Traditional features construction

Feature extraction from EACH image from the training data using image feature descriptor SIFT (Scale-invariant feature transform) (see https://docs.opencv.org/4.x/da/df5/ tutorial_py_sift_intro.html) with 128 dimensions for each keypoint (Note that you should obtain many keypoints from each image).Plot the keypoints on one image from your training dataset (see Fig 1 for one such figure). (0.5 point)

```
import numpy as np
import cv2 as cv
from IPython.display import Image, display
# Read the image
img = cv.imread('/content/drive/MyDrive/TrainingImages/Apple Golden 1/0_100.jpg')
# Convert the image to grayscale
gray = cv.cvtColor(img, cv.COLOR_BGR2GRAY)
# Initialize the SIFT detector with 180 keypoints
sift = cv.SIFT_create(nfeatures=180)
# Detect keypoints
kp = sift.detect(gray, None)
# Draw keypoints on the original image
img_with_keypoints = cv.drawKeypoints(gray, kp, img)
# Save the image with keypoints
cv.imwrite('sift_keypoints.jpg', img_with_keypoints)
# Display the image
display(Image('sift_keypoints.jpg'))
```



Create a new keypoints dataset K P that consists of all the keypoints from all the training images.

Perform K-mean clustering such that K = 100 on KP.

Use the learned K-mean clusters (seehttps://scikit-learn.org/stable/modules/clustering. html) to construct a 100-D vector for each image.

Example: For Image A, we have 20 keypoints in Cluster 1 and 10 keypoints in cluster 2, and no keypoints in the other clusters, then the vector representing Image A is (20,10,0,0,0,...,0,0)

Create a new 100-D dataset D consisting of vectors constructed from the training images.

```
from sklearn.cluster import KMeans
import cv2 as cv
import numpy as np
import os
# Initialize an empty list to store keypoints and labels
labels = []
# List of directories containing your training images
dir paths = ['/content/drive/MvDrive/TrainingImages/Apple Golden 1'.
             '/content/drive/MyDrive/TrainingImages/Apple Red 1',
             '/content/drive/MyDrive/TrainingImages/Apple Red Delicious',
             '/content/drive/MyDrive/TrainingImages/Avocado',
             '/content/drive/MyDrive/TrainingImages/Ginger Root',
             '/content/drive/MyDrive/TrainingImages/Limes',
             '/content/drive/MyDrive/TrainingImages/Maracuja',
             '/content/drive/MyDrive/TrainingImages/Papaya',
             '/content/drive/MyDrive/TrainingImages/Potato Red',
             '/content/drive/MyDrive/TrainingImages/Rambutan'
             # Add more directory paths as needed
            1
# Initialize SIFT detector
sift = cv.SIFT_create()
# Initialize a list to store the image vectors
image_vectors = []
# Initialize KMeans with the desired number of clusters
kmeans = KMeans(n clusters=100)
# Loop through each directory
for label, dir path in enumerate(dir paths):
   # Loop through each image file in the directory
   for filename in os.listdir(dir path):
       # Read the image
        img = cv.imread(os.path.join(dir path. filename))
       # Convert the image to grayscale
       gray = cv.cvtColor(img, cv.COLOR_BGR2GRAY)
       # Detect keypoints
        kp = sift.detect(gray, None)
        # Append keypoints to list
       KP.append(kp)
       # Append label to the labels list
        labels.append(label)
# Fit KMeans to keypoints
kmeans.fit(np.concatenate([np.array([kp.pt for kp in kp]) for kp in KP]))
# Loop through each directory again to extract features
for label, dir_path in enumerate(dir_paths):
   # Loop through each image file in the directory
   for filename in os.listdir(dir path):
       # Read the image
```

