

Lab 7

Statistics, Machine Learning, Deep Learning

1. Write a Python program that computes the value of the Gaussian distribution at a given vector **X**. Hence, plot the effect of varying mean and variance to the normal distribution.

Code:-

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

def gaussian_distribution(x, mean, variance):

    sigma = np.sqrt(variance)
    return (1 / (sigma * np.sqrt(2 * np.pi))) * np.exp(-0.5 * ((x -
mean) ** 2) / variance)

# Define range for x
x = np.linspace(-10, 10, 1000)

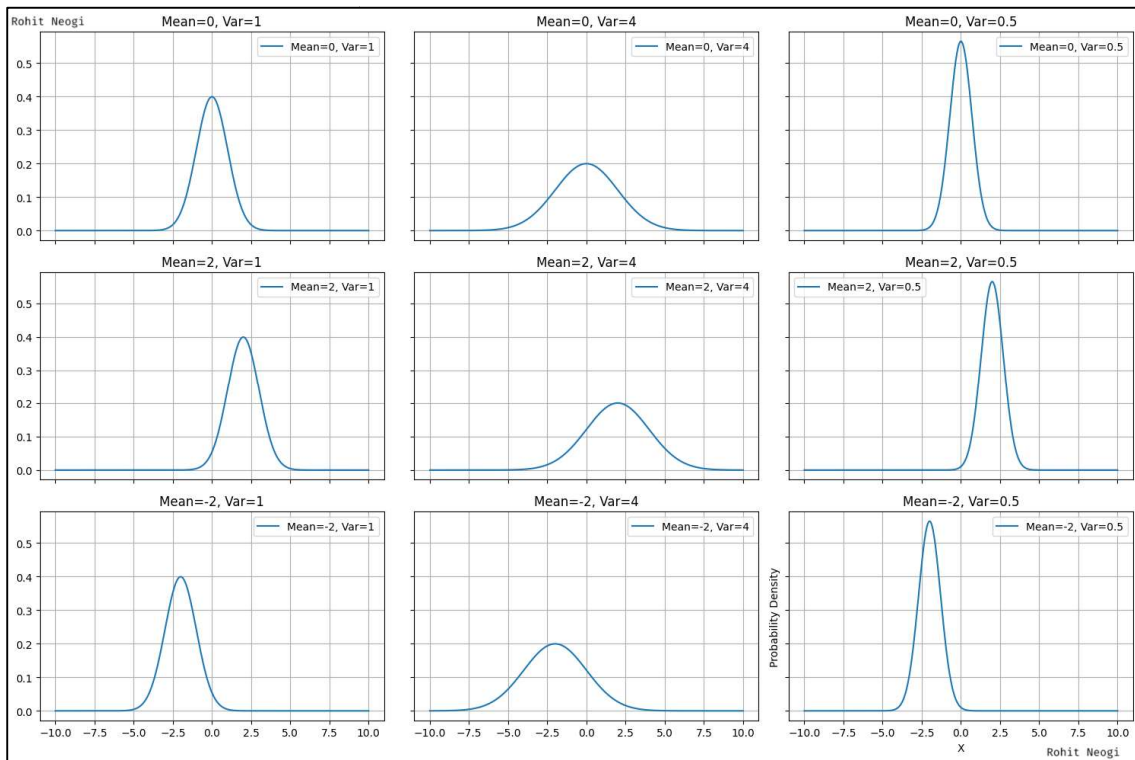
# Parameters to vary
means = [0, 2, -2]
variances = [1, 4, 0.5]

# Create subplots
fig, axes = plt.subplots(len(means), len(variances), figsize=(15,
10), sharex=True, sharey=True)

# Plot for varying means
for i, mean in enumerate(means):
    for j, variance in enumerate(variances):
        ax = axes[i, j]
        y = gaussian_distribution(x, mean, variance)
        ax.plot(x, y, label=f'Mean={mean}, Var={variance}')
        ax.set_title(f'Mean={mean}, Var={variance}')
        ax.legend()
        ax.grid(True)

