# Fit the model on the training data
model.fit(X train, y train)
# Predict on the test set
y pred = model.predict(X test)
# Mean Squared Error (MSE)
mse = mean_squared_error(y_test, y_pred)
print(f"Mean Squared Error (MSE): {mse}")
# R-squared score (R2)
r2 = r2 score(y test, y pred)
print(f"R-squared (R2): {r2}")
# Visualize actual vs predicted values
plt.scatter(y_test, y_pred)
plt.title("Actual vs Predicted BMI")
plt.xlabel("Actual BMI")
plt.ylabel("Predicted BMI")
plt.show()
```

OUTPUT:

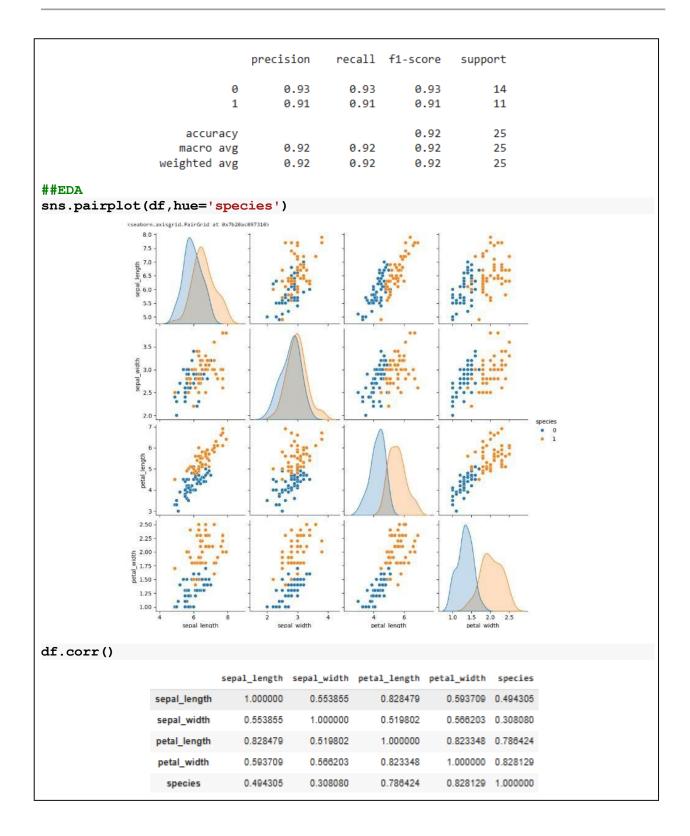


Aim:Implement Logistic Regression(iris dataset)

```
import seaborn as sns
import pandas as pd
import numpy as np
df=sns.load_dataset('iris')
df.head()
                    sepal_length sepal_width petal_length petal_width species
                 0
                             5.1
                                        3.5
                                                    1.4
                                                                0.2
                                                                     setosa
                 1
                             4.9
                                        3.0
                                                    1.4
                                                                0.2
                                                                     setosa
                 2
                             4.7
                                        3.2
                                                    1.3
                                                                     setosa
                 3
                             4.6
                                                    1.5
                                        3.1
                                                                0.2
                                                                     setosa
                                                    1.4
                                                                     setosa
df['species'].unique()
            array(['setosa', 'versicolor', 'virginica'], dtype=object)
df.isnull().sum()
                                                  0
                                        sepal_length 0
                                         sepal width 0
                                         petal_length 0
                                         petal_width 0
                                          species
                                        dtype: int64
df=df[df['species']!='setosa']
df.head()
                     sepal_length sepal_width petal_length petal_width species
                 50
                             7.0
                                         3.2
                                                      4.7
                                                                  1.4 versicolor
                 51
                             6.4
                                         3.2
                                                      4.5
                                                                  1.5 versicolor
                 52
                             6.9
                                         3.1
                                                      4.9
                                                                  1.5 versicolor
                 53
                             5.5
                                         2.3
                                                      4.0
                                                                  1.3 versicolor
                             6.5
                                         2.8
                                                      4.6
                                                                  1.5 versicolor
df['species']=df['species'].map({'versicolor':0,'virginica':1})
df.head()
```

```
sepal_length sepal_width petal_length petal_width species
                     50
                                 7.0
                                             3.2
                                                          4.7
                                                                      1.4
                     51
                                 6.4
                                             3.2
                                                          4.5
                                                                      1.5
                                                                                0
                                 6.9
                                             3.1
                                                          4.9
                                                                      1.5
                     53
                                 5.5
                                             2.3
                                                          4.0
                                                                      1.3
                                                                                0
                                 6.5
                                             2.8
                                                          4.6
                                                                      1.5
### Split dataset into independent and dependent features
X=df.iloc[:,:-1]
y=df.iloc[:,-1]
                              sepal_length sepal_width petal_length petal_width
                          50
                                     7.0
                                                3.2
                                                           4.7
                                                                      1.4
                                     6.4
                                                3.2
                                                           4.5
                                                                      1.5
                           52
                                     6.9
                                                3.1
                                                           4.9
                                                                      1.5
                                                2.3
                                                           4.0
                                                                      1.3
                                                           4.6
                                                3.0
                                                                      2.3
                                     6.3
                                                2.5
                                                           5.0
                                                                      1.9
                          146
                          147
                                     6.5
                                                                      2.0
                                     6.2
                                                3.4
                                                           5.4
                                                                      2.3
                          148
                          149
                                     5.9
                                                           5.1
                                                                      1.8
                         100 rows × 4 columns
                                           species
                                        50
                                        149
                                       100 rows × 1 columns
                                       dtype: int64
from sklearn.model_selection import train_test_split
X_train, X_test, y_train, y_test = train_test_split(
     X, y, test_size=0.25, random_state=42)
```

