

I N D E X

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21/03/24

Lab 1. Iris data Set.

① import pandas as pd
url = "https://archive.ics.uci.edu/ml/
machine-learning-database/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()

sepal-length in cm	sepal length in cm	petal length in cm
5.1	3.5	3.5 1.4

petal width in cm	class
0.4	iris-setosa

ins. data to csv "cleaned-ins. data"

Data Set is done in CSV

Due
24/11

① Telch

→ To load

diff

pat

OS.

sube

→ housing

Name

→ housing

Sho

→ %

impe

hou

glt.

→ (

med-inis - data.csv
CSV

Week-02

① Fetch the Data of housing:

→ To load the data:

```
def load_housing_data(housing-path = housing-path):  
    csv-path = os.path.join(housing-path, "housing.csv")  
    return pd.read_csv(csv-path).
```

→ housing["Ocean-proximity"].value_counts()

Name: ocean-proximity, dtype: int64

→ housing.describe()
Shows all the output

→ %matplotlib inline
import matplotlib.pyplot as plt
housing.hist(bins=50, figsize=(20,15))
plt.show.

Output → histogram for each numerical attribute.

→ Create a test case:

```
housing["income-cat"] = pd.cut(housing["medium-income"],
```

```
bins = [0, 1.5, 3.0, 4.5, 6., np.inf],  
labels = [1, 2, 3, 4, 5])
```

→ histogram of income categories.

→ strat-test-set["income-cat"].value_counts
not len(strat-test-set).

⇒ income-cat, dtype: float64

→ To create a copy of main dataset
housing = strat-train-set.copy()

→ Visualizing Geographical Data.

housing.plot(kind="scatter", x="longitude",
y="latitude")

Correlations.

corr_matrix = housing.corr().

→>> corr_matrix["median-house-value"].
sort_values(ascending=False)

Name: median-house-value, dtype: float64

Scatter Matrix

housing.plot(kind="scatter",
x="median-income", y=
"median-house-value",
alpha=0.1)

→ Median income versus median
house value.

Analyze +

→ feature

feature
array()

Ans
41

value-counts

label
py()

"longitude"

value"]

: float 64

7

Analyze the Best Models and their Errors
→ feature-importance = grid-search-best-estimator.feature-importances

feature-importance

array([7.33, 6.29-02 ... 2.8564-03])

Av
4/14/24

Week-03

part-02: Linear and Multiple
Regression.

import cv
→ Salary file

① Import pandas as pd.

d = pd.read_csv('salary.csv')
df.head()

② import libraries.

③ df.sal.describe()

④ plt.title('Salary Distribution Plot')
sns.distplot(df.sal['Salary'])
plt.show()

→ plot of Salary Distribution Plot
v/s Salary.

⑤ plt.scatter(df.sal['Years Experience'],
df.sal['Salary'], colour = 'light coral')

plt.title('Salary vs Experience')

plt.xlabel('Years of Experience')

plt.ylabel('Salary')

plt.box(False)

plt.show()

→

⑥ x = df.sal.
y = df.sal

⑦ x_train, x_test,
split(x, y)

⑧ regressor =
regressor.

// Start with

① Import

② Import

③ df

④ plt.
sns.
plt.

⑤ plot

⑥

⑦

⑧

⑨

and Multiple
Regression.

d.
v ('salary.csv')

tribution Plot
['Salary']

tribution Plot

s Experience',
light color')

ci")

u')

me.

⑥ x = df_sal.iloc[:, 1:]
y = df_sal.iloc[:, 2:]

⑦ x_train, x_test, y_train, y_test = train_test_split(x, y, test_size = 0.2, random_state = 0)

⑧ regressor = LinearRegression()
regressor.fit(x_train, y_train)

Startup file

① Import file.

② Import libraries.

③ df_start.describe()

④ plt.title('Profit Distribution Plot')
sns.distplot(df_start['profit'])
plt.show()

⑤ plot Profit vs Rand D spend graph.

⑥ x = df_start.iloc[:, :-1], values
y = df_start.iloc[:, :-1].values

④ x_train, x_test, y_train, y_test =
train_test_split(x, y, test_size = 0.2,
random_state = 0)

⑤ regressor = LinearRegression()
regressor.fit(x_train, y_train)

⑥ y_pred = regressor.predict(x_test)

Week 4. Decision Tree

```
> import pandas as pd
import matplotlib.pyplot as plt
from sklearn.datasets import load_iris
import seaborn as sns

from sklearn.datasets import load_iris
iris_data = load_iris()
print(iris_data.DESCR)
X = iris_data.data
y = iris_data.target
print("Shape of X:", X.shape)
print("Shape of y:", y.shape)

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 DecisionTree
Classifier.
```

```
tree_model = DecisionTreeClassifier()
tree_model.fit(X_train, y_train)
```

Decision Tree Classifier()

```
from sklearn.tree import DecisionTree
Classifier, plot_tree
import matplotlib.pyplot as plt.
```


- Libraries
- Importing Boston Housing Data.
 - Loading the Data
 - Preparing Data
 - Splitting Data into training and Testing Sets.
 - Creating and Training the Linear Regression Model.
 - Evaluating Model performance.
 - Plotting the error.

