

```
In [1]: import numpy as np
import matplotlib.pyplot as plt
from sklearn import datasets, linear_model
```

```
In [2]: diabetes = datasets.load_diabetes()    # dataset loaded from sklearn library.
diabetes
```

```
Out[2]: {'data': array([[ 0.03807591,  0.05068012,  0.06169621, ..., -0.00259226,
    0.01990749, -0.01764613],
 [-0.00188202, -0.04464164, -0.05147406, ..., -0.03949338,
   -0.06833155, -0.09220405],
 [ 0.08529891,  0.05068012,  0.04445121, ..., -0.00259226,
   0.00286131, -0.02593034],
 ...,
 [ 0.04170844,  0.05068012, -0.01590626, ..., -0.01107952,
  -0.04688253,  0.01549073],
 [-0.04547248, -0.04464164,  0.03906215, ...,  0.02655962,
   0.04452873, -0.02593034],
 [-0.04547248, -0.04464164, -0.0730303 , ..., -0.03949338,
  -0.00422151,  0.00306441]]),
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  145.,  74.,  45., 115., 264.,  87., 202., 127., 182., 241.,  66.,
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114., 142., 109., 180., 144., 163., 147., 97., 220., 190., 109.,
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263., 178., 113., 200., 139., 139., 88., 148., 88., 243., 71.,
77., 109., 272., 60., 54., 221., 90., 311., 281., 182., 321.,
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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.]),
'frame': None,
'DESCR': '.. _diabetes_dataset:\n\nDiabetes dataset\n-----\n\nTen baseline variables, age, sex, body mass index, average blood\npressur
e, and six blood serum measurements were obtained for each of n =\n442 diabetes patients, as well as the response of interest, a\nquantitative measu
re of disease progression one year after baseline.\n\n**Data Set Characteristics:**\n\n :Number of Instances: 442\n\n :Number of Attributes: First
10 columns are numeric predictive values\n\n :Target: Column 11 is a quantitative measure of disease progression one year after baseline\n\n :Attr
ibute Information:\n      - age      age in years\n      - sex\n      - bmi      body mass index\n      - bp      average blood pressure\n      - s1
tc, total serum cholesterol\n      - s2      ldl, low-density lipoproteins\n      - s3      hdl, high-density lipoproteins\n      - s4      tch, tot
al cholesterol / HDL\n      - s5      ltcg, possibly log of serum triglycerides level\n      - s6      glu, blood sugar level\n\nNote: Each of these
10 feature variables have been mean centered and scaled by the standard deviation times the square root of `n_samples` (i.e. the sum of squares of e
ach column totals 1).\n\nSource URL:\nhttps://www4.stat.ncsu.edu/~boos/var.select/diabetes.html\n\nFor more information see:\nBradley Efron, Trevor
Hastie, Iain Johnstone and Robert Tibshirani (2004) "Least Angle Regression," Annals of Statistics (with discussion), 407-499.\n(https://web.stanfor
d.edu/~hastie/Papers/LARS/LeastAngle_2002.pdf)\n',
'feature_names': ['age',
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'bmi',
'bp',
's1',
's2',
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's4',
's5',
's6'],
'data_filename': 'diabetes_data_raw.csv.gz',
'target_filename': 'diabetes_target.csv.gz',
'data_module': 'sklearn.datasets.data'}

```

In [3]: diabetes.target

Out[3]: array([151., 75., 141., 206., 135., 97., 138., 63., 110., 310., 101.,  
69., 179., 185., 118., 171., 166., 144., 97., 168., 68., 49.,  
68., 245., 184., 202., 137., 85., 131., 283., 129., 59., 341.,  
87., 65., 102., 265., 276., 252., 90., 100., 55., 61., 92.,  
259., 53., 190., 142., 75., 142., 155., 225., 59., 104., 182.,  
128., 52., 37., 170., 170., 61., 144., 52., 128., 71., 163.,  
150., 97., 160., 178., 48., 270., 202., 111., 85., 42., 170.,  
200., 252., 113., 143., 51., 52., 210., 65., 141., 55., 134.,  
42., 111., 98., 164., 48., 96., 90., 162., 150., 279., 92.,  
83., 128., 102., 302., 198., 95., 53., 134., 144., 232., 81.,  
104., 59., 246., 297., 258., 229., 275., 281., 179., 200., 200.,  
173., 180., 84., 121., 161., 99., 109., 115., 268., 274., 158.,  
107., 83., 103., 272., 85., 280., 336., 281., 118., 317., 235.,  
60., 174., 259., 178., 128., 96., 126., 288., 88., 292., 71.,  
197., 186., 25., 84., 96., 195., 53., 217., 172., 131., 214.,  
59., 70., 220., 268., 152., 47., 74., 295., 101., 151., 127.,  
237., 225., 81., 151., 107., 64., 138., 185., 265., 101., 137.,  
143., 141., 79., 292., 178., 91., 116., 86., 122., 72., 129.,  
142., 90., 158., 39., 196., 222., 277., 99., 196., 202., 155.,  
77., 191., 70., 73., 49., 65., 263., 248., 296., 214., 185.,  
78., 93., 252., 150., 77., 208., 77., 108., 160., 53., 220.,  
154., 259., 90., 246., 124., 67., 72., 257., 262., 275., 177.,  
71., 47., 187., 125., 78., 51., 258., 215., 303., 243., 91.,  
150., 310., 153., 346., 63., 89., 50., 39., 103., 308., 116.,  
145., 74., 45., 115., 264., 87., 202., 127., 182., 241., 66.,  
94., 283., 64., 102., 200., 265., 94., 230., 181., 156., 233.,  
60., 219., 80., 68., 332., 248., 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.]