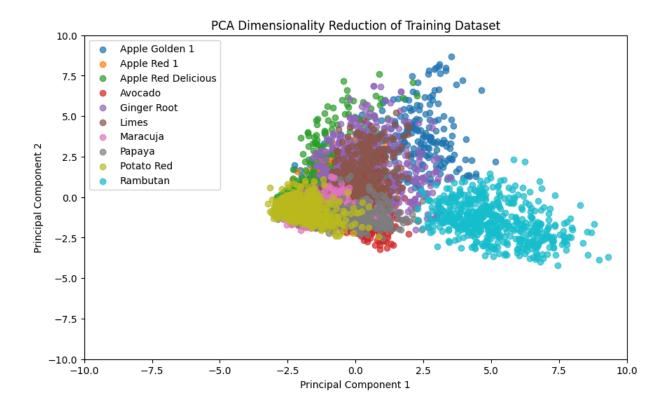
```
img = cv.imread(os.path.join(dir_path, filename))
        # Convert the image to grayscale
        gray = cv.cvtColor(img, cv.COLOR BGR2GRAY)
        # Detect keypoints
        kp = sift.detect(gray, None)
        # Get cluster labels for the keypoints of this image
        image labels = kmeans.predict(np.array([kp.pt for kp in kp]))
        # Count the occurrences of each cluster label
        cluster_counts = np.bincount(image_labels, minlength=100)
        # Append the cluster counts to the list of image vectors
        image vectors.append(cluster counts)
# Convert the list of image vectors to a numpy array
image vectors array = np.array(image vectors)
# Ensure each vector follows the specified structure
for i in range(len(image vectors array)):
    # Find the indices where the count is nonzero
   nonzero_indices = np.nonzero(image_vectors_array[i])[0]
    # Create a new vector with counts only at those indices
    new vector = np.zeros like(image vectors arrav[i])
    new vector[nonzero indices] = image vectors array[i][nonzero indices]
    # Replace the original vector with the new one
    image vectors array[i] = new vector
# Convert the list of labels to a numpy array
labels array = np.array(labels)
# Print an example vector
example vector = image vectors array[0]
print("Example vector representing Image A:", example_vector)
# Print the shape of the dataset D
print("Shape of dataset D:", image vectors array.shape)
     /usr/local/lib/python3.10/dist-packages/sklearn/cluster/ kmeans.py:870: FutureWarning: The default value of `n_init` will change from 10 to 'auto' in 1.4. Set
      warnings.warn(
     Example vector representing Image A: [1 0 0 0 1 0 0 0 1 1 0 0 0 1 0 0 2 2 0 3 0 2 1 0 0 0 1 0 0 1 1 1 0 0 0 0
     \begin{smallmatrix} 0 & 0 & 1 & 1 & 3 & 2 & 0 & 1 & 2 & 0 & 1 & 0 & 0 & 2 & 0 & 0 & 0 & 0 & 1 & 4 & 1 & 1 & 0 & 0 & 0 & 2 & 2 & 2 & 0 & 1 & 2 & 1 & 0 & 0 & 3 & 1 \\ \end{smallmatrix}
     Shape of dataset D: (4665, 100)
```

Dimensionality reduction(using Principal Component Analysis,PCA)(seehttps://scikit-learn.org/stable/modules/generated/sklearn.decomposition.PCA.html for PCA. https:// scikit-learn.org/stable/auto_examples/decomposition/plot_pca_iris.html for code example.)

