```
In [2]: import os
        # os.environ["CUDA_VISIBLE_DEVICES"] = "-1" # To use local CPU instead of local GPU
        import numpy as np, pandas as pd
        import matplotlib.pyplot as plt
        import matplotlib.image as implt
        from PIL import Image
        import glob
        from sklearn.utils import shuffle
        from sklearn import metrics
        from sklearn.model_selection import train_test_split, StratifiedKFold, cross_val_pr
        from sklearn.metrics import confusion_matrix, classification_report
        from sklearn.datasets import load files
        from io import BytesIO
        import seaborn as sns
        import xml.etree.ElementTree as ET
        from pathlib import Path
        from sklearn.neural_network import MLPClassifier
        from sklearn.naive_bayes import GaussianNB, MultinomialNB, BernoulliNB
        from sklearn.ensemble import RandomForestClassifier
        from skimage import exposure
        from sklearn.preprocessing import StandardScaler
        from sklearn.metrics import ConfusionMatrixDisplay
In [3]: os.listdir()
Out[3]: ['.git',
          '.ipynb_checkpoints',
          'anno',
          'Assignment2.ipynb',
          'Cropped',
          'Desktop',
          'DM1_ProgrammingAssignment1-Fall2024_712178661.pdf',
          'id_rsa_dond',
          'id_rsa_dond.pub',
          'image',
          'image.zip',
          'stanford-dogs-transfer-crop-stack.html',
          'stanford-dogs-transfer-crop-stack.ipynb',
          'stanford-dogs-transfer-crop-stack.pdf',
          'stanford-dogs-transfer-crop-stack_files',
          'student_8']
In [3]: %matplotlib inline
        %config InlineBackend.figure_format = 'svg'
        dog_images = glob.glob('./image/*/*')
        breeds = glob.glob('./anno/*')
        annotations = glob.glob('./anno/*/*')
        cropped = "./Cropped/"
        img_size = 299 # For Xception input
        train_dir = './Cropped' # './Images'
        batch_size_training = 256
        batch size validation = 256
        input_shape = (img_size,img_size,3)
```

```
####### Read X and Y coordinate ranges from an annotation #######
def get_bounding_boxes(annot):
   xml = annot
   tree = ET.parse(xml)
   root = tree.getroot()
   objects = root.findall('object')
   bbox = []
   for o in objects:
        bndbox = o.find('bndbox')
        xmin = int(bndbox.find('xmin').text)
       ymin = int(bndbox.find('ymin').text)
        xmax = int(bndbox.find('xmax').text)
        ymax = int(bndbox.find('ymax').text)
        bbox.append((xmin,ymin,xmax,ymax))
   return bbox
####### Get image path from annotation path #######
def get_image(annot):
   img_path = './image/'
   file = annot.split('\\')
   img_filename = img_path + file[-2] +'/'+file[-1]+'.jpg'
   return img_filename
####### Fill image with black to make a square (not used) #######
def make_square(im, min_size=100, fill_color=(0, 0, 0, 0)):
   x, y = im.size
   size = max(min_size, x, y)
   new_im = Image.new('RGB', (size, size), fill_color)
   new_im_paste(im, (int((size - x) / 2), int((size - y) / 2)))
   return new_im
print(len(dog_images), len(breeds), len(annotations))
```

767 4 767

```
In [4]: plt.figure(figsize=(10,6))
for i in range(8):
    plt.subplot(2,4,i+1)
    plt.axis("off")
    dog = get_image(annotations[i])
    im = Image.open(dog)
    im = im.resize((256,256), Image.Resampling.LANCZOS)
    plt.imshow(im)
```













```
In [5]:
    plt.figure(figsize=(10,6))
    for i in range(len(dog_images)):
        bbox = get_bounding_boxes(annotations[i])
        dog = get_image(annotations[i])
        im = Image.open(dog)
        for j in range(len(bbox)):
            im2 = im.crop(bbox[j])
            im2 = im2.resize((128,128), Image.Resampling.LANCZOS)
            new_path = dog.replace('./image/','./Cropped/')
            new_path = new_path.replace('.jpg','-' + str(j) + '.jpg')
            im2=im2.convert('RGB')
            head, tail = os.path.split(new_path)
            Path(head).mkdir(parents=True, exist_ok=True)
            im2.save(new_path)
```

