



21/03/2024

WEEK - 1

Write a python program to import and export data using pandas library functions.

```
import pandas as pd
```

```
url = "https://archive.ics.uci.edu/ml/machine-learning-databases/iris/iris.data"
```

```
col_names = ["sepal-length-in-cm", "sepal-width-in-cm",  
             "petal-length-in-cm", "petal-width-in-cm", "class"]
```

```
iris_data = pd.read_csv(url, names=col_names)
```

```
iris_data.head()
```

OUTPUT

	sepal-length-in-cm	sepal-width-in-cm	<del>pet</del> sepal-length-in-cm	petal-width-in-cm	class
0	5.1	3.5	1.4	0.2	Iris-setosa

```
iris_data.to_csv("cleaned-iris-data.csv")
```

```
df2 = pd.read_csv("cleaned-iris-data.csv")
```

	sepal-length-in-cm	sepal-width-in-cm	petal-length-in-cm	petal-width-in-cm	class
0	5.1	3.5	1.4	0.2	Iris-setosa

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## End-to-End Machine Learning Project

1. Select a Performance Measure

2. Get the Data

```
HOUSING_PATH = 'https://raw.githubusercontent.com/content.com/housing.csv'
```

```
housing = pd.read_csv(HOUSING_PATH)
```

```
housing.head()
```

```
housing.info()
```

```
housing['ocean-proximity'].value_counts()
```

```
housing.describe()
```

```
housing.groupby(by=['longitude', 'latitude']).count()  
['total_rooms'].sort_values()
```

```
from sklearn.model_selection import train_test_split
```

```
train_set, test_set = train_test_split(housing, test_size=0.2,  
                                        random_state=42)
```

```
housing['income_cat'] = pd.cut(x=housing['median_income'],  
                               bins=[0, 1.5, 3, 4.5, 6, np.inf], labels=[1, 2, 3, 4, 5])
```

```
housing['income_cat'].hist()
```



### 3. Discover and Visualize the Data to Gain Insights

plot  
housing.plot(kind='scatter', x='longitude', y='latitude')  
plt.show()

housing.plot(kind='scatter', x='longitude', y='latitude',  
alpha=0.1)  
plt.show()

#### Correlation

housing[['population', 'median-house-value']].corr()

corr\_matrix = housing.corr()

corr\_matrix['median-house-value'].sort\_values(ascending=False)

scatter-matrix() for correlation

from pandas.plotting import scatter\_matrix

attributes = ['median-house-value', 'median-income',  
'total-rooms', 'housing-median-age']

scatter\_matrix(frame = housing[attributes], figsize=(12, 8))

Experimenting with attribute combinations

housing['rooms-per-household'] = housing['total-rooms'] /  
housing['households']

housing['bedrooms-per-room'] = housing['total-bedrooms'] /  
housing['total-rooms']

housing['population-per-household', sort-values (ascending =  
= housing['population'] / housing['households']

#### 4. Prepare the Data for Machine Learning Algorithms

##### Data cleaning

```
from sklearn.impute import SimpleImputer
imputer = SimpleImputer(strategy='median')
housing-num = housing.drop("ocean-proximity", axis=1)
imputer.fit(housing-num)
imputer.statistics_
housing-num.median().values
```

##### One Hot Encoding

```
from sklearn.preprocessing import OneHotEncoder
one-hot-encoder = OneHotEncoder()
housing-cat-1hot = one-hot-encoder.fit_transform(housing-cat-1hot)
housing-cat-1hot
housing-cat-1hot.toarray()
one-hot-encoder.categories_
```

##### Transformation pipelines

```
from sklearn.pipeline import Pipeline
from sklearn.preprocessing import StandardScaler
num_pipeline = Pipeline([('imputer', SimpleImputer(
    strategy='median')), ('attribute-adder',
    CombinedAttributesAdder())], ('std-scaler', StandardScaler())
])
```

```
housing-num-tr = num_pipeline.fit_transform(housing-num)
housing-num-tr.shape
```

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20/3/24

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## 5. Select and Train a model.

```
from sklearn.linear_model import LinearRegression  
lin_reg = LinearRegression()  
lin_reg.fit(X=housing_prepared, y=housing_labels)
```

Try to modelling on a few instances from the training set:

```
some_data = housing.iloc[:5]  
some_labels = housing_labels.iloc[:5]  
some_data_prepared = full_pipeline.transform(some_data)  
print("Predictions:", lin_reg.predict(some_data_prepared))  
print("Labels:", some_labels.tolist())
```