- i. Perform Principal Component Analysis (PCA) dimensionality reduction Dataset D to 2 dimensions. (Note: You should not use the class labels)
- ii. Plot the 2D points using 10 different colors/symbols for data from the 10 classes (see Figure 2 for an example of the plot without normalization).(1 point)

```
from sklearn.decomposition import PCA
import matplotlib.pyplot as plt
import os
# List of directories containing your training images
train dir paths = ['/content/drive/MyDrive/TrainingImages/Apple Golden 1',
                   '/content/drive/MyDrive/TrainingImages/Apple Red 1',
                   '/content/drive/MyDrive/TrainingImages/Apple Red Delicious',
                   '/content/drive/MyDrive/TrainingImages/Avocado',
                   '/content/drive/MyDrive/TrainingImages/Ginger Root',
                   '/content/drive/MvDrive/TrainingImages/Limes'.
                   '/content/drive/MyDrive/TrainingImages/Maracuja',
                   '/content/drive/MyDrive/TrainingImages/Papaya',
                   '/content/drive/MyDrive/TrainingImages/Potato Red',
                   '/content/drive/MvDrive/TrainingImages/Rambutan'l
# Initialize a list to store the image vectors
train image vectors = []
# Initialize a list to store the corresponding class labels
train labels = []
# Loop through each directory
for label, dir path in enumerate(train dir paths):
   # Loop through each image file in the directory
    for filename in os.listdir(dir_path):
        # Read the image
        img = cv.imread(os.path.join(dir path, filename))
        # Convert the image to gravscale
        gray = cv.cvtColor(img, cv.COLOR_BGR2GRAY)
       # Detect keypoints
        kp = sift.detect(grav. None)
       # Get cluster labels for the keypoints of this image
        image labels = kmeans.predict(np.array([kp.pt for kp in kp]))
        # Count the occurrences of each cluster label
        cluster counts = np.bincount(image labels, minlength=100)
        # Append the cluster counts to the list of image vectors
       train_image_vectors.append(cluster_counts)
        # Assign class label based on folder name
       train_labels.append(os.path.basename(dir_path))
# Convert the lists to numpy arrays
train_image_vectors_array = np.array(train_image_vectors)
train labels array = np.array(train labels)
# Perform PCA dimensionality reduction to 2 dimensions
pca = PCA(n components=2)
D pca = pca.fit transform(train image vectors array)
# Plot the 2D points with different colors/symbols for each class
plt.figure(figsize=(10, 6))
for label in np.unique(train_labels_array):
    class_indices = np.where(train_labels_array == label)[0]
    plt.scatter(D pca[class indices, 0], D pca[class indices, 1], label=f'{label}', alpha=0.7)
plt.xlabel('Principal Component 1')
plt.ylabel('Principal Component 2')
```

```
plt.title('PCA Dimensionality Reduction of Training Dataset')
plt.legend()
plt.xlim(-10, 10)
plt.ylim(-10, 10)
plt.show()
```



"Traditional" Machine Learning Model - Support Vector Machine (SVM)

- (a) We perform model selection to learn a linear SVM (see https://scikit-learn.org/stable/ modules/generated/sklearn.svm.SVC.html) prediction model (i.e., kernel = 'linear').
- (b) Train the SVM model using different C parameters. In particular, you will use the following values: 0.01, 0.1, 1.0, 100.
- (c) Plot a graph to show the SVM performance on the training data and test data similar to
- (d) Repeat Step (b) and (c) for kernel = 'rbf', 'poly', 'sigmoid'. (2 points include 0.5 for linear kernel.)
- (e) Plot a graph to compare the best performance for SVM using the four kernels on the training data and test data similar to Figure 4 (note that you identify the best results for each kernel from results in Step (d))

First, performing the same steps on testing data to create testing dataset

Combine all the keypoints from all the testing images into a single dataset.

Use the 100 cluster centers obtained from the training data to create a 100D vector for each test image.

Process the test images into the test dataset.