```
from sklearn.linear model import LogisticRegression
classifier=LogisticRegression()
from sklearn.model selection import GridSearchCV
parameter={'penalty':['11','12','elasticnet'],'C':[1,2,3,4,5,6,10,20,30,
40,50], 'max iter':[100,200,300]}
classifier regressor=GridSearchCV(classifier,param grid=parameter,scorin
g='accuracy',cv=5)
classifier regressor.fit(X_train,y_train)
           /usr/local/lib/python3.10/dist-packages/sklearn/model_selection/_validation.py:540: FitFailedWarning:
           330 fits failed out of a total of 495.
           The score on these train-test partitions for these parameters will be set to nan.
          If these failures are not expected, you can try to debug them by setting error_score='raise'.
          Below are more details about the failures:
          165 fits failed with the following error:
                          warnings.warn(
                                   GridSearchCV
                         ▶ best_estimator_: LogisticRegression
                                ▶ LogisticRegression
print(classifier regressor.best params )
                    {'C': 1, 'max iter': 100, 'penalty': 'l2'}
print(classifier regressor.best score )
                                0.9733333333333334
##prediction
y pred=classifier regressor.predict(X test)
## accuracy score
from sklearn.metrics import accuracy score, classification report
score=accuracy_score(y_pred,y_test)
print(score)
print(classification report(y pred,y test))
```

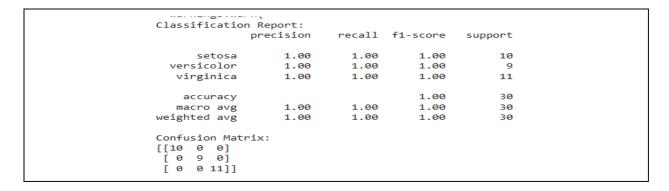


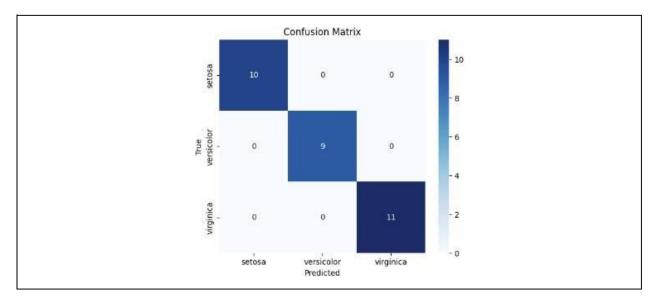
Aim: Implements Multinomial Logistic Regression (Iris Dataset)

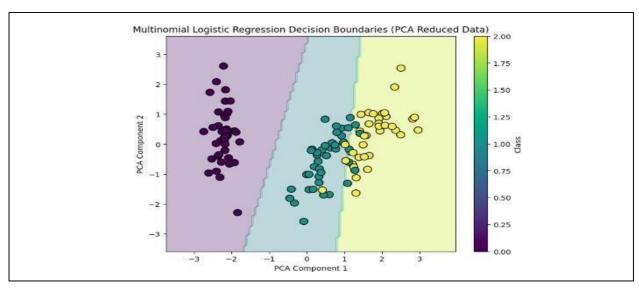
```
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns
from sklearn.datasets import load iris
from sklearn.linear model import LogisticRegression
from sklearn.model selection import train test split
from sklearn.preprocessing import StandardScaler
from sklearn.metrics import classification report, confusion matrix
from sklearn.decomposition import PCA
data = load iris()
X = data.data
y = data.target
df = pd.DataFrame(X, columns=data.feature names)
df['species'] = pd.Categorical.from codes(y, data.target names)
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2,
random state=42)
scaler = StandardScaler()
X train scaled = scaler.fit transform(X train)
X test scaled = scaler.transform(X test)
model = LogisticRegression(multi class='multinomial', solver='lbfgs',
max iter=200)
model.fit(X train scaled, y train)
y pred = model.predict(X test scaled)
print("Classification Report:")
print(classification report(y test, y pred,
target names=data.target names))
print("Confusion Matrix:")
cm = confusion matrix(y test, y pred)
print(cm)
plt.figure(figsize=(6, 5))
sns.heatmap(cm, annot=True, fmt='d', cmap='Blues',
xticklabels=data.target names, yticklabels=data.target names)
```

