Worksheet 14

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Topics

- Naive Bayes
- Model Evaluation

Naive Bayes

Attribute A	Attribute B	Attribute C	Class	
Yes	Single	High	No	
No	Married	Mid	No	
No	Single	Low	No	
Yes	Married	High	No	
No	Divorced	Mid	Yes	
No	Married	Low	No	
Yes	Divorced	High	No	
No	Single	Mid	Yes	
No	Married	Low	No	
No	Single	Mid	Yes	

- a) Compute the following probabilities:
 - P(Attribute A = Yes | Class = No)
 - P(Attribute B = Divorced | Class = Yes)
 - P(Attribute C = High | Class = No)
 - P(Attribute C = Mid | Class = Yes)

P(Attribute A = Yes | Class = No) = 3/7

P(Attribute B = Divorced | Class = Yes) = 1/3

P(Attribute C = High | Class = No) = 3/7

P(Attribute C = Mid | Class = Yes) = 3/3 = 1

- b) Classify the following unseen records:
 - (Yes, Married, Mid)
 - (No, Divorced, High)
 - (No, Single, High)
 - (No, Divorced, Low)

```
Under Naive Bayes' assumption of independence,
P(Yes, Married, Mid | Class = Yes)
= P(Yes | Class = Yes) * P(Married | Class = Yes) * P(Mid | Class = Yes)
= 0
P(Yes, Married, Mid | Class = No)
= P(Yes | Class = No) * P(Married | Class = No) * P(Mid | Class = No)
= 3/7 * 4/7 * 1/7
= 0.034985
  1. (Yes, Married, Mid) belongs to class No because 0.034985 > 0 so we will predict class No
P(No, Divorced, High | Class = Yes)
= P(No | Class = Yes) * P(Divorced | Class = Yes) * P(High | Class = Yes)
= 1 * 1/3 * 0
= 0
P(No, Divorced, High | Class = No)
= P(No | Class = No) * P(Divorced | Class = No) * P(High | Class = No)
= 4/7 * 1/7 * 3/7
= 0.034985
 2. (No, Divorced, High) belongs to class No because 0.034985 > 0 so we will predict class No
P(No, Single, High | Class = Yes)
= P(No | Class = Yes) * P(Single | Class = Yes) * P(High | Class = Yes)
= 1 * 2/3 * 0
= 0
P(No, Single, High | Class = No)
= P(No | Class = No) * P(Single | Class = No) * P(High | Class = No)
= 4/7 * 2/7 * 3/7
= 0.069971
 3. (No, Single, High) belongs to class No because 0.069971 > 0 so we will predict class No
P(No, Divorced, Low | Class = Yes)
= P(No | Class = Yes) * P(Divorced | Class = Yes) * P(Low | Class = Yes)
```

```
= 1 * 1/3 * 0

= 0

P(No, Divorced, Low | Class = No)

= P(No | Class = No) * P(Divorced | Class = No) * P(Low | Class = No)

= 4/7 * 1/7 * 3/7

= 0.034985
```

4. (No, Divorced, Low) belongs to class No because 0.034985 > 0 so we will predict class No

Model Evaluation

a) Write a function to generate the confusion matrix for a list of actual classes and a list of predicted classes

```
In [ ]: actual_class = ["Yes", "No", "No", "Yes", "No", "Yes", "No", "No", "No"]
        predicted_class = ["Yes", "No", "Yes", "No", "No", "No", "Yes", "Yes", "Yes", "No"]
        def confusion_matrix(actual, predicted):
            trueP = 0
            trueN = 0
            falseP = 0
            falseN = 0
            for x in range(len(actual)):
              if actual[x] == "Yes" and actual[x] == predicted[x]:
                trueP += 1
              elif actual[x] == "No" and actual[x] == predicted[x]:
                trueN += 1
              elif actual[x] == "No" and actual[x] != predicted[x]:
                falseP += 1
              else:
                falseN += 1
            matrix = [[trueP, falseN],[falseP, trueN]]
            return matrix
        print(confusion_matrix(actual_class, predicted_class))
```

[[2, 1], [3, 4]]

b) Assume you have the following Cost Matrix:

	predicted = Y	predicted = N
actual = Y	-1	5
actual = N	10	0

What is the cost of the above classification?