This code will create the test_image_vectors_array with the features obtained using the cluster centers obtained from the training data.

```
# List of directories containing your testing images
test_dir_paths = ['/content/drive/MyDrive/TestingImages/Apple Golden 1',
                  '/content/drive/MyDrive/TestingImages/Apple Red 1',
                  '/content/drive/MyDrive/TestingImages/Apple Red Delicious',
                  '/content/drive/MyDrive/TestingImages/Avocado',
                  '/content/drive/MyDrive/TestingImages/Ginger Root',
                  '/content/drive/MyDrive/TestingImages/Limes',
                  '/content/drive/MvDrive/TestingImages/Maracuia'.
                  '/content/drive/MyDrive/TestingImages/Papaya',
                  '/content/drive/MyDrive/TestingImages/Potato Red',
                  '/content/drive/MvDrive/TestingImages/Rambutan'.
                  # Add more directory paths as needed
# Initialize an empty list to store keypoints for testing data
test KP = []
# Initialize a list to store the image vectors for testing data
test image vectors = []
# Initialize a list to store the labels for testing data
test_labels = []
# Loop through each directory for testing data
for label, dir_path in enumerate(test_dir_paths):
   # Loop through each image file in the directory
   for filename in os.listdir(dir_path):
       # Read the image
        img = cv.imread(os.path.join(dir_path, filename))
       # Convert the image to grayscale
       gray = cv.cvtColor(img, cv.COLOR_BGR2GRAY)
        # Detect keypoints
        kp = sift.detect(grav. None)
        # Append keypoints to the test_KP dataset
        test KP.extend(kp)
       # Get cluster labels for the keypoints of this image for testing data
        image_labels = kmeans.predict(np.array([kp.pt for kp in kp]))
        # Count the occurrences of each cluster label
        cluster counts = np.bincount(image labels, minlength=100)
       # Append the cluster counts to the list of image vectors for testing data
        test image vectors.append(cluster counts)
       # Append label to the test_labels list
       test labels.append(label)
# Convert the list of keypoints to an array for testing data
test KP array = np.array([kp.pt for kp in test KP])
# Convert the list of image vectors to a numpy array for testing data
```

```
test_image_vectors_array = np.array(test_image_vectors)
# Convert the list of labels to a numpy array for testing data
test labels array = np.array(test labels)
# Ensure each vector follows the specified structure for testing data
for i in range(len(test image vectors array)):
   # Find the indices where the count is nonzero
   nonzero_indices = np.nonzero(test_image_vectors_array[i])[0]
   # Create a new vector with counts only at those indices
   new_vector = np.zeros_like(test_image_vectors_array[i])
   new vector[nonzero indices] = test image vectors array[i][nonzero indices]
   # Replace the original vector with the new one for testing data
   test image vectors array[i] = new vector
# Print an example vector for testing data
example vector test = test image vectors array[0] # Assuming the first image is Image A
print("Example vector representing Image A for testing data:", example vector test)
# Print the shape of the testing dataset D
print("Shape of testing dataset D:", test_image_vectors_array.shape)
    Example vector representing Image A for testing data: [0 2 1 0 0 0 0 0 0 0 1 1 0 0 0 0 0 0 1 0 0 0 0 0 3 1 0 0 1 1 1 0 4 1 0
     4 0 2 1 2 0 0 2 2 0 0 3 1 1 0 1 0 1 0 0 0 0 2 0 0 0]
    Shape of testing dataset D: (1542, 100)
```

Now, we train the model and test it (using classes from the training images as labels)

The below code uses the default kernel function for the Support Vector Classifier (SVC) in scikit-learn, which is the Radial Basis Function (RBF) kernel.

```
from sklearn.svm import SVC
from sklearn.metrics import accuracy_score
# Define C values
C_values = [0.01, 0.1, 1.0, 10, 100]
# Initialize dictionaries to store results
train scores = {}
test_scores = {}
# Train SVM models for different C values
for C in C_values:
   # Initialize SVM model
   svm_model = SVC(C=C)
   # Train the model
    svm_model.fit(image_vectors_array, labels_array)
   # Predictions on training data
    train_predictions = svm_model.predict(image_vectors_array)
   # Accuracy on training data
    train_accuracy = accuracy_score(labels_array, train_predictions)
    train_scores[C] = train_accuracy
    # Predictions on testing data
    test_predictions = svm_model.predict(test_image_vectors_array)
    # Accuracy on testing data
    test_accuracy = accuracy_score(test_labels_array, test_predictions)
    test_scores[C] = test_accuracy
# Print the results
for C in C_values:
   print(f"C Value: {C}")
   print(f"Training Accuracy: {train_scores[C]:.4f}")
   print(f"Testing Accuracy: {test_scores[C]:.4f}")
    print()
    C Value: 0.01
    Training Accuracy: 0.2114
    Testing Accuracy: 0.2095
    C Value: 0.1
    Training Accuracy: 0.8845
    Testing Accuracy: 0.6589
    C Value: 1.0
    Training Accuracy: 0.9908
    Testing Accuracy: 0.7309
    C Value: 10
    Training Accuracy: 1.0000
    Testing Accuracy: 0.7302
    C Value: 100
```

Training Accuracy: 1.0000 Testing Accuracy: 0.7302

In the context of Support Vector Machines (SVM), C values represent the regularization parameter.