# Set common labels
plt.xlabel('X')
plt.ylabel('Probability Density')
plt.tight_layout()
plt.show()
```

Output:-



2. Write a python program to implement linear regression.

Code:-

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

# Generate synthetic data
np.random.seed(0)
X = 2 * np.random.rand(100, 1)
y = 4 + 3 * X + np.random.randn(100, 1) # y = 4 + 3*X + noise

# Add bias term (x0 = 1) to each instance
X_b = np.c_[np.ones((100, 1)), X] # Add a column of ones to X

# Compute the optimal parameters using the Normal Equation
theta_best = np.linalg.inv(X_b.T.dot(X_b)).dot(X_b.T).dot(y)

# Extract the parameters
intercept, slope = theta_best

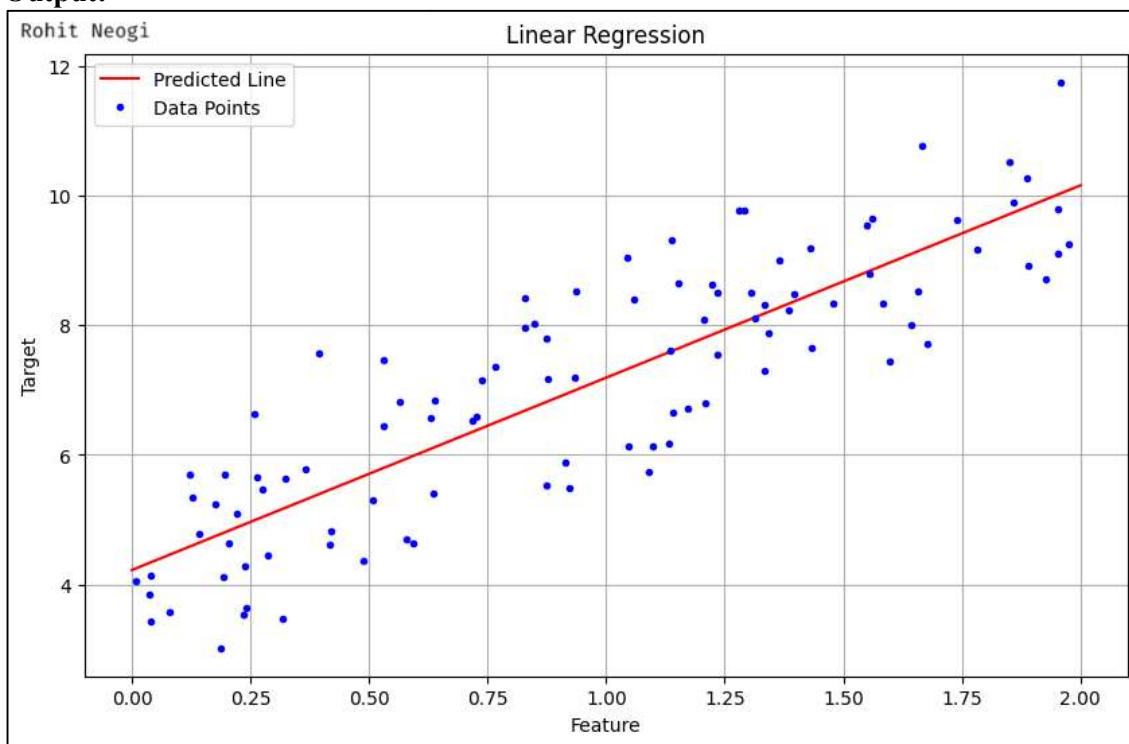
# Predict using the fitted model
X_new = np.linspace(0, 2, 100).reshape(100, 1)
X_new_b = np.c_[np.ones((100, 1)), X_new] # Add bias term
y_predict = X_new_b.dot(theta_best)

# Plotting
```

```
plt.figure(figsize=(10, 6))
plt.plot(X_new, y_predict, "r-", label="Predicted Line")
plt.plot(X, y, "b.", label="Data Points")
plt.xlabel("Feature")
plt.ylabel("Target")
plt.title("Linear Regression")
plt.legend()
plt.grid(True)
plt.show()

# Output the parameters
print(f"Intercept: {intercept[0]}")
print(f"Slope: {slope[0]}")
```

Output:-



3. Write a python program to implement gradient descent.

Code:-

```
# Import necessary libraries
import pandas as pd
import matplotlib.pyplot as plt
from sklearn.linear_model import LinearRegression
import numpy as np
import math

# Load and prepare the data
data_url = "/content/clean_weather.csv" # Path to your dataset
```

```

data = pd.read_csv(data_url, index_col=0).ffill() # Forward fill
missing values

# Display the first few rows of the data
print("First few rows of the data:")
print(data.head())

# Plot the relationship between tmax and tmax_tomorrow
plt.figure(figsize=(10, 6))
plt.scatter(data["tmax"], data["tmax_tomorrow"], color='blue',
label='Data Points')
plt.plot([30, 120], [30, 120], color='green', label='Regression
Line')
plt.xlabel('tmax')
plt.ylabel('tmax_tomorrow')
plt.title('Scatter Plot of tmax vs. tmax_tomorrow')
plt.legend()
plt.show()

# Train a linear regression model
lr = LinearRegression()
lr.fit(data[["tmax"]], data["tmax_tomorrow"])

# Display the model parameters
print(f"\nLinear Regression Model Parameters:")
print(f"Weight: {lr.coef_[0]:.2f}")
print(f"Bias: {lr.intercept_:.2f}")

# Loss calculation
loss = lambda w, y: ((w * 80 + 11.99) - y) ** 2
ws = np.arange(-1, 3, 0.1)
losses = loss(ws, 81)

# Plot the loss function
plt.figure(figsize=(10, 6))
plt.scatter(ws, losses, color='blue', label='Loss Values')
plt.plot(1, loss(1, 81), 'ro', label='Loss at Weight=1')
plt.xlabel('Weight')
plt.ylabel('Loss')
plt.title('Loss Function vs. Weight')
plt.legend()
plt.show()

# Gradient calculation
gradient = lambda w, y: ((w * 80 + 11.99) - y) * 2
gradients = gradient(ws, 81)

# Plot the gradient

