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Market Basket Insights

TEAM: 10

Market Basket Insights

Important note:

This notebook has become very large, and I could not add more information to it, so I will divide it into a number of notebooks. So you can understand the content And be an excellent reference for you

Table of Content:

Machine Learning and Types

Application of Machine Learning

Steps of Machine Learning

Factors help to choose algorithm

Algorithm

Evaluate Algorithms

Linear Regression

Logistic Regression

Support Vector Machine

Naive Bayes Algorithm

KNN

Perceptron

Random Forest

Decision Tree

Extra Tree

Gradient Boosting

Light GBM

XGBoost

Catboost

Stochastic Gradient Descent

Lasso

Kernel Ridge Regression

Bayesian Ridge

Elastic Net Regression

LDA

K-Means Algorithm

CNN

LSTM

PCA

Apriori

Prophet

ARIMA

Machine Learning:

Linear Regression:

It is a basic and commonly used type of predictive analysis. These regression estimates are used to explain the relationship between one dependent variable and one or more independent variables. Y = a + bX where

- Y Dependent Variable
- a intercept Bise
- X Independent variable
- b Slope -Weights

Hypothesis:
$$h_{\theta}(x) = \theta_0 + \theta_1 x$$

Parameters:
$$\theta_0, \theta_1$$

Parameters:
$$\theta_0, \theta_1$$
 Cost Function: $J(\theta_0, \theta_1) = \frac{1}{2m} \sum_{i=1}^m \left(h_\theta(x^{(i)}) - y^{(i)}\right)^2$

Goal:
$$\min_{\theta_0,\theta_1} \text{minimize } J(\theta_0,\theta_1)$$

Example: University GPA' = (0.675)(High School GPA) + 1.097

import numpy as np

import pandas as pd

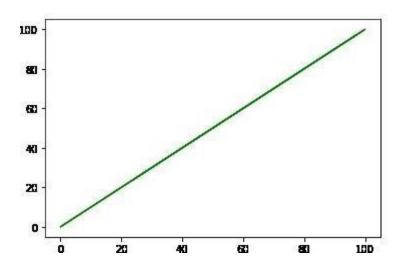
from matplotlib import pyplot as plt

```
from sklearn.linear_model import LinearRegression
from sklearn.metrics import mean_squared_error, r2_score
train = pd.read_csv("../input/random-linear-regression/train.csv")
test = pd.read_csv("../input/random-linear-regression/test.csv")
train = train.dropna()
test = test.dropna()
train.head()
OUTPUT:
      X
             y
      24.0 21.549452
0
1
      50.0 47.464463
2
      15.0 17.218656
3
      38.0 36.586398
4
      87.0 87.288984
Model with plots and accuracy:
X_train = np.array(train.iloc[:, :-1].values)
y_train = np.array(train.iloc[:, 1].values)
X_test = np.array(test.iloc[:, :-1].values)
y_test = np.array(test.iloc[:, 1].values)
model = LinearRegression(fit_intercept=True,
normalize=True,copy_X=True,n_jobs=-1)
model.fit(X_train, y_train)
```

y_pred = model.predict(X_test)

accuracy = model.score(X_test, y_test)

plt.plot(X_train, model.predict(X_train), color='green')
plt.show()
print(accuracy)

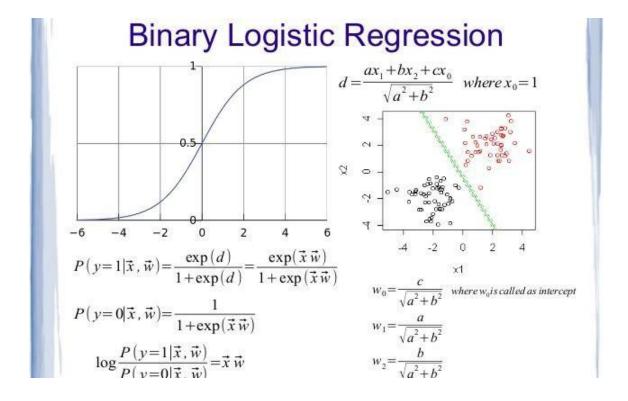


0.9888014444327563

Logistic Regression:

It's a classification algorithm, that is used where the response variable is categorical. The idea of Logistic Regression is to find a relationship between features and probability of particular outcome.

odds=p(x)/(1-p(x)) = probability of event occurrence / probability of not event occurrence



Libraries and data:

import sklearn

from sklearn.model_selection import train_test_split from sklearn.linear_model import LogisticRegression from sklearn.metrics import r2_score from statistics import mode

train = pd.read_csv("../input/titanic/train.csv")
test = pd.read_csv('../input/titanic/test.csv')
train.head()
OUTPUT:

	Passengerld	Survived	Pclass	Name	Sex	Age	SibSp	Parch	Ticket	Fare	Cabin	Embarked
0	1	0	3	Braund, Mr. Owen Harris	male	22.0	1	0	A/5 21171	7.2500	NaN	S
1	2	1	1	Cumings, Mrs. John Bradley (Florence Briggs Th	female	38.0	1	0	PC 17599	71.2833	C85	С
2	3	1	3	Heikkinen, Miss. Laina	female	26.0	0	0	STON/02. 3101282	7.9250	NaN	S
3	4	1	1	Futrelle, Mrs. Jacques Heath (Lily May Peel)	female	35.0	1	0	113803	53.1000	C123	S
4	5	0	3	Allen, Mr. William Henry	male	35.0	0	0	373450	8.0500	NaN	S

Model and Accuracy:

```
from sklearn.model_selection import train_test_split
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.25, random_state=5)
from sklearn.linear_model import LogisticRegression
#linear_model.LogisticRegression(penalty='12', dual=False, tol=0.0001, C=1.0, fit
_intercept=True, intercept_scaling=1,
# class_weight=None, random_state=None, solver='warn', max_iter=100,
# multi_class='warn', verbose=0, warm_start=False, n_jobs=None)

model = LogisticRegression(max_iter = 500000)
model.fit(X_train, y_train)
y_pred = model.predict(X_test)
accuracy = model.score(X_test, y_test)
print(accuracy)
OUTPUT:
0.8251121076233184
```

Support Vector Machine:

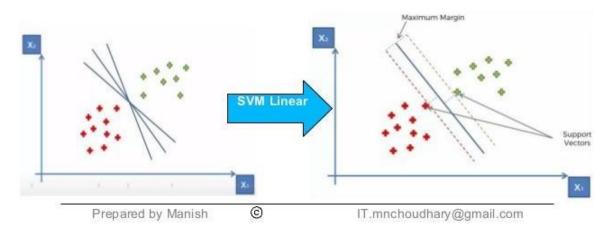
Data Science



Classification Model: SVM - Linear

Linearly separate the data points

Support Vector Machine" (SVM) is a **supervised machine** learning algorithm which can be used for **both classification or regression challenges**. However, it is mostly used in classification problems. In this algorithm, we plot each data item as a point in n-dimensional space (where n is number of features you have) with the value of each feature being the value of a particular coordina. Then, we perform classification by finding the **hyper-plane that differentiate the two classes** very well (look at the below snapshot).