The cost is -1(-1) + 5(100) + 10(1) + 0 = 511.

c) Write a function that takes in the actual values, the predictions, and a cost matrix and outputs a cost. Test it on the above example.

```
In [ ]: def outputCost(actual, predicted, costMatrix):
    cost = 0
    cost += costMatrix[0][0] * -1
    cost += costMatrix[0][1] * 100
    cost += costMatrix[1][0] * 1
    return cost

print(outputCost(actual_class, predicted_class, confusion_matrix(actual_class, predicted_class))
```

d) Implement functions for the following:

- accuracy
- precision
- recall
- f-measure

and apply them to the above example.

```
In [ ]: | def accuracy(actual, predicted):
          total = len(actual)
          correct = 0
          for x in range(len(actual)):
            if actual[x] == predicted[x]:
              correct += 1
          return correct/total
        def precision(actual, predicted):
            trueP = 0
            trueN = 0
            falseP = 0
            falseN = 0
            for x in range(len(actual)):
              if actual[x] == "Yes" and actual[x] == predicted[x]:
                trueP += 1
              elif actual[x] == "No" and actual[x] == predicted[x]:
                trueN += 1
              elif actual[x] == "No" and actual[x] != predicted[x]:
                falseN += 1
              else:
                falseP += 1
            return trueP / (trueP + falseP)
        def recall(actual, predicted):
            trueP = 0
            trueN = 0
            falseP = 0
            falseN = 0
            for x in range(len(actual)):
              if actual[x] == "Yes" and actual[x] == predicted[x]:
                trueP += 1
              elif actual[x] == "No" and actual[x] == predicted[x]:
              elif actual[x] == "No" and actual[x] != predicted[x]:
                falseN += 1
              else:
                falseP += 1
            return trueP / (trueP + falseN)
```

```
def fMeasure(actual, predicted):
  rec = recall(actual, predicted)
  pre = precision(actual, predicted)
  return 2 * rec * pre / (rec + pre)
print(accuracy(actual_class, predicted_class))
print(precision(actual_class, predicted_class))
print(recall(actual_class, predicted_class))
print(fMeasure(actual_class, predicted_class))
```

0.6

0.4

0.5

Challenge (Midterm prep part 2)

In this exercise you will update your submission to the titanic competition.

- a) First let's add new numerical features / columns to the datasets that might be related to the survival of individuals.
 - has cabin should have a value of 0 if the cabin feature is nan and 1 otherwise
 - family members should have the total number of family members (by combining SibSp and Parch)
 - title_type: from the title extracted from the name, we will categorize it into 2 types: common for titles that many passengers have, rare for titles that few passengers have. Map common to 1 and rare to 0. Describe what threshold you used to define common and rare titles and how you found it.
 - fare_type: using Kmeans clustering on the fare column, find an appropriate number of clusters / groups of similar fares. Using the clusters you created, fare_price should be an ordinal variable that represents the expensiveness of the fare. For example if you split fare into 3 clusters (0 - 15, 15 - 40, and 40+) then the fare_price value should be 0 for fare values 0 - 15, 1 for 15 -40, and 2 for 40+.
 - Create an addition two numerical features of your invention that you think could be relevant to the survival of individuals.

Note: The features must be numerical because the sklearn DecisionTreeClassifier can only take on numerical features.

```
In [ ]:
        import pandas as pd
        import numpy as np
        import csv
        pdTitanic = pd.read csv("Titanic.csv")
        pdTitanicTwo = pd.read csv("Titanic2.csv")
        df = pd.concat([pdTitanic, pdTitanicTwo], ignore_index=True)
        df.head(5)
```

Out[]:	Passenger	ld	Survived	Pclass	Name	Sex	Age	SibSp	Parch	Ticket	Fare	Cabin
	0	1	0.0	3	Braund, Mr. Owen Harris	male	22.0	1	0	A/5 21171	7.2500	NaN
	1	2	1.0	1	Cumings, Mrs. John Bradley (Florence Briggs Th	female	38.0	1	0	PC 17599	71.2833	C85
	2	3	1.0	3	Heikkinen, Miss. Laina	female	26.0	0	0	STON/O2. 3101282	7.9250	NaN
	3	4	1.0	1	Futrelle, Mrs. Jacques Heath (Lily May Peel)	female	35.0	1	0	113803	53.1000	C123
	4	5	0.0	3	Allen, Mr. William Henry	male	35.0	0	0	373450	8.0500	NaN
In []:	df.isnull().	SUI	m()									
Out[]:												