1/1/20

Week
 → import
 import
 from sklearn
 import

- from
 iris_data
 print (
 X = ir
 Y = ir
 print (
 print

from
 train
 X_train
 from
 ran

from
 Clas
 In
 tre
 D
 J
 u


```
clf = DecisionTreeClassifier()
clf.fit(X, y)
```

```
plt.figure(figsize=(12, 8))
plot_tree(clf, filled=True, feature_names=iris.data.feature_names, class_names=iris.data.target_names)
```

```
petal length <= 2.45
gini = 0.667
samples = 150
value = [50, 50, 50]
class = setosa
```

```
gini = 0.0
samples = 50
value = [50, 0, 0]
class = setosa
```

```
petal width (cm) <= 1.7
gini = 0.5
samples = 100
value = [0, 50, 50]
class = versicolour
```

```
import from sklearn.model_selection
import cross_val_score
scores = cross_val_score(clf, X, y, cv=5)
```

```
accuracy = scores.mean()
```

```
print("Mean Accuracy", accuracy)
```

```
Mean Accuracy = 0.960000
```


- ① import pandas as pd.
from matplotlib import pyplot as plt.
- ② read the csv file.

```

③ plt.scatter(df['age'], df['bought'],
               marker='+', color='red')
from sklearn.model_selection import
train_test_split
x_train, x_test, y_train = train_test
split(df[['age']], df['bought'],
       train_size=0.8)

```

```

print(x_test)
from sklearn.linear_model
import LogisticRegression
model = LogisticRegression()
model.fit(x_train, y_train)

```

print(x_test)

```

y_predicted = model.predict(x_test)
model.predict_proba(x_test)
model.score(x_test, y_test)

```



```
print(y_predicted)
print(x_test)
import math
```

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

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

```
def prediction_function(x):
```

```
    z = 0.042 * age - 1.53
```

```
    y = sigmoid(z)
```

```
    return y
```

```
age = 35
```

```
prediction_function(age)
```

```
y:- 0.485044983805899
```

```
age: 43
```

```
prediction_function(age)
```

```
y:- 0.568565299705
```

K.S.M.

Lab 4: KNN

04.05.24

sklearn KNN

classifiers

① importing libraries

② printing

insurance_data

insurance_data

③ plt.scatter (df.loc[:, 0], df.loc[:, 1],

c = df.loc[:, -1], cmap = viridis)

plt.xlabel ('feature')

plt.ylabel ('feature')

plt.title ('scatterplot of the dataset')

plt.show()

④ X = df.loc[:, :-1]

y = df.loc[:, -1]

⑤ X_train, X_test, y_train, y_test =

train_test_split (X, y, test_size = 0.3,

random_state = 42)

⑥ k_nn = KNeighborsClassifier (n_neighbors = 3)

⑦ k_nn.fit (X_train, y_train)

⇒ 15 Neighbors Classifier (n_neighbors = 3)

⑧ y_pred = k_nn.predict (X_test)

⑨ accuracy = accuracy_score (y_test, y_pred)

print ('Accuracy:', accuracy)

Accuracy = 0.888888888

Week 1: SVM. sklearn.svm

① import libraries

② plt.scatter (df[df['target']==0]['sepal length (cm)'], df[df['target']==0]['petal width (cm)'], label='setosa')

df['target']==1 df['target']==2

label='Virginica'

label='Virginica'

plt.xlabel ('sepal length (cm)')

plt.ylabel ('petal width (cm)')

plt.title ('sepal width vs length')

plt.legend()

plt.show()

X_train, X_test, y_train, y_test =

train_test_split (iris_data, iris_target,

test_size=0.2, random_state=42)

svm_classifier = SVC(kernel='linear')

svm_classifier.fit (X_train, y_train)

SVC(kernel='linear')

sum

$y = 0$ [input]

$[target] = 0$

label = 'setosa']

avgf = 2

color

area

gth (cm)

(cm)

length

nt =

, iris-target,

- state = 42)

$l = 'linear'$

y-train

8 - pred = sum accuracy (pred, y-train)
accuracy = accuracy (pred, y-train)
print ('Accuracy = ', accuracy)

Accuracy = 1.0

(1.0, 1.0, 1.0, 1.0, 1.0)

(1.0, 1.0, 1.0, 1.0, 1.0)

(1.0, 1.0, 1.0, 1.0, 1.0)

(1.0, 1.0, 1.0, 1.0, 1.0)