<Figure size 1000x600 with 0 Axes>

```
import random
import matplotlib.pyplot as plt
import numpy as np
from skimage import filters
from skimage.color import rgb2gray

breeds = glob.glob('./Cropped/*')

## Getting all images from each class

def allimages(breeds):
```

```
images = {}
            for folder in breeds:
                image_files = [f for f in os.listdir(folder)]
                images[folder] = image_files
            return images
        images_files_list = allimages(breeds)
        # """Calculate the angles between horizontal and vertical operators."""
        def pathtoimage(image_path):
            return rgb2gray(np.array(Image.open(image_path).convert('RGB')))
        def angle(dx, dy):
            return np.mod(np.arctan2(dy, dx), np.pi)
        def compute_edge_histogram(image):
            angle_sobel = angle(filters.sobel_h(image),filters.sobel_v(image))
            hist, hist_center = exposure.histogram(angle_sobel, nbins=36)
            return hist
        edge_hist_dict = {}
        for file_path_name,file_path_list in images_files_list.items():
            edge_histograms = []
            for file_path in file_path_list:
                edge_histogram = compute_edge_histogram(pathtoimage(file_path_name+ '\\'+fi
                edge_histograms.append(edge_histogram)
            edge_hist_dict[file_path_name] = edge_histograms
In [ ]:
In [7]: # Convert the images to edge histograms. (Assignment 1 - These will be the vector r
        # images). This will be your dataset for Part 3. (0.25 point)
        # images_files_list.keys,
```

```
# Assuming edge_hist_dict is already defined, containing histograms for each class
        # Initialize lists for training and test sets for each class
        X train dict = {}
        X_test_dict = {}
        y_train_dict = {}
        y_test_dict = {}
        # Split each class into training and test sets
        for class name, histograms in edge hist dict.items():
            # Convert histograms to a numpy array
            histograms_array = np.array(histograms)
            # Create class labels for the current class
            labels = np.array([class_name] * len(histograms_array))
            # Perform an 80/20 split for this class
            X_train_class, X_test_class, y_train_class, y_test_class = train_test_split(
                histograms_array, labels, test_size=0.2, random_state=42
            # Store in separate dictionaries
            X_train_dict[class_name] = X_train_class
            X_test_dict[class_name] = X_test_class
            y_train_dict[class_name] = y_train_class
            y_test_dict[class_name] = y_test_class
        # Print sizes of the datasets for each class
        # for class_name in edge_hist_dict.keys():
              print(f"Class: {class_name}")
              print("Training set size:", X_train_dict[class_name].shape)
              print("Test set size:", X_test_dict[class_name].shape)
        # Combine all training data and labels for standardization
        # Standardization
        scaler = StandardScaler()
        for X_train_name,X_train in X_train_dict.items():
            X_train_dict[X_train_name] = scaler.fit_transform(X_train)
            X_test_dict[X_train_name] = scaler.transform(X_test_dict[X_train_name])
In [8]: ## References:
        # https://scikit-learn.org/1.5/modules/neural_networks_supervised.html
        # https://scikit-learn.org/1.5/modules/generated/sklearn.ensemble.RandomForestClass
In [9]: ### Classifiers
```

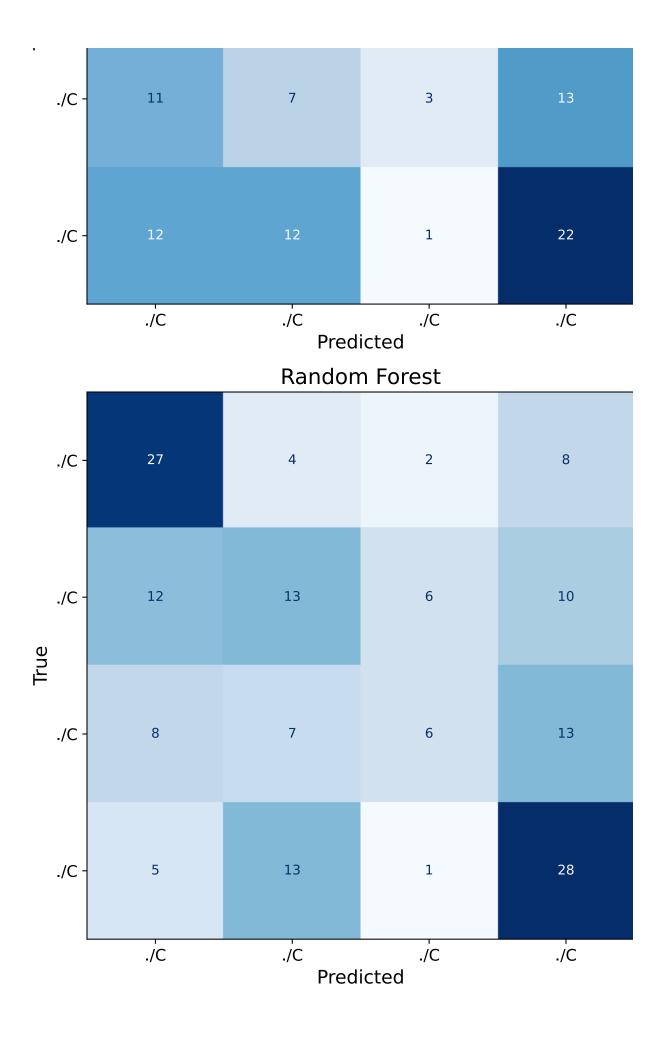
```
Classifiers = {
  'GausianDB':GaussianNB(),
  'Neuarl Network':MLPClassifier(hidden_layer_sizes=(10, 10, 10), random_state=42),
  'Random Forest':RandomForestClassifier(random_state=42)
}
```