In []: df.info()

```
Data columns (total 12 columns):
       #
           Column
                        Non-Null Count Dtype
           PassengerId 1309 non-null
       0
                                       int64
       1
           Survived 891 non-null
                                       float64
                        1309 non-null int64
           Pclass
       2
       3
           Name
                       1309 non-null object
                       1309 non-null object
           Sex
       4
       5
                       1046 non-null float64
           Age
                       1309 non-null int64
       6
           SibSp
       7
           Parch
                       1309 non-null int64
                       1309 non-null object
       8
           Ticket
       9
           Fare
                        1308 non-null
                                       float64
       10 Cabin
                        295 non-null
                                        object
       11 Embarked
                       1307 non-null
                                        object
      dtypes: float64(3), int64(4), object(5)
      memory usage: 122.8+ KB
In [ ]: from sklearn.cluster import KMeans
        from sklearn.preprocessing import StandardScaler
        from collections import Counter
        df['has_cabin'] = df['Cabin'].apply(lambda x: 0 if pd.isna(x) else 1)
        df['family members'] = df['SibSp'] + df['Parch']
        listName = df['Name'].tolist()
        listTitle = []
        for x in listName:
          listTitle.append(x.split()[1])
        df['title'] = listTitle
        title_counts = Counter(listTitle)
        threshold = len(listTitle) / 100
        results = {}
        for item, item count in title counts.items():
            results[item] = 1 if item_count > threshold else 0
        df['title_type'] = df['title'].map(results)
        df.drop(['title'], axis=1, inplace=True)
        df = df.dropna(subset=['Fare'])
In [ ]: fare_scaled = StandardScaler().fit_transform(df[['Fare']])
        kmeans = KMeans(n clusters=3, random state=42).fit(fare scaled)
        df['fare_cluster'] = kmeans.labels_
        ordered_clusters = df.groupby('fare_cluster')['Fare'].mean().sort_values().index
        cluster_mapping = {old: new for new, old in enumerate(ordered_clusters)}
        df['fare_price'] = df['fare_cluster'].map(cluster_mapping)
        df.drop(['fare_cluster'], axis=1, inplace=True)
      /usr/local/lib/python3.10/dist-packages/sklearn/cluster/_kmeans.py:870: FutureWarning: Th
      e default value of `n_init` will change from 10 to 'auto' in 1.4. Set the value of `n_ini
      t` explicitly to suppress the warning
        warnings.warn(
In []: df.head(5)
```

<class 'pandas.core.frame.DataFrame'>
RangeIndex: 1309 entries, 0 to 1308

	PassengerId	Survived	Pclass	Name	Sex	Age	SibSp	Parch	Ticket	Fare	Cabin
0	1	0.0	3	Braund, Mr. Owen Harris	male	22.0	1	0	A/5 21171	7.2500	NaN
1	2	1.0	1	Cumings, Mrs. John Bradley (Florence Briggs Th	female	38.0	1	0	PC 17599	71.2833	C85
2	3	1.0	3	Heikkinen, Miss. Laina	female	26.0	0	0	STON/O2. 3101282	7.9250	NaN
\$ 3	4	1.0	1	Futrelle, Mrs. Jacques Heath (Lily May Peel)	female	35.0	1	0	113803	53.1000	C123
4	5	0.0	3	Allen, Mr. William Henry	male	35.0	0	0	373450	8.0500	NaN

Out[]:

b) Using a method covered in class, tune the parameters of a decision tree model on the titanic dataset (containing all numerical features including the ones you added above). Evaluate this model locally and report it's performance.

Note: make sure you are not tuning your parameters on the same dataset you are using to evaluate the model. Also explain how you know you are not overfitting to the training set.

```
In []: from sklearn.model_selection import train_test_split, GridSearchCV
    from sklearn.tree import DecisionTreeClassifier
    from sklearn.metrics import accuracy_score, classification_report
    import pandas as pd

X = df.drop('Survived', axis=1)
y = df['Survived']

X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42
dt = DecisionTreeClassifier()

grid_search = GridSearchCV(estimator=dt, param_grid=param_grid, cv=5, scoring='accuracy'
grid_search.fit(X_train, y_train)

print("Best parameters:", grid_search.best_params_)
print("Best score:", grid_search.best_score_)

predictions = grid_search.predict(X_test)
print("Accuracy on test set:", accuracy_score(y_test, predictions))
print("Classification Report:\n", classification_report(y_test, predictions))
```

c) Try reducing the dimension of the dataset and create a Naive Bayes model. Evaluate this model.

```
In [ ]: from sklearn.decomposition import PCA
        from sklearn.model selection import train test split
        from sklearn.preprocessing import StandardScaler
        from sklearn.naive_bayes import GaussianNB
        from sklearn.metrics import accuracy score
        import pandas as pd
        X = df.drop('Survived', axis=1)
        y = df['Survived']
        scaler = StandardScaler()
        X scaled = scaler.fit transform(X)
        pca = PCA(n_components=0.95)
        X_pca = pca.fit_transform(X_scaled)
        X_train, X_test, y_train, y_test = train_test_split(X_pca, y, test_size=0.3, random_stat
        model = GaussianNB()
        model.fit(X_train, y_train)
        y_pred = model.predict(X_test)
        accuracy = accuracy_score(y_test, y_pred)
        print(f"Accuracy: {accuracy:.4f}")
```