In [4]: diabetes.DESCR

Out[4]: '.. \_diabetes\_dataset:\n\nDiabetes dataset\n-----\n\nTen baseline variables, age, sex, body mass index, average blood\npressure, and six blood serum measurements were obtained for each of n =\n442 diabetes patients, as well as the response of interest, a\nquantitative measure of disease progression one year after baseline.\n\n\*\*Data Set Characteristics:\*\*\n\n :Number of Instances: 442\n\n :Number of Attributes: First 10 columns are numeric predictive values\n\n :Target: Column 11 is a quantitative measure of disease progression one year after baseline\n\n :Attribute Information:\n\n - age age in years\n\n - sex\n\n - bmi body mass index\n\n - bp average blood pressure\n\n - s1 tc, total serum cholesterol\n\n - s2 ldl, low-density lipoproteins\n\n - s3 hdl, high-density lipoproteins\n\n - s4 tch, total cholesterol / HDL\n\n - s5 ltg, possibly log of serum triglycerides level\n\n - s6 glu, blood sugar level\n\nNote: Each of these 10 feature variables have been mean centered and scaled by the standard deviation times the square root of `n\_samples` (i.e. the sum of squares of each column totals 1).\n\nSource URL:\nhttps://www4.stat.ncsu.edu/~boos/var.select/diabetes.html\n\nFor more information see:\nBradley Efron, Trevor Hastie, Iain Johnstone and Robert Tibshirani (2004) "Least Angle Regression," Annals of Statistics (with discussion), 407-499.\n(https://web.stanford.edu/~hastie/Papers/LARS/LeastAngle\_2002.pdf)\n'

In [5]: diabetes.data

Out[5]: array([[ 0.03807591, 0.05068012, 0.06169621, ..., -0.00259226,  
 0.01990749, -0.01764613],  
 [-0.00188202, -0.04464164, -0.05147406, ..., -0.03949338,  
 -0.06833155, -0.09220405],  
 [ 0.08529891, 0.05068012, 0.04445121, ..., -0.00259226,  
 0.00286131, -0.02593034],  
 ...,  
 [ 0.04170844, 0.05068012, -0.01590626, ..., -0.01107952,  
 -0.04688253, 0.01549073],  
 [-0.04547248, -0.04464164, 0.03906215, ..., 0.02655962,  
 0.04452873, -0.02593034],  
 [-0.04547248, -0.04464164, -0.0730303 , ..., -0.03949338,  
 -0.00422151, 0.00306441]])

```
In [6]: diabetes_x = diabetes.data[:, np.newaxis, 2] # if we remove the list from this expression then mean squared error changes.  
print(diabetes_x) # and plots cannot be drawn.
```

```
[-0.04285156]  
[ 0.00564998]  
[-0.03530688]  
[ 0.02397278]  
[-0.01806189]  
[ 0.04229559]  
[-0.0547075 ]  
[-0.00297252]  
[-0.06656343]  
[-0.01267283]  
[-0.04177375]  
[-0.03099563]  
[-0.00512814]  
[-0.05901875]  
[ 0.0250506 ]  
[-0.046085 ]  
[ 0.00349435]  
[ 0.05415152]  
[-0.04500719]  
[-0.05791093]
```

```
In [7]: diabetes_x_train = diabetes_x[:-30]  
diabetes_x_test = diabetes_x[-30:]  
diabetes_y_train = diabetes.target[:-30]  
diabetes_y_test = diabetes.target[-30:]
```

```
In [13]: model = linear_model.LinearRegression()  
model.fit(diabetes_x_train, diabetes_y_train)  
diabetes_y_predicted = model.predict(diabetes_x_test)
```

```
In [14]: from sklearn.metrics import mean_squared_error
```

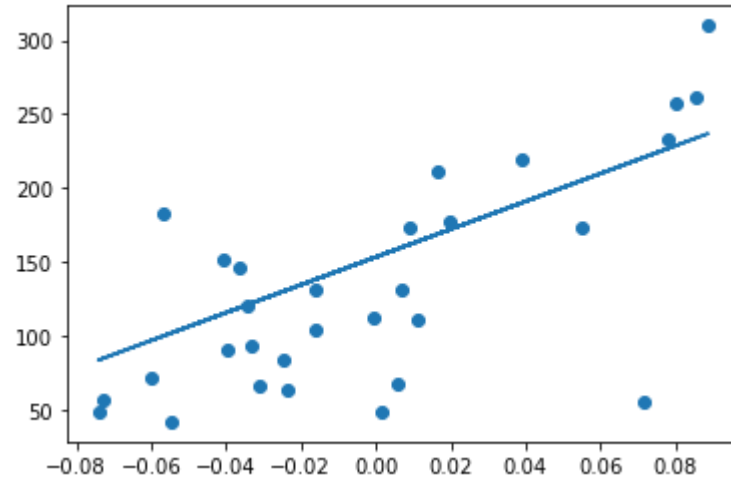
```
In [15]: print("Mean Squared Error:", mean_squared_error(diabetes_y_test, diabetes_y_predicted))
```

Mean Squared Error: 3035.0601152912695

```
In [16]: print("weights: ", model.coef_)  
print('intercept: ', model.intercept_)
```

```
weights: [941.43097333]  
intercept: 153.39713623331644
```

```
In [19]: plt.scatter(diabetes_x_test, diabetes_y_test)  
plt.plot(diabetes_x_test, diabetes_y_predicted)  
plt.show()
```



```
In [ ]:
```