Regularization is a technique used to prevent overfitting by adding a penalty term to the loss function. In SVM, the parameter C controls the trade-off between maximizing the margin (decision boundary) and minimizing the classification error.

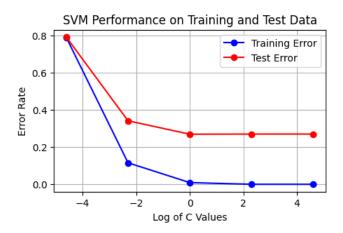
A small C value allows for a larger margin but may misclassify some points (soft margin). A large C value penalizes misclassifications heavily, potentially resulting in a smaller margin (hard margin).

In essence, C values determine the balance between achieving a low training error and generalizing well to unseen data.

```
import matplotlib.pyplot as plt

# Calculate error rates
train_errors = [1 - train_scores[C] for C in C_values]
test_errors = [1 - test_scores[C] for C in C_values]

# Plotting
plt.figure(figsize=(5, 3))
plt.plot(np.log(C_values), train_errors, marker='o', label='Training Error', color='blue')
plt.plot(np.log(C_values), test_errors, marker='o', label='Test Error', color='red')
plt.xlabel('Log of C Values')
plt.ylabel('Error Rate')
plt.title('SVM Performance on Training and Test Data')
plt.grid(True)
plt.legend()
plt.show()
```

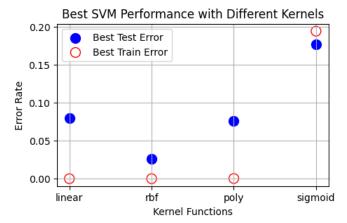


An error rate of 0.3 means that 30% of the samples in the test data are classified incorrectly by the model.

We repeat for kernel = 'linear', 'RBF', 'poly', 'sigmoid'

```
import numpy as np
import matplotlib.pyplot as plt
from sklearn.svm import SVC
from sklearn.metrics import accuracy score
from sklearn.model selection import train test split
# Define C values
C \text{ values} = [0.01, 0.1, 1.0, 10, 100]
# Split the data into training and testing sets
train image vectors array, test image vectors array, train labels array, test labels array = \
    train test split(image vectors array, labels array, test size=0.2, random state=42)
# Define kernel functions
kernel_functions = ['linear', 'rbf', 'poly', 'sigmoid']
# Initialize dictionaries to store results
train errors = {kernel: [] for kernel in kernel functions}
test_errors = {kernel: [] for kernel in kernel_functions}
best test errors = {kernel: float('inf') for kernel in kernel functions}
best train errors = {kernel: float('inf') for kernel in kernel functions}
best_test_C_values = {kernel: None for kernel in kernel_functions}
best_train_C_values = {kernel: None for kernel in kernel_functions}
# Train SVM models for different kernel functions and C values
for kernel in kernel functions:
   for C in C_values:
        # Initialize SVM model
        svm_model = SVC(kernel=kernel, C=C)
        # Train the model
        svm_model.fit(train_image_vectors_array, train_labels_array)
       # Predictions on training data
       train predictions = svm model.predict(train image vectors array)
       # Error rate on training data
       train_error = 1.0 - accuracy_score(train_labels_array, train_predictions)
       # Predictions on testing data
       test_predictions = svm_model.predict(test_image_vectors_array)
       # Error rate on testing data
       test_error = 1.0 - accuracy_score(test_labels_array, test_predictions)
       # Store errors
       train errors[kernel].append(train error)
       test_errors[kernel].append(test_error)
       # Update best test error for this kernel
       if test error < best test errors[kernel]:</pre>
            best test errors[kernel] = test error
```

```
best_test_C_values[kernel] = C
        # Update best train error for this kernel
        if train error < best train errors[kernel]:</pre>
            best_train_errors[kernel] = train_error
            best train C values[kernel] = C
# Plotting
plt.figure(figsize=(5, 3))
# Plot testing data errors
for kernel in kernel functions:
   plt.scatter([kernel], [best test errors[kernel]], color='blue', marker='o', s=100)
# Plot training data errors
for kernel in kernel functions:
   plt.scatter([kernel], [best_train_errors[kernel]], color='none', edgecolor='red', s=100)
# Add labels outside the loop
plt.scatter([], [], color='blue', marker='o', label='Best Test Error', s=100)
plt.scatter([], [], color='none', edgecolor='red', label='Best Train Error', s=100)
# Add legend
plt.legend()
plt.xlabel('Kernel Functions')
plt.ylabel('Error Rate')
plt.title('Best SVM Performance with Different Kernels')
plt.grid(True)
plt.legend()
plt.show()
# Find the best C value and kernel
best kernel = None
best C value = None
best_test_error = float('inf')
best_train_error = float('inf')
for kernel in kernel_functions:
   if best_test_errors[kernel] < best_test_error:</pre>
       best_test_error = best_test_errors[kernel]
       best_kernel = kernel
       best C value = best test C values[kernel]
   if best_train_errors[kernel] < best_train_error:</pre>
        best_train_error = best_train_errors[kernel]
# Print the results
print("Best C value:", best_C_value)
print("Best kernel:", best_kernel)
print("Testing error with the best parameters:", best test error)
print("Training error with the best parameters:", best_train_error)
```



