

数据科学基础

作业报告



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任务 1: numpy 创建数组，数组形状修改结果截图

对应关键代码段：

```
import numpy as np

a = np.arange(9)

print('原始数组：')

print(a)

print('\n')

b = a.reshape(3,3)

print('修改后的数组：')


print(b)
```

运行结果截图：

```
C:\Users\徐豆豆\PycharmProjects\pythonProject\venv\Scripts\python.exe C:/Users/徐豆豆/PycharmProjects/pythonProject/main.py
原始数组：
[0 1 2 3 4 5 6 7 8]

修改后的数组：
[[0 1 2]
 [3 4 5]
 [6 7 8]]

Process finished with exit code 0
```



任务 2: 输出糖尿病数据集所有变量值及其数组形状

对应关键代码段：

```
import matplotlib.pyplot as plt

import numpy as np

from sklearn import datasets

from sklearn.datasets import load_diabetes

diabetes = datasets.load_diabetes()

print(diabetes.data)
```

运行结果截图：

```
[[ 0.03887591  0.05068012  0.06169621 ... -0.00259226  0.01990842
 -0.01764613]
 [-0.00188202 -0.04464164 -0.05147406 ... -0.03949338 -0.06832974
 -0.09220405]
 [ 0.08529891  0.05068012  0.04445121 ... -0.00259226  0.00286377
 -0.02593834]
 ...
 [ 0.04170844  0.05068012 -0.01590626 ... -0.01107952 -0.04687948
  0.01549073]
 [-0.04547248 -0.04464164  0.03906215 ...  0.02655962  0.04452837
 -0.02593834]
 [-0.04547248 -0.04464164 -0.0730303 ... -0.03949338 -0.00421986
  0.00386441]]

Process finished with exit code 0
```

任务 3：输出糖尿病数据所有样本真实标签及其数组形状

对应关键代码段：

```
import matplotlib.pyplot as plt

import numpy as np

from sklearn import datasets

from sklearn.datasets import load_diabetes

diabetes = datasets.load_diabetes()

print(diabetes.data)

print(diabetes.target)
```

运行结果截图：

```
45. 115. 264.  87. 202. 127. 182. 241.  66.  94. 203.  64. 102. 200.
265.  94. 230. 181. 156. 233.  60. 219.  80.  60. 332. 240.  84. 200.
55.  85.  89.  31. 129.  83. 275.  65. 198. 236. 253. 124.  44. 172.
114. 142. 109. 180. 144. 163. 147.  97. 220. 190. 109. 191. 122. 230.
242. 248. 249. 192. 131. 237.  78. 135. 244. 199. 270. 164.  72.  96.
306.  91. 214.  95. 216. 263. 178. 113. 200. 139. 139.  88. 148.  88.
243.  71.  77. 109. 272.  60.  54. 221.  90. 311. 281. 182. 321.  58.
262. 206. 233. 242. 123. 167.  63. 197.  71. 168. 140. 217. 121. 235.
245.  40.  52. 104. 132.  88.  69. 219.  72. 201. 110.  51. 277.  63.
118.  69. 273. 258.  43. 198. 242. 232. 175.  93. 168. 275. 293. 281.
72. 140. 189. 181. 209. 136. 261. 113. 131. 174. 257.  55.  84.  42.
146. 212. 233.  91. 111. 152. 120.  67. 310.  94. 183.  66. 173.  72.
49.  64.  48. 178. 104. 132. 220.  57.]

Process finished with exit code 0
```

