

Machine Learning Lab 1

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```
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
import math
import os
```

```
/tmp/ipykernel_9435/4209716058.py:2: DeprecationWarning:
Pyarrow will become a required dependency of pandas in the next major
release of pandas (pandas 3.0),
(to allow more performant data types, such as the Arrow string type,
and better interoperability with other libraries)
but was not found to be installed on your system.
If this would cause problems for you,
please provide us feedback at
https://github.com/pandas-dev/pandas/issues/54466
```

```
import pandas as pd

train_dirs=[]
test_dirs=[]
for dir in os.listdir("./"):
    if(dir.find("5-fold")!=-1):
        train_dirs.append("./"+dir+"/train/")
        test_dirs.append("./"+dir+"/test/")

def cal_header_val(file_path):
    with open(file_path, "r") as file:
        lines=file.readlines()
    return lines.index('@data\n')+1

headers=[]
for dir in train_dirs:
    file_path=dir+os.listdir (dir)[0]
    headers.append(cal_header_val(file_path))

headers=np.array(headers)
```

SVM

```
def Support_Vector_Regression(train_file, test_file, header):
    train_df = pd.read_csv(train_file, header=header, delimiter=",")
    test_df = pd.read_csv(test_file, header=header, delimiter=",")

    X_train = train_df.iloc[:, :-1].values
    y_train = train_df.iloc[:, -1].values
```

```

X_test = test_df.iloc[:, :-1].values
y_test = test_df.iloc[:, -1].values

from sklearn.svm import SVR
width=np.array([2**i for i in range(-18, 21, 2)])

best_mse, best_w=float('inf'), None

for w in width:
    regressor=SVR(kernel="rbf", gamma=w)
    regressor.fit(X_train, y_train)
    y_pred=regressor.predict(X_test)
    from sklearn.metrics import mean_squared_error
    mse=mean_squared_error(y_pred, y_test)
    if(mse<best_mse):
        best_mse, best_w=mse, w

return np.array([best_mse, best_w])

for train_dir, test_dir, header in zip(train_dirs, test_dirs,
headers):
    train_files=os.listdir(train_dir)
    test_files=os.listdir(test_dir)
    val=np.zeros(2)
    for train, test in zip(train_files, test_files):
        val+=Support_Vector_Regression(train_dir+train, test_dir+test,
header)
    print(train_dir)
    val/=len(train_files)
    val[0]=math.sqrt(val[0])
    val=pd.DataFrame(val, index=["RMSE", "Alpha"], columns=["Values"])
    print(val)
    print("-----\n")

```

/usr/lib/python3/dist-packages/scipy/__init__.py:146: UserWarning: A NumPy version >=1.17.3 and <1.25.0 is required for this version of SciPy (detected version 1.26.3

warnings.warn(f"A NumPy version >={np_minversion} and <{np_maxversion}")

```

./ele-1-5-fold/train/
      Values
RMSE    1139.563045
Alpha      0.000015
-----

```

```

./diabetes-5-fold/train/
      Values
RMSE      0.295963
Alpha     4.450000

```

```

-----
./quake-5-fold/train/
      Values
RMSE      0.136809
Alpha    4313.800000
-----

./laser-5-fold/train/
      Values
RMSE     23.374021
Alpha     0.000098
-----

./plastic-5-fold/train/
      Values
RMSE     1.490691
Alpha     0.010937
-----

```

SVM / MAE

```

def Support_Vector_Regression(train_file, test_file, header):
    train_df = pd.read_csv(train_file, header=header, delimiter=",")
    test_df = pd.read_csv(test_file, header=header, delimiter=",")

    X_train = train_df.iloc[:, :-1].values
    y_train = train_df.iloc[:, -1].values

    X_test = test_df.iloc[:, :-1].values
    y_test = test_df.iloc[:, -1].values

    from sklearn.svm import SVR
    width=np.array([2**i for i in range(-18, 21, 2)])

    best_mse, best_w=float('inf'), None

    for w in width:
        regressor=SVR(kernel="rbf", gamma=w)
        regressor.fit(X_train, y_train)
        y_pred=regressor.predict(X_test)
        from sklearn.metrics import mean_absolute_error
        mse=mean_absolute_error(y_pred, y_test)
        if(mse<best_mse):
            best_mse, best_w=mse, w

    return np.array([best_mse, best_w])

```

```

for train_dir, test_dir, header in zip(train_dirs, test_dirs,
headers):
    train_files=os.listdir(train_dir)
    test_files=os.listdir(test_dir)
    val=np.zeros(2)
    for train, test in zip(train_files, test_files):
        val+=Support_Vector_Regression(train_dir+train, test_dir+test,
header)
    print(train_dir)
    val/=len(train_files)
    val=pd.DataFrame(val, index=["MAE", "Alpha"], columns=["Values"])
    print(val)
    print("-\n")

```

```

./ele-1-5-fold/train/
      Values
MAE      792.005959
Alpha    0.000015
-----

```

```

./diabetes-5-fold/train/
      Values
MAE      0.183959
Alpha    4.450000
-----

```

```

./quake-5-fold/train/
      Values
MAE      0.103932
Alpha   1228.800002
-----

```

```

./laser-5-fold/train/
      Values
MAE     12.226322
Alpha   0.000244
-----

```

```

./plastic-5-fold/train/
      Values
MAE     1.117342
Alpha   0.071875
-----

```

SVM / MSE

```

def Support_Vector_Regression(train_file, test_file, header):
    train_df = pd.read_csv(train_file, header=header, delimiter=",")
    test_df = pd.read_csv(test_file, header=header, delimiter=",")

```

```

X_train = train_df.iloc[:, :-1].values
y_train = train_df.iloc[:, -1].values

X_test = test_df.iloc[:, :-1].values
y_test = test_df.iloc[:, -1].values

from sklearn.svm import SVR
width=np.array([2**i for i in range(-18, 21, 2)])

best_mse, best_w=-float('inf'), None

for w in width:
    regressor=SVR(kernel="rbf", gamma=w)
    regressor.fit(X_train, y_train)
    y_pred=regressor.predict(X_test)
    from sklearn.metrics import r2_score
    mse=r2_score(y_pred, y_test)
    if(mse>best_mse):
        best_mse, best_w=mse, w

    return np.array([best_mse, best_w])

for train_dir, test_dir, header in zip(train_dirs, test_dirs,
headers):
    train_files=os.listdir(train_dir)
    test_files=os.listdir(test_dir)
    val=np.zeros(2)
    for train, test in zip(train_files, test_files):
        val+=Support_Vector_Regression(train_dir+train, test_dir+test,
header)
        print(train_dir)
        val/=len(train_files)
        val=pd.DataFrame(val, index=["R2 score", "Alpha"],
columns=["Values"])
        print(val)
        print("-----\n")

./ele-1-5-fold/train/
          Values
R2 score -312.579759
Alpha      0.000015
-----

./diabetes-5-fold/train/
          Values
R2 score   0.755725
Alpha      55.450000
-----

```

```
./quake-5-fold/train/
```

```
Values
```

```
R2 score  -1.180579
```

```
Alpha      460.806250
```

```
-----
```

```
./laser-5-fold/train/
```

```
Values
```

```
R2 score  0.417075
```

```
Alpha      0.000098
```

```
-----
```

```
./plastic-5-fold/train/
```

```
Values
```

```
R2 score  0.782111
```

```
Alpha      0.015625
```

```
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```