```
plt.title('Confusion Matrix')
plt.xlabel('Predicted')
plt.ylabel('True')
plt.show()
pca = PCA(n components=2)
X train pca = pca.fit transform(X train scaled)
X test pca = pca.transform(X test scaled)
model pca = LogisticRegression(multi class='multinomial',
solver='lbfgs', max iter=200)
model pca.fit(X train pca, y train)
plt.figure(figsize=(8, 6))
xx, yy = np.meshgrid(np.linspace(X train pca[:, 0].min() - 1,
X \text{ train pca}[:, 0].max() + 1, 100),
                      np.linspace(X train pca[:, 1].min() - 1,
X \text{ train pca}[:, 1].max() + 1, 100))
Z = model pca.predict(np.c [xx.ravel(), yy.ravel()])
Z = Z.reshape(xx.shape)
plt.contourf(xx, yy, Z, alpha=0.3, cmap='viridis')
plt.scatter(X train pca[:, 0], X train pca[:, 1], c=y train,
cmap='viridis', edgecolors='k', s=100)
plt.title("Multinomial Logistic Regression Decision Boundaries (PCA
Reduced Data)")
plt.xlabel('PCA Component 1')
plt.ylabel('PCA Component 2')
plt.colorbar(label='Class')
plt.show()
```

Output:







AIM: Implement SVM Classifier.

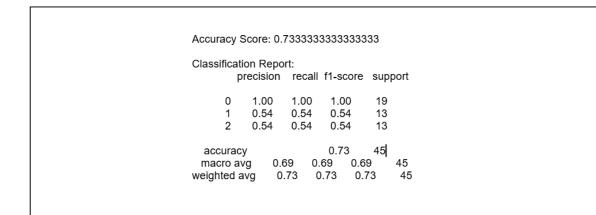
```
import numpy as np
import matplotlib.pyplot as plt
from sklearn import datasets
from sklearn.model_selection import train_test_split
from sklearn.preprocessing import StandardScaler
from sklearn.svm import SVC
from sklearn.metrics import classification report, accuracy score
# Load the Iris dataset
iris = datasets.load_iris()
X = iris.data[:, :2] # We only take the first two features (sepal length and sepal width)
y = iris.target
# Split the dataset into training and testing sets
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.3, random_state=42)
# Standardize the features
scaler = StandardScaler()
X train = scaler.fit transform(X train)
X \text{ test} = \text{scaler.transform}(X \text{ test})
# Create and train the SVM classifier
svm = SVC(kernel='linear', random_state=42)
svm.fit(X_train, y_train)
# Make predictions
y_pred = svm.predict(X_test)
# Evaluate performance
print("Accuracy Score:", accuracy_score(y_test, y_pred))
print("\nClassification Report:\n", classification_report(y_test, y_pred))
# Visualization of decision boundary
x_{min}, x_{max} = X_{train}[:, 0].min() - 1, X_{train}[:, 0].max() + 1
y_{min}, y_{max} = X_{train}[:, 1].min() - 1, X_{train}[:, 1].max() + 1
xx, yy = np.meshgrid(np.arange(x_min, x_max, 0.01),
             np.arange(y_min, y_max, 0.01))
Z = \text{sym.predict(np.c_[xx.ravel(), yy.ravel()])}
Z = Z.reshape(xx.shape)
```

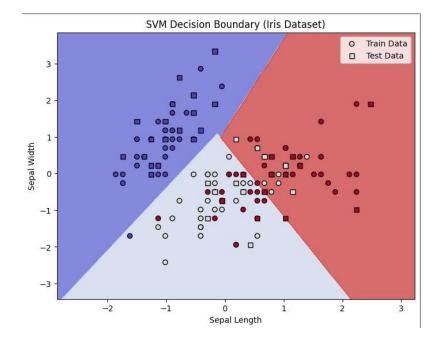
```
# Plotting the decision boundary plt.figure(figsize=(8, 6)) plt.contourf(xx, yy, Z, alpha=0.75, cmap=plt.cm.coolwarm)

# Plot the training points plt.scatter(X_train[:, 0], X_train[:, 1], c=y_train, cmap=plt.cm.coolwarm, marker='o', edgecolors='k', label="Train Data") plt.scatter(X_test[:, 0], X_test[:, 1], c=y_test, cmap=plt.cm.coolwarm, marker='s', edgecolors='k', label="Test Data")

plt.title('SVM Decision Boundary (Iris Dataset)') plt.xlabel('Sepal Length') plt.ylabel('Sepal Width') plt.legend() plt.show()
```

Output-





Aim: Train and fine-tune a Decision Tree for the Moons Dataset