```

```

plt.figure(figsize=(10, 6))
plt.scatter(ws, gradients, color='blue', label='Gradient Values')
plt.plot(1, gradient(1, 81), 'ro', label='Gradient at Weight=1')
plt.xlabel('Weight')
plt.ylabel('Gradient')
plt.title('Gradient Function vs. Weight')
plt.legend()
plt.show()

# Initialize model parameters
def init_params(predictors):
    k = math.sqrt(1 / predictors)
    np.random.seed(0)
    weights = np.random.rand(predictors, 1) * 2 * k - k
    biases = np.ones((1, 1)) * 2 * k - k
    return [weights, biases]

# Forward pass to make predictions
def forward(params, x):
    weights, biases = params
    return x @ weights + biases

# Mean Squared Error calculation
def mse(actual, predicted):
    return np.mean((actual - predicted) ** 2)

# Backward pass to update parameters
def backward(params, x, lr, grad):
    w_grad = (x.T / x.shape[0]) @ grad
    b_grad = np.mean(grad, axis=0)
    params[0] -= w_grad * lr
    params[1] -= b_grad * lr
    return params

# Gradient Descent
lr = 1e-4
epochs = 50000
params = init_params(train_x.shape[1])

for i in range(epochs):
    predictions = forward(params, train_x)
    grad = mse_grad(train_y, predictions)
    params = backward(params, train_x, lr, grad)

    if i % 10000 == 0:
        predictions = forward(params, valid_x)
        valid_loss = mse(valid_y, predictions)
        print(f"Epoch {i} validation loss: {valid_loss:.2f}")

```

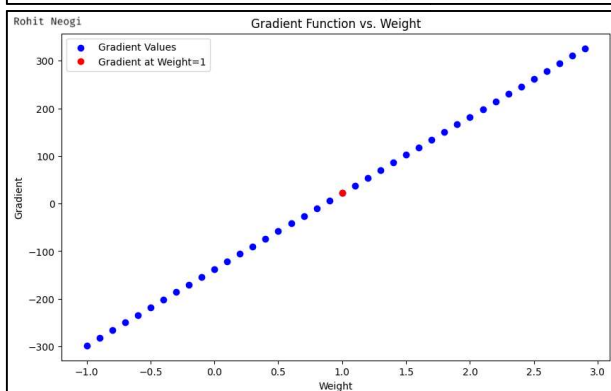
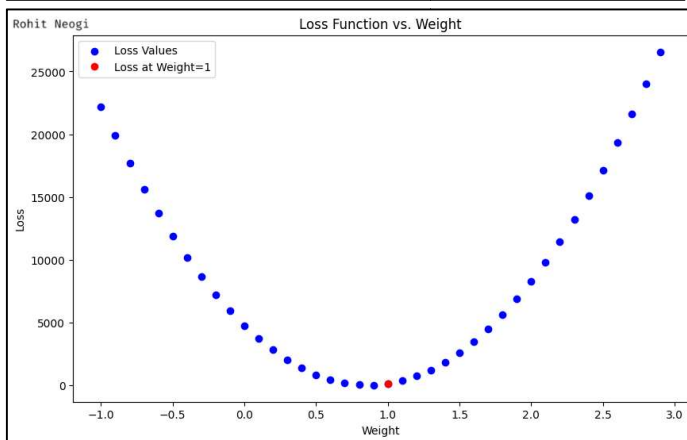
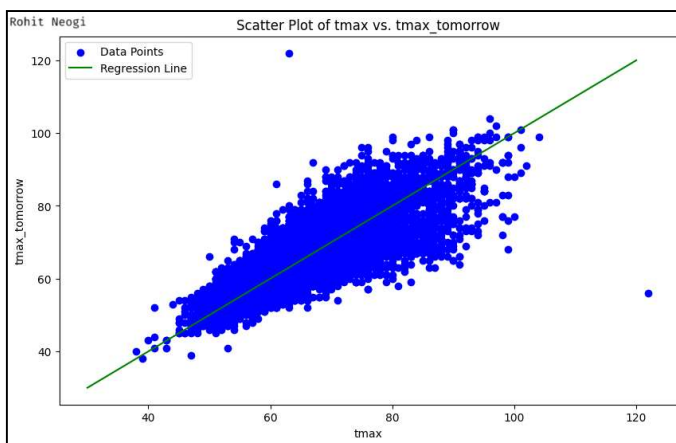
```
# Evaluate the model on the test set
predictions = forward(params, test_x)
test_loss = mse(test_y, predictions)
print(f"\nTest MSE: {test_loss:.2f}")
```

Output:-



First few rows of the data:

	tmax	tmin	rain	tmax_tomorrow
1970-01-01	60.0	35.0	0.0	52.0
1970-01-02	52.0	39.0	0.0	52.0
1970-01-03	52.0	35.0	0.0	53.0
1970-01-04	53.0	36.0	0.0	52.0
1970-01-05	52.0	35.0	0.0	50.0



```
Epoch 0 validation loss: 297.28
Epoch 10000 validation loss: 22.65
Epoch 20000 validation loss: 22.61
Epoch 30000 validation loss: 22.58
Epoch 40000 validation loss: 22.55

Test MSE: 23.34
```