Best C value: 10 Best kernel: rbf

Testing error with the best parameters: 0.025723472668810254

Training error with the best parameters: 0.0

This code plots the best performance (lowest error rate) for each kernel for both the testing and training datasets. It identifies the best performing C value for each kernel based on the lowest error rate achieved on the respective dataset.

Let's test and see what label our trained model will predict for one of our testing images

```
import cv2
import numpy as np
from sklearn.svm import SVC
from sklearn.metrics import accuracy_score
# Define C value
C = 1.0
# Initialize SVM model
svm_model = SVC(C=C, kernel='linear')
# Train the SVM model
svm_model.fit(image_vectors_array, labels_array)
# Read the testing image
image_path = '/content/drive/MyDrive/TestingImages/Apple Red Delicious/13_100.jpg'
testing_image = cv2.imread(image_path)
# Check if the image is loaded successfully
if testing_image is not None:
   # Convert the testing image to grayscale
   gray_testing_image = cv2.cvtColor(testing_image, cv2.COLOR_BGR2GRAY)
```

```
# Detect keypoints
    kp = sift.detect(gray_testing_image, None)
    # Get cluster labels for the keypoints of this image
    image_labels = kmeans.predict(np.array([kp.pt for kp in kp]))
   # Count the occurrences of each cluster label
    cluster_counts = np.bincount(image_labels, minlength=100)
    # Reshape the cluster counts to match the shape of the training data
    input_vector = cluster_counts.reshape(1, -1)
    # Use the trained SVM model to predict the label of the testing image
    predicted_label = svm_model.predict(input_vector)
# Define the class labels
    class_labels = ['Apple Golden 1', 'Apple Red 1', 'Apple Red Delicious', 'Avocado', 'Ginger Root',
                    'Limes', 'Maracuja', 'Papaya', 'Potato Red', 'Rambutan']
   # Map the predicted index to the corresponding class label
    predicted_class_label = class_labels[predicted_label[0]]
   # Print the predicted class label
   print(f"Predicted Class Label: {predicted_class_label}")
   # Display the testing image
    plt.imshow(cv2.cvtColor(testing_image, cv2.COLOR_BGR2RGB))
    plt.axis('off')
    nl+ chow()
```