```
import numpy as np
import matplotlib.pyplot as plt
from sklearn.datasets import make moons
from sklearn.model selection import train test split, GridSearchCV
from sklearn.tree import DecisionTreeClassifier
from sklearn.metrics import classification report, accuracy score
# Generate the moons dataset
X, y = make moons(n samples=1000, noise=0.2, random state=42)
# Split the data into training and testing sets
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.3, random_state=42)
# Create a DecisionTreeClassifier
dt = DecisionTreeClassifier(random_state=42)
# Hyperparameter tuning using GridSearchCV
param grid = {
  'max depth': [3, 5, 10, 20, None], # Depth of the tree
  'min_samples_split': [2, 5, 10], # Minimum samples required to split an internal node
  'min_samples_leaf': [1, 2, 4], # Minimum samples required to be at a leaf node
  'criterion': ['gini', 'entropy'] # The function to measure the quality of a split
# Apply GridSearchCV to find the best hyperparameters
grid search = GridSearchCV(estimator=dt, param grid=param grid, cv=5, n jobs=-1,
verbose=2)
grid_search.fit(X_train, y_train)
# Best parameters from GridSearchCV
print(f"Best parameters: {grid search.best params }")
# Use the best model from GridSearchCV
best_dt = grid_search.best_estimator_
# Predict on the test set
y pred = best dt.predict(X test)
# Evaluate the model
print("\nAccuracy Score:", accuracy score(y test, y pred))
print("\nClassification Report:\n", classification_report(y_test, y_pred))
# Visualization of the decision boundary
# Create a mesh grid to plot the decision boundaries
x_{min}, x_{max} = X[:, 0].min() - 1, X[:, 0].max() + 1
y_{min}, y_{max} = X[:, 1].min() - 1, X[:, 1].max() + 1
xx, yy = np.meshgrid(np.arange(x_min, x_max, 0.01),
             np.arange(y min, y max, 0.01))
Z = best_dt.predict(np.c_[xx.ravel(), yy.ravel()])
Z = Z.reshape(xx.shape)
# Plotting the decision boundary
plt.figure(figsize=(8, 6))
plt.contourf(xx, yy, Z, alpha=0.75, cmap=plt.cm.coolwarm)
```

```
# Plot the points from the moons dataset
plt.scatter(X_train[:, 0], X_train[:, 1], c=y_train, cmap=plt.cm.coolwarm, marker='o',
edgecolors='k', label="Train Data")
plt.scatter(X_test[:, 0], X_test[:, 1], c=y_test, cmap=plt.cm.coolwarm, marker='s',
edgecolors='k', label="Test Data")
plt.title('Decision Tree Classifier (Moons Dataset)')
plt.xlabel('Feature 1')
plt.ylabel('Feature 2')
plt.legend()
plt.show()
```

Output:

Fitting 5 folds for each of 90 candidates, totalling 450 fits

Best parameters: {'criterion': 'gini', 'max_depth': 10, 'min_samples_leaf': 1,

'min_samples_split': 10}

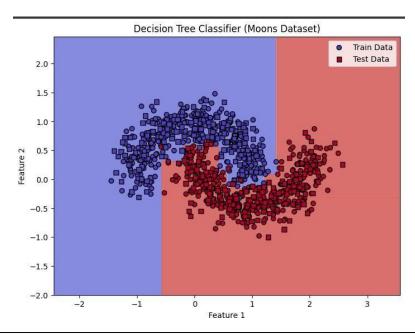
Accuracy Score: 0.96

Classification Report:

precision recall f1-score support

0 0.94 0.99 0.96 156 1 0.99 0.93 0.96 144

accuracy 0.96 300 macro avg 0.96 0.96 0.96 300 weighted avg 0.96 0.96 0.96 300



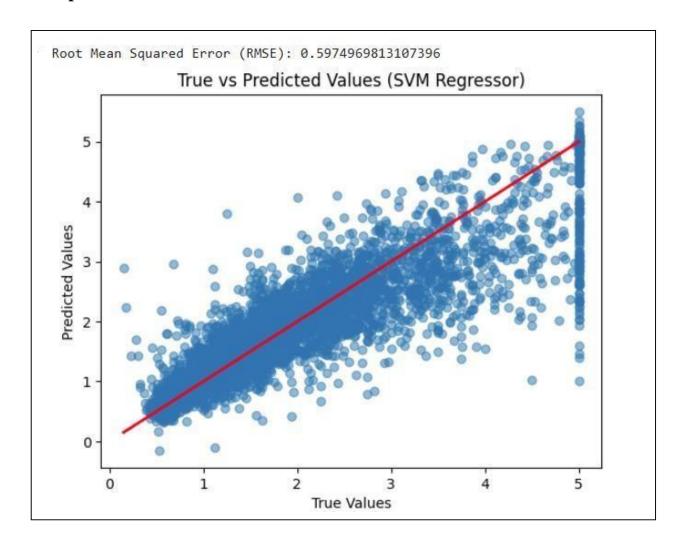
Aim: Train an SVM regressor on the California Housing Dataset

```
import numpy as np
import pandas as pd
from sklearn.datasets import fetch_california_housing
from sklearn.model selection import train test split
from sklearn.preprocessing import StandardScaler
from sklearn.svm import SVR
from sklearn.metrics import mean squared error
import matplotlib.pyplot as plt
# Load the California Housing Dataset
california housing = fetch california housing()
X = california housing.data
y = california_housing.target
# Split the dataset into training and testing sets
X train, X test, y train, y test = train test split(X, y, test size=0.2,
random state=42)
# Feature scaling (important for SVM)
scaler = StandardScaler()
X train scaled = scaler.fit transform(X train)
X test scaled = scaler.transform(X test)
# Train the SVM regressor
svm regressor = SVR(kernel='rbf') # You can experiment with different
kernels like 'linear', 'poly', etc.
svm_regressor.fit(X_train_scaled, y train)
# Predict on the test set
y_pred = svm_regressor.predict(X_test_scaled)
# Evaluate the model
mse = mean_squared_error(y_test, y_pred)
```

```
rmse = np.sqrt(mse)
print(f"Root Mean Squared Error (RMSE): {rmse}")