4. Write a python program to classify different flower images using MLP.

Code: -

```
import numpy as np
import matplotlib.pyplot as plt
from sklearn.datasets import load_iris
from sklearn.model_selection import train_test_split
from sklearn.preprocessing import StandardScaler
from sklearn.neural_network import MLPClassifier
from sklearn.metrics import accuracy_score, classification_report,
confusion_matrix
import seaborn as sns

def load_and_preprocess_data():
    # Load the Iris dataset
    iris = load_iris()
    X = iris.data
    y = iris.target
    feature_names = iris.feature_names
    class_names = iris.target_names

    # 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)

    # Standardize features
    scaler = StandardScaler()
    X_train = scaler.fit_transform(X_train)
    X_test = scaler.transform(X_test)

    return X_train, X_test, y_train, y_test, class_names

def build_and_train_model(X_train, y_train):
    # Create an MLP model
    model = MLPClassifier(hidden_layer_sizes=(10, 10), max_iter=500,
random_state=42)

    # Train the model
    model.fit(X_train, y_train)
```

```

    return model

def evaluate_model(model, X_test, y_test, class_names):
    # Make predictions
    y_pred = model.predict(X_test)

    # Calculate accuracy
    accuracy = accuracy_score(y_test, y_pred)
    print(f"Accuracy: {accuracy:.2f}")

    # Print classification report
    print("\nClassification Report:")
    print(classification_report(y_test, y_pred,
target_names=class_names))

    # Print confusion matrix
    conf_matrix = confusion_matrix(y_test, y_pred)
    print("\nConfusion Matrix:")
    print(conf_matrix)

    # Plot confusion matrix
    plt.figure(figsize=(8, 6))
    sns.heatmap(conf_matrix, annot=True, fmt='d', cmap='Blues',
xticklabels=class_names, yticklabels=class_names)
    plt.title('Confusion Matrix')
    plt.xlabel('Predicted')
    plt.ylabel('Actual')
    plt.show()

def main():
    X_train, X_test, y_train, y_test, class_names =
load_and_preprocess_data()
    model = build_and_train_model(X_train, y_train)
    evaluate_model(model, X_test, y_test, class_names)

if __name__ == "__main__":
    main()

```

Output :-


```

/usr/local/lib/python3.10/dist-packages/sklearn/neural_network/
warnings.warn(
Accuracy: 1.00

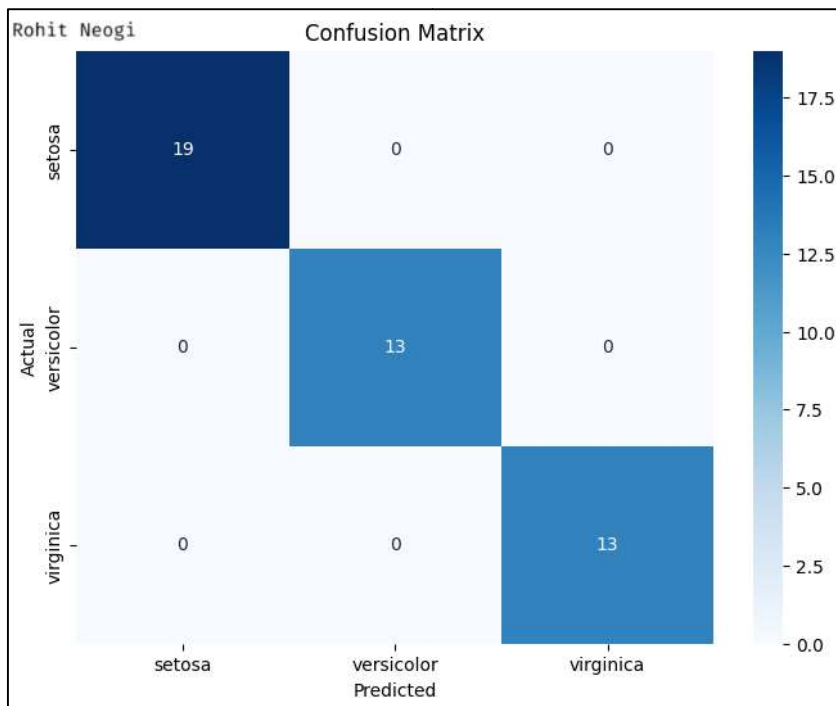
Classification Report:
      precision    recall  f1-score   support

   setosa         1.00      1.00      1.00        19
  versicolor         1.00      1.00      1.00        13
   virginica         1.00      1.00      1.00        13

   accuracy         1.00      1.00      1.00        45
  macro avg         1.00      1.00      1.00        45
 weighted avg         1.00      1.00      1.00        45

Confusion Matrix:
[[19  0  0]
 [ 0 13  0]
 [ 0  0 13]]

```



5. Write a python program to classify different flower images using the SVM classifier.

Code:-

```

import numpy as np
import tensorflow as tf
import tensorflow_datasets as tfds
from sklearn import svm
from sklearn.model_selection import train_test_split
from sklearn.preprocessing import StandardScaler
from sklearn.metrics import accuracy_score, classification_report,
confusion_matrix
import seaborn as sns
import matplotlib.pyplot as plt

# Load flower dataset
def load_flower_data():

