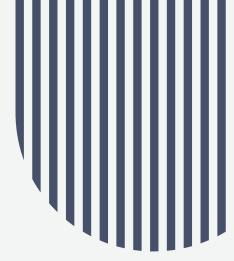


supervized ML



Supervised learning is a type of machine learning where the computer is taught using examples. These examples have both the input (what you know) and the output (what you want to predict). The goal is for the computer to learn how to predict the output based on the input.

process:



- Data Collection: Gather a labeled dataset, where each example consists of an input-output pair.
- 2. Data Preprocessing: Clean the data, handle missing values, normalize or scale features, and possibly perform feature engineering.
- 3. Train-Test Split: Split the dataset into training and testing sets to evaluate the model's performance.
- 4. Model Selection: Choose a suitable algorithm for the problem at hand (e.g., linear regression for regression tasks, decision trees for classification tasks).
- 5. Training: Use the training data to train the model, adjusting the model's parameters to minimize the error between predicted and actual outputs.
- 6. Evaluation: Assess the model's performance on the test set using appropriate metrics (e.g., accuracy, precision, recall for classification; mean squared error, R-squared for regression).
- 7. Hyperparameter Tuning: Optimize the model's hyperparameters to improve performance.
- 8. Prediction: Use the trained model to make predictions on new, unseen data.

Algorithm used in this project:

for classifying images of cats and dogs, we extract features from the images and then use traditional machine learning algorithms such as Decision tree and k-Nearest Neighbors (k-NN). here is a steps for what we do:

1- import libraries:

```
import os
import numpy as np
from tensorflow.keras.preprocessing.image import load_img, img_to_array
from sklearn.preprocessing import LabelEncoder
from sklearn.model_selection import train_test_split
from sklearn.neighbors import KNeighborsClassifier
from sklearn.tree import DecisionTreeClassifier
from sklearn.metrics import accuracy_score, classification_report, confusion_matrix
import matplotlib.pyplot as plt

Your paragraph text
```

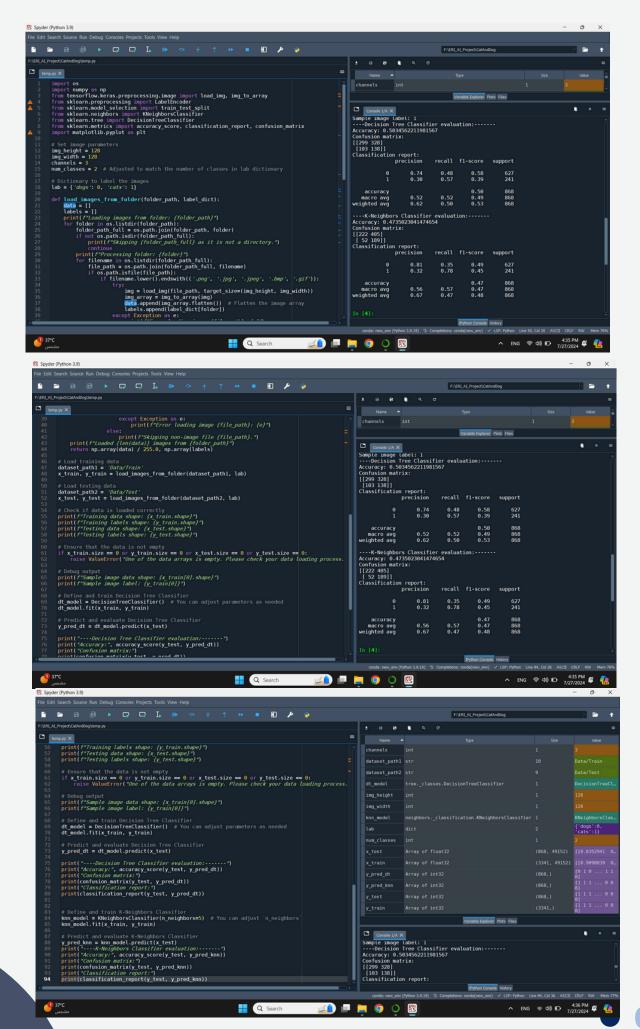
2- Load and Preprocess Images:

```
img height = 128
img width = 128
channels = 3
num classes = 2 # Adjusted to match the number of classes in lab dictionary
# Dictionary to label the images
lab = { 'dogs': 0, 'cats': 1}
def load_images_from_folder(folder_path, label_dict):
    data = []
    labels = []
    print(f"Loading images from folder: {folder_path}")
    for folder in os.listdir(folder_path):
        folder_path_full = os.path.join(folder_path, folder)
        if not os.path.isdir(folder_path_full):
            print(f"Skipping {folder_path_full} as it is not a directory.")
        print(f"Processing folder: {folder}")
for filename in os.listdir(folder_path_full):
            file_path = os.path.join(folder_path_full, filename)
            if os.path.isfile(file_path):
                 if filename.lower().endswith(('.png', '.jpg', '.jpeg', '.bmp', '.gif')):
                         img = load_img(file_path, target_size=(img_height, img_width))
                         img_array = img_to_array(img)
                         data.append(img_array.flatten()) # Flatten the image array
                         labels.append(label_dict[folder])
                    except Exception as e:
```