~~to~~ Measuring performance using RMSE metric.

```
from sklearn.metrics import mean_squared_error  
housing_predictions = lin_reg.predict(housing_prepared)  
lin_mse = mean_squared_error(housing_labels,  
                             housing_predictions)
```

```
lin_rmse = np.sqrt(lin_mse)
```

```
lin_rmse
```



## 6. Fine-Tune Your Model

### Grid search

Mentioning hyper-parameters and values to test and it will test out all combinations of hyper-parameters and use cross-validation for evaluation.

```
from sklearn.model_selection import GridSearchCV  
param_grid = [ {'n_estimators': [3, 10, 30], 'max_features': [25, 50, 75],  
                {'bootstrap': [False], 'n_estimators': [3, 10], 'max_features':  
                  [2, 3, 4]} ]
```

```
forest_reg = RandomForestRegressor()
```

```
grid_search = GridSearchCV(estimator=forest_reg,  
                             param_grid=param_grid, scoring='neg-mean-squared-  
                             error',  
                             cv=5, return_train_score=True, n_jobs=-1)
```

```
grid_search.fit(X=housing_prepared, y=housing_labels)
```

```
grid_search.best_params_
```

```
grid_search.best_estimator_
```

```
cv_results = grid_search.cv_results_
```

```
for mean_score, params in zip(cv_results['mean_test_score'],  
                               cv_results['params']):
```

```
    print(np.sqrt(-mean_score), params)
```

## 7. Launch, Monitor, & Maintain your system

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## Simple Linear Regression

```
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
import seaborn as sns
from sklearn.model_selection import train_test_split
from pandas.core.common import import random_state
from sklearn.linear_model import LinearRegression
```

```
df_sal = pd.read_csv('content/salary-Data.csv')
df_sal.head()
```

```
df_sal.describe()
```

```
plt.title('Salary Distribution Plot')
sns.distplot(df_sal['salary'])
plt.show()
```

```
plt.scatter(df_sal['Years Experience'], df_sal['salary'],
            color='lightcoral')
```

```
plt.title('Salary vs Experience')
plt.xlabel('Years of Experience')
plt.ylabel('Salary')
plt.box(False)
plt.show()
```

## Split data

```
x = df_sal.iloc[:, 1:]
y = df_sal.iloc[:, 2:]
```



## Split into Train test split sets

```
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=0)
```

## Train model

```
regressor = LinearRegression()  
regressor.fit(X_train, y_train)
```

## Predict results

```
y_pred_test = regressor.predict(X_test)  
y_pred_train = regressor.predict(X_train)
```

## Visualize predictions

```
plt.scatter(X_train, y_train, color='lightcoral')  
plt.plot(X_train, y_pred_train, color='firebrick')  
plt.title('Salary vs Experience (Training Set)')  
plt.xlabel('Years of Experience')  
plt.ylabel('Salary')  
plt.legend(['X_train/Pred(y_test)', 'X_train/y_train'],  
           title='sal/exp', loc='best', facecolor='white')  
plt.box(False)  
plt.show()
```

```
plt.scatter(X_test, y_test, color='lightcoral')  
plt.plot(X_train, y_pred_train, color='firebrick')  
plt.scattertitle('Salary vs Experience (Test Set)')  
plt.xlabel('Years of Experience')  
plt.ylabel('Salary')  
plt.show()
```

Coefficient and Intercept:

print ('Coefficient: regressor.coef - y')

print ('Intercept: regressor.intercept - y')

Coefficient: [59312.57]

Intercept: [26780.097]

## Multiple Linear Regression

```
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
import seaborn as sns
from sklearn.model_selection import train_test_split
from sklearn.linear_model import LinearRegression
```

```
df_start = pd.read_csv('/content/startup.csv')
df_start.head()
```

```
df_start.describe()
```

### Distribution

```
plt.title('Profit Distribution Plot')
sns.distplot(df_start['Profit'])
plt.show()
```

### Relation between Profit and R&D Spend

```
plt.scatter(df_start['R&D Spend'], df_start['Profit'],
            color='lightcoral')
```

```
plt.title('Profit vs R&D Spend')
```

```
plt.xlabel('R&D Spend')
```

```
plt.ylabel('Profit')
```

```
plt.box(False)
```

```
plt.show()
```



Split into Independent/Dependent variables

`x = df.start.iloc[:, :-1].values`

`y = df.start.iloc[:, -1].values`

One-hot encoding

`ct = ColumnTransformer(transformers = ['encoder', OneHotEncoder,  
[3]], remainder = 'passthrough')`

