projeto 4

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Stochastic Processes - class 1/2024 - University of Brasília Computational work 4 - classification Gabriel Tambara Rabelo - 241106461

O presente projeto visa classificar de diferentes métodos para

h[0],h[1],h[2],h[3],h[4],h[5],h[6],h[7],class 58419,11150,9625,13219,13300,450,20,17,0 52196, 7232, 6733, 7406, 9320, 9624, 12177, 1512, 0

Para uma rodada dos testes do código, foram geradas as matrizes de confusão e suas análises indicam alguns pontos de interesse quanto ao método e o conjunto específico de dados utilizado. Esses resultados obtidos não necessariamente indicam que um algoritmo é melhor que o outro, contudo, para o presente conjunto de dados e amostras, podemos definir sua classificabilidade.

A seguir, têm-se os resultados de um dos testes da classificação, seguido pela sua análise.
CASE1
Bayes Confusion Matrix: $[[11\ 0\ 0\ 0\ 0]\ [\ 0\ 1\ 0\ 1\ 0]\ [\ 0\ 0\ 3\ 0\ 0]\ [\ 0\ 0\ 0\ 6\ 0]\ [\ 0\ 0\ 0\ 0\ 4]]$ Bayes Accuracy per Class: $[1.\ 0.5\ 1.\ 1.\ 1.\]$ Bayes Accuracy: 0.9615384615384616
QDA Confusion Matrix: [[11 0 0 0 0] [0 2 0 0 0] [0 0 3 0 0] [0 0 0 6 0] [0 0 0 0 4]] QDA Accuracy per Class: [1. 1. 1. 1.] QDA Accuracy: 1.0
LDA Confusion Matrix: [[11 0 0 0 0] [0 0 0 0 2] [0 0 3 0 0] [0 0 0 6 0] [0 0 2 0 2]] LDA Accuracy per Class: [1. 0. 1. 1. 0.5] LDA Accuracy: 0.8461538461538461 ———————————————————————————————————
Bayes Confusion Matrix: $[[11\ 0\ 0\ 0\ 0]\ [\ 0\ 1\ 0\ 1\ 0]\ [\ 0\ 0\ 3\ 0\ 0]\ [\ 0\ 1\ 0\ 4\ 1]\ [\ 0\ 0\ 0\ 0\ 4]]$ Bayes Accuracy per Class: $[1.\ 0.5\ 1.\ 0.666666667\ 1.\]$ Bayes Accuracy: 0.8846153846153846
QDA Confusion Matrix: [[11 0 0 0 0] [0 1 0 1 0] [0 0 3 0 0] [0 0 0 4 2] [0 0 0 0 4]] QDA Accuracy per Class: [1. 0.5 1. 0.666666667 1.] QDA Accuracy: 0.8846153846153846

LDA Confusion Matrix: [[11 0 0 0 0] [0 0 1 1 0] [0 0 3 0 0] [0 0 0 6 0] [0 0 1 2 1]] LDA Accuracy per Class: [1. 0. 1. 1. 0.25] LDA Accuracy: 0.8076923076923077

Para o caso 1:

O classificador Naive Bayes teve um bom desempenho para a maioria das classes, com uma precisão perfeita para 4 das 5 classes. A classe 2 teve um desempenho menor com precisão de 50%, indicando que há espaço para melhorias na diferenciação dessa classe. A precisão geral é alta, mostrando que o classificador funciona bem com esses dados, mas pode ter limitações em situações onde as classes não são bem separadas. Esse problema seria resolvido com um espaço amostral maior, o que poderia gerar melhores acurácias também para a classe problemática, porém, também poderia reduzir a acurácia da classificação das outras classes.

O QDA obteve uma precisão perfeita em todas as classes. Isso indica que o QDA foi capaz de capturar muito bem as características dos dados, provavelmente devido à sua capacidade de modelar distribuições não lineares. No entanto, precisão perfeita pode às vezes indicar overfitting, especialmente com um conjunto de dados de teste pequeno.

O LDA teve uma boa performance para a maioria das classes, mas falhou completamente para a classe 2, e teve uma precisão de 50% para a classe 4. A precisão geral é inferior ao Naive Bayes e QDA.

Para o caso 2:

O Naive Bayes apresentou uma redução na precisão geral e por classe comparado ao caso 1. A classe 2 ainda tem uma precisão de 50%, e a classe 4 reduziu para 66.67%.

O QDA também apresentou uma redução na precisão geral e por classe comparado ao caso 1. A precisão para a classe 4 diminuiu para 66.67%, e a classe 2 manteve a precisão de 50%...

O LDA teve uma redução significativa na precisão geral e para a classe 4 comparado ao Case 1. A precisão para a classe 2 é nula e a precisão para a classe 4 é de apenas 25%, indicando dificuldades significativas em separar essas classes com as características usadas. A precisão geral é a mais baixa entre os classificadores, indicando que o LDA é o menos eficaz com as características do caso 2.

A seguir, segue o código equivalente.

```
import os
import csv
import cv2 as cv
import numpy as np
import matplotlib.pyplot as plt
from collections import Counter
from sklearn.preprocessing import StandardScaler

img_size = 300
classes = ['Alzheimer', 'COVID', 'Brazilian_seeds', 'Brazilian_leaves',
'skin_cancer']
data_dir = './image_dataset/'
case1_csv_filename = 'case1_statistics.csv'
case2_csv_filename = 'case2_statistics.csv'
reg_param = 1e-5  # Adjusted regularization parameter
```

```
def load_images_from_folder(folder_path):
    image_files = [f for f in os.listdir(folder_path) if os.path.isfile(os.path.

→join(folder_path, f))]
    images = []
   for image_file in image_files:
        img = cv.imread(os.path.join(folder_path, image_file), cv.
 →IMREAD_GRAYSCALE)
       ratio = img_size / img.shape[1]
        img_resized = cv.resize(img, (img_size, int(img.shape[0] * ratio)), cv.
 →INTER_AREA)
       images.append(img_resized)
   return images
def makeHist(image, bits):
   hist, bins = np.histogram(image.flatten(), bins=np.arange(0, 257, 2 ** (8 -
 ⇔bits)))
   return hist, bins
def normalizeHist(hist):
   total_pixels = np.sum(hist)
   normalized_hist = hist / total_pixels
   return normalized_hist
def bin_centers(bin_borders):
   bin_center_list = (bin_borders[:-1] + bin_borders[1:]) / 2
   bin_center_list[-1] = 255
   return bin_center_list
def expectancy(hist, bin_centers):
   return np.sum(hist * bin_centers)
def median(hist, bin centers):
    cumulative_freq = 0
   for i, freq in enumerate(hist):
        cumulative_freq += freq
        if cumulative_freq >= 0.5:
            return bin_centers[i]
def mode(hist, bin_centers):
   return bin_centers[np.argmax(hist)]
def moment(hist, bin_centers, order, expectancy):
   return np.sum(((bin_centers - expectancy) ** order) * hist)
```

```
def entropy(hist):
   epsilon = 1e-27
   return -np.sum(hist * np.log2(hist + epsilon))
def calculate_statistics(image, bins):
   hist, bin_edges = makeHist(image, bins)
   normalized hist = normalizeHist(hist)
    centered_bins = bin_centers(bin_edges)
   var_exp = expectancy(normalized_hist, centered_bins)
   var median = median(normalized hist, centered bins)
   var_mode = mode(normalized_hist, centered_bins)
   var_variance = moment(normalized_hist, centered_bins, 2, var_exp)
   var_skewness = moment(normalized_hist, centered_bins, 3, var_exp)
   var_kurtosis = moment(normalized_hist, centered_bins, 4, var_exp)
   var_entropy = entropy(normalized_hist)
   return hist, var_exp, var_mode, var_median, var_variance, var_skewness, u
 →var_kurtosis, var_entropy
def save statistics to csv(images, class label, case1 csv filename,
 ⇔case2 csv filename):
   with open(case1_csv_filename, mode='a', newline='') as case1_file, __
 →open(case2_csv_filename, mode='a', newline='') as case2_file:
        case1 writer = csv.writer(case1 file)
       case2_writer = csv.writer(case2_file)
        for image in images:
            hist, var_exp, var_mode, var_median, var_variance, var_skewness,

¬var_kurtosis, var_entropy = calculate_statistics(image, bins=8)

            case_1_data = list(hist) + [class_label]
            case_2_data = [var_exp, var_mode, var_median, var_variance,__
 →var_skewness, var_kurtosis, var_entropy, class_label]
            case1_writer.writerow(case_1_data)
            case2_writer.writerow(case_2_data)
def load_data_from_csv(filename):
   data = []
   labels = []
   with open(filename, mode='r') as file:
       reader = csv.reader(file)
       next(reader)
       for row in reader:
            data.append([float(x) for x in row[:-1]])
            labels.append(int(row[-1]))
```

```
return np.array(data), np.array(labels)
def train_test_split(data, labels, test_size=0.1):
   np.random.seed(np.random.randint(0,99))
   indices = np.arange(len(data))
   np.random.shuffle(indices)
   split_idx = int(len(data) * (1 - test_size))
   train_indices = indices[:split_idx]
   test indices = indices[split idx:]
   return data[train_indices], data[test_indices], labels[train_indices],
 ⇔labels[test indices]
with open(case1_csv_filename, mode='w', newline='') as case1_file,__
 →open(case2_csv_filename, mode='w', newline='') as case2_file:
   case1 writer = csv.writer(case1 file)
   case2_writer = csv.writer(case2_file)
   case1_writer.writerow([f'h[{i}]' for i in range(8)] + ['class'])
   case2_writer.writerow(['expectancy', 'mode', 'median', 'variance',
 for class_id, class_name in enumerate(classes):
   folder path = os.path.join(data dir, class name)
   images = load_images_from_folder(folder_path)
   save_statistics_to_csv(images, class_id, case1_csv_filename,_
⇔case2_csv_filename)
data_case1, labels_case1 = load_data_from_csv(case1_csv_filename)
data_case2, labels_case2 = load_data_from_csv(case2_csv_filename)
X_train_case1, X_test_case1, y_train_case1, y_test_case1 =
→train_test_split(data_case1, labels_case1, test_size=0.1)
X train case2, X test case2, y train case2, y test case2 = 1
 class NaiveBayesClassifier:
   def fit(self, X, y):
       self.classes = np.unique(y)
       self.mean = \{\}
       self.var = \{\}
       self.priors = {}
       for cls in self.classes:
           X_c = X[y == cls]
           self.mean[cls] = np.mean(X c, axis=0)
           self.var[cls] = np.var(X_c, axis=0) + reg_param
```

```
self.priors[cls] = X_c.shape[0] / X.shape[0]
    def predict(self, X):
        predictions = [self._predict(x) for x in X]
        return np.array(predictions)
    def _predict(self, x):
        posteriors = []
        for cls in self.classes:
            prior = np.log(self.priors[cls])
            posterior = np.sum(np.log(self._pdf(cls, x)))
            posterior = prior + posterior
            posteriors.append(posterior)
        return self.classes[np.argmax(posteriors)]
    def _pdf(self, cls, x):
        mean = self.mean[cls]
        var = self.var[cls]
        numerator = np.exp(-(x - mean) ** 2 / (2 * var))
        denominator = np.sqrt(2 * np.pi * var)
        results = numerator / denominator
        for i in range(len(results)):
            if results[i] == 0:
                results[i] = reg_param
        return results
class QuadraticDiscriminantAnalysis:
    def fit(self, X, y):
        self.classes = np.unique(y)
        self.mean = \{\}
        self.covariance = {}
        self.priors = {}
        for cls in self.classes:
            X c = X[y == cls]
            self.mean[cls] = np.mean(X_c, axis=0)
            self.covariance[cls] = np.cov(X_c, rowvar=False) + np.eye(X.
 ⇒shape[1]) * reg_param
            self.priors[cls] = X_c.shape[0] / X.shape[0]
        #print("Means:", self.mean)
        #print("Covariances:", self.covariance)
        #print("Priors:", self.priors)
```

```
def predict(self, X):
       predictions = [self._predict(x) for x in X]
        return np.array(predictions)
   def _predict(self, x):
       discriminants = []
       for cls in self.classes:
            G = self._quadratic_discriminant(cls, x)
            discriminants.append(G)
       return self.classes[np.argmax(discriminants)]
   def _quadratic_discriminant(self, cls, x):
       mean = self.mean[cls]
       covariance = self.covariance[cls]
        inv_covariance = np.linalg.inv(covariance)
        det_cov = np.linalg.det(covariance)
        if det_cov == 0:
            det_cov = reg_param
       W k = -0.5 * inv covariance
        w_k = np.dot(inv_covariance, mean)
        w_k0 = -0.5 * np.dot(mean.T, np.dot(inv_covariance, mean)) - 0.5 * np.
 →log(det_cov) + np.log(self.priors[cls])
        discriminant = np.dot(x.T, np.dot(W_k, x)) + np.dot(w_k.T, x) + w_k0
        return discriminant
class LinearDiscriminantAnalysis:
   def fit(self, X, y):
       self.classes = np.unique(y)
       self.mean = {}
       self.covariance = {}
       self.priors = {}
       for cls in self.classes:
            X_c = X[y == cls]
            self.mean[cls] = np.mean(X_c, axis=0)
            self.covariance[cls] = np.cov(X_c, rowvar=False) + np.eye(X.
 ⇒shape[1]) * reg_param
            self.priors[cls] = X_c.shape[0] / X.shape[0]
        #print("Means:", self.mean)
```

```
#print("Covariances:", self.covariance)
        #print("Priors:", self.priors)
   def predict(self, X):
       predictions = [self._predict(x) for x in X]
       return np.array(predictions)
   def _predict(self, x):
       discriminants = []
       for cls in self.classes:
            G = self._linear_discriminant(cls, x)
            discriminants.append(G)
       return self.classes[np.argmax(discriminants)]
   def _linear_discriminant(self, cls, x):
       mean = self.mean[cls]
        covariance = self.covariance[cls]
        inv_covariance = np.linalg.inv(covariance)
        w_k = np.dot(inv_covariance, mean)
        w_k0 = -0.5 * np.dot(mean.T, np.dot(inv_covariance, mean)) + np.
 →log(self.priors[cls])
        discriminant = np.dot(w_k.T, x) + w_k0
        return discriminant
def confusion_matrix(y_true, y_pred, num_classes):
   cm = np.zeros((num_classes, num_classes), dtype=int)
   for i in range(len(y_true)):
       cm[y_true[i]][y_pred[i]] += 1
   return cm
def accuracy_per_class(cm):
   return np.diag(cm) / np.sum(cm, axis=1)
def overall_accuracy(cm):
   return np.sum(np.diag(cm)) / np.sum(cm)
data, labels = load_data_from_csv(case1_csv_filename)
X_train, X_test, y_train, y_test = train_test_split(data, labels)
# Standardizing the features
scaler = StandardScaler()
```

```
X_train = scaler.fit_transform(X_train)
X_test = scaler.transform(X_test)
nb = NaiveBayesClassifier()
nb.fit(X_train, y_train)
y_pred_nb = nb.predict(X_test)
cm_nb = confusion_matrix(y_test, y_pred_nb, len(classes))
acc_nb = accuracy_per_class(cm_nb)
overall_acc_nb = overall_accuracy(cm_nb)
qda = QuadraticDiscriminantAnalysis()
qda.fit(X_train, y_train)
y_pred_qda = qda.predict(X_test)
cm_qda = confusion_matrix(y_test, y_pred_qda, len(classes))
acc_qda = accuracy_per_class(cm_qda)
overall_acc_qda = overall_accuracy(cm_qda)
lda = LinearDiscriminantAnalysis()
lda.fit(X_train, y_train)
y_pred_lda = lda.predict(X_test)
cm_lda = confusion_matrix(y_test, y_pred_lda, len(classes))
acc_lda = accuracy_per_class(cm_lda)
overall_acc_lda = overall_accuracy(cm_lda)
print("\n----\n")
print("Bayes Confusion Matrix:\n", cm_nb)
print("Bayes Accuracy per Class:\n", acc_nb)
print("Bayes Accuracy:", overall_acc_nb)
print("\n----\n")
print("QDA Confusion Matrix:\n", cm_qda)
print("QDA Accuracy per Class:\n", acc_qda)
print("QDA Accuracy:", overall_acc_qda)
print("\n----\n")
print("LDA Confusion Matrix:\n", cm lda)
print("LDA Accuracy per Class:\n", acc_lda)
print("LDA Accuracy:", overall_acc_lda)
data, labels = load_data_from_csv(case2_csv_filename)
X_train, X_test, y_train, y_test = train_test_split(data, labels)
# Standardizing the features
```

```
scaler = StandardScaler()
X train = scaler.fit_transform(X_train)
X_test = scaler.transform(X_test)
nb = NaiveBayesClassifier()
nb.fit(X_train, y_train)
y_pred_nb = nb.predict(X_test)
cm_nb = confusion_matrix(y_test, y_pred_nb, len(classes))
acc_nb = accuracy_per_class(cm_nb)
overall_acc_nb = overall_accuracy(cm_nb)
qda = QuadraticDiscriminantAnalysis()
qda.fit(X_train, y_train)
y_pred_qda = qda.predict(X_test)
cm_qda = confusion_matrix(y_test, y_pred_qda, len(classes))
acc_qda = accuracy_per_class(cm_qda)
overall_acc_qda = overall_accuracy(cm_qda)
lda = LinearDiscriminantAnalysis()
lda.fit(X_train, y_train)
y_pred_lda = lda.predict(X_test)
cm_lda = confusion_matrix(y_test, y_pred_lda, len(classes))
acc_lda = accuracy_per_class(cm_lda)
overall_acc_lda = overall_accuracy(cm_lda)
print("\n----\n")
print("Bayes Confusion Matrix:\n", cm_nb)
print("Bayes Accuracy per Class:\n", acc_nb)
print("Bayes Accuracy:", overall_acc_nb)
print("\n----\n")
print("QDA Confusion Matrix:\n", cm_qda)
print("QDA Accuracy per Class:\n", acc_qda)
print("QDA Accuracy:", overall_acc_qda)
print("\n----\n")
print("LDA Confusion Matrix:\n", cm_lda)
print("LDA Accuracy per Class:\n", acc lda)
print("LDA Accuracy:", overall_acc_lda)
```