# Plot the true vs predicted values
plt.scatter(y_test, y_pred, alpha=0.5)
plt.plot([min(y_test), max(y_test)], [min(y_test), max(y_test)],
color='red', lw=2)
plt.xlabel('True Values')
plt.ylabel('Predicted Values')
plt.title('True vs Predicted Values (SVM Regressor)')
plt.show()
```

Output-



Aim: Implement Batch Gradient Descent with early stopping for Softmax Regression

Code:-

#Let's import IRIS data

from sklearn import datasets

iris = datasets.load_iris()

list(iris.keys())

Output:-

['data',

'target',

'frame',

'target_names',

'DESCR',

'feature names',

'filename',

'data module']

Code:-

print(iris.DESCR)

Output:-

.. iris dataset:

Iris plants dataset

Data Set Characteristics:

- :Number of Instances: 150 (50 in each of three classes)
- :Number of Attributes: 4 numeric, predictive attributes and the class
- :Attribute Information:
 - sepal length in cm
 - sepal width in cm
 - petal length in cm
 - petal width in cm
 - class:
 - Iris-Setosa
 - Iris-Versicolour
 - Iris-Virginica

:Summary Statistics:

```
Min Max Mean SD Class Correlation

sepal length: 4.3 7.9 5.84 0.83 0.7826

sepal width: 2.0 4.4 3.05 0.43 -0.4194

petal length: 1.0 6.9 3.76 1.76 0.9490 (high!)

petal width: 0.1 2.5 1.20 0.76 0.9565 (high!)
```

```
Code:-
print('iris.data.shape = ',iris.data.shape)
print('iris.target.shape = ',iris.target.shape)

Output:-
iris.data.shape = (150, 4)
iris.target.shape = (150,)
```

type(iris.data)

Output:-

numpy.ndarray

Code:-

from sklearn.linear_model import LogisticRegression

Code:-

```
X = iris.data[:, (2,3)]
```

y = iris.target

 $softmax_reg = LogisticRegression(multi_class = 'multinomial', solver = 'lbfgs', C = 10)$ $softmax_reg.fit(X,y)$

Output:-

/usr/local/lib/python3.10/dist-packages/sklearn/linear_model/_logistic.py:1247: FutureWarning: 'multi_class' was deprecated in version 1.5 and will be removed in 1.7. From then on, it will always use 'multinomial'. Leave it to its default value to avoid this warning. warnings.warn(

LogisticRegression LogisticRegression(C=10, multi_class='multinomial')

softmax_reg.predict([[5,2]])

Output:-

array([2])

Code:-

softmax_reg.predict_proba([[5,2]])

Output:-

array([[6.21626375e-07, 5.73689803e-02, 9.42630398e-01]])

Code:-

import numpy as np

np.bincount(y)

Output:-

array([50, 50, 50])

Code:-

Add a bias term in X

 $X_{\text{with_bias}} = \text{np.c}_{\text{[np.ones}([len(X), 1]), X]}$

Code:-

```
# Dividing into train-val-test
```

```
test ratio = 0.2
```

validation ratio = 0.2

 $total_size = len(X_with_bias)$

```
test size = int(total size * test ratio)
```

validation_size = int(total_size * validation_ratio)

train size = total size - test size - validation size

rnd_indices = np.random.permutation(total_size)

```
X_train = X_with_bias[rnd_indices[:train_size]]
y_train = y[rnd_indices[:train_size]]
X_valid = X_with_bias[rnd_indices[train_size:-test_size]]
y_valid = y[rnd_indices[train_size:-test_size]]
X_test = X_with_bias[rnd_indices[-test_size:]]
y_test = y[rnd_indices[-test_size:]]
```

```
Code:-
def to_one_hot(y):
    n_classes = y.max() + 1
    m = len(y)
    Y_one_hot = np.zeros((m, n_classes))
    Y_one_hot[np.arange(m), y] = 1
    return Y_one_hot
```

```
Y_train_one_hot = to_one_hot(y_train)
Y_valid_one_hot = to_one_hot(y_valid)
Y_test_one_hot = to_one_hot(y_test)
```

```
Code:-
def softmax(logits):
    exps = np.exp(logits)
    exp_sums = np.sum(exps, axis=1, keepdims=True)
    return exps / exp_sums
```

```
Code:-
n_inputs = X_train.shape[1] # == 3 (2 features plus the bias term)
n_outputs = len(np.unique(y_train)) # == 3 (3 iris classes)
```

```
Code:-
eta = 0.01
n iterations = 5001
m = len(X train)
epsilon = 1e-7
Theta = np.random.randn(n_inputs, n_outputs)
for iteration in range(n iterations):
  logits = X train.dot(Theta)
  Y proba = softmax(logits)
  loss = -np.mean(np.sum(Y train one hot *np.log(Y proba + epsilon), axis=1))
  error = Y proba - Y train one hot
  if iteration \% 500 == 0:
    print(iteration, loss)
  gradients = 1/m * X train.T.dot(error)
  Theta = Theta - eta * gradients
Output:-
0 1.4562135108859078
500 0.8945648262383888
1000 0.7241274446686337
1500 0.620840316123756
2000 0.5547103842019515
2500 0.508972430657483
3000 0.4750987442917988
3500 0.44863047810584006
4000 0.4270897793972089
4500 0.409009301588904
5000 0.3934688215211218
```

Theta

Output:-