`x = np.array(ct.fit_transform(x))`

Split into Train/Test sets

`x_train, x_test, y_train, y_test = train_test_split(x, y,  
test_size = 0.2, random_state = 0)`

Train model

`regressor = LinearRegression()`

`regressor.fit(x_train, y_train)`

Predict results

`y_pred = regressor.predict(x_test)`

Compare predictions

`np.set_printoptions(precision = 2)`

`result = np.concatenate((y_pred.reshape(len(y_pred), 1),  
y_test.reshape(len(y_test), 1)), 1)`

`result`

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## Decision tree (ID3)

```
import pandas as pd
from sklearn.tree import DecisionTreeClassifier,
plot_tree
```

```
import matplotlib.pyplot as plt
import math
```

```
df = pd.read_csv('content/drive/MyDrive/Iris.csv')
df.head()
```

```
from sklearn import datasets
```

```
iris = datasets.load_iris()
```

```
iris_df = pd.DataFrame(data=iris.data, columns=iris.
feature_names)
```

```
iris_df['species'] = iris.target
```

```
iris_df['species'] = iris_df['species'].map(0: 'setosa', 1:
'versicolor', 2: 'virginica')
```

```
iris_df.head()
```

```
y = iris_df['species']
```

```
x = iris_df.drop(['species'], axis=1)
```

```
from sklearn.model_selection import train_test_split
```

```
x_train, x_test, y_train, y_test =
```

```
train_test_split(x, y, test_size=0.33, random_state=42)
```

```
from sklearn.tree import DecisionTreeClassifier
```

```
clf = DecisionTreeClassifier (criterion="gini", random_state=100,  
                             max_depth=5, min_samples_leaf=8)
```

```
clf.fit (X_train, y_train)
```

```
y_pred = clf.predict (X_test)
```

```
from sklearn.metrics import accuracy_score
```

```
accuracy = accuracy_score (y_pred, y_test)
```

```
print ("Accuracy : {accuracy %}")
```

Accuracy : 0.98

```
from sklearn import tree
```

```
plt.figure (figsize = (12, 8))
```

```
tree.plot_tree (clf, filled = True, feature_names = ['sepal length (cm)',  
                                                      'sepal width (cm)', 'petal length (cm)', 'petal width (cm)'],
```

```
class_names = ['setosa', 'versicolor', 'virginica'])
```

```
plt.show()
```



petal length (cm)  $\leq 2.45$   
gini = 0.666  
samples = 100  
value = [3, 35, 34]  
class = versicolor

gini = 0.0  
samples = 3  
value = [3, 0, 0]  
class = setosa

petal width (cm)  $\leq 1.75$   
gini = 0.5  
samples = 69  
value = [0, 35, 34]  
class = versicolor

petal length (cm)  $\leq 4.45$   
gini = 0.188  
samples = 38  
value = [0, 34, 4]  
class = versicolor

petal width (cm)  $\leq 1.85$   
gini = 0.062  
samples = 31  
value = [0, 1, 35]  
class = virginica

gini = 0.0  
samples = 23  
value = [0, 24, 0]  
class = versicolor

gini = 0.391  
samples = 15  
value = [0, 11, 4]  
class = versicolor

gini = 0.198  
samples = 9  
value = [0, 1, 8]  
class = virginica

gini = 0.0  
samples = 22  
value = [0, 0, 22]  
class = virginica

Link  
18/4/24

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## Insurance System

```
import pandas as pd.  
from matplotlib.pyplot import plt.  
from sklearn.model_selection  
import train_test_split  
from sklearn.linear_model import LogisticRegression  
  
df = pd.read_csv('id.csv')  
  
X_train, X_test, y_train, y_test = train_test_split(df[['age']],  
df['bought_insurance'], test_size=0.2)  
  
model = LogisticRegression()  
model.fit(X_train, y_train)  
y_predicted = model.predict(X_test)  
print(y_predicted)  
  
print(model.predict_proba(X_test))  
print(model.score(X_test, y_test))
```

import Math

```
def sigmoid(z):  
    return 1/(1 + Math.exp(-z))
```

```
def pradi(age):  
    z = 0.042 * age - 1.53  
    y = sigmoid(z)  
    return y
```

```
print(pradi(35))
```

```
print(pradi(43))
```

Output

Prediction: array([1, 0, 1, 0, 0, 0, 0, 1, 0])

Score: 0.8888

Linear Reg score: 0.584321

Predictions: 0.485

0.5685.