6. (Performance Comparison) Perform stratified 5-fold cross-validation on the 4-class classification problem using the three classification methods (available on canvas) assigned to you. Plot the (3) confusion matrices for using three approaches (clearly label the classes) on the test set (See Figure 1). (If you use code from any website, please do proper referencing. You will get 0 point for this assignment without proper referencing) (3.75 points)

```
In [10]: # Assuming edge_hist_dict is already defined, containing histograms for each class
         # Combine all training data and labels for standardization
         X_test_combined = np.vstack(list(X_test_dict.values()))
         y_test_combined = np.hstack(list(y_test_dict.values()))
         X_train_combined = np.vstack(list(X_train_dict.values()))
         y_train_combined = np.hstack(list(y_train_dict.values()))
         # Perform stratified 5-fold cross-validation and collect predictions
         cv = StratifiedKFold(n_splits=5, shuffle=True, random_state=42)
         predictions = {}
         predictions = {}
         for name, clf in Classifiers.items():
             # Train the classifier on the training set
             clf.fit(X_train_combined, y_train_combined)
             # Predict on the test set
             preds = clf.predict(X_test_combined)
             predictions[name] = preds
         # Plot confusion matrices with increased size
         fig, axes = plt.subplots(3, 1, figsize=(18, 20)) # Increase figsize here
         for ax, (name, preds) in zip(axes, predictions.items()):
             cm = confusion_matrix(y_test_combined, preds, labels=list(edge_hist_dict.keys())
             display_labels = [label[:3] for label in edge_hist_dict.keys()] # Adjust this
             disp = ConfusionMatrixDisplay(confusion_matrix=cm, display_labels=display_label
             # Plot confusion matrix
             disp.plot(ax=ax, cmap=plt.cm.Blues, colorbar=False)
             # Set titles and other parameters
             ax.set_title(name, fontsize=16)
             ax.set_xlabel("Predicted", fontsize=14)
             ax.set_ylabel("True", fontsize=14)
             ax.tick_params(axis='both', which='major', labelsize=12)
```

plt.tight_layout()
plt.show()

GausianDB ./C -./C ./C ./C ./C ./C ./C /C Predicted Neuarl Network ./C -./C -True



```
In [ ]:
```

By visually comparing (e.g., looking at the color on the diagonal values, etc.) the three confusion matrices (on the test set), which do you think is the best method? Why? (0.50 point) by images Neural Network is best method

```
In [11]: from sklearn.model_selection import cross_val_score

mean_accuracies = {}

for name, clf in Classifiers.items():
    scores = cross_val_score(clf, X_train_combined, y_train_combined, cv=cv, scorin mean_accuracies[name] = scores.mean()

print("Mean Validation Accuracies:")
for name, mean_accuracy in mean_accuracies.items():
    print(f"{name}: {mean_accuracy:.4f}")
```

Mean Validation Accuracies:

GausianDB: 0.1502 Neuarl Network: 0.2602 Random Forest: 0.3451

Based on the mean validation accuracies (from the 5-fold cross-validation) for the three methods. Which is the best method?