```

```

    # Load the Flowers dataset from TensorFlow Datasets
    dataset, info = tfds.load('tf_flowers', with_info=True,
as_supervised=True, split=['train[:80%]', 'train[80%:]'],
shuffle_files=True)

    train_data, test_data = dataset

    # Preprocess and extract features
    def preprocess(image, label):
        image = tf.image.resize(image, [64, 64]) # Resize image to
64x64
        image = tf.cast(image, tf.float32) / 255.0 # Normalize
pixel values
        return image, label

    train_data = train_data.map(preprocess).batch(32).prefetch(1)
    test_data = test_data.map(preprocess).batch(32).prefetch(1)

    return train_data, test_data, info

# Extract features and labels
def extract_features_labels(data):
    features = []
    labels = []

    for images, batch_labels in data:
        features.extend(images.numpy().reshape(images.shape[0], -1))
    # Flatten images
        labels.extend(batch_labels.numpy())

    return np.array(features), np.array(labels)

def main():
    # Load and preprocess data
    train_data, test_data, info = load_flower_data()

    # Extract features and labels
    X_train, y_train = extract_features_labels(train_data)
    X_test, y_test = extract_features_labels(test_data)

    # Standardize features
    scaler = StandardScaler()
    X_train = scaler.fit_transform(X_train)
    X_test = scaler.transform(X_test)

    # Create and train SVM model
    model = svm.SVC(kernel='linear', C=1.0, random_state=42)
    model.fit(X_train, y_train)

```

```

# Make predictions
y_pred = model.predict(X_test)

# Evaluate the model
accuracy = accuracy_score(y_test, y_pred)
print(f"Accuracy: {accuracy:.2f}")

print("\nClassification Report:")
print(classification_report(y_test, y_pred,
target_names=info.features['label'].names))

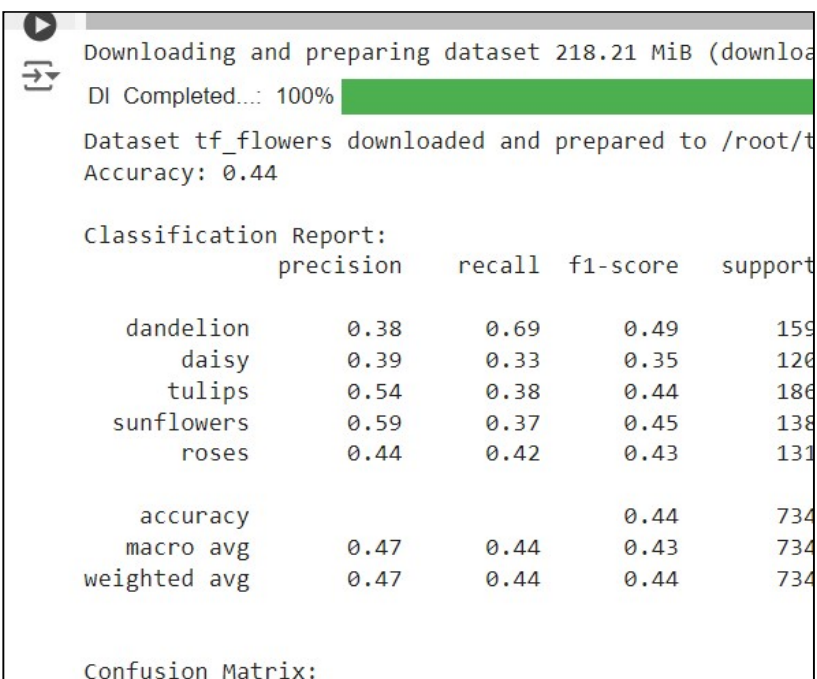
conf_matrix = confusion_matrix(y_test, y_pred)
print("\nConfusion Matrix:")
print(conf_matrix)

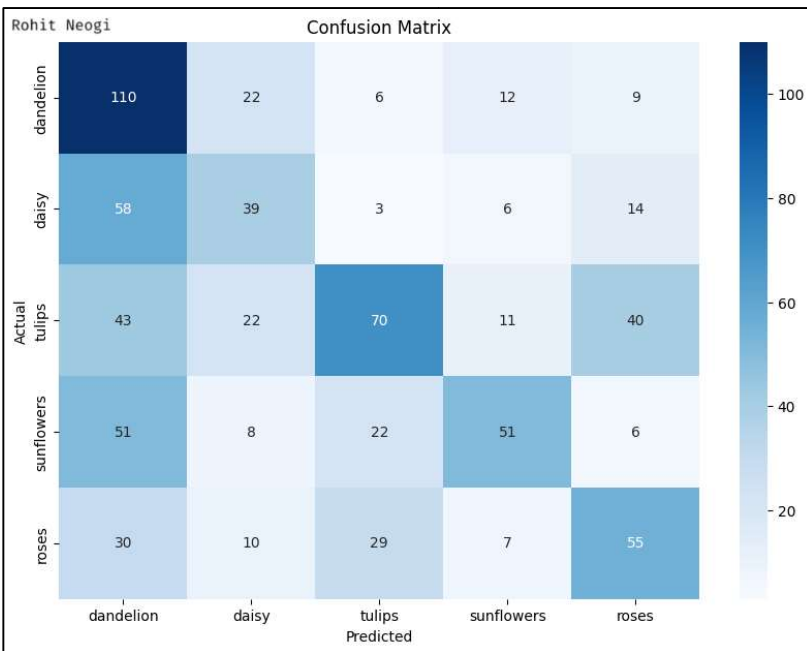
# Plot confusion matrix
plt.figure(figsize=(10, 7))
sns.heatmap(conf_matrix, annot=True, fmt='d', cmap='Blues',
xticklabels=info.features['label'].names,
yticklabels=info.features['label'].names)
plt.title('Confusion Matrix')
plt.xlabel('Predicted')
plt.ylabel('Actual')
plt.show()

if __name__ == "__main__":
    main()

