

Linear Discriminant Analysis

February 23, 2022

1 Linear Discremenrt Analyss

```
[1]: import numpy as np
import pandas as pd
from sklearn import datasets
```

```
[2]: path_in = "/Users/saeedkhalili/Desktop/IntroML_Grant/iris.csv"
```

```
[3]: df = pd.read_csv(path_in)
```

```
[4]: df.head()
```

```
[4]:   sepal_length  sepal_width  petal_length  petal_width  species
0           5.1           3.5           1.4           0.2   setosa
1           4.9           3.0           1.4           0.2   setosa
2           4.7           3.2           1.3           0.2   setosa
3           4.6           3.1           1.5           0.2   setosa
4           5.0           3.6           1.4           0.2   setosa
```

```
[5]: df.tail()
```

```
[5]:   sepal_length  sepal_width  petal_length  petal_width  species
145           6.7           3.0           5.2           2.3  virginica
146           6.3           2.5           5.0           1.9  virginica
147           6.5           3.0           5.2           2.0  virginica
148           6.2           3.4           5.4           2.3  virginica
149           5.9           3.0           5.1           1.8  virginica
```

```
[6]: df.shape
```

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[6]: (150, 5)
```

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[7]: df.size
```

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[7]: 750
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```
[8]: df.describe()
```

```
[8]:      sepal_length  sepal_width  petal_length  petal_width
count      150.000000    150.000000    150.000000    150.000000
mean         5.843333      3.057333      3.758000      1.199333
std          0.828066      0.435866      1.765298      0.762238
min          4.300000      2.000000      1.000000      0.100000
25%          5.100000      2.800000      1.600000      0.300000
50%          5.800000      3.000000      4.350000      1.300000
75%          6.400000      3.300000      5.100000      1.800000
max          7.900000      4.400000      6.900000      2.500000
```

```
[9]: y = df['species']
```

```
[10]: y
```

```
[10]: 0      setosa
1      setosa
2      setosa
3      setosa
4      setosa
...
145    virginica
146    virginica
147    virginica
148    virginica
149    virginica
Name: species, Length: 150, dtype: object
```

```
[11]: from sklearn.preprocessing import LabelEncoder
y = LabelEncoder().fit_transform(y)
```

```
[12]: y
```

```
[12]: array([0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0,
0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0,
0, 0, 0, 0, 0, 0, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1,
1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1,
1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2,
2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2,
2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2])
```

```
[13]: X = df.drop(['species'], axis = 1)
```

```
[14]: from sklearn.discriminant_analysis import LinearDiscriminantAnalysis
lda = LinearDiscriminantAnalysis()
```

```
[15]: lda.fit(X, y)
```

```
[15]: LinearDiscriminantAnalysis()
```

```
[16]: yhat = lda.predict(X)
```

```
[17]: yhat
```

```
[17]: array([[0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0,
          0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0,
          0, 0, 0, 0, 0, 0, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1,
          1, 1, 1, 1, 2, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 2, 1, 1, 1, 1, 1,
          1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2,
          2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2,
          2, 1, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2])
```

```
[18]: from sklearn import metrics
```

```
[19]: metrics.accuracy_score(y, yhat)
```

```
[19]: 0.98
```

```
[20]: from sklearn.metrics import confusion_matrix
      confusion_matrix(y, yhat)
```

```
[20]: array([[50,  0,  0],
          [ 0, 48,  2],
          [ 0,  1, 49]])
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