3-Train decision tree Model and k-nn model and evaluate them:

```
dt model = DecisionTreeClassifier() # You can adjust parameters as needed
      dt_model.fit(x_train, y_train)
      # Predict and evaluate Decision Tree Classifier
      y_pred_dt = dt_model.predict(x_test)
      print("----Decision Tree Classifier evaluation:-----")
      print("Accuracy:", accuracy_score(y_test, y_pred_dt))
     print("Confusion matrix:")
print(confusion_matrix(y_test, y_pred_dt))
      print("Classification report:")
      print(classification_report(y_test, y_pred_dt))
      # Define and train K-Neighbors Classifier
      knn_model = KNeighborsClassifier(n_neighbors=5) # You can adjust `n_neighbors
84
      knn_model.fit(x_train, y_train)
      # Predict and evaluate K-Neighbors Classifier
      y pred knn = knn model.predict(x test)
      print("----K-Neighbors Classifier evaluation:-----")
     print("Accuracy:", accuracy_score(y_test, y_pred_knn))
     print( "Confusion matrix: ")
      print(confusion_matrix(y_test, y_pred_knn))
      print("Classification report:")
94
     print(classification report(y test, y pred knn))
```

```
Console 1/A X
Sample image label: 1
----Decision Tree Classifier evaluation:-----
Accuracy: 0.5034562211981567
Confusion matrix:
[[299 328]
 [103 138]]
Classification report:
                           recall f1-score
              precision
                                               support
                   0.74
                             0.48
                                        0.58
                                                   627
           1
                   0.30
                             0.57
                                        0.39
                                                   241
                                        0.50
                                                   868
   accuracy
                                        0.49
  macro avg
                   0.52
                             0.52
                                                   868
                             0.50
                                        0.53
                                                   868
weighted avg
                   0.62
----K-Neighbors Classifier evaluation:-----
Accuracy: 0.4735023041474654
Confusion matrix:
[[222 405]
 [ 52 189]]
Classification report:
              precision
                            recall f1-score
                                               support
           0
                   0.81
                             0.35
                                        0.49
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           1
                   0.32
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                                        0.47
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    accuracy
                   0.56
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                                        0.47
                                                   868
  macro avg
                                        0.48
weighted avg
                   0.67
                             0.47
                                                   868
```



Unsupervized ML

Used Histogram of Oriented Gradients (HOG) for feature extraction, Principal Component Analysis (PCA) for dimensionality reduction, and KMeans clustering for grouping similar images. The project evaluates the clustering results against the ground truth labels and visualizes the clusters and classification results. (How ever because the images are complex this approach wasn't the best for this case)

Data Preprocessing

Image Loading and Preprocessing

Images are loaded from separate folders for training and testing datasets. Each image is resized to 64x64 pixels and normalized by dividing pixel values by 255.0. The labels are assigned as 0 for dogs and 1 for cats.

HOG Feature Extraction

HOG features are extracted from the grayscale images. The HOG descriptor captures the edge and gradient structure of the images, which is useful for distinguishing between different object categories. python

Standardization and Scaling

The extracted HOG features are standardized using StandardScaler and optionally scaled to a [0, 1] range using MinMaxScaler.

```
60 # Standardize the HOG features
61 scaler_standard = StandardScaler()
62 train_hog_features_standard_scaled = scaler_standard.fit_transform(train_hog_features)
63 test_hog_features_standard_scaled = scaler_standard.transform(test_hog_features)
64
65 # Apply MinMaxScaler
66 scaler_minmax = MinMaxScaler(feature_range=(0, 1))
67 train_hog_features_minmax_scaled = scaler_minmax.fit_transform(train_hog_features)
68 test_hog_features_minmax_scaled = scaler_minmax.transform(test_hog_features)
69
```