Random Forest is best method

```
for name, clf in Classifiers.items():
    clf.fit(X_train_combined, y_train_combined) # Fit on all training data
    test_preds = clf.predict(X_test_combined) # Use scaled test data
    accuracy = np.mean(test_preds == y_test_combined) # Calculate accuracy
    test_accuracies[name] = accuracy

print("Test Set Accuracies:")
for name, accuracy in test_accuracies.items():
    print(f"{name}: {accuracy:.4f}")
```

Test Set Accuracies: GausianDB: 0.2883 Neuarl Network: 0.3374 Random Forest: 0.4540

Compute the accuracies for the three methods on the test set. Which is the best method? (0.25 point)

Neural Network is best method

```
In [13]: from sklearn.metrics import f1_score
f_measures = {}
```

```
for name, clf in Classifiers.items():
    clf.fit(X_train_combined, y_train_combined) # Fit on all training data
    test_preds = clf.predict(X_test_combined) # Use scaled test data
    f_measure = f1_score(y_test_combined, test_preds, average='weighted') # Calcul
    f_measures[name] = f_measure

print("F-measures on Test Set:")
for name, f_measure in f_measures.items():
    print(f"{name}: {f_measure:.4f}")
```

F-measures on Test Set: GausianDB: 0.1291 Neuarl Network: 0.3133 Random Forest: 0.4333

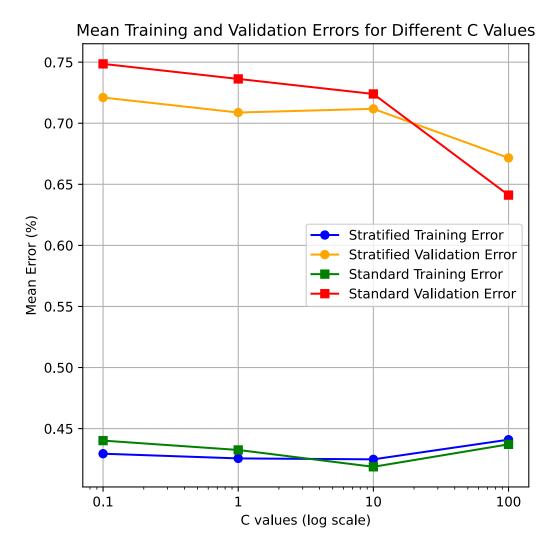
Compute the F-measure for the three methods on the test set. Which is the best method? (0.25 point) Neural Network is best method

7. (Model Selection) Use images from TWO classes. Perform a standard 5-fold cross-validation and a stratified 5-fold cross-validation on the training set (i.e., the standardized edge histogram dataset obtained from the training set) for Support Vector Classifiers using LinearSVC such that parameter C = 0.1, 1, 10, 100 and other parameters set as default. (2.5 points) • Plot a graph (x-axis: C; y-axis: mean validation/training error (%)) containing four error curves (2 validation error curves and 2 training error curves - label them clearly using a legend to define the curves). Which C has/have the lowest mean error for each curve? Comment about (1) the model complexity for SVM in relation to C, and (2) when/whether there is overfitting/underfitting. (1.5 points) • Use the C value with the lowest mean validation error for your SVM classifier from the stratified 5-fold cross-validation. What is the error for the test dataset (i.e., the standardized edge histogram dataset obtained from the test set)? (0.25 point)

```
In [14]: import numpy as np
         import matplotlib.pyplot as plt
         from sklearn.model_selection import StratifiedKFold, KFold
         from sklearn.svm import LinearSVC
         from sklearn.metrics import accuracy_score
         import warnings
         # Suppress warnings
         warnings.filterwarnings("ignore")
         # Assuming X_train_dict, y_train_dict, X_test_dict, and y_test_dict are already def
         # Replace 'Class_A' and 'Class_B' with your actual class labels
         names = list(X_train_dict.keys())
         selected_classes = [names[0], names[1]] # Selecting two classes
         # Filter the training and testing data for the selected classes
         X_train_filtered = np.vstack([X_train_dict[key] for key in X_train_dict if key in s
         y_train_filtered = np.hstack([y_train_dict[key] for key in y_train_dict if key in s
         X_test_filtered = np.vstack([X_test_dict[key] for key in X_test_dict if key in sele
         y_test_filtered = np.hstack([y_test_dict[key] for key in y_test_dict if key in sele
```

```
# Values of C to evaluate
C_{values} = [0.1, 1, 10, 100]
train_errors_stratified = []
val_errors_stratified = []
train_errors_standard = []
val_errors_standard = []
# Stratified 5-Fold Cross-Validation
stratified_cv = StratifiedKFold(n_splits=5, shuffle=True, random_state=42)
standard_cv = KFold(n_splits=5, shuffle=True, random_state=42)
for C in C values:
   svc = LinearSVC(C=C, max_iter=10000)
   # Stratified Cross-Validation for validation error
   stratified_train_errors = []
   stratified_val_errors = []
   for train_index, val_index in stratified_cv.split(X_train_filtered, y_train_fil
       X_train_split, X_val_split = X_train_filtered[train_index], X_train_filtere
       y_train_split, y_val_split = y_train_filtered[train_index], y_train_filtere
       # Fit the model
       svc.fit(X_train_split, y_train_split)
        # Calculate training error
       train_preds = svc.predict(X_train_split)
        stratified_train_errors.append(1 - accuracy_score(y_train_split, train_pred
       # Calculate validation error
       val_preds = svc.predict(X_val_split)
        stratified_val_errors.append(1 - accuracy_score(y_val_split, val_preds))
   # Store mean errors for stratified
   train_errors_stratified.append(np.mean(stratified_train_errors))
   val_errors_stratified.append(np.mean(stratified_val_errors))
   # Standard Cross-Validation for training and validation error
   standard_train_errors = []
   standard_val_errors = []
   for train_index, test_index in standard_cv.split(X_train_filtered):
       X_train_split, X_test_split = X_train_filtered[train_index], X_train_filter
       y_train_split, y_test_split = y_train_filtered[train_index], y_train_filter
       # Fit the model
       svc.fit(X_train_split, y_train_split)
       # Calculate training error
       train_preds = svc.predict(X_train_split)
        standard_train_errors.append(1 - accuracy_score(y_train_split, train_preds)
        # Calculate validation error
        val_preds = svc.predict(X_test_split)
        standard_val_errors.append(1 - accuracy_score(y_test_split, val_preds))
```

```
# Store mean errors for standard
   train_errors_standard.append(np.mean(standard_train_errors))
   val_errors_standard.append(np.mean(standard_val_errors))
# Plotting the errors
plt.figure(figsize=(6, 6))
# Stratified Training and Validation Errors
plt.plot(C_values, train_errors_stratified, label='Stratified Training Error', mark
plt.plot(C_values, val_errors_stratified, label='Stratified Validation Error', mark
# Standard Training and Validation Errors
plt.plot(C_values, train_errors_standard, label='Standard Training Error', marker='
plt.plot(C_values, val_errors_standard, label='Standard Validation Error', marker='
plt.xscale('log')
plt.xticks(C_values, C_values)
plt.xlabel('C values (log scale)')
plt.ylabel('Mean Error (%)')
plt.title('Mean Training and Validation Errors for Different C Values')
plt.legend()
plt.grid()
plt.show()
# Determine the C with the lowest validation error for both methods
```



As before, the model complexity can be understood through the relationship between the \mathcal{C} C parameter and the resulting errors. Lower values of \mathcal{C} C may lead to underfitting, while higher values may lead to overfitting.

The model is underfitting as the gap is training and validaition error is high both in standard and stratified

```
In [15]: best_C_stratified_index = np.argmin(val_errors_stratified)
best_C_stratified = C_values[best_C_stratified_index]

best_C_standard_index = np.argmin(val_errors_standard)
best_C_standard = C_values[best_C_standard_index]

print(f"Best C value based on stratified validation error: {best_C_stratified}")
print(f"Best C value based on standard validation error: {best_C_standard}")

# Now use the best C value from stratified to compute the error on the test dataset
final_model_stratified = LinearSVC(C=best_C_stratified, max_iter=10000)
final_model_stratified.fit(X_train_filtered, y_train_filtered)
test_preds_stratified = final_model_stratified.predict(X_test_filtered)
test_error_stratified = 1 - accuracy_score(y_test_filtered, test_preds_stratified)

print(f"Test_Error_for_best_C_value ({best_C_stratified}) from Stratified: {test_er
```

```
# Now use the best C value from standard to compute the error on the test dataset
final_model_standard = LinearSVC(C=best_C_standard, max_iter=10000)
final_model_standard.fit(X_train_filtered, y_train_filtered)
test_preds_standard = final_model_standard.predict(X_test_filtered)
test_error_standard = 1 - accuracy_score(y_test_filtered, test_preds_standard)

print(f"Test Error for best C value ({best_C_standard}) from Standard: {test_error_
Best C value based on stratified validation error: 100
Best C value based on standard validation error: 100
Test Error for best C value (100) from Stratified: 0.4878
Test Error for best C value (100) from Standard: 0.4268
```

In []:	
In []:	
In []:	