```

Output:-





6. Write a python program to classify different flower images using CNN.

Code:-

```
import numpy as np
import tensorflow as tf
import tensorflow_datasets as tfds
from tensorflow.keras.models import Sequential
from tensorflow.keras.layers import Conv2D, MaxPooling2D, Flatten, Dense
from tensorflow.keras.preprocessing.image import ImageDataGenerator
from tensorflow.keras.callbacks import EarlyStopping
from sklearn.metrics import accuracy_score, classification_report, confusion_matrix
import seaborn as sns
import matplotlib.pyplot as plt

# Load and preprocess flower dataset
def load_flower_data():
    # Load the Flowers dataset
    dataset, info = tfds.load('tf_flowers', with_info=True,
as_supervised=True, split=['train[:80%]', 'train[80%:]'],
shuffle_files=True)
    train_data, test_data = dataset

    # Define preprocessing function
    def preprocess(image, label):
        image = tf.image.resize(image, [128, 128]) # Resize to
128x128
        image = tf.cast(image, tf.float32) / 255.0 # Normalize
pixel values
        return image, label

    # Apply preprocessing
```

```

train_data = train_data.map(preprocess).batch(32).prefetch(1)
test_data = test_data.map(preprocess).batch(32).prefetch(1)

return train_data, test_data, info

# Build CNN model
def build_cnn_model(num_classes):
    model = Sequential([
        Conv2D(32, (3, 3), activation='relu', input_shape=(128, 128,
3)),
        MaxPooling2D((2, 2)),
        Conv2D(64, (3, 3), activation='relu'),
        MaxPooling2D((2, 2)),
        Conv2D(128, (3, 3), activation='relu'),
        MaxPooling2D((2, 2)),
        Flatten(),
        Dense(512, activation='relu'),
        Dense(num_classes, activation='softmax')
    ])

    model.compile(optimizer='adam',
                  loss='sparse_categorical_crossentropy',
                  metrics=['accuracy'])

    return model

# Evaluate model performance
def evaluate_model(model, test_data, info):
    # Make predictions
    y_true = []
    y_pred = []

    for images, labels in test_data:
        predictions = model.predict(images)
        y_true.extend(labels.numpy())
        y_pred.extend(np.argmax(predictions, axis=1))

    # Convert lists to arrays
    y_true = np.array(y_true)
    y_pred = np.array(y_pred)

    # Evaluate the model
    accuracy = accuracy_score(y_true, y_pred)
    print(f"Accuracy: {accuracy:.2f}")

    print("\nClassification Report:")
    print(classification_report(y_true, y_pred,
target_names=info.features['label'].names))

```

```

conf_matrix = confusion_matrix(y_true, y_pred)
print("\nConfusion Matrix:")
print(conf_matrix)

# Plot confusion matrix
plt.figure(figsize=(10, 7))
sns.heatmap(conf_matrix, annot=True, fmt='d', cmap='Blues',
xticklabels=info.features['label'].names,
yticklabels=info.features['label'].names)
plt.title('Confusion Matrix')
plt.xlabel('Predicted')
plt.ylabel('Actual')
plt.show()

def main():
    # Load and preprocess data
    train_data, test_data, info = load_flower_data()

    # Build and train CNN model
    num_classes = len(info.features['label'].names)
    model = build_cnn_model(num_classes)

    # Use EarlyStopping to prevent overfitting
    early_stopping = EarlyStopping(monitor='val_loss', patience=3)

    history = model.fit(train_data,
                        epochs=20,
                        validation_data=test_data,
                        callbacks=[early_stopping])

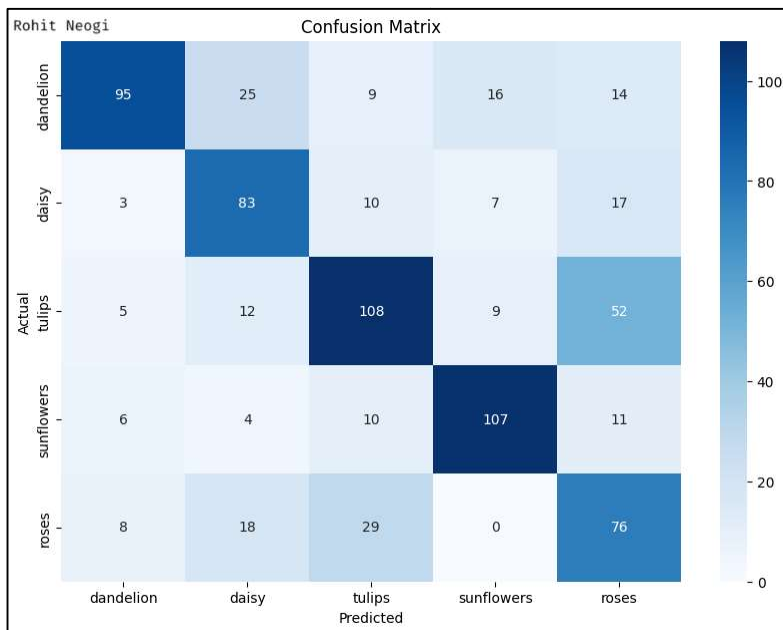
    # Evaluate model performance
    evaluate_model(model, test_data, info)

if __name__ == "__main__":
    main()