Dwa  
9-5-2024



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Program 6

KNN - K Nearest Neighbours

```
import pandas as pd
import seaborn as sns
import matplotlib.pyplot as plt
import numpy as np

df = pd.read_csv ("content/drive/My Drive/Iris.csv")
df.head()

classes = df["Species"].unique()
colors = ['r', 'g', 'b']

for i, cls in enumerate(classes):
    class_data = df[df["Species"] == cls]
    plt.scatter(class_data["sepal.length.cm"], class_data["Petal
        Length.cm"], c=colors[i], label=cls)

plt.xlabel('Sepal length [cm]')
plt.ylabel('Petal length [cm]')
plt.legend()
plt.show()

y = df["Species"]
X = df.drop(["Species"], axis=1)

from sklearn.model_selection import train_test_split
X_train, X_test, y_train, y_test =
    train_test_split(X, y, test_size=0.3, random_state=0)
```

```

from sklearn.neighbors import KNeighborsClassifier
knn = KNeighborsClassifier(n_neighbors=3)
knn.fit(X_train, y_train)
y_pred = knn.predict(X_test)
y2 = knn.predict(X_train)

```

```

from sklearn.metrics import accuracy_score
score = accuracy_score(y_pred, y_test)
print(f"Testing Accuracy: {score}")
score2 = accuracy_score(y2, y_train)
print(f"Training Accuracy: {score2}")

```

Testing Accuracy: 1.0  
Training Accuracy: 1.0

Program 7

SVM - Support Vector Machine

```

from sklearn.svm import SVC
model = SVC(kernel='linear', random_state=0, C=1.0)
model.fit(X_train, y_train)
y_pred = model.predict(X_test)
y2 = model.predict(X_train)

```

```

score = accuracy_score(y_pred, y_test)
print(f"Testing Accuracy: {score}")
score2 = accuracy_score(y2, y_train)
print(f"Training Accuracy: {score2}")

```

Testing Accuracy: 1.0

Training Accuracy: 1.0

*Wah*  
9-5-24

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## Program 8.

### Implementation of ANN using Back propagation for given values

```
import numpy as np
```

```
X = np.array([2, 9], [1, 5], [3, 6]), dtype=float)
```

```
y = np.array([92], [86], [89]), dtype=float)
```

```
X = X/np.array(X, axis=0)
```

```
y = y/100
```

```
epoch = 5000
```

```
lr = 0.1
```

```
input_layer_neurons = 2
```

```
hidden_layer_neurons = 3
```

```
output_neurons = 1
```

```
w1 = np.random.uniform(size=(input_layer_neurons, hidden_layer_neurons))
```

```
b1 = np.random.uniform(size=(1, hidden_layer_neurons))
```

```
w2 = np.random.uniform(size=(hidden_layer_neurons, output_neurons))
```

```
b2 = np.random.uniform(size=(1, output_neurons))
```

```
def sigmoid(x):
```

```
    return 1/(1+np.exp(-x))
```

```
def derivative_sigmoid(x):
```

```
    return x*(1-x)
```



```
for i in range(epochs):
```

```
    hinp1 = np.dot(X, wh)
```

```
    hinp = hinp1 + bh
```

```
    hlayer_act = sigmoid(hinp)
```

```
    outinp1 = np.dot(hlayer_act, wout)
```

```
    outinp = outinp1 + bout
```

```
    output = sigmoid(outinp)
```

```
    E0 = y - output
```

```
    outgrad = derivatives_sigmoid(output)
```

```
    d_output = E0 * outgrad
```

```
    EH = d_output * dot(wout.T)
```

```
    hiddengrad = derivatives_sigmoid(hlayer_act)
```

```
    d_hidden_layer = EH * hiddengrad
```

```
    wout += hlayer_act.T * dot(d_output) * lr
```

```
    wh += X.T * dot(d_hidden_layer) * lr
```

```
    print("Input: \n" + str(X))
```

```
    print("Actual Output: \n" + str(y))
```

```
    print("Predicted Output: \n", output)
```

## Output

Input:

```
[ [0.6667  1.    ]  
  [0.3334  0.556 ]  
  [1.     0.6667]]
```

Actual output:

```
[ [0.92]  
  [0.86]  
  [0.89]]
```

predicted Output:

```
[ [0.935]  
  [0.923]  
  [0.939]]
```

## Program 9a

### Random Forest Algorithm

```
import pandas as pd
import seaborn as sns
import matplotlib.pyplot as plt
import numpy as np
```

```
df = pd.read_csv ("content/drive/My Drive/Irrs.csv")
df.head()
```

```
y = df["species"]
```

```
X = df.drop(["species"], axis=1)
```

```
from sklearn.model_selection import train_test_split
X_train, X_test, y_train, y_test = train_test_split
(X, y, test_size=0.3, random_state=0)
```

~~Random~~

```
from sklearn.ensemble import RandomForestClassifier
clf = RandomForestClassifier(n_estimators=100)
```

```
clf.fit(X_train, y_train)
```

```
y_pred = clf.predict(X_test)
```

~~q2~~

```
from sklearn.metrics import accuracy_score
```

```
score = accuracy_score(y_pred, y_test)
```

```
print(f"Accuracy: {score}")
```

Output

Accuracy: 1.0.

## AdaBoost (with default parameters)

```
from sklearn.ensemble import AdaBoostClassifier.
```

```
adb = AdaBoostClassifier()
```

```
adb_model = adb.fit(X_train, y_train)
```

```
y_pred = adb_model.predict(X_test)
```

```
score = accuracy_score(y_pred, y_test)
```

```
print(f"Accuracy : {score}")
```

Accuracy : 0.977

## AdaBoost (with Hyper parameters)

```
from sklearn.linear_model import LogisticRegression
```

```
lrmodel = LogisticRegression()
```

```
adthp = AdaBoostClassifier(n_estimators=150, estimator=lrmodel,  
learning_rate=1)
```

```
model = adthp.fit(X_train, y_train)
```

```
y_pred = model.predict(X_test)
```

```
score = accuracy_score(y_pred, y_test)
```

```
print(f"Accuracy : {score}")
```

OUTPUT

Accuracy : 1.0.

*Shake*  
23/05/24



30/05/2024

## Lap. program - 10

Build K-Means algorithm to cluster a set of data stored in a .csv file.

```
import matplotlib.pyplot as plt
from sklearn import datasets
from sklearn.cluster import KMeans
import pandas as pd
import numpy as np
```

```
iris = datasets.load_iris()
```

```
X = pd.DataFrame(iris.data)
```

```
X.columns = ['Sepal_Length', 'Sepal_Width', 'Petal_Length',  
             'Petal_Width']
```

```
y = pd.DataFrame(iris.target)
```

```
y.columns = ['Targets']
```

```
model = KMeans(n_clusters=3)
```

```
model.fit(X)
```

```
plt.figure(figsize=(14,14))
```

```
colormap = np.array(['red', 'lime', 'black'])
```


```
plt.subplot(2,2,1)
```

```
plt.scatter(X.Petal_Length, X.Petal_Width, c=colormap[y.Targets],  
            s=40)
```

```
plt.title('Real clusters')
```

```
plt.xlabel('Petal Length')
```

```
plt.ylabel('Petal Width')
```



```
plt.subplot(2,2,2)
```

```
plt.scatter(X.Petal.Length, X.Petal.Width,
```

```
c=colormap[model.labels-], s=40)
```

```
plt.title('K-Means Clustering')
```

```
plt.xlabel('Petal.Length')
```

```
plt.ylabel('Petal.Width')
```

✓

1-1

## Lab program - II

Implement Dimensionality reduction using Principle Component Analysis (PCA) method.

```
import matplotlib.pyplot as plt
import pandas as pd
import numpy as np
import seaborn as sns
%matplotlib inline
```

```
from sklearn.datasets import load_breast_cancer()
cancer = load_breast_cancer()
cancer.keys()
```

```
print(cancer['DESCR'])
```

```
df = pd.DataFrame(cancer['data'], columns=cancer['feature_names'])
df.head()
```

```
from sklearn.preprocessing import StandardScaler
scaler = StandardScaler()
scaler.fit(df)
```

```
scaled_data = scaler.transform(df)
```

```
from sklearn.decomposition import PCA
pca = PCA(n_components=2)
pca.fit(scaled_data)
```

```
scaled_data x_pca = pca.transform(scaled_data)
```

```
scaled_data.shape
```

```
(569, 30)
```



x\_pca.shape

(569, 2)

plt.figure(figsize=(8,6))

plt.scatter(x\_pca[:,0], x\_pca[:,1],  
c=cancer['target'], cmap='plasma')

plt.xlabel('First Principal Component')

plt.ylabel('Second Principal Component')

✓  
3-15h