Support Vector Machines are perhaps one of the most popular and talked about machine learning algorithms. It is primarily a classier method that performs classification tasks by constructing hyperplanes in a multidimensional space that separates cases of different class labels. SVM supports both regression and classification tasks and can handle multiple continuous and categorical variables

Example: One class is linearly separable from the others like if we only had two features like Height and Hair length of an individual, we'd first plot these two variables in two dimensional space where each point has two co-ordinates

Libraries and Data:

from sklearn.model_selection import train_test_split from sklearn.model_selection import cross_val_score from sklearn.svm import SVC

data_svm = pd.read_csv("../input/svm-classification/UniversalBank.csv") data svm.head()

	ID	Age	Experience	Income	ZIP Code	Family	CCAvg	Education	Mortgage	Personal Loan	Securities Account	CD Account	Online	CreditCard
0	1	25	1	49	91107	4	1.6	1	0	0	1	0	0	0
1	2	45	19	34	90089	3	1.5	1	0	0	1	0	0	0
2	3	39	15	11	94720	1	1.0	1	0	0	0	0	0	0
3	4	35	9	100	94112	1	2.7	2	0	0	0	0	0	0
4	5	35	8	45	91330	4	1.0	2	0	0	0	0	0	1

Model and Accuracy:

OUTPUT:

Naive Bayes Algorithm:

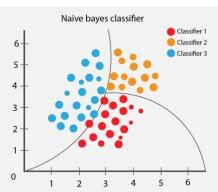
Naive Bayes



In machine learning, naive Bayes classifiers are a family of simple "probabilistic classifiers" based on applying Bayes' theorem with strong (naive) independence assumptions between the features.

$$P(A|B) = \frac{P(B|A) P(A)}{P(B)}$$

using Bayesian probability terminology, the above equation can be written as



A naive Bayes classifier is not a single algorithm, but a family of machine learning algorithms which use probability theory to classify data with an assumption of independence between predictors It is easy to build and particularly useful for very large data sets. Along with simplicity, Naive Bayes is known to outperform even highly sophisticated classification methods

Example: Emails are given and we have to find the spam emails from that. A spam filter looks at email messages for certain key words and puts them in a spam folder if they match.

Libraries and Data:

from sklearn.naive_bayes import GaussianNB from sklearn.preprocessing import StandardScalerfrom sklearn.metrics import accuracy_score

data =
pd.read_csv('../input/classification-suv-dataset/Social_Network_Ads.
csv')

data_nb = data
data_nb.head()

	User ID	Gender	Age	EstimatedSalary	Purchased
0	15624510	Male	19	19000	0
1	15810944	Male	35	20000	0
2	15668575	Female	26	43000	0
3	15603246	Female	27	57000	0
4	15804002	Male	19	76000	0

Model and Accuracy:

```
X = data_nb.iloc[:,
[2,3]].valuesy = data_nb.iloc[:,
4].values
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size = 0.20, random_state = 0)
sc_X = StandardScaler()
X_train = sc_X.fit_transform(X_train)
X_test = sc_X.transform(X_test)

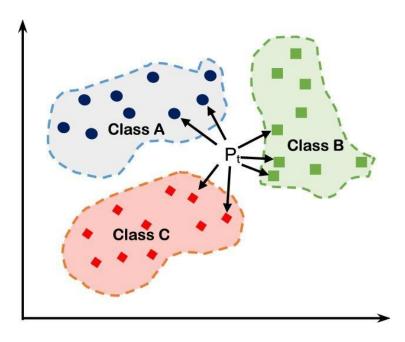
#sklearn.naive_bayes.GaussianNB(priors=None, var_smoothing=1e-09)
""
```

```
classifier=GaussianNB()
classifier.fit(X_train,y_train)
y_pred=classifier.predict(X_test)
acc=accuracy_score(y_test, y_pred)
```

0.9125

KNN:

KNN does not learn any model. and stores the entire training data set which it uses as its representation. The output can be calculated as the class with the highest frequency from the K-most similar instances. Each instance in essence votes for their class and the class with the most votes is taken as the prediction



Libraries and Data:

from sklearn.neighbors import KNeighborsClassifier knn = pd.read_csv("../input/iris/Iris.csv") knn.head()

	Id	SepalLengthCm	SepalWidthCm	PetalLengthCm	PetalWidthCm	Species
0	1	5.1	3.5	1.4	0.2	Iris-setosa
1	2	4.9	3.0	1.4	0.2	Iris-setosa
2	3	4.7	3.2	1.3	0.2	Iris-setosa
3	4	4.6	3.1	1.5	0.2	Iris-setosa
4	5	5.0	3.6	1.4	0.2	Iris-setosa

Model and Accuracy:

```
X = \text{knn.iloc}[:, [1,2,3,4]].\text{values}
```

y = knn.iloc[:, 5].values

X_train, X_test, y_train, y_test = train_test_split(X, y, test_size = 0.20, random_state = 0)

sc_X = StandardScaler()

X_train = sc_X.fit_transform(X_train)

 $X_{test} = sc_X.transform(X_{test})$

,,,

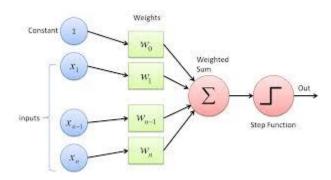
sklearn.neighbors.KNeighborsClassifier(n_neighbors=5, weights='uniform', algorithm='auto', leaf_size=30,

p=2, metric='minkowski', metric_params=None,n_jobs=None)

classifier=KNeighborsClassifier(n_neighbors=5,metric='minkowski',p=2)
classifier.fit(X_train,y_train)
y_pred=classifier.predict(X_test)
acc=accuracy_score(y_test, y_pred)
print(acc)

1.0

Perceptron:



It is single layer neural network and used for classification

from sklearn.linear_model import Perceptron from sklearn.neighbors import KNeighborsClassifier p = pd.read_csv("../input/iris/Iris.csv") p.head()

	ld	SepalLengthCm	SepalWidthCm	PetalLengthCm	PetalWidthCm	Species
0	1	5.1	3.5	1.4	0.2	Iris-setosa
1	2	4.9	3.0	1.4	0.2	Iris-setosa
2	3	4.7	3.2	1.3	0.2	Iris-setosa
3	4	4.6	3.1	1.5	0.2	Iris-setosa
4	5	5.0	3.6	1.4	0.2	Iris-setosa

X = p.iloc[:, [1,2,3,4]].values
y = p.iloc[:, 5].values
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size = 0.20, random_state = 0)
sc_X = StandardScaler()
X_train = sc_X.fit_transform(X_train)
X_test = sc_X.transform(X_test)
""

classifier=Perceptron(penalty=None, alpha=0.0001, fit_intercept=True, max_iter=None, tol=None, shuffle=True,

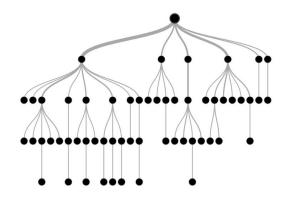
verbose=0, eta0=1.0, n_jobs=None, random_state=0, early_stopping=False, validation_fraction=0.1, n_iter_no_change=5, class_weight=None, warm_start=False, n_iter=None)

classifier=Perceptron()
classifier.fit(X_train,y_train)
y_pred=classifier.predict(X_test)
acc=accuracy_score(y_test, y_pred)
print(acc)

0.966666666666667

Decision Tree:

Decision tree algorithm is classification algorithm under supervised machine learning and it is simple to understand and use in data. The idea of Decision tree is to split the big data(root) into smaller(leaves)



from sklearn.tree import DecisionTreeClassifier
dt = data
dt.head()

	User ID	Gender	Age	EstimatedSalary	Purchased
0	15624510	Male	19	19000	0
1	15810944	Male	35	20000	0
2	15668575	Female	26	43000	0
3	15603246	Female	27	57000	0
4	15804002	Male	19	76000	0

```
X = dt.iloc[:, [2,3]].values
y = dt.iloc[:, 4].values
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size = 0.20, random_state = 0)
sc_X = StandardScaler()
X_train = sc_X.fit_transform(X_train)
X_test = sc_X.transform(X_test)
```

...