```
array([[ 3.6450864 , -0.07748292, -2.11002678], [-0.40702617, 1.02826592, 0.5528866 ], [-1.53608619, -1.39115264, 1.42070209]])
```

```
Code:-
logits = X_valid.dot(Theta)

Y_proba = softmax(logits)

y_predict = np.argmax(Y_proba, axis=1)

accuracy_score = np.mean(y_predict == y_valid)
accuracy_score

Output:-
0.8
```

```
Code:-
eta = 0.1
n iterations = 5001
m = len(X train)
epsilon = 1e-7
alpha = 0.1 # regularization hyperparameter
Theta = np.random.randn(n inputs, n outputs)
for iteration in range(n iterations):
  logits = X train.dot(Theta)
  Y proba = softmax(logits)
  xentropy loss = -np.mean(np.sum(Y train one hot *np.log(Y proba + epsilon), axis=1))
  12 loss = 1/2 * np.sum(np.square(Theta[1:]))
  loss = xentropy loss + alpha * 12 loss
  error = Y proba - Y train one hot
  if iteration \% 500 == 0:
    print(iteration, loss)
  gradients = 1/m * X train.T.dot(error) + np.r [np.zeros([1, n outputs]), alpha * Theta[1:]]
  Theta = Theta - eta * gradients
```

Output:-

0 4.19763586247358 500 0.5561026466477075 1000 0.5190516378835687

```
1500 0.5087624410064894

2000 0.5050276975465959

2500 0.5035487641201934

3000 0.5029375815310052

3500 0.5026788293930453

4000 0.502567673305409

4500 0.5025194842171282

5000 0.5024984706307303
```

```
Code:-
logits = X_valid.dot(Theta)

Y_proba = softmax(logits)

y_predict = np.argmax(Y_proba, axis=1)

accuracy_score = np.mean(y_predict == y_valid)
accuracy_score

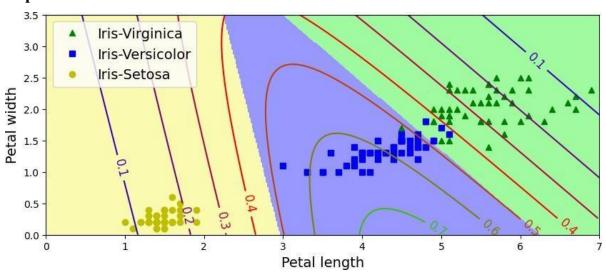
Output:-
0.933333333333333333
```

```
%matplotlib inline
import matplotlib.pyplot as plt
x0, x1 = np.meshgrid(
    np.linspace(0, 8, 500).reshape(-1, 1),
    np.linspace(0, 3.5, 200).reshape(-1, 1),
)
X_new = np.c_[x0.ravel(), x1.ravel()]
X_new_with_bias = np.c_[np.ones([len(X_new), 1]), X_new]

logits = X_new_with_bias.dot(Theta)
Y_proba = softmax(logits)
y_predict = np.argmax(Y_proba, axis=1)
```

```
zz1 = Y_proba[:, 1].reshape(x0.shape)
zz = y_predict.reshape(x0.shape)
plt.figure(figsize=(10, 4))
plt.plot(X[y==2, 0], X[y==2, 1], "g^", label="Iris-Virginica")
plt.plot(X[y==1, 0], X[y==1, 1], "bs", label="Iris-Versicolor")
plt.plot(X[y==0, 0], X[y==0, 1], "yo", label="Iris-Setosa")
from matplotlib.colors import ListedColormap
custom_cmap = ListedColormap(['#fafab0','#9898ff','#a0faa0'])
plt.contourf(x0, x1, zz, cmap=custom_cmap)
contour = plt.contour(x0, x1, zz1, cmap=plt.cm.brg)
plt.clabel(contour, inline=1, fontsize=12)
plt.xlabel("Petal length", fontsize=14)
plt.ylabel("Petal width", fontsize=14)
plt.legend(loc="upper left", fontsize=14)
plt.axis([0, 7, 0, 3.5])
plt.show()
```

Output:-



Aim: Implement MLP for classification of handwritten digits (MNIST Dataset)

```
import tensorflow as tf
from tensorflow.keras import layers, models
from tensorflow.keras.datasets import mnist
from tensorflow.keras.utils import to categorical
# 1. Load and preprocess the dataset
(x train, y train), (x test, y test) = mnist.load data()
# Normalize the pixel values to be between 0 and 1
x train, x test = x train / 255.0, x test / 255.0
# Flatten the images into vectors of size 784 (28 * 28)
x train = x train.reshape(-1, 28 * 28)
x_{test} = x_{test.reshape(-1, 28 * 28)}
# One-hot encode the labels
y_train = to_categorical(y_train, 10)
y_test = to_categorical(y_test, 10)
# 2. Build the MLP model
model = models.Sequential([
    layers.InputLayer(input_shape=(28 * 28,)), # Input layer (flattened
28x28 images)
    layers.Dense(128, activation='relu'),  # First hidden layer
with 128 neurons
    layers.Dense(64, activation='relu'),
                                               # Second hidden layer
with 64 neurons
    layers.Dense(10, activation='softmax') # Output layer with 10
classes (digits 0-9)
1)
# 3. Compile the model
model.compile(
    optimizer='adam',
                                     # Adam optimizer
    loss='categorical_crossentropy',  # Categorical cross-entropy loss
    metrics=['accuracy']
                                      # Track accuracy during training
# 4. Train the model
model.fit(x train, y train, epochs=10, batch size=32,
```

