Curve Fitting

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- 1 import pandas as pd
- 2 import numpy as np
- 3 import matplotlib.pyplot as plt
- 4 import seaborn as sns
- 5 from sklearn.model_selection import train_test_split
- 6 from sklearn import datasets
- 7 from sklearn.linear_model import LinearRegression
- 8 from sklearn import metrics
- 1 # Data Preparation
- 2 # importing data set from sklearn diabetes
- 3 db_data = datasets.load_diabetes()
- 1 diabetes_data = pd.DataFrame(db_data.data, columns=db_data.feature_names)
- 2 diabetes_data.describe()

	age	sex	bmi	bp	s1	s2
count	4.420000e+02	4.420000e+02	4.420000e+02	4.420000e+02	4.420000e+02	4.420000e+02
mean	-3.634285e- 16	1.308343e-16	-8.045349e- 16	1.281655e-16	-8.835316e- 17	1.327024e-16
std	4.761905e-02	4.761905e-02	4.761905e-02	4.761905e-02	4.761905e-02	4.761905e-02
min	-1.072256e- 01	-4.464164e- 02	-9.027530e- 02	-1.123996e- 01	-1.267807e- 01	-1.156131e- 01
25%	-3.729927e- 02	-4.464164e- 02	-3.422907e- 02	-3.665645e- 02	-3.424784e- 02	-3.035840e- 02
50%	5.383060e-03	-4.464164e- 02	-7.283766e- 03	-5.670611e- 03	-4.320866e- 03	-3.819065e- 03

- 1 # checking its rows and columns
- 2 diabetes_data.shape

(442, 10)

- 1 # target refers to y-axis. setting target as the data of disease progression
- 2 diabetes_data['disease progression'] = db_data.target
- 3 diabetes_data['disease progression'].head()
 - 0 151.0
 - 1 75.0
 - 2 141.0
 - 3 206.0
 - 4 135.0

Name: disease progression, dtype: float64

- 1 #cheking the updated rows and columns of the data
- 2 diabetes_data.shape

(442, 11)

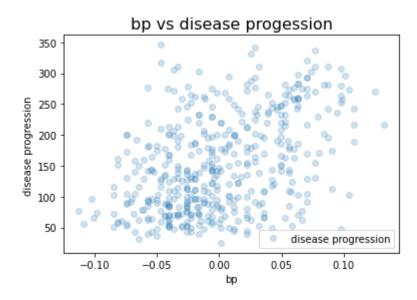
Visualization of percentage related on the following bp = average blood pressure

```
s4= total cholesterol / HDL (tch)
```

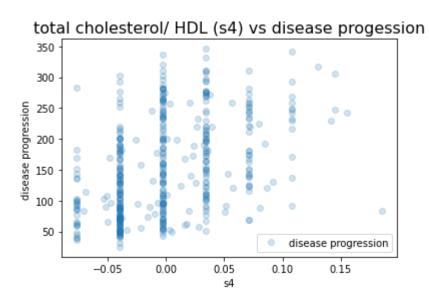
s6 = blood sugar level

which measure of disease progression one year after baseline [1]

```
1 # Visualizing the average blood pressure and disease progression
2 diabetes_data.plot(x='bp', y='disease progression', style='o', alpha=0.2)
3 plt.title('bp vs disease progession', fontsize=16)
4 plt.xlabel('bp')
5 plt.ylabel('disease progression ')
6 plt.show()
```

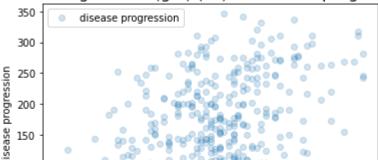


```
1 diabetes_data.plot(x='s4', y='disease progression', style='o', alpha=0.2)
2 plt.title('total cholesterol/ HDL (s4) vs disease progession', fontsize=16)
3 plt.xlabel('s4')
4 plt.ylabel('disease progression ')
5 plt.show()
```



```
1 diabetes_data.plot(x='s6', y='disease progression', style='o', alpha=0.2)
2 plt.title('blood sugar level (glu) (s6) vs disease progession', fontsize=16)
3 plt.xlabel('s6')
4 plt.ylabel('disease progression ')
5 plt.show()
```

blood sugar level (glu) (s6) vs disease progession



Multiple Linear Regression

```
-0.15
                -0.10
                       -0.05
                               0 00
                                      0.05
                                              0 10
1 # Multiple Regreesion
2 X = pd.DataFrame(np.c_[diabetes_data['s4'], diabetes_data['bp'], diabetes_data['s6']], col
3 y = diabetes_data['disease progression']
4 X_train, X_test, y_train, y_test = train_test_split(X, y, test_size = 0.2, random_state=1)
1 # Fitting a Multiple Linear Model
2 l_reg= LinearRegression()
3 l_reg.fit(X_train, y_train)
    LinearRegression(copy_X=True, fit_intercept=True, n_jobs=None, normalize=False)
1 # Summary of modelling
2 l_reg_sum = pd.DataFrame(X.columns, columns=['Features'])
3 l_reg_sum['Weights Raw'] = l_reg.coef_.reshape(3,1)
4 l_reg_sum = l_reg_sum.append({'Features':'Intercept', 'Weights Raw':float(l_reg.intercept_
5 l_reg_sum
```

	Features	Weights Raw
0	s4	476.371697
1	bp	520.717573
2	s6	223.540217
3	Intercept	152.003079

```
1 # prediction model
2 preds = l_reg.predict(X_test)
3 res = pd.DataFrame({'Actual': y_test, 'Predicted': preds})
4 res
```

Actual Predicted

```
1 plt.figure(figsize=(10,10))
2 plt.title('Predictions', fontsize=20)
3 plt.scatter(y_test, preds, s = 150, alpha=0.5)
4 plt.xlabel('Ground Truth', fontsize=20)
5 plt.ylabel('Prediction', fontsize=20)
6 plt.show()
```

Predictions 250 225 200 Prediction 120 120 125 100 75 50 150 200 250 350 100 300 Ground Truth

```
1 # Metrics
2 # adjusted r squared as per lecture
4 def adjr2(r2,x):
5
     n = x.shape[0]
6
     p = x.shape[1]
7
     adjusted_r2 = 1-(1-r2)*(n-1)/(n-p-1)
8
     return adjusted_r2
1 MSE = metrics.mean_squared_error(y_test, preds)
2 RMSE = np.sqrt(MSE)
3 R2 = metrics.r2_score(y_test, preds)
4 AR2 = adjr2(R2,X_train)
5 model_metrics = pd.DataFrame([['MSE'],['RMSE'],['R^2'],
6
                                ['Adjusted R^2']],
7
                               columns=['Metrics'])
8 model_metrics['Multiple Regression'] = MSE, RMSE, R2, AR2
9 model_metrics
```

	Metrics	Multiple Regression
0	MSE	3834.449487
1	RMSE	61.922932

▼ Polynomial Regression

```
1 # POLYNOMIAL REGRESSION
2 from sklearn.preprocessing import PolynomialFeatures
3 X = pd.DataFrame(np.c_[diabetes_data['s4'], diabetes_data['bp'], diabetes_data['s6'],diabe
4 y = diabetes_data['disease progression']
5 X_train, X_test, y_train, y_test = train_test_split(X, y, test_size = 0.2, random_state=1)
1 poly_reg = PolynomialFeatures(degree=2)
2 X_poly = poly_reg.fit_transform(X_train)
1 print(X_poly)
    [[ 1.00000000e+00 3.43088589e-02 -6.76422830e-02 ... 1.97895780e-03
      [ 1.00000000e+00 -2.59226200e-03 4.25295792e-02 ... 2.72463581e-05
     -3.72467191e-04 5.09175604e-03]
    [ 1.00000000e+00 -3.94933829e-02 -2.28849640e-02 ... 1.17063562e-03
      2.17217379e-03 4.03057866e-03]
    [ 1.00000000e+00 1.29062088e-02 5.85963092e-02 ... 2.36464207e-03
      -2.88648387e-03 3.52348849e-03]
    [ 1.00000000e+00 -3.94933829e-02 -5.73136710e-02 ... 2.72463581e-05
      1.30287145e-04 6.23009508e-04]]
1 # Fitting a Multiple Linear Model
2 l_reg2 = LinearRegression()
3 l_reg2.fit(X_poly, y_train)
   LinearRegression(copy_X=True, fit_intercept=True, n_jobs=None, normalize=False)
1 preds = l_reg2.predict(poly_reg.fit_transform(X_test))
1 poly_MSE = metrics.mean_squared_error(y_test,preds)
2 poly_RMSE = np.sqrt(poly_MSE)
3 poly_R2 = metrics.r2_score(y_test,preds)
4 poly_AR2 = adjr2(poly_R2,X_train)
5 model_metrics = pd.DataFrame([['MSE'],['RMSE'],['R^2'],
                              ['Adjusted R^2']],
6
                             columns=['Metrics'])
8 model_metrics['Polynomial Regression'] = poly_MSE, poly_RMSE, poly_R2, poly_AR2
9 model_metrics
```

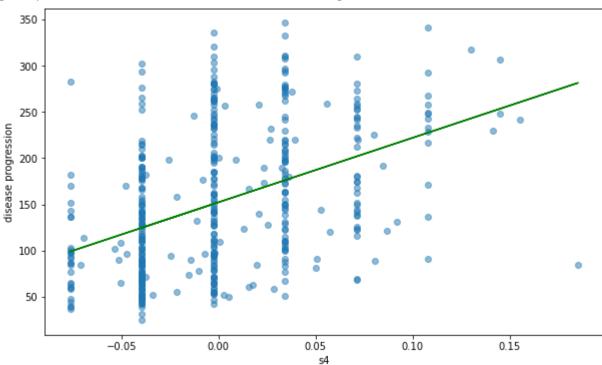
Metrics Polynomial Regression

0	MSE	4219.197941
1	RMSE	64.955353
2	R^2	0.208253
3	Adjusted R^2	0.199152

Normal Equation

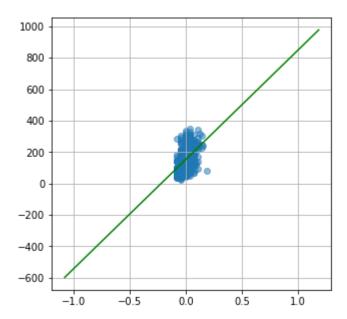
```
1 X = diabetes data['s4'].values.reshape(-1,1)
2 y = diabetes_data['disease progression'].values.reshape(-1,1)
3
4 n = len(diabetes_data['s4']) #no. rows
5 bias = np.ones((n,1)) #column-1 of Matrix X
6 xnew = np.reshape(X,(n,1)) #reshaping the data
7 xnew1 =np.append(bias,xnew,axis=1) #forming Matrix X
8 xnew1_t = np.transpose(xnew1) #transpose
9 xnew_t_dot_x_new = xnew1_t.dot(xnew1) #matrix multiplication
10 temp_1 = np.linalg.inv(xnew_t_dot_x_new) #inverse of a matrix
11 \text{ temp}_2 = \text{xnew1}_t.dot(y)
1 #Finding coefficients:
3 theta = temp_1.dot(temp_2)
4 Intercept = theta[0]
5 Slope = theta[1]
6 print("Intercept:",Intercept)
7 print("Slope:",Slope)
     Intercept: [152.13348416]
     Slope: [696.88303009]
1 #Predicting values:
2 def predict_value(input_feature,slope,intercept):
       return slope*input_feature+intercept
1 s4 = 4
2 prediction =predict_value(s4,Slope,Intercept)
3 print(prediction)
     [2939.66560453]
1 #Plotting the regression Line:
2 plt.figure(figsize=(5*2,3*2))
3 plt.scatter(X,y, alpha=0.5)
4 plt.xlabel('s4')
5 plt.ylabel('disease progression')
6 plt.plot(X,Slope*X+Intercept, color="green")
```

[<matplotlib.lines.Line2D at 0x7f1ed5b34710>]



▼ Polynomial Curve

```
1 def linear_regressor(X,y):
   X = np.array(X)
   y = np.array(y)
 4
    n = X.size
    w0 = (y.mean()*np.sum(X**2)-X.mean()*np.sum(X*y)) / (np.sum(X**2) - n*X.mean()**2)
  w1 = (np.sum(X*y) - X.mean()*np.sum(y)) / (np.sum(X**2) - n*X.mean()**2)
    return w0,w1
 8 w0,w1 = linear_regressor(X,y)
 9 print("Linear Regression Equation: y = {:.3f}x + {:.3f}".format(w1, w0))
     Linear Regression Equation: y = 696.883x + 152.133
 1 def show_regline(X,y,w1,w0):
   x_{min}, x_{max} = X.min() - 1, X.max() + 1
 3
   linex = np.linspace(x_min, x_max)
   liney = w1*linex+w0
 5 plt.figure(figsize=(5,5))
  plt.grid()
 7
    plt.scatter(X,y, alpha=0.5)
    plt.plot(linex, liney, c='green')
    plt.show()
10 show_regline(X,y,w1,w0)
```



```
1 def lin_reg(val,w0,w1):
2    return w1*val + w0 #model
3 print(lin_reg(10, w0, w1))
4 X_new, y_new = X.copy(), y.copy()
5 for i in range(10,16):
6    X_new = np.insert(X_new,-1, i)
7    y_new = np.insert(y_new,-1, lin_reg(i,w0,w1))
8 show_regline(X_new, y_new, w1, w0)
```

```
7120.963785085329
```

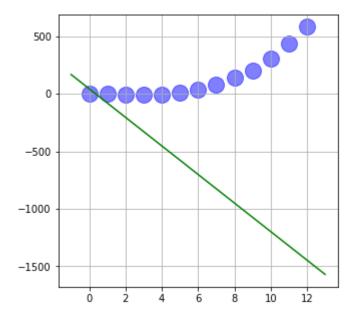
```
10000
     8000
1 X_1 = np.arange(0, 13,1)
2 y_1 = X_1 - 2 * (X_1 ** 2) + 0.5 * (X_1 ** 3) + np.random.normal(-3, 3, 13)
1 def show_regline(X,y,w1,w0):
   x_{min}, x_{max} = X.min() - 1, X.max() + 1
3
   linex = np.linspace(x_min, x_max)
  liney = w1*linex+w0
4
   plt.figure(figsize=(5,5))
5
6
   plt.grid()
7
   plt.scatter(X_1,y_1, s = 256, color='blue', alpha=0.5)
8 plt.plot(linex, liney, c='green')
   plt.show()
1 def linear_regressor(X,y):
  X = np.array(X)
3
  y = np.array(y)
  n = X.size
   w0 = (y.mean()*np.sum(X**2)-X.mean()*np.sum(X*y)) / (np.sum(X**2) - n*X.mean()**2)
```

w1 = (np.sum(X*y) - X.mean()*np.sum(y)) / (np.sum(X**2) - n*X.mean()**2)

```
1 w0_q,w1_q = linear_regressor(X_1, y_1)
2 show_regline(X_1,y_1,w0_q,w1_q)
```

8 w0,w1 = linear_regressor(X,y)

return w0,w1



1

References:

[1] "7.1. Toy datasets — scikit-learn 0.24.2 documentation." https://scikit-learn.org/stable/datasets/toy_dataset.html (accessed May 05, 2021).

[2] "numeth2021/NuMeth_4_5_Applied_Linear_Regression.ipynb at main · dyjdlopez/numeth2021 · GitHub." https://github.com/dyjdlopez/numeth2021/blob/main/Week%209-13%20-%20Curve%20Fitting%20Techniques/NuMeth_4_5_Applied_Linear_Regression.ipynb (accessed May 05, 2021).

[3] "numeth2021/NuMeth_4_Curve_Fitting.ipynb at main · dyjdlopez/numeth2021 · GitHub." https://github.com/dyjdlopez/numeth2021/blob/main/Week%209-13%20-%20Curve%20Fitting%20Techniques/NuMeth_4_Curve_Fitting.ipynb (accessed May 05, 2021).

[4] "Implementing and Visualizing Linear Regression in Python with SciKit Learn | by Sthitaprajna Mishra | Becoming Human: Artificial Intelligence Magazine." https://becominghuman.ai/implementing-and-visualizing-linear-regression-in-python-with-scikit-learn-a073768dc688 (accessed May 05, 2021).

[5] "Implementation of Simple Linear Regression Using Normal Equation(Matrices) | by Pratik Shukla | Medium." https://medium.com/@shuklapratik22/implementation-of-simple-linear-regression-using-normal-equation-matrices-f9021c3590da (accessed May 05, 2021).

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