sklearn.tree.DecisionTreeClassifier(criterion='gini', splitter='best', max_depth=None,min_samples_split=2,

min_samples_leaf=1,min_weight_fraction_leaf=0.0,max_features=None, random_state=None, max_leaf_nodes=None,min_impurity_decrease=0.0, min_impurity_split=None, class_weight=None,presort=False)

ш

classifier=DecisionTreeClassifier(criterion="entropy",random_state=0) classifier.fit(X_train,y_train) y_pred=classifier.predict(X_test) acc=accuracy_score(y_test, y_pred) print(acc)

0.9

Extra Tree:

from sklearn.ensemble import ExtraTreesClassifier et = data et.head()

	User ID	Gender	Age	EstimatedSalary	Purchased
0	15624510	Male	19	19000	0
1	15810944	Male	35	20000	0
2	15668575	Female	26	43000	0
3	15603246	Female	27	57000	0
4	15804002	Male	19	76000	0

```
X = et.iloc[:, [2,3]].values
y = et.iloc[:, 4].values
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size = 0.20, random_state = 0)
sc_X = StandardScaler()
X_train = sc_X.fit_transform(X_train)
X_test = sc_X.transform(X_test)
```

classifier=ExtraTreesClassifier(criterion="entropy",random_state=0) classifier.fit(X_train,y_train)

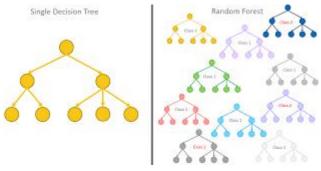
[&]quot;ExtraTreesRegressor(n_estimators=10, max_features=32, random_state=0)"

y_pred=classifier.predict(X_test)
acc=accuracy_score(y_test, y_pred)
print(acc)

0.9

Random Forest:

Random forest is collection of tress(forest) and it builds multiple decision trees and merges them together to get a more accurate and stable prediction. It can be used for both classification and regression problems.



Example: Suppose we have a bowl of 100 unique numbers from 0 to 99. We want to select a random sample of numbers from the bowl. If we put the number back in the bowl, it may be selected more than once.

Libraries and Data:

from sklearn.ensemble import RandomForestClassifier
rf = pd.read_csv("../input/mushroom-classification/mushrooms.csv")
rf.head()

	class	cap- shape	cap- surface	cap- color	bruises	odor	gill- attachment	gill- spacing	gill- size	gill- color	***	stalk- surface- below- ring	stalk- color- above- ring	stalk- color- below- ring	veil- type	veil- color	ring- number	ring- type	spore- print- color	popul
	р	х	s	n	t	р	f	С	n	k	***	S	W	W	р	W	0	р	k	S
	е	X	s	У	t	а	f	С	b	k	***	S	w	w	р	w	0	р	n	n
	е	b	S	W	t	1	f	С	b	n	***	S	w	w	р	W	0	р	n	n
	р	х	У	w	t	р	f	С	n	n		S	w	w	р	w	0	р	k	s
	е	×	s	g	f	n	f	w	b	k		S	W	w	р	W	0	е	n	а
ë	_				_	_														•

X = rf.drop('class', axis=1)

y = rf['class']

X = pd.get_dummies(X)

y = pd.get_dummies(y)

X_train, X_test, y_train, y_test = train_test_split(X, y)

ш

ensemble.RandomForestClassifier(n_estimators='warn', criterion='gini', max_depth=None, min_samples_split=2, min_samples_leaf=1,min_weight_fraction_leaf=0.0, max_features='auto',max_leaf_nodes=None,min_impurity_decrease=0.0, min_impurity_split=None, bootstrap=True,oob_score=False, n_jobs=None,

```
random_state=None, verbose=0,warm_start=False, class_weight=None)
```

```
model = RandomForestClassifier (n_estimators=100, max_depth=10, random\_state=1) \\ model.fit(X_train, y_train) \\ model.score(X_test, y_test)
```

1.0