Project 2 Breast Cancer Wisconsin Data Set

by Yuheng Lin for B455 Principles of Machine Learning

Import the Packages

```
import pandas as pd
import matplotlib.pyplot as plt
from sklearn.svm import SVC
from sklearn.model_selection import KFold
from sklearn.model_selection import train_test_split
from sklearn.model_selection import ShuffleSplit
from sklearn.neural_network import MLPClassifier
from sklearn.metrics import classification_report
from sklearn.model_selection import cross_val_score, learning_curve
```

Read Breast Cancer Wisconsin Data From UCI Learning

Check the Data

By using head()

patient data.head()

₽		ID	diagnosis	radius_mean	texture_mean	perimeter_mean	area- mean	${\tt smoothness}_{_}$
	0	842302	М	17.99	10.38	122.80	1001.0	0.1
	1	842517	М	20.57	17.77	132.90	1326.0	0.0
	2	84300903	М	19.69	21.25	130.00	1203.0	0.1
	3	84348301	М	11.42	20.38	77.58	386.1	0.1
	4	84358402	М	20.29	14.34	135.10	1297.0	0.1

5 rows × 32 columns

Check the Info of Our Data

```
print(patient_data.info())
```



```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 569 entries, 0 to 568
Data columns (total 32 columns):
                           569 non-null int64
diagnosis
                           569 non-null object
                           569 non-null float64
radius mean
                           569 non-null float64
texture mean
perimeter_mean
                           569 non-null float64
area-mean
                           569 non-null float64
                           569 non-null float64
smoothness mean
                           569 non-null float64
compactness mean
                           569 non-null float64
concavity_mean
                           569 non-null float64
concave points mean
symmetry mean
                           569 non-null float64
fractal dimension mean
                           569 non-null float64
radius se
                           569 non-null float64
                           569 non-null float64
texture_se
                           569 non-null float64
perimeter se
area-se
                           569 non-null float64
                           569 non-null float64
smoothness se
compactness se
                           569 non-null float64
                           569 non-null float64
concavity se
                           569 non-null float64
concave points se
symmetry_se
                           569 non-null float64
fractal_dimension_se
                           569 non-null float64
radius worst
                           569 non-null float64
                           569 non-null float64
texture worst
                           569 non-null float64
perimeter worst
area-worst
                           569 non-null float64
                           569 non-null float64
smoothness worst
                           569 non-null float64
compactness worst
                           569 non-null float64
concavity worst
                           569 non-null float64
concave_points_worst
                           569 non-null float64
symmetry_worst
fractal dimension worst
                           569 non-null float64
dtypes: float64(30), int64(1), object(1)
memory usage: 142.3+ KB
```

Set Our Inputs and Outputs

We drop the ID and diagnosis in our data as our inputs

Diagnosis will be our outputs

```
inputs = patient_data.drop('ID', axis=1).drop('diagnosis', axis=1)
outputs = patient_data['diagnosis']
```

Randomly Select Train and Test Inputs and Outputs

```
inputs_train, inputs_test, outputs_train, outputs_test = train_test_split(inputs, outputs)
```

Train the Model With 5 Hidden Layer With 2000 Max Iterations

```
BC_mlp = MLPClassifier(hidden_layer_sizes=(30, 30, 30, 30, 30), max_iter=2000) BC_mlp.fit(inputs_train, outputs_train)
```

Check the Accuracy of Our MLP by Random selections of Inputs and Outputs

```
mlp_prediction = BC_mlp.predict(inputs_test)
print(classification_report(outputs_test, mlp_prediction))
```

₽			precision	recal1	f1-score	support
		В	0.94	0.95	0.94	95
		M	0.89	0.88	0.88	48
	micro	avg	0.92	0.92	0.92	143
	macro	avg	0.92	0.91	0.91	143
	weighted	avg	0.92	0.92	0.92	143

Check the Accuracy of Our MLP by Using 5-Fold Cross Validation

```
five_fold_mlp_accuracy = cross_val_score(BC_mlp, inputs, outputs, cv=5)
print(five_fold_mlp_accuracy)
```

[0.93043478 0.93043478 0.9380531 0.92035398 0.91150442]

Conclusion for Model Parameters for Our MLP

In the mlp model, I used 5 hidden layers with 2000 iterations. The accuracy of the mlp model is good and around 0.92. And the learning curve of mlp model converfent. As result, using 5 hidden layers for the mlp model is valid and suitable.

Plot the Learning Curve of the Model

```
train_sizes = [1, 50, 115, 230, 335, 454]
def plot_learning_curve(estimator, title, inputs, outputs, train_sizes, cv):
  plt.figure()
  plt.title(title)
  train_sizes, train_error, validation_error = learning_curve(estimator, inputs, outputs, train_sizes, train_error_mean = train_error.mean(axis=1)
  validation_error_mean = validation_error.mean(axis=1)
  plt. xlabel('Training set size')
  plt. ylabel('Error')
  plt. plot(train_sizes, train_error_mean, color = 'b', label = 'Training error')
  plt. plot(train_sizes, validation_error_mean, color = 'r', label = 'Validation_error')
  plt. legend(loc="best")
  return plt

plot_learning_curve(BC_mlp, "Learning Curve of MLP Model", inputs, outputs, train_sizes, 5)
  plt. show()
```



Learning Curve Conclusion

From the graph above, we can see that the train and validation error both converge and they are around 0.92 to 0.95. And training error is higher than validation error.

Build the SVM for the Same Train and Test Data

Using SVC from the sklearn package

```
BC_svm = SVC(kernel='linear')
BC_svm.fit(inputs_train, outputs_train)
```

Check the Accuracy of Our SVM Model by Using Random Selections of Inputs

svm_prediction = BC_svm.predict(inputs_test)
print(classification_report(outputs_test, svm_prediction))

₽		precision	recall	f1-score	support
	В	0.95	0.98	0.96	95
	M	0.96	0.90	0.92	48
micro	avø	0.95	0.95	0.95	143
macro	_	0.95	0.94	0.94	143
weighted	avg	0.95	0.95	0.95	143

Check the Accuracy of Our SVM by Using 5-Fold Cross Validation

```
five_fold_svm_accuracy = cross_val_score(BC_svm, inputs, outputs, cv=5)
print(five_fold_svm_accuracy)
```

[0.94782609 0.93043478 0.97345133 0.92035398 0.95575221]

Conclusion

We can see that the average accuracy of mlp is around 0.92. And the average accuracy of svm is around 0.95. So, the accuracy of svm is higher. By running the code, svm takes a little longer to run than mlp.