

MLP Classifier

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1 Multi-Layer Perceptron Implementation

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```
[1]: import sklearn
import pandas as pd
import numpy as np
import seaborn as sns

from sklearn.model_selection import train_test_split
from sklearn.neural_network import MLPClassifier
from sklearn.metrics import accuracy_score, confusion_matrix, precision_score, \
    balanced_accuracy_score

from warnings import filterwarnings    #to ignore MLP warnings for no convergence
filterwarnings('ignore')
```

```
[2]: from sklearn.datasets import load_wine
```

```
[3]: data = load_wine()
columns = data.feature_names
targets = np.array(data.target)

data = pd.DataFrame(data.data, columns=data.feature_names)
targets = pd.DataFrame(targets, columns=['class'])
frames = [data, targets]
dataset = pd.concat(frames, axis = 1)
```

```
[4]: print(dataset.head())
```

	alcohol	malic_acid	ash	alcalinity_of_ash	magnesium	total_phenols	\
0	14.23	1.71	2.43	15.6	127.0	2.80	
1	13.20	1.78	2.14	11.2	100.0	2.65	
2	13.16	2.36	2.67	18.6	101.0	2.80	
3	14.37	1.95	2.50	16.8	113.0	3.85	
4	13.24	2.59	2.87	21.0	118.0	2.80	

	flavanoids	nonflavanoid_phenols	proanthocyanins	color_intensity	hue	\
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0	3.06	0.28	2.29	5.64	1.04
1	2.76	0.26	1.28	4.38	1.05
2	3.24	0.30	2.81	5.68	1.03
3	3.49	0.24	2.18	7.80	0.86
4	2.69	0.39	1.82	4.32	1.04

	od280/od315_of_diluted_wines	proline	class
0	3.92	1065.0	0
1	3.40	1050.0	0
2	3.17	1185.0	0
3	3.45	1480.0	0
4	2.93	735.0	0

```
[5]: print("Number of Samples:", len(dataset))
      print("\nNumber of Classes:")
      print(dataset['class'].value_counts())
```

Number of Samples: 178

Number of Classes:

```
1    71
0    59
2    48
```

Name: class, dtype: int64

```
[6]: pd.set_option('float_format', '{:.2f}'.format)
      dataset.describe()
```

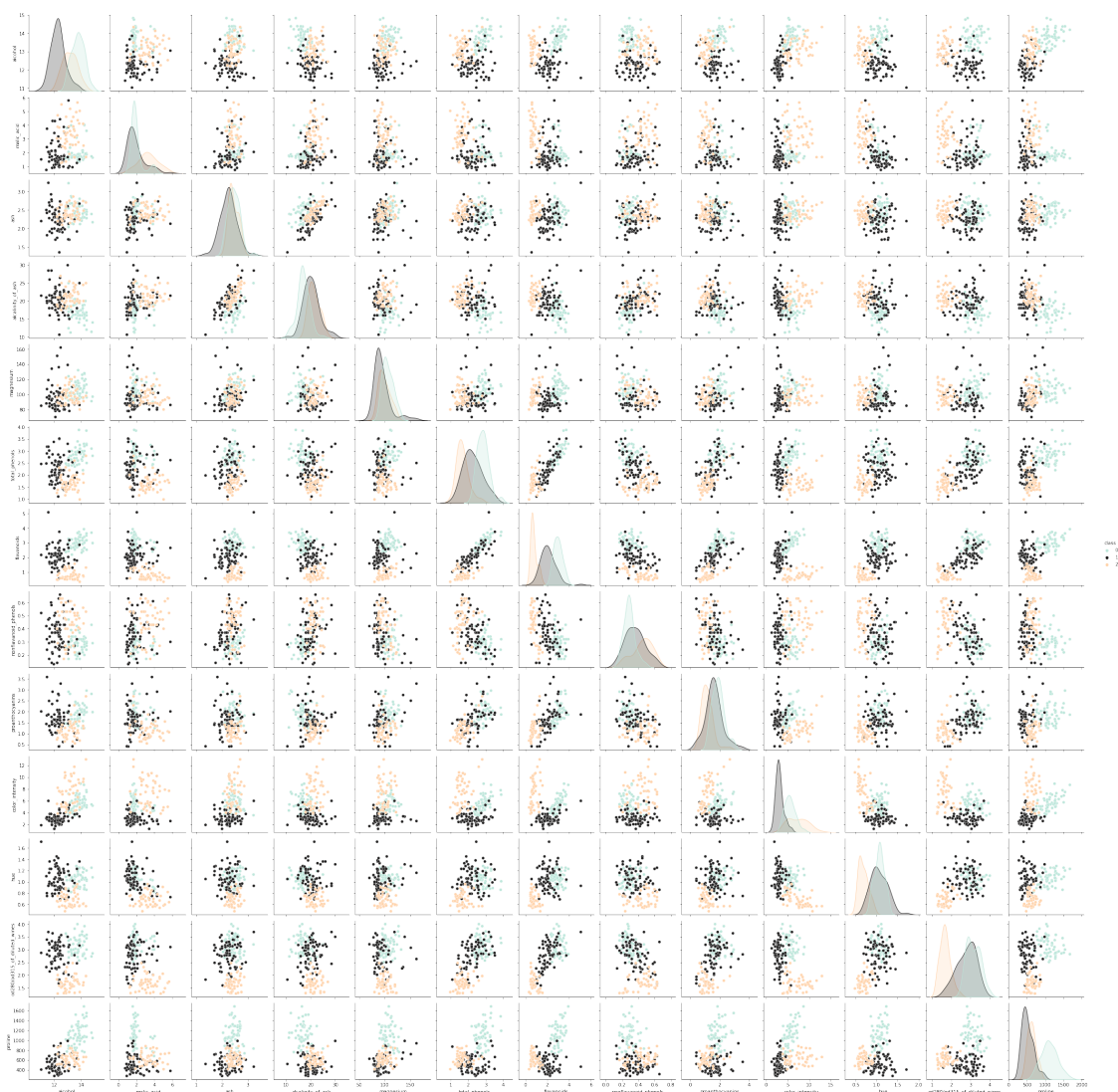
	alcohol	malic_acid	ash	alcalinity_of_ash	magnesium	\
count	178.00	178.00	178.00	178.00	178.00	
mean	13.00	2.34	2.37	19.49	99.74	
std	0.81	1.12	0.27	3.34	14.28	
min	11.03	0.74	1.36	10.60	70.00	
25%	12.36	1.60	2.21	17.20	88.00	
50%	13.05	1.87	2.36	19.50	98.00	
75%	13.68	3.08	2.56	21.50	107.00	
max	14.83	5.80	3.23	30.00	162.00	

	total_phenols	flavanoids	nonflavanoid_phenols	proanthocyanins	\
count	178.00	178.00	178.00	178.00	
mean	2.30	2.03	0.36	1.59	
std	0.63	1.00	0.12	0.57	
min	0.98	0.34	0.13	0.41	
25%	1.74	1.20	0.27	1.25	
50%	2.35	2.13	0.34	1.56	
75%	2.80	2.88	0.44	1.95	
max	3.88	5.08	0.66	3.58	

	color_intensity	hue	od280/od315_of_diluted_wines	proline	class
count	178.00	178.00	178.00	178.00	178.00
mean	5.06	0.96	2.61	746.89	0.94
std	2.32	0.23	0.71	314.91	0.78
min	1.28	0.48	1.27	278.00	0.00
25%	3.22	0.78	1.94	500.50	0.00
50%	4.69	0.96	2.78	673.50	1.00
75%	6.20	1.12	3.17	985.00	2.00
max	13.00	1.71	4.00	1680.00	2.00

```
[7]: sns.pairplot(dataset, hue='class', palette='icefire')
```

```
[7]: <seaborn.axisgrid.PairGrid at 0x7f9fd6fd80d0>
```



```
[8]: print("Columns:\n",columns)
```

Columns:

```
['alcohol', 'malic_acid', 'ash', 'alcalinity_of_ash', 'magnesium',  
'total_phenols', 'flavanoids', 'nonflavanoid_phenols', 'proanthocyanins',  
'color_intensity', 'hue', 'od280/od315_of_diluted_wines', 'proline']
```

```
[9]: x = np.array(dataset[['alcohol', 'flavanoids', 'ash']])  
y = np.array(dataset['class'])
```

```
[10]: x_train, x_test, y_train, y_test = train_test_split(x, y, test_size=0.3,  
→random_state=0)
```

```
[11]: mlp = MLPClassifier(activation='tanh', solver='sgd')  
mlp.fit(x_train, y_train)  
y_pred = mlp.predict(x_test)
```

```
[12]: print("Accuracy: {0} %".format(round(100 * float(accuracy_score(y_test,y_pred)),  
→2)))  
print("Precision: {0} %".format(round(100 * float(precision_score(y_test,  
→y_pred, average="weighted")), 2)))
```

Accuracy: 83.33 %

Precision: 84.56 %

```
[13]: confusion_matrix(y_test, y_pred)
```

```
[13]: array([[18,  1,  0],  
          [ 6, 15,  1],  
          [ 0,  1, 12]])
```

```
[14]: bas = round(balanced_accuracy_score(y_test, y_pred) * 100, 2)  
  
print("Balanced Accuracy Score:", bas, "%")
```

Balanced Accuracy Score: 85.08 %