```

Output:-

Classification Report:				
	precision	recall	f1-score	support
dandelion	0.81	0.60	0.69	159
daisy	0.58	0.69	0.63	120
tulips	0.65	0.58	0.61	186
sunflowers	0.77	0.78	0.77	138
roses	0.45	0.58	0.50	131
accuracy			0.64	734
macro avg	0.65	0.65	0.64	734
weighted avg	0.66	0.64	0.64	734
Confusion Matrix:				
[[95 25 9 16 14]				
[3 83 10 7 17]				
[5 12 108 9 52]				
[6 4 10 107 11]				
[8 18 29 0 76]]				



7. Write a python program to classify different handwritten character images using the SVM classifier.

Code:-

```
import numpy as np
from sklearn import datasets
from sklearn.model_selection import train_test_split
from sklearn.svm import SVC
from sklearn.metrics import accuracy_score, classification_report
import matplotlib.pyplot as plt

# Load the dataset
digits = datasets.load_digits()
X = digits.images
y = digits.target

# Flatten the images
n_samples = len(X)
X = X.reshape((n_samples, -1))

# Split the dataset
X_train, X_test, y_train, y_test = train_test_split(X, y,
test_size=0.5, random_state=42)

# Train the SVM classifier
clf = SVC(gamma=0.001, C=100.)
clf.fit(X_train, y_train)

# Predict and evaluate
y_pred = clf.predict(X_test)
accuracy = accuracy_score(y_test, y_pred)
```

```

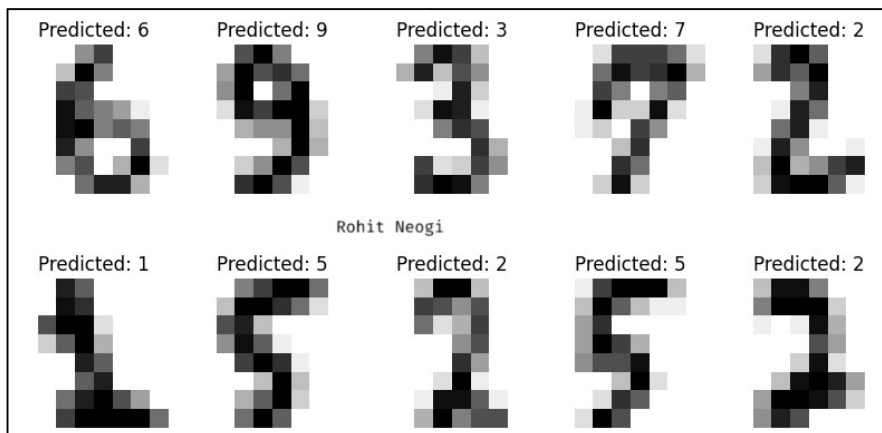
print(f'Accuracy: {accuracy:.2f}')
print('Classification Report:')
print(classification_report(y_test, y_pred))

# Visualize some predictions
plt.figure(figsize=(10, 5))
for i in range(10):
    plt.subplot(2, 5, i + 1)
    plt.imshow(X_test[i].reshape(8, 8), cmap=plt.cm.gray_r,
interpolation='nearest')
    plt.title(f'Predicted: {y_pred[i]}')
    plt.axis('off')
plt.show()

```

Output:-

Accuracy: 0.99				
Classification Report:				
	precision	recall	f1-score	support
0	0.99	1.00	0.99	82
1	1.00	1.00	1.00	89
2	1.00	1.00	1.00	83
3	0.99	0.97	0.98	93
4	1.00	1.00	1.00	93
5	0.99	0.98	0.98	99
6	1.00	0.98	0.99	98
7	0.98	0.99	0.98	87
8	0.97	1.00	0.98	83
9	0.97	0.97	0.97	92
accuracy			0.99	899
macro avg	0.99	0.99	0.99	899
weighted avg	0.99	0.99	0.99	899



8. Write a python program to classify different face images using CNN.

Code:-

```

import tensorflow as tf
from tensorflow.keras import datasets, layers, models
import matplotlib.pyplot as plt
import numpy as np
(X_train, y_train), (X_test, y_test) =
datasets.cifar10.load_data()
X_train.shape

```



```

X_test.shape

y_train.shape
y_train[:5]

y_train = y_train.reshape(-1,)
y_train[:5]
y_test = y_test.reshape(-1,)
classes =
["airplane","automobile","bird","cat","deer","dog","frog","horse",
"ship","truck"]

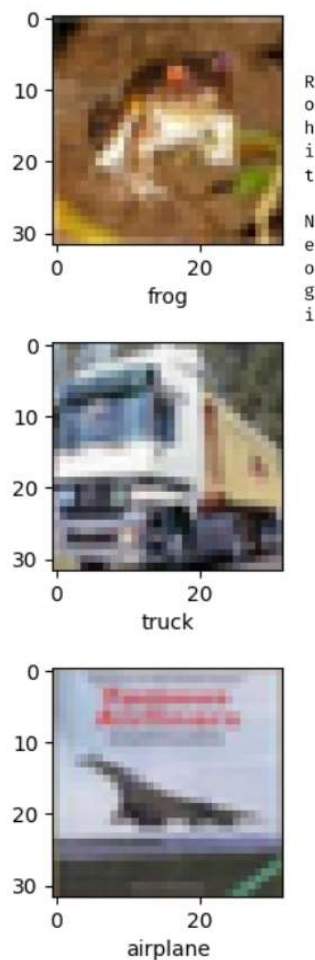
def plot_sample(X, y, index):
    plt.figure(figsize = (15,2))
    plt.imshow(X[index])
    plt.xlabel(classes[y[index]])

plot_sample(X_train, y_train, 0)
plot_sample(X_train, y_train, 1)
plot_sample(X_test, y_test,3)

X_train = X_train / 255.0
X_test = X_test / 255.0

```

Output:-




9. Write a python program to identify a person from the walking style (gait recognition) using convolutional recurrent neural network.