(1.0, 1.0, 1.0, 1.0, 1.0)

(1.0, 1.0, 1.0, 1.0, 1.0)

(1.0, 1.0, 1.0, 1.0, 1.0)

(1.0, 1.0, 1.0, 1.0, 1.0)

(1.0, 1.0, 1.0, 1.0, 1.0)

(1.0, 1.0, 1.0, 1.0, 1.0)

(1.0, 1.0, 1.0, 1.0, 1.0)

(1.0, 1.0, 1.0, 1.0, 1.0)

(1.0, 1.0, 1.0, 1.0, 1.0)

(1.0, 1.0, 1.0, 1.0, 1.0)

(1.0, 1.0, 1.0, 1.0, 1.0)

(1.0, 1.0, 1.0, 1.0, 1.0)

Task 8

ANN using

Backpropagation

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.amax(X, axis=0)

Y = Y / 100

epoch = 5000

lr = 0.1

input_layer_neurons = 2

hidden_layer_neurons = 3

output_neurons = 1

wh = np.random.uniform(size=(input_layer,

hidden_layer_neurons))

bh = np.random.uniform(size=(1, hidden_layer,

neurons))

wout = np.random.uniform(size=(hidden_layer,

neurons, output_neurons))

bout = np.random.uniform(size=(1, output_neurons))


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

```
def deriv_sigmoid(x):  
    return x * (1 - x)
```

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

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

```
    h_inpt = h_inpt + bh
```

```
    hlayer_act = sigmoid(h_inpt)
```

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

```
    out_inpt = out_inpt + bout
```

```
    output = sigmoid(out_inpt)
```

```
    d_hidden_grad = deriv_sigmoid(hlayer_act)
```

```
    d_hidden_layer = E * h * d_hidden_grad
```

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

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

Input:

```
[[0.66667 1.0.33333 0.55556  
 1. 0.66667]]
```

AO:

```
[[92.] [86.] [89.]]
```

PO: [[0.84284073]

[0.83454433]

[0.8418144]]

Week 9a

Random Forest Algorithm

```
import pandas as pd
from sklearn.model_selection import
    train_test_split
from sklearn.ensemble import
    Random Forest Classifier.
```

```
iris = load_iris()
```

```
data = pd.DataFrame(data = iris.data,
    column = Iris.feature_name)
```

```
data['target'] = iris.target
```

```
X = data.drop('target', axis = 1)
```

```
Y = data['target']
```

```
X_train, X_test, Y_train, Y_test =
```

```
train_test_split(X, Y, test_size = 0.2,
```

```
random_state = 42)
```

```
rf_classifier = Random Forest Classifier()
```

```
rf_classifier.fit(X_train, Y_train)
```

```
Y_pred = rf_classifier.predict(X_test)
```

```
accuracy = accuracy_score(Y_test, Y_pred)
```

```
print('Accuracy: ', accuracy)
```

Accuracy = 1.0

Week 26.

ADA boost

```
import libraries
```

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

```
df = read_csv('iris.csv')
```

```
df.head()
```

```
X = df.drop('species', axis=1)
```

```
Y = df['species']
```

```
X_train, X_test, Y_train, Y_test =
```

```
train_test_split(X, Y, test_size=0.1
```

```
random_state=56)
```

```
X_train.shape, X_test.shape,
```

```
Y_train.shape, Y_test.shape
```

```
(135, 5), (15, 5), (135, 1), (15, 1)
```

```
my_logreg_model = LogisticRegression()
```

```
adaboost = AdaBoostClassifier(n_estimators=100,
```

```
base_estimator=my_logreg_model,
```

```
learning_rate=1)
```

```
model = adaboost.fit(X_train, Y_train)
```

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

```
print('Accuracy score', accuracy_score(Y_test,
```

```
Y_pred))
```

```
Accuracy = 1.0
```


K-Means

```
→ 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  
plt.xlabel  
plt.ylabel
```

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plt title ('K-Means clustering')
plt.xlabel ('Petal Length')
plt.ylabel ('Petal width')

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Sepal - Width
width']
target)

)]

'time'

petal.

ts], s=40)

- Width,
s=40)

11. Principal Analysis

```
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()
```

```
dict_keys(['DESCR', 'data',  
           'feature_names', 'target_names',  
           'target'])
```

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

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

```
df.head()
```

```
from sklearn.preprocessing import  
StandardScaler.
```

```
scaler = StandardScaler()
```

```
scaler.fit(df).
```

```
StandardScaler(copy=True, with_mean=True,  
                with_std=True)
```


scaled_data = scaler.transform(df)
from sklearn.decomposition import PCA
pca = PCA(n_components=2)
pca.fit(scaled_data)
pca(copy=True, n_components=2, whiten=False)

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')~~

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5/5/22

port

au cloth - mean