d) Create an ensemble classifier using a combination of KNN, Decision Trees, and Naive Bayes models. Evaluate this classifier.

```
In [ ]: from sklearn.neighbors import KNeighborsClassifier
        from sklearn.metrics import accuracy_score
        X = df[['Age', 'Sex']]
        y = df['Survived']
        X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=1)
        train_accuracy_lst = []
        test_accuracy_lst = []
        for K in range(1, 51):
          knn = KNeighborsClassifier(n neighbors=K)
          knn.fit(X_train, y_train)
          y_train_pred = knn.predict(X_train)
          y_test_pred = knn.predict(X_test)
          train_accuracy = accuracy_score(y_train, y_train_pred)
          test_accuracy = accuracy_score(y_test, y_test_pred)
          train_accuracy_lst.append(train_accuracy)
          test_accuracy_lst.append(test_accuracy)
        plt.figure(figsize=(12, 8))
        plt.plot(range(1, 51), train_accuracy_lst, label='Training Accuracy')
        plt.plot(range(1, 51), test_accuracy_lst, label='Testing Accuracy')
        plt.xlabel('K')
        plt.ylabel('Accuracy')
        plt.title('Training and Testing Accuracy for Different K Values')
        plt.legend()
        plt.xticks(range(1, 51))
        plt.show()
        knn = KNeighborsClassifier(n_neighbors=7)
        knn.fit(X_train, y_train)
        y_train_pred = knn.predict(X_train)
```

```
y_test_pred = knn.predict(X_test)
train_accuracy = accuracy_score(y_train, y_train_pred)
test_accuracy = accuracy_score(y_test, y_test_pred)
print(train_accuracy, test_accuracy)
```

e) Update your kaggle submission using the best model you created (best model means the one that performed the best on your local evaluation)

https://www.kaggle.com/competitions/titanic/submissions

Some useful code for the midterm

```
In [ ]: import seaborn as sns
        from sklearn.svm import SVC
        import matplotlib.pyplot as plt
        from sklearn.decomposition import PCA
        from sklearn.pipeline import make pipeline
        from sklearn.metrics import confusion_matrix, accuracy_score
        from sklearn.datasets import fetch lfw people
        from sklearn.ensemble import BaggingClassifier
        from sklearn.model_selection import GridSearchCV, train_test_split
        sns.set()
        # Get face data
        faces = fetch_lfw_people(min_faces_per_person=60)
        # plot face data
        fig, ax = plt.subplots(3, 5)
        for i, axi in enumerate(ax.flat):
            axi.imshow(faces.images[i], cmap='bone')
            axi.set(xticks=[], yticks=[],
                    xlabel=faces.target_names[faces.target[i]])
        plt.show()
        # split train test set
        Xtrain, Xtest, ytrain, ytest = train_test_split(faces.data, faces.target, random_state=4
        pca = PCA(n_components=150, whiten=True)
        svc = SVC(kernel='rbf', class_weight='balanced')
        svcpca = make pipeline(pca, svc)
        # Tune model to find best values of C and gamma using cross validation
        param_grid = \{ 'svc_C' : [1, 5, 10, 50], \}
                      'svc__gamma': [0.0001, 0.0005, 0.001, 0.005]}
        kfold = 10
        grid = GridSearchCV(svcpca, param_grid, cv=kfold)
        grid.fit(Xtrain, ytrain)
        print(grid.best_params_)
        # use the best params explicitly here
        pca = PCA(n_components=150, whiten=True)
        svc = SVC(kernel='rbf', class_weight='balanced', C=10, gamma=0.005)
        svcpca = make_pipeline(pca, svc)
        model = BaggingClassifier(svcpca, n estimators=100).fit(Xtrain, ytrain)
        yfit = model.predict(Xtest)
```

```
fig, ax = plt.subplots(6, 6)
for i, axi in enumerate(ax.flat):
    axi.imshow(Xtest[i].reshape(62, 47), cmap='bone')
    axi.set(xticks=[], yticks=[])
    axi.set ylabel(faces.target names[yfit[i]].split()[-1],
                   color='black' if yfit[i] == ytest[i] else 'red')
fig.suptitle('Predicted Names; Incorrect Labels in Red', size=14)
plt.show()
mat = confusion_matrix(ytest, yfit)
sns.heatmap(mat.T, square=True, annot=True, fmt='d', cbar=False,
            xticklabels=faces.target_names,
            yticklabels=faces.target_names)
plt.xlabel('true label')
plt.ylabel('predicted label')
plt.show()
print("Accuracy = ", accuracy_score(ytest, yfit))
```





eorge W Blushichiro Koiz Ginenoirge W BushTony Blair Ariel Sharon

eorge W Bloshald Rums Geeldrge W Bucsehorge W Bucsehorge W Bus

{'svc__C': 10, 'svc__gamma': 0.005}

Predicted Names; Incorrect Labels in Red