任务 4：输出测试数据散点图（学号尾号为基数散点图为红色方形，学号尾号为偶数散点图为蓝色圆形）

对应关键代码段：

```
import matplotlib.pyplot as plt

import numpy as np

from sklearn import datasets
```

```
from sklearn.datasets import load_diabetes

diabetes = datasets.load_diabetes()

print(diabetes.data)

print(diabetes.target)

diabetes = datasets.load_diabetes()

diabetes_X = diabetes.data[:, 2]

diabetes_X_train = diabetes_X[:-20]

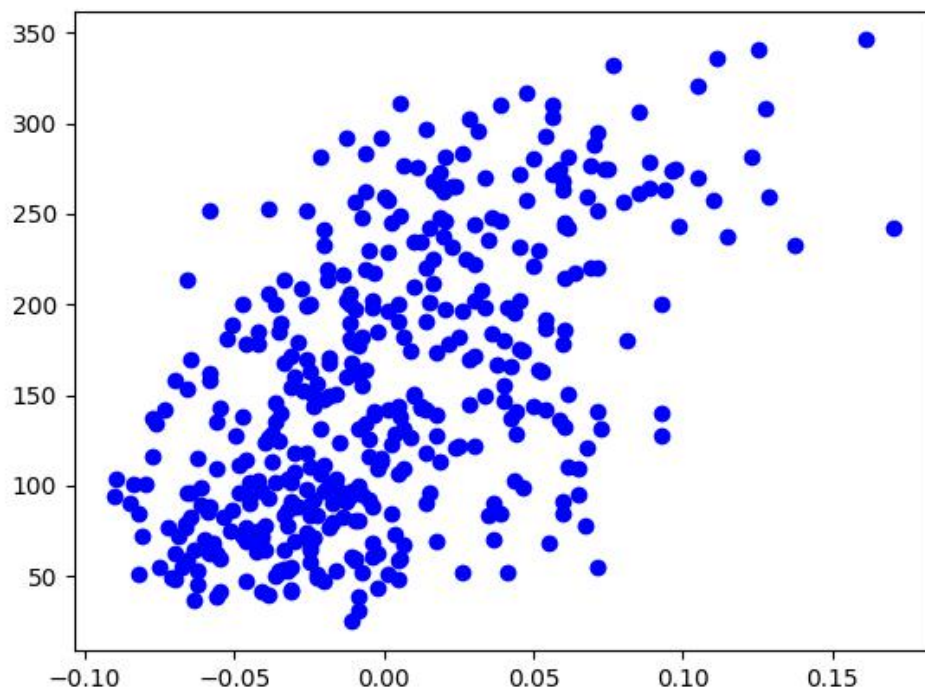
diabetes_y_train = diabetes.target[:-20]

plt.scatter(diabetes_X_train, diabetes_y_train,

color='blue', marker='o')

plt.show()
```

运行结果截图：



任务 5: `diabetes_X_train=np.array(diabetes_X_train).reshape(-1, 1)` 句的意义?

答: 训练数据集转换成 1 列, numpy 根据剩下的维度计算行数。

任务 6: 线性回归回归系数计算

对应关键代码段:

```
import matplotlib.pyplot as plt

import numpy as np

from sklearn.linear_model import LinearRegression

from sklearn import datasets, linear_model

from sklearn.datasets import load_diabetes

from sklearn.datasets import load_diabetes

from sklearn.metrics import mean_squared_error, r2_score

diabetes = datasets.load_diabetes()

diabetes_X = diabetes.data[:, 2]

diabetes_X_train = diabetes_X[:-20]

diabetes_X_test = diabetes_X[-20:]

diabetes_y_train = diabetes.target[:-20]

diabetes_y_test = diabetes.target[-20:]

regr = linear_model.LinearRegression()

regr.fit(np.array(diabetes_X_train).reshape(-1, 1),

np.array(diabetes_y_train).reshape(-1, 1))

diabetes_y_pred = regr.predict(np.array(diabetes_X_test).reshape(-1, 1))

print('Coefficients: ', regr.coef_)

print("Mean squared error: %.2f"
```

```
% mean_squared_error(diabetes_y_test,diabetes_y_pred))

print('Variance score: %.2f' % r2_score(diabetes_y_test,diabetes_y_pred))
```

运行结果截图：

```
C:\Users\徐豆豆\PycharmProjects\pythonProject\venv\Scripts\python.exe C:/Users/徐豆豆/PycharmProjects/pythonProject/main.py
Coefficients:  [[938.23786125]]
Mean squared error: 2548.07
Variance score: 0.47

Process finished with exit code 0
```