```
validation_data=(x_test, y_test))
# 5. Evaluate the model
test_loss, test_acc = model.evaluate(x_test, y_test)
print(f"Test Loss: {test_loss}")
print(f"Test Accuracy: {test_acc}")
```

Output:

```
Downloading data from https://storage.googleapis.com/tensorflow/tf-keras-
datasets/mnist.npz
11490434/11490434 -
                                -----Os Ous/step
/usr/local/lib/python3.11/dist-packages/keras/src/layers/core/input layer.py:27:
UserWarning: Argument `input shape` is deprecated. Use `shape` instead.
  warnings.warn(
Epoch 1/10
1875/1875 -
                               -15s 7ms/step - accuracy: 0.8770 - loss: 0.4147 -
val accuracy: 0.9604 - val loss: 0.1320
1875/1875 -
                               -18s 5ms/step - accuracy: 0.9684 - loss: 0.1028 -
val accuracy: 0.9688 - val loss: 0.0958
Epoch 3/10
1875/1875 -
                                -17s 4ms/step - accuracy: 0.9794 - loss: 0.0673 -
val_accuracy: 0.9689 - val_loss: 0.1022
Epoch 4/10
1875/1875 -
                               -11s 4ms/step - accuracy: 0.9834 - loss: 0.0520 -
val_accuracy: 0.9759 - val loss: 0.0784
Epoch 5/10
                               -- 3s 4ms/step - accuracy: 0.9874 - loss: 0.0398 -
val accuracy: 0.9794 - val loss: 0.0752
Epoch 6/10
1875/1875 -
                               -7s 4ms/step - accuracy: 0.9898 - loss: 0.0311 -
val_accuracy: 0.9769 - val_loss: 0.0810
Epoch 7/10
1875/1875 -
                               -8s 4ms/step - accuracy: 0.9929 - loss: 0.0222 -
val accuracy: 0.9761 - val loss: 0.0950
Epoch 8/10
1875/1875 -
                            ---- 6s 3ms/step - accuracy: 0.9922 - loss: 0.0229 -
val accuracy: 0.9773 - val loss: 0.0898
Epoch 9/10
1875/1875 -
                            val accuracy: 0.9761 - val loss: 0.0898
Epoch 10/10
1875/1875 -
                               -9s 3ms/step - accuracy: 0.9945 - loss: 0.0165 -
val_accuracy: 0.9767 - val_loss: 0.0925
                              -ds 2ms/step - accuracy: 0.9729 - loss: 0.1090
Test Loss: 0.09251588582992554
Test Accuracy: 0.9767000079154968
```

Aim: Classification of images of clothing using Tensorflow (Fashion MNIST dataset)

```
import tensorflow as tf
from tensorflow.keras import layers, models
from tensorflow.keras.datasets import fashion_mnist
from tensorflow.keras.utils import to_categorical
# 1. Load and preprocess the dataset
(x_train, y_train), (x_test, y_test) = fashion_mnist.load_data()
# Normalize the images to be between 0 and 1
x_{train}, x_{test} = x_{train} / 255.0, x_{test} / 255.0
# Flatten the images to vectors of size 784 (28 * 28)
x_{train} = x_{train.reshape}(-1, 28 * 28)
x_{test} = x_{test.reshape}(-1, 28 * 28)
# One-hot encode the labels
y_train = to_categorical(y_train, 10)
y_test = to_categorical(y_test, 10)
# 2. Build the MLP model
model = models.Sequential([
  layers.InputLayer(input_shape=(28 * 28,)), # Input layer (flattened 28x28 images)
  layers.Dense(128, activation='relu'),
                                            # First hidden layer with 128 neurons
  layers.Dense(64, activation='relu'),
                                            # Second hidden layer with 64 neurons
  layers.Dense(10, activation='softmax')
                                              # Output layer with 10 classes (clothing types)
1)
# 3. Compile the model
model.compile(
  optimizer='adam',
                                # Adam optimizer
  loss='categorical_crossentropy', # Categorical cross-entropy loss
  metrics=['accuracy']
                               # Track accuracy during training
# 4. Train the model
model.fit(x_train, y_train, epochs=10, batch_size=32, validation_data=(x_test, y_test))
# 5. Evaluate the model
test_loss, test_acc = model.evaluate(x_test, y_test)
```

```
print(f"Test Loss: {test_loss}")
print(f"Test Accuracy: {test_acc}")
```

OUTPUT:

```
Downloading data from https://storage.googleapis.com/tensorflow/tf-
keras-datasets/train-labels-idx1-ubyte.gz
29515/29515 -
                                  - 0us/step
Downloading data from https://storage.googleapis.com/tensorflow/tf-
keras-datasets/train-images-idx3-ubyte.gz
26421880/26421880 -
                                        20us/step
Downloading data from https://storage.googleapis.com/tensorflow/tf-
keras-datasets/t10k-labels-idx1-ubyte.gz
                                - 1us/step
Downloading data from https://storage.googleapis.com/tensorflow/tf-
keras-datasets/t10k-images-idx3-ubyte.gz
4422102/4422102 -
                                     ─s Ous/step
/usr/local/lib/python3.11/dist-
packages/keras/src/layers/core/input layer.py:27: UserWarning: Argument
`input shape` is deprecated. Use `shape` instead.
 warnings.warn(
Epoch 1/10
1875/1875 -
                               ──13 6ms/step - accuracy: 0.7770 - loss:
0.6334 - val accuracy: 0.8391 - val loss: 0.4358
Epoch 2/10
1875/1875 <del>---</del>
                                ─19 5ms/step - accuracy: 0.8615 - loss:
0.3781 - val accuracy: 0.8554 - val loss: 0.3999
Epoch 3/10
Test Loss: 0.34715157747268677
Test Accuracy: 0.8815000057220459
```

Aim: Implement Regression to predict fuel efficiency using Tensorflow (Auto MPG dataset)

```
import tensorflow as tf
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
from tensorflow.keras.models import Sequential
from tensorflow.keras.layers import Dense, Normalization
from sklearn.model_selection import train_test_split
from sklearn.preprocessing import StandardScaler
dataset_url = "http://archive.ics.uci.edu/ml/machine-learning-databases/auto-mpg/auto-mpg.data"
column names = ['MPG', 'Cylinders', 'Displacement', 'Horsepower', 'Weight', 'Acceleration', 'Model
Year', 'Origin']
dataset = pd.read_csv(dataset_url, names=column_names, na_values="?", comment='\t', sep=" ",
skipinitialspace=True)
dataset = dataset.dropna()
dataset['Origin'] = dataset['Origin'].astype(int)
dataset = pd.get_dummies(dataset, columns=['Origin'], prefix=", prefix_sep=")
X = dataset.drop('MPG', axis=1)
y = dataset['MPG']
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42)
scaler = StandardScaler()
X_train = scaler.fit_transform(X_train)
X_{test} = scaler.transform(X_{test})
model = Sequential([
  Dense(64, activation='relu', input_shape=(X_train.shape[1],)),
  Dense(64, activation='relu'),
  Dense(1) # Output layer for regression
1)
model.compile(optimizer=tf.keras.optimizers.Adam(learning rate=0.01), loss='mse', metrics=['mae'])
history = model.fit(X_train, y_train, epochs=100, batch_size=32, validation_data=(X_test, y_test),
verbose=1)
loss, mae = model.evaluate(X_test, y_test, verbose=2)
print(f'\nTest Mean Absolute Error: {mae:.2f} MPG')
```

Output:

Facab 4/400	
Epoch 1/100	
10/10	8s 123ms/step - loss: 492.4525 - mae: 20.6579 - val_loss:
49.8234 - val_mae: 5.6433	
Epoch 2/100	
10/10	
18.9818 - val_mae: 3.3336	
Epoch 3/100	
10/10	1s 42ms/step - loss: 23.4737 - mae: 3.8814 - val_loss:
17.4990 - val_mae: 3.4165	<u>-</u>
Epoch 4/100	
10/10	1s 65ms/step - loss: 17.4230 - mae: 3.2979 - val_loss:
15.5191 - val mae: 3.1137	
Epoch 5/100	
10/10	
10.8163 - val_mae: 2.3958	05 301113/3(ep - 1033, 12.0310 - 111ae, 2.0130 - Val_1033,
Epoch 6/100	00 26moleton loop 0 2770 mag 2 2 420 val loop 0 2000
10/10	
val_mae: 2.1948	
Epoch 7/100	
10/10	1s 28ms/step - loss: 8.2256 - mae: 2.0704 - val_loss: 7.6242 -
val_mae: 1.9974	
Epoch 8/100	
10/10	
- val_mae: 2.2980	
Epoch 9/100	
10/10	Os 18ms/step - loss: 8.6890 - mae: 2.2575 - val_loss: 6.8007 -
val_mae: 1.8021	
Epoch 10/100	
10/10	
val_mae: 1.8900	·
Epoch 11/100	
10/10	
val_mae: 1.9310	13 22 2.5p 1222 2222 1ac. 1.0000 1.a1000 1.10100
Epoch 12/100	
10/10	Os 10ms/step - loss: 7.8139 - mae: 2.0588 - val_loss: 6.5023 -
val_mae: 1.8036	55 Total Glop 1000. 1.0 Tot 111do. 2.0000 Val_1000. 0.0020
Epoch 13/100	
10/10	
10/10	03 101113/31cp - 1033. 0.7020 - 111ac. 1.0420 - Val_1033. 7.4070 -
vai_mae. 1.9773 Epoch 14/100	
10/10 ——————————————————————————————————	0c 0mc/cton_loce: 7 0916_map: 2 0727_val_loce: 7 2709
val_mae: 1.9083	
Epoch 15/100	0.40 // 1. 2004
10/10	
val_mae: 1.9725	
Epoch 16/100	
10/10	Os 10ms/step - loss: 8.4659 - mae: