## Aprendizagem 2023 Homework II – Group 28

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Part I: Pen and Paper

Consider the following dataset  $(y_3 - y_5)$  are all categorical variables and the domain of  $y_2$  is [0, 1]:

D	$y_1$	$y_2$	$y_3$	$y_4$	$y_5$	$y_6$
<b>X</b> 1	0.24	0.36	1	1	0	Α
<b>X</b> 2	0.16	0.48	1	0	1	Α
<b>X</b> 3	0.32	0.72	0	1	2	Α
<b>X</b> 4	0.54	0.11	0	0	1	В
<b>X</b> 5	0.66	0.39	0	0	0	В
<b>X</b> 6	0.76	0.28	1	0	2	В
<b>X</b> 7	0.41	0.53	0	1	1	В
<b>X</b> 8	0.38	0.52	0	1	0	Α
<b>X</b> 9	0.42	0.59	0	1	1	В

1. Consider  $x_1$  -  $x_7$  to be training observations,  $x_8$  -  $x_9$  to be testing observations,  $y_1$  -  $y_5$  to be input variables and  $y_6$  to be the target variable.

Hint: you can use scipy.stats.multivariate\_normal for multivariate distribution calculus

(a) Learn a Bayesian classifier assuming: i)  $\{y_1, y_2\}$ ,  $\{y_3, y_4\}$  and  $\{y_5\}$  sets of independent variables (e.g.,  $y_1 \perp y_3$  yet  $y_1 \perp y_2$ ), and ii)  $y_1 \times y_2 \in \mathbb{R}^2$  is normally distributed. Show all parameters (distributions and priors for subsequent testing).

Gonçalo

(b) Under a MAP assumption, classify each testing observation showing all your calculus.

Gonçalo

(c) Consider that the default decision threshold of  $\theta = 0.5$  can be adjusted according to

$$f(\mathbf{x}|\theta) = \begin{cases} A, & P(A|\mathbf{x}) > \theta \\ B, & \text{otherwise} \end{cases}$$

Under a maximum likelihood assumption, what thresholds optimize testing accuracy?

Raquel

- 2. Let  $y_1$  be the target numeric variable,  $y_2$   $y_6$  be the input variables where  $y_2$  is binarized under an equal-width (equal-range) discretization. For the evaluation of regressors, consider a 3-fold cross-validation over the full dataset  $(x_1 x_9)$  without shuffling the observations.
  - (a) Identify the observations and features per data fold after the binarization procedure.

To do the **binarization procedure** with an **equal-width discretization**, we need to find the range of values for  $y_2$  and then divide it into two intervals. Bellow are the calculations of the intervals:

 $y_2$  ranges from 0.11 to 0.72

Therefore, the width shall be:

$$width = \frac{0.72 - 0.11}{2} = 0.305$$

**Now,** we can finally separate  $y_2$  into two intervals:

$$interval_1 = [0.11, 0.11 + 0.305] = [0.11, 0.415]$$

$$interval_2 = [0.72 - 0.305, 0.72] = [0.415, 0.72]$$

**Here is the binazation of**  $y_2$  based on those intervals:

D	У1	У2	У3	У4	<i>y</i> 5	У6
$x_1$	0.24	0	1	1	0	A
$x_2$	0.16	1	1	0	1	A
<i>x</i> <sub>3</sub>	0.32	1	0	1	2	A
$x_4$	0.54	0	0	0	1	В
<i>x</i> <sub>5</sub>	0.66	0	0	0	0	В
$x_6$	0.76	0	1	0	2	В
<i>x</i> <sub>7</sub>	0.41	1	0	1	1	В
$x_8$	0.38	1	0	1	0	A
<i>x</i> <sub>9</sub>	0.42	1	0	1	1	В

The next step is identifying our folds, which will be:

Fold 
$$1 = x_1 x_2 x_3$$
  
Fold  $2 = x_4 x_5 x_6$   
Fold  $3 = x_7 x_8 x_9$ 

So our datasets will be:

Fold 1								
D	y <sub>1</sub>	<i>y</i> <sub>2</sub>	<b>у</b> 3	У4	<i>y</i> <sub>5</sub>	У6		
$x_1$	0.24	0	1	1	0	A		
$x_2$	0.16	1	1	0	1	A		
$x_3$	0.24 0.16 0.32	1	0	1	2	A		

$ \begin{array}{c ccccccccccccccccccccccccccccccccccc$								
D	У1	У2	У3	У4	<i>y</i> <sub>5</sub>	У6		
<i>x</i> <sub>4</sub>	0.54	0	0	0	1	В		
<i>x</i> <sub>5</sub>	0.66	0	0	0	0	В		
$x_6$	0.54 0.66 0.76	0	1	0	2	В		

Fold 3								
D	У1	<i>y</i> <sub>2</sub>	У3	У4	<i>y</i> 5	У6		
<i>x</i> <sub>7</sub>	0.41	1	0	1	1	В		
$x_8$	0.41 0.38 0.42	1	0	1	0	A		
<i>x</i> <sub>9</sub>	0.42	1	0	1	1	В		

(b) Consider a distance-weighted kNN with k = 3, Hamming distance (d), and 1/d weighting. Compute the MAE of this kNN regressor for the  $1^{st}$  iteration of the cross-validation (i.e. train observations have the lower indices).

Raquel

## Part II: Programming and critical analysis

Considering the column\_diagnosis.arff dataset available at the course webpage's homework tab. Using sklearn, apply a 10-fold stratified cross-validation with shuffling (random\_state=0) for the assessment of predictive models along this section.

- 1. Compare the performance of kNN with k = 5 and Naïve Bayes with Gaussian assumption (consider all remaining parameters for each classifier as sklearn's default):
  - (a) Plot two boxplots with the fold accuracies for each classifier.

```
import matplotlib.pyplot as plt, pandas as pd
2 from sklearn.model_selection import StratifiedKFold, cross_val_score
3 from sklearn.neighbors import KNeighborsClassifier
4 from sklearn.naive_bayes import GaussianNB
5 from scipy.io.arff import loadarff
_{7} # Read the ARFF file and prepare data
8 data = loadarff("./data/column_diagnosis.arff")
9 df = pd.DataFrame(data[0])
10 df["class"] = df["class"].str.decode("utf-8")
11 X, y = df.drop("class", axis=1), df["class"]
13 # Define cross-validation strategy
14 folds = StratifiedKFold(n_splits=10, shuffle=True, random_state=0)
16 # Initialize classifiers
17 knn_predictor = KNeighborsClassifier(n_neighbors=5)
18 nb_predictor = GaussianNB()
20 # Evaluate classifiers
21 knn_accs = cross_val_score(knn_predictor, X, y, cv=folds, scoring="accuracy")
22 nb_accs = cross_val_score(nb_predictor, X, y, cv=folds, scoring="accuracy")
24 # Plot boxplots
25 plt.figure(figsize=(7, 5))
26 b_plot = plt.boxplot(
      [knn_accs, nb_accs], patch_artist=True, labels=["kNN", "Naive Bayes"]
28
30 colors = ["#f8766d", "#00bfc4"]
for patch, color in zip(b_plot["boxes"], colors):
      patch.set_facecolor(color)
33 for median in b_plot["medians"]:
      median.set_color("black")
36 plt.ylabel("Accuracy")
37 plt.grid(axis="y")
38 plt.show()
```

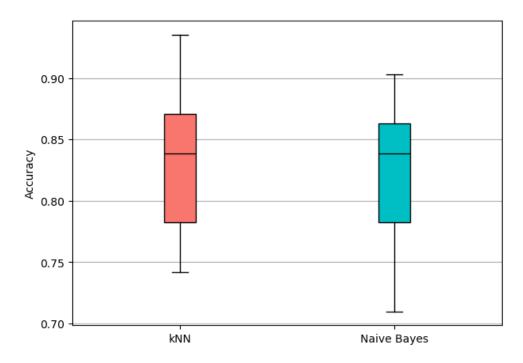


Figure 1: Boxplots with the fold accuracies of kNN (k = 5) and Naïve Bayes

(b) Using scipy, test the hypothesis "kNN is statistically superior to Naïve Bayes regarding accuracy", asserting whether is true.

We'll consider the null hypothesis and alternate hypothesis below and perform a single-tailed test using the accuracies obtained in the previous answer,

```
H_0: accuracy<sub>kNN</sub> = accuracy<sub>Naïve Bayes</sub>

H_1: accuracy<sub>kNN</sub> > accuracy<sub>Naïve Bayes</sub>
```

```
from scipy.stats import ttest_rel

# Is knn better than naive bayes?
res = ttest_rel(knn_accs, nb_accs, alternative="greater")
print("Is knn > naive bayes? pval =", res.pvalue)
```

Using scipy we get a p-value of, approximately, 0.190428 = 19.0428 %.

This means we cannot reject the hypothesis  $H_0$  at common significance levels (1%, 5% and 10%).

**Therefore,** we cannot assert that kNN is statistically superior to Naïve Bayes. We also cannot state that the hypothesis on the statement is outright false without checking other statistical tests.

2. Consider two kNN predictors with k=1 and k=5 (uniform weights, Euclidean distance, all remaining parameters as default). Plot the differences between the two cumulative confusion matrices of the predictors. Comment.

```
import numpy as np, matplotlib.pyplot as plt, pandas as pd, seaborn as sns
from sklearn.model_selection import StratifiedKFold
from sklearn.neighbors import KNeighborsClassifier
from sklearn.metrics import confusion_matrix
from scipy.io.arff import loadarff
```

```
_{7} # Read the ARFF file and prepare data
8 data = loadarff("./data/column_diagnosis.arff")
9 df = pd.DataFrame(data[0])
10 df["class"] = df["class"].str.decode("utf-8")
II X, y = df.drop("class", axis=1), df["class"]
13 # Initialize StratifiedKFold with 10 folds and shuffling
14 folds = StratifiedKFold(n_splits=10, shuffle=True, random_state=0)
16 # Create kNN classifiers with k=1 and k=5
17 knn_1 = KNeighborsClassifier(n_neighbors=1)
18 knn_5 = KNeighborsClassifier(n_neighbors=5)
20 labels = ["Hernia", "Normal", "Spondylolisthesis"]
21 \text{ cm}_1, \text{ cm}_5 = \text{np.zeros}((3, 3)), \text{np.zeros}((3, 3))
22 for train_k, test_k in folds.split(X, y):
      X_train, X_test = X.iloc[train_k], X.iloc[test_k]
      y_train, y_test = y.iloc[train_k], y.iloc[test_k]
24
25
      # Fit kNN classifiers and assess
      knn_1.fit(X_train, y_train)
27
      knn_5.fit(X_train, y_train)
      knn_1_pred, knn_5_pred = knn_1.predict(X_test), knn_5.predict(X_test)
      cm_1 += np.array(confusion_matrix(y_test, knn_1_pred, labels=labels))
      cm_5 += np.array(confusion_matrix(y_test, knn_5_pred, labels=labels))
33 # Calculate cumulative confusion matrices
34 \text{ cm\_diff} = \text{cm\_1} - \text{cm\_5}
35 cm_diff_df = pd.DataFrame(cm_diff, index=labels, columns=labels)
37 # Plot the differences
38 plt.figure(figsize=(9, 7))
39 sns.heatmap(
      cm_diff_df, cmap="Purples", annot=True, annot_kws={"fontsize": 14}, fmt="g"
41 )
42 plt.xlabel("Predicted")
43 plt.ylabel("Real")
44 plt.show()
```

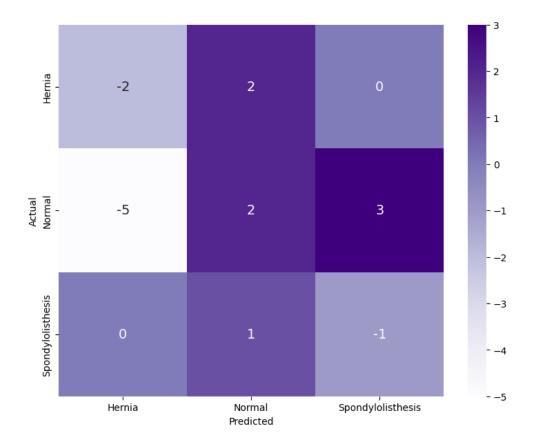


Figure 2: Confusion Matrix Differences Between k=1 and k=5 k-Nearest Neighbors (kNN) Classifiers

Blah

## 3. Considering the unique properties of column\_diagnosis, identify three possible difficulties of Naïve Bayes when learning from the given dataset.

Here are three possible difficulties of Naïve Bayes when learning from the given dataset, in no particular order:

- Variable dependencies (inadequacy of independence assumption).
- Variables not normally distributed (inadequacy of Gaussian assumption). Probability estimates from a limited number of observations (e.g., inadequate estimates, null probabilities).
- Imbalanced class creating biases in MAP estimates via priors.