任务 7：线性回归的回归结果折线图及散点图展示

对应关键代码段：

```
import matplotlib.pyplot as plt

import numpy as np

from sklearn.linear_model import LinearRegression

from sklearn import datasets, linear_model

from sklearn.datasets import load_diabetes

from sklearn.datasets import load_diabetes

from sklearn.metrics import mean_squared_error, r2_score

diabetes = datasets.load_diabetes()

diabetes_X = diabetes.data[:, 2]

diabetes_X_train = diabetes_X[:-20]

diabetes_X_test = diabetes_X[-20:]

diabetes_y_train = diabetes.target[:-20]

diabetes_y_test = diabetes.target[-20:]

regr = linear_model.LinearRegression()
```

```

regr.fit(np.array(diabetes_X_train).reshape(-1, 1),
np.array(diabetes_y_train).reshape(-1, 1))

diabetes_y_pred = regr.predict(np.array(diabetes_X_test).reshape(-1, 1))

print('Coefficients: ', regr.coef_)

print("Mean squared error: %.2f"

      % mean_squared_error(diabetes_y_test, diabetes_y_pred))

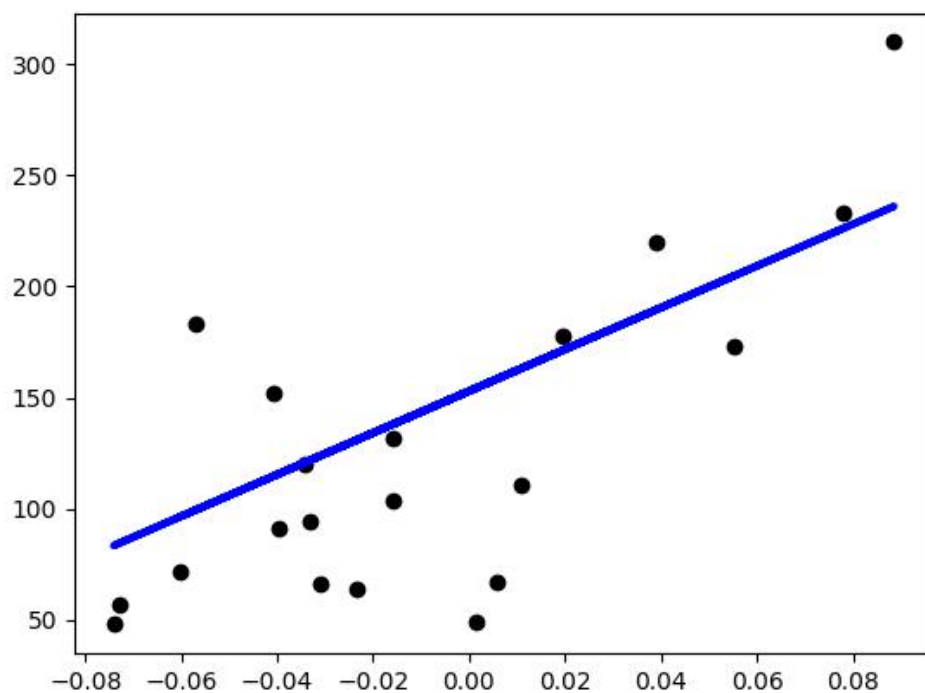
plt.scatter(diabetes_X_test, diabetes_y_test, color='black')

plt.plot(diabetes_X_test, diabetes_y_pred, color='blue', linewidth=3)

plt.show()

```

运行结果截图：



任务 8：逻辑回归回归系数计算

对应关键代码段：

```
import matplotlib.pyplot as plt

import numpy as np

from sklearn.datasets import load_iris

from sklearn.linear_model import LogisticRegression

iris = load_iris()

X = iris.data[:, :2]

Y = iris.target

lr = LogisticRegression(C=1e5)


lr.fit(X, Y)

print('Coefficients: ', lr.coef_)
```

运行结果截图：

```
C:\Users\徐豆豆\PycharmProjects\pythonProject\venv\Scripts\python.exe C:/Users/徐豆豆/PycharmProjects/pythonProject/main.py
Coefficients:  [[-36.45485824  30.74798948]
 [ 17.27627299 -15.57630379]
 [ 19.17858526 -15.17160568]]

Process finished with exit code 0
```