Code:-

```
import tensorflow as tf
from tensorflow.keras import layers, models
model = models.Sequential([
    layers.Conv2D(64, (3, 3), activation='relu', input_shape=(64,
64, 1)),
    layers.MaxPooling2D((2, 2)),
    layers.Conv2D(128, (3, 3), activation='relu'),
    layers.MaxPooling2D((2, 2)),
    layers.Flatten(),
    layers.RepeatVector(10),
    layers.LSTM(64, return_sequences=True),
    layers.TimeDistributed(layers.Dense(1, activation='sigmoid'))
])
7
model.compile(optimizer='adam',
loss='binary_crossentropy', metrics=['accuracy'])
print("Model summary for gait recognition:")
model.summary()
```

Output:

 /usr/local/lib/python3.10/dist-packages/keras/src/layers/convolutional/base_conv.py:107: UserWarning: super().__init__(activity_regularizer=activity_regularizer, **kwargs)

Model summary for gait recognition:
Model: "sequential"

Layer (type)	Output Shape	Param #
conv2d (Conv2D)	(None, 62, 62, 64)	640
max_pooling2d (MaxPooling2D)	(None, 31, 31, 64)	0
conv2d_1 (Conv2D)	(None, 29, 29, 128)	73,856
max_pooling2d_1 (MaxPooling2D)	(None, 14, 14, 128)	0
flatten (Flatten)	(None, 25088)	0
repeat_vector (RepeatVector)	(None, 10, 25088)	0
lstm (LSTM)	(None, 10, 64)	6,439,168
time_distributed (TimeDistributed)	(None, 10, 1)	65

Total params: 6,513,729 (24.85 MB)
Trainable params: 6,513,729 (24.85 MB)
Non-trainable params: 0 (0.00 B)

Rohit Neogi

10. Write a python program to classify breast cancer from histopathological images using VGG-16 and DenseNet-201 CNN architectures.

Code:-

```
import tensorflow as tf
from tensorflow.keras.applications import VGG16, DenseNet201
from tensorflow.keras import layers, models
from tensorflow.keras.preprocessing.image import ImageDataGenerator

# Load pre-trained VGG16 and DenseNet201 models without the top
layer
vgg16_model = VGG16(weights='imagenet', include_top=False,
input_shape=(224, 224, 3))
densenet_model = DenseNet201(weights='imagenet', include_top=False,
input_shape=(224, 224, 3))

# Function to create a model with a base pre-trained model
def create_model(base_model):
    model = models.Sequential([
        base_model,
        layers.Flatten(),
        layers.Dense(256, activation='relu'),
        layers.Dropout(0.5),
        layers.Dense(1, activation='sigmoid')
    ])
    model.compile(optimizer='adam', loss='binary_crossentropy',
metrics=['accuracy'])
    return model

# Create models using VGG16 and DenseNet201 as base models
vgg16_cancer_model = create_model(vgg16_model)
densenet_cancer_model = create_model(densenet_model)

# Print the summary of both models
print("VGG-16 model summary:")
vgg16_cancer_model.summary()

print("DenseNet-201 model summary:")
densenet_cancer_model.summary()
```

Output:

Downloading data from https://storage.googleapis.com/tensorflow/keras-applications/vgg16/vgg16_weights_tf_58889256/58889256 56s 1us/step
Downloading data from https://storage.googleapis.com/tensorflow/keras-applications/densenet/densenet201_weights_tf_74836368/74836368 59s 1us/step
VGG-16 model summary:
Model: "sequential"

Layer (type)	Output Shape	Param #
vgg16 (Functional)	(None, 7, 7, 512)	14,714,688
flatten (Flatten)	(None, 25088)	0
dense (Dense)	(None, 256)	6,422,784
dropout (Dropout)	(None, 256)	0
dense_1 (Dense)	(None, 1)	257

Total params: 21,137,729 (80.63 MB)
Trainable params: 21,137,729 (80.63 MB)
Non-trainable params: 0 (0.00 B)
DenseNet-201 model summary:
Model: "sequential_1"

Layer (type)	Output Shape	Param #
densenet201 (Functional)	(None, 7, 7, 1920)	18,321,984
flatten_1 (Flatten)	(None, 94080)	0
dense_2 (Dense)	(None, 256)	24,084,736
dropout_1 (Dropout)	(None, 256)	0
dense_3 (Dense)	(None, 1)	257

Total params: 42,406,977 (161.77 MB)
Trainable params: 42,177,921 (160.90 MB)
Non-trainable params: 229,056 (894.75 KB)

Rohit Neogi