Dimensionality Reduction

PCA is applied to the standardized HOG features to reduce dimensionality and retain 50 principal components. This step helps in reducing the computational cost and noise while preserving the essential features for clustering.

```
70 # Apply PCA for feature extraction
71 pca = PCA(n_components=50)
72 train_hog_features_pca = pca.fit_transform(train_hog_features_standard_scaled)
73 test_hog_features_pca = pca.transform(test_hog_features_standard_scaled)
74 print(f"Training PCA shape: {train_hog_features_pca.shape}")
75 print(f"Testing PCA shape: {test_hog_features_pca.shape}")
76
```

Clustering

Determining Optimal Number of Clusters

The elbow method is used to determine the optimal number of clusters by plotting the within-cluster sum of squares (inertia) for different values of k.

KMeans Clustering

Based on the elbow graph, the optimal number of clusters is chosen, and KMeans clustering is applied to the PCA-transformed features.

```
92 # Choose the optimal number of clusters based on the elbow graph
93 optimal_clusters = 2 # Set this to the number of clusters identified by the elbow method
94
95 # Apply KMeans clustering with the optimal number of clusters
96 kmeans = KMeans(n_clusters=optimal_clusters, init='k-means++', n_init=10, random_state=0)
97 train_clusters = kmeans.fit_predict(train_hog_features_pca)
98 test_clusters = kmeans.predict(test_hog_features_pca)
99
```

Evaluation

The clustering results are evaluated against the ground truth labels using confusion matrices and classification reports. Clusters are relabeled to align with the actual labels for better interpretability.

```
100 # Print evaluation metrics
101 print("Ground truth clusters:\n", test_clusters)
102 print("Estimated clusters:\n", test_clusters)
103 print("Confusion matrix:\n", confusion_matrix(test_labels, test_clusters))
104 print("Classification report:\n", classification_report(test_labels, test_clusters))
105
106 # Relabel clusters
107 relabel_clusters = [1 if cluster == 0 else 0 for cluster in test_clusters]
108
109 print("Relabel clusters:\n", relabel_clusters)
110 print("Confusion matrix (relabelled):\n", confusion_matrix(test_labels, relabel_clusters))
111 print("Classification report (relabelled):\n", classification_report(test_labels, relabel_clusters))
112
```

PCA Component Plot

The PCA components are plotted to visualize the clustering results

```
113 # Function to plot PCA components
114 def plot_pca_components(images_pca, labels, title):
115    plt.figure(figsize=(8, 6))
116    scatter = plt.scatter(images_pca[:, 0], images_pca[:, 1], c=labels, cmap='viridis',
        alpha=0.5)
117    plt.title(title)
118    plt.xlabel('PCA Component 1')
119    plt.ylabel('PCA Component 2')
120    plt.colorbar(scatter, label='Cluster')
121    plt.show()
122
123 # Plot PCA components of training data
124 plot_pca_components(train_hog_features_pca, train_clusters, 'KMeans Clustering Results
        (Training Data)')
125
```

Image Plot with Cluster Labels

A subset of test images is plotted with their predicted cluster labels.

```
126 # Function to plot images with cluster labels
127 def plot_images(images, labels, title, image_size=(64, 64)):
128     fig, axes = plt.subplots(2, 5, figsize=(12, 6))
129     fig.suptitle(title)
130     for i, ax in enumerate(axes.flat):
131         if i < len(images):
132             ax.imshow(images[i].reshape(image_size + (3,)))
133             ax.set_title(f'Cluster {labels[i]}')
134             ax.axis('off')
135         plt.show()
136
137 # Plot images with predicted clusters
138 plot_images(test_images, test_clusters, 'KMeans Clustering Results (Test Data)')
139</pre>
```

2D PCA Visualization

PCA is performed again to reduce the dimensionality to 2D for visualization of clustering results and ground truth labels.

```
140 # Perform PCA to reduce dimensionality to 2D for visualization

141 pca_2d = PCA(n_components=2)

142 pca_result = pca_2d.fit_transform(test_hog_features) # Use HOG features for PCA
visualization

143

144 # Visualize the clustering results

145 colormap = np.array(['green', 'red'])

146

147 plt.figure(figsize=(12, 6))

148

149 plt.subplot(1, 2, 1)

150 plt.title('Classification K-means')

151 plt.scatter(pca_result[:, 0], pca_result[:, 1], c=colormap[relabel_clusters],
label='Clusters')

152 plt.xlabel('Principal Component 1')

153 plt.ylabel('Principal Component 2')

154 plt.legend()

155

156 plt.subplot(1, 2, 2)

157 plt.title('Ground Truth')

158 plt.scatter(pca_result[:, 0], pca_result[:, 1], c=colormap[test_labels], label='Ground
Truth')

159 plt.xlabel('Principal Component 1')

160 plt.ylabel('Principal Component 2')

161 plt.legend()

162

163 plt.show()
```

Output:

0.49

0.49

weighted avg

0.49

0.49

0.49

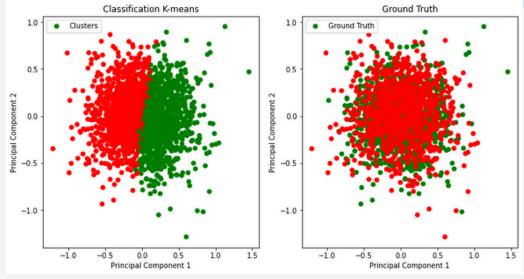
0.49

2023

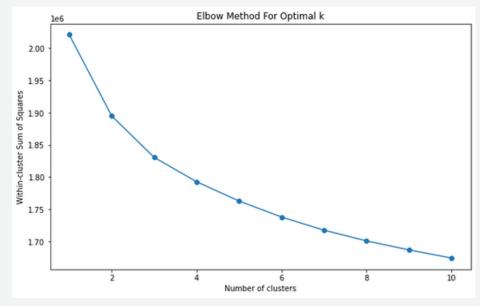
2023

```
1]: runfile('C:/Users/Salma/SpyderProjects/AIcourse/Unsupervised ML/scripts/main.py', wdir='C:/Users/Salma/SpyderProjects/
 AIcourse/Unsupervised ML/scripts')
Training images shape: (8005, 64, 64, 3)
Training labels shape: (8005,)
Testing images shape: (2023, 64, 64, 3)
Testing labels shape: (2023,)
Training PCA shape: (8005, 50)
Testing PCA shape: (8003, 50)
Testing PCA shape: (2023, 50)
                 Figures are displayed in the Plots pane by default. To make them also appear inline in the console, you need to uncheck "Mute inline plotting" under the options menu of Plots.
 Ground truth clusters:
[0 0 0 ... 1 1 1]
Estimated clusters:
 [1 1 0 ... 1 1 1]
 Confusion matrix:
 [[552 460]
[537 474]]
 lassification report:
                                precision
                                                             recall f1-score
                                                                                                      support
                                                                                     0.53
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 Classification report (relabelled):
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                                                                                                           1012
                                          0.49
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                                                                                                           1011
                                                                                                            2023
         accuracy
```

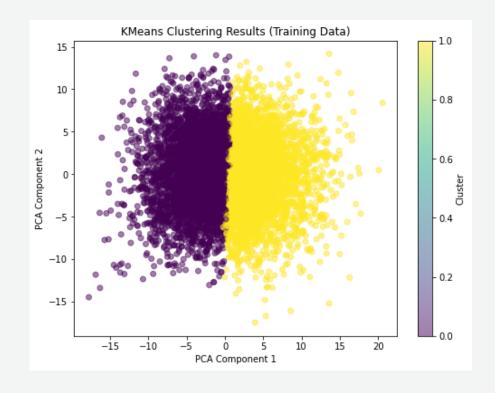
Classfication KMeans and Ground Truth



Elbow method for optimal K



KMeans Clustering Results (Trainging Data)



KMeans Clustering Results (Test Data)



Conclusion

The application of unsupervised learning techniques for image classification. The combination of HOG features, PCA, and KMeans clustering provides a method for grouping similar images without the need for labeled training data. The results, evaluated through confusion matrices and visualizations, show effectiveness of this approach in distinguishing between images cats. **Future** improvements of and could include dogs experimenting with different feature extraction methods. clustering algorithms, and further hyperparameter tuning to enhance performance.

Deep Learning

Deep learning is a subset of machine learning that uses multilayered neural networks, called deep neural networks, to simulate the complex decision-making power of the human brain..

Data Preprocessing

Image Loading and Preprocessing

Images are loaded from separate folders for training and testing datasets. Each image is resized to 64x64 pixels and normalized by dividing pixel values by 255.0. The labels are assigned as 0 for dogs and 1 for cats.

```
# Load training data
train_data = []
train_labels = []
train_base_path = 'C:/Users/t460/Downloads/archive/training_set/training_set'

for folder in os.listdir(train_base_path):
    folder_path = os.path.join(train_base_path, folder)
    if os.path.isdir(folder_path):
        for filename in os.listdir(folder_path):
            img_path = os.path.join(folder_path, filename)
            try:
            img = load_img(img_path, target_size=(img_height, img_width))
            img_array = img_to_array(img)
            train_data.append(img_array)
            train_labels.append(lab[folder])
            except Exception as e:
            print(f"Error Loading image {img_path}: {e}")
```

This section of the code focuses on preparing and normalizing the training data for a machine learning model. It converts the training data into a NumPy array and normalizes the pixel values by dividing by 255.0, which scales the values to a range between 0 and 1. This is a common preprocessing step for image data to ensure that the model trains more effectively. Additionally, the labels for the training data are also converted into a NumPy array. The code then proceeds to load the test data and labels from the specified directory path, preparing them for evaluation purposes.

```
30  x_train = np.array(train_data) / 255.0 # Normalize pixel values
31  y_train = np.array(train_labels)
32
33  # Load test data
34  test_data = []
35  test_labels = []
36  test_base_path = 'C:/Users/t460/DownLoads/archive/test_set/test_set'
```

This section of the code is responsible for loading and preprocessing the test data for the machine learning model. It iterates through each folder in the specified test data directory. For each folder, it checks if the path is a directory and then iterates through each image file within that directory. Each image is loaded and resized to the specified dimensions ('img_height' and 'img_width'). The image is then converted into an array and appended to the 'test_data' list. Correspondingly, the label for the image, determined by the folder name, is appended to the 'test_labels' list. If there is an error loading an image, it prints an error message. Finally, the test data and labels are converted into NumPy arrays, and the pixel values of the test data are normalized by dividing by 255.0, ensuring they are in the range of 0 to 1.

This section of the code defines, trains, and evaluates a Convolutional Neural Network (CNN) model for image classification:

1. Model Definition:

- The model is built using convolutional (Conv2D) and pooling (MaxPooling2D) layers to extract features from images.
- A flattening layer (Flatten) is used to convert the 2D matrix data to a vector, followed by dense (Dense) layers for classification.

2. Model Compilation:

 The model is compiled with the sparse_categorical_crossentropy loss function and adam optimizer.

3. Model Training:

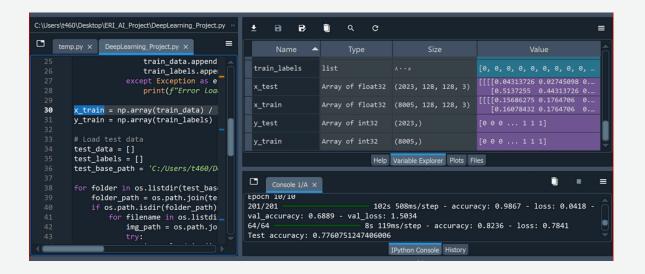
 The model is trained on the training data for 10 epochs with a batch size of 32, using 20% of the data for validation.

4. Model Evaluation:

 The model's accuracy is evaluated on the test data, and the test accuracy is printed.

```
# Define the model
model = Sequential()
model.add(Conv2D(32, (3, 3), activation='relu', input_shape=(img_height, img_width, 3)))
model.add(MaxPooling2D((2, 2)))
model.add(Conv2D(64, (3, 3), activation='relu'))
model.add(MaxPooling2D((2, 2)))
model.add(Conv2D(128, (3, 3), activation='relu'))
model.add(MaxPooling2D((2, 2)))
model.add(Flatten())
model.add(Dense(128, activation='relu'))
model.add(Dense(num_classes, activation='softmax'))
model.compile(loss='sparse_categorical_crossentropy', optimizer='adam', metrics=['accurac
# Print model summary
model.summary()
model.fit(x_train, y_train, epochs=10, batch_size=32, validation_split=0.2)
# Evaluate the model
test_loss, test_acc = model.evaluate(x_test, y_test)
print(f"Test accuracy: {test_acc}")
```

The accuracy



Our

Team

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