任务 9：逻辑回归回归散点图展示

对应关键代码段：

```
import matplotlib.pyplot as plt

import numpy as np

from sklearn.datasets import load_iris

from sklearn.linear_model import LogisticRegression

iris = load_iris()

X = iris.data[:, :2]
```



```

Y = iris.target

lr = LogisticRegression(C=1e5)

lr.fit(X, Y)

h = .02

x_min, x_max = X[:, 0].min() - .5, X[:, 0].max() + .5

y_min, y_max = X[:, 1].min() - .5, X[:, 1].max() + .5

xx, yy = np.meshgrid(np.arange(x_min, x_max, h),
np.arange(y_min, y_max, h))

Z = lr.predict(np.c_[xx.ravel(), yy.ravel()])

Z = Z.reshape(xx.shape)

plt.figure(1, figsize=(8, 6))

plt.pcolormesh(xx, yy, Z, cmap=plt.cm.Paired)

plt.scatter(X[:50, 0], X[:50, 1], color = 'red', marker =
'o', label = 'setosa')

plt.scatter(X[50:100, 0], X[50:100, 1], color = 'blue',
marker = 'x', label = 'versicolor')

plt.scatter(X[100:, 0], X[100:, 1], color = 'green', marker
= 's', label = 'Virginica')

plt.xlabel('Sepal length')

plt.ylabel('Sepal width')

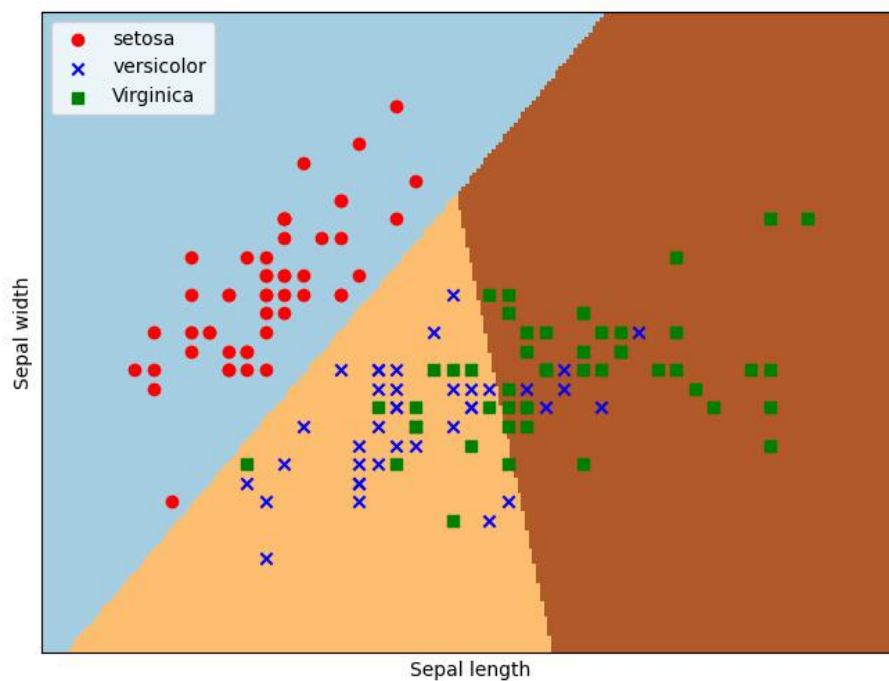
plt.xlim(xx.min(),xx.max())

plt.ylim(yy.min(),yy.max())

```

```
plt.xticks()  
  
plt.yticks()  
  
plt.legend(loc=2)  
  
plt.show()
```

运行结果截图：



任务 10：对鸢尾花数据进行 K-means 聚类，绘制聚类中心为 3 的聚类结果图

对应关键代码段：

```
import matplotlib.pyplot as plt  
  
import numpy as np  
  
from sklearn.cluster import KMeans  
  
from sklearn import datasets  
  
iris = datasets.load_iris()  
  
X = iris.data[:, :4]
```

```
print(X.shape)

plt.scatter(X[:, 0], X[:, 1], c="red", marker='o',
            label='see')

plt.xlabel('sepal length')
plt.ylabel('sepal width')

plt.legend(loc=2)

plt.show()

estimator = KMeans(n_clusters=3)

estimator.fit(X)

label_pred = estimator.labels_

x0 = X[label_pred == 0]
x1 = X[label_pred == 1]
x2 = X[label_pred == 2]

plt.scatter(x0[:, 0], x0[:, 1], c="red", marker='o',
            label='label0')

plt.scatter(x1[:, 0], x1[:, 1], c="green", marker='*',
            label='label1')

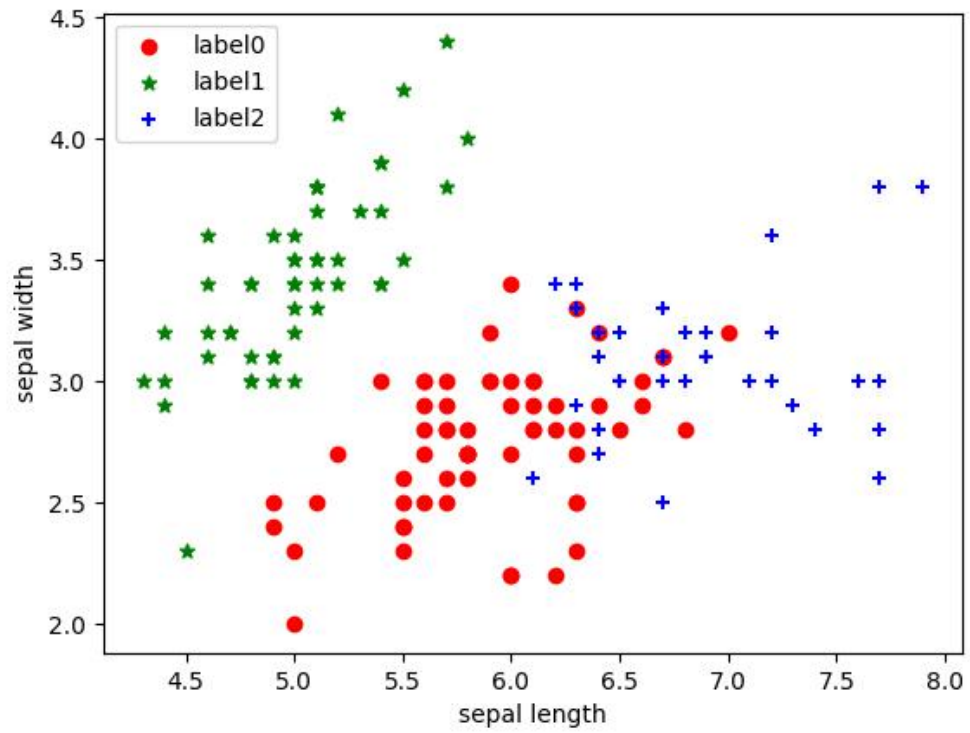
plt.scatter(x2[:, 0], x2[:, 1], c="blue", marker='+',
            label='label2')

plt.xlabel('sepal length')
plt.ylabel('sepal width')
```

```
plt.legend(loc=2)
```

```
plt.show()
```

运行结果截图：



附加题：

中国男足在亚洲水平的聚类实验

关键代码段：

```
from sklearn.cluster import KMeans

from sklearn import preprocessing

import pandas as pd

import numpy as np

# 输入数据

data = pd.read_csv('data.csv', encoding='gbk')

train_x = data[["2006 年世界杯", "2010 年世界杯 ", "2007 年亚洲杯 "]]

df = pd.DataFrame(train_x)

kmeans = KMeans(n_clusters=3)

# 规范化到 [0,1] 空间

min_max_scaler=preprocessing.MinMaxScaler()

train_x=min_max_scaler.fit_transform(train_x)

# kmeans 算法

kmeans.fit(train_x)

predict_y = kmeans.predict(train_x)

# 合并聚类结果，插入到原数据中

result = pd.concat((data,pd.DataFrame(predict_y)),axis=1)

result.rename({0:'聚类'},axis=1,inplace=True)

print(result)
```

组别	聚类	中心点
1	日本，韩国，伊朗，沙特	(0.21, 0.41, 0.16)
2	乌兹别克斯坦，巴林，朝鲜	(0.7, 0.7333, 0.4167)
3	中国，伊拉克，卡塔尔，阿联酋，泰国，越南，阿曼，印尼	(1, 0.94, 0.40625)

按照中心点位置（数值越小，排名越前），可以将聚类结果划分为三档：

亚洲一流：日本，韩国，伊朗，沙特

亚洲二流：乌兹别克斯坦，巴林，朝鲜

亚洲三流：中国，伊拉克，卡塔尔，阿联酋，泰